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(54) Title: SYSTEMS AND METHODS FOR TARGETING CANCER CELLS

(57) Abstract: The present disclosure provides an immune cell genetically modified to produce two antigentriggered polypeptides, each recognizing a different cell surface antigen. The present disclosure provides a system two antigen-triggered polypeptides, each recognizing a different cell surface antigen. The present disclosure provides a method of killing a target cancer cell, using a genetically modified immune cell or a system of the present disclosure. The present disclosure provides a computational method to identify target antigen pairs on a cancer cell.

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SYSTEMS AND METHODS FOR TARGETING CANCER CELLS

CROSS-REFERENCE

[0001] This application claims the benefit of U.S. Provisional Patent Application No. 62/333,106, filed May 6, 2016, which application is incorporated herein by reference in its entirety.

STATEMENT REGARDING FEDERALLY SPONSORED RESEARCH

[0002] This invention was made with government support under Grant No. GM071966 and GM071508 awarded by the National Institutes of Health/NIGMS and Grant No. HL117798 awarded by the National Institutes of Health/NHLBI. This invention was also made with government support under Grant No. R01 CA196277 awarded by the National Institutes of Health. The government has certain rights in the invention.

INCORPORATION BY REFERENCE OF SEQUENCE LISTING PROVIDED AS A TEXT FILE

[0003] A Sequence Listing is provided herewith as a text file, "UCSF-537WO_SeqList_ST25.txt" created on May 5, 2017 and having a size of 37,675 KB. The contents of the text file are incorporated by reference herein in their entirety.

INTRODUCTION

[0004] Immune cell activation can be engineered *ex vivo* through expression of various designer antigen-triggered immune cell receptors including, e.g., synthetic chimeric antigen receptors (CAR), synthetic Notch polypeptides (synNotch), inhibitory CARs (iCARs), split CARs, and engineered T cell Receptors (TCR). A goal of such immune cell activation is targeting and killing of cancer cells in a patient, while avoiding or at least minimizing killing of non-cancerous cells.

SUMMARY

[0005] The present disclosure provides an immune cell genetically modified to produce two antigen-triggered polypeptides, each recognizing a different cell surface antigen. The present disclosure provides a system comprising two antigen-triggered polypeptides (or nucleic acids encoding same), each recognizing a different cell surface antigen. The present disclosure provides a method of killing a target cancer cell, using a genetically modified immune cell or a system of the present disclosure. The present disclosure provides a computational method to identify target antigen pairs on a cancer cell.

BRIEF DESCRIPTION OF THE DRAWINGS

[0006] **FIG. 1** provides antigen combinations for various cancers.

[0007] **FIG. 2** provides amino acid sequences of antigens within antigen combinations.

[0008] **FIG. 3A-3D** provide schematic depictions of various antigen-triggered polypeptides; and AND and AND-NOT logic gates using the antigen-triggered polypeptides.

[0009] **FIG. 4** provides cancer-associated target antigens in clinical trials.

[0010] **FIG. 5A-5G** provide schematic depictions of exemplary synNotch receptor Notch regulatory regions.

[0011] **FIG. 6A-6F** provide schematic depictions of exemplary split CARs.

[0012] **FIG. 7** provides an example of a decision tree for identifying AND gate antigen pairs.

[0013] **FIG. 8** provides an example of a decision tree for identifying AND-NOT gate antigen pairs.

[0014] **FIG. 9** provides clinical antigen AND-gate combinations for various exemplary cancers.

[0015] **FIG. 10** provides clinical antigen AND NOT-gate combinations for various exemplary cancers.

[0016] **FIG. 11** provides novel antigen AND-gate combinations for various exemplary cancers.

[0017] **FIG. 12** provides novel antigen AND NOT-gate combinations for various exemplary cancers.

[0018] **FIG. 13** provides clinical-clinical antigen combinations for various exemplary cancers.

[0019] **FIG. 14** provides single antigens useful for various exemplary cancers.

[0020] **FIG. 15** provides three antigen combinations for various exemplary cancers.

DEFINITIONS

[0021] The terms "polynucleotide" and "nucleic acid," used interchangeably herein, refer to a polymeric form of nucleotides of any length, either ribonucleotides or deoxyribonucleotides. Thus, this term includes, but is not limited to, single-, double-, or multi-stranded DNA or RNA, genomic DNA, cDNA, DNA-RNA hybrids, or a polymer comprising purine and pyrimidine bases or other natural, chemically or biochemically modified, non-natural, or derivatized nucleotide bases.

[0022] The terms "polypeptide," "peptide," and "protein", used interchangeably herein, refer to a polymeric form of amino acids of any length, which can include genetically coded and non-genetically coded amino acids, chemically or biochemically modified or derivatized amino acids, and polypeptides having modified peptide backbones. The term includes fusion proteins, including, but not limited to, fusion proteins with a heterologous amino acid sequence, fusions with heterologous and homologous leader sequences, with or without N-terminal methionine residues; immunologically tagged proteins; and the like.

[0023] The terms "chimeric antigen receptor" and "CAR", used interchangeably herein, refer to artificial multi-module molecules capable of triggering or inhibiting the activation of an immune cell which generally but not exclusively comprise an extracellular domain (e.g., a ligand/antigen binding domain), a

transmembrane domain and one or more intracellular signaling domains. The term CAR is not limited specifically to CAR molecules but also includes CAR variants. CAR variants include split CARs wherein the extracellular portion (e.g., the ligand binding portion) and the intracellular portion (e.g., the intracellular signaling portion) of a CAR are present on two separate molecules. CAR variants also include ON-switch CARs which are conditionally activatable CARs, e.g., comprising a split CAR wherein conditional hetero-dimerization of the two portions of the split CAR is pharmacologically controlled. CAR variants also include bispecific CARs, which include a secondary CAR binding domain that can either amplify or inhibit the activity of a primary CAR. CAR variants also include inhibitory chimeric antigen receptors (iCARs) which may, e.g., be used as a component of a bispecific CAR system, where binding of a secondary CAR binding domain results in inhibition of primary CAR activation. CAR molecules and derivatives thereof (i.e., CAR variants) are described, e.g., in PCT Application No. US2014/016527; Fedorov et al. *Sci Transl Med* (2013) ;5(215):215ra172; Glienke et al. *Front Pharmacol* (2015) 6:21; Kakarla & Gottschalk 52 *Cancer J* (2014) 20(2):151-5; Riddell et al. *Cancer J* (2014) 20(2):141-4; Pegram et al. *Cancer J* (2014) 20(2):127-33; Cheadle et al. *Immunol Rev* (2014) 257(1):91-106; Barrett et al. *Annu Rev Med* (2014) 65:333-47; Sadelain et al. *Cancer Discov* (2013) 3(4):388-98; Cartellieri et al., *J Biomed Biotechnol* (2010) 956304; the disclosures of which are incorporated herein by reference in their entirety.

[0024] As used herein, the term "immune cells" generally includes white blood cells (leukocytes) which are derived from hematopoietic stem cells (HSC) produced in the bone marrow. "Immune cells" includes, e.g., lymphocytes (T cells, B cells, natural killer (NK) cells) and myeloid-derived cells (neutrophil, eosinophil, basophil, monocyte, macrophage, dendritic cells).

[0025] "T cell" includes all types of immune cells expressing CD3 including T-helper cells (CD4+ cells), cytotoxic T-cells (CD8+ cells), T-regulatory cells (Treg) and gamma-delta T cells.

[0026] A "cytotoxic cell" includes CD8+ T cells, natural-killer (NK) cells, and neutrophils, which cells are capable of mediating cytotoxicity responses.

[0027] As used herein, the terms "treatment," "treating," and the like, refer to obtaining a desired pharmacologic and/or physiologic effect. The effect may be prophylactic in terms of completely or partially preventing a disease or symptom thereof and/or may be therapeutic in terms of a partial or complete cure for a disease and/or adverse effect attributable to the disease. "Treatment," as used herein, covers any treatment of a disease in a mammal, e.g., in a human, and includes: (a) preventing the disease from occurring in a subject which may be predisposed to the disease but has not yet been diagnosed as having it; (b) inhibiting the disease, i.e., arresting its development; and (c) relieving the disease, i.e., causing regression of the disease.

[0028] The terms "individual," "subject," "host," and "patient," used interchangeably herein, refer to a mammal, including, but not limited to, murines (e.g., rats, mice), lagomorphs (e.g., rabbits), non-human

primates, humans, canines, felines, ungulates (e.g., equines, bovines, ovines, porcines, caprines), etc. In some cases, the individual is a human.

[0029] A "therapeutically effective amount" or "efficacious amount" refers to the amount of an agent, or combined amounts of two agents, that, when administered to a mammal or other subject for treating a disease, is sufficient to effect such treatment for the disease. The "therapeutically effective amount" will vary depending on the agent(s), the disease and its severity and the age, weight, etc., of the subject to be treated.

[0030] Before the present invention is further described, it is to be understood that this invention is not limited to particular embodiments described, as such may, of course, vary. It is also to be understood that the terminology used herein is for the purpose of describing particular embodiments only, and is not intended to be limiting, since the scope of the present invention will be limited only by the appended claims.

[0031] Where a range of values is provided, it is understood that each intervening value, to the tenth of the unit of the lower limit unless the context clearly dictates otherwise, between the upper and lower limit of that range and any other stated or intervening value in that stated range, is encompassed within the invention. The upper and lower limits of these smaller ranges may independently be included in the smaller ranges, and are also encompassed within the invention, subject to any specifically excluded limit in the stated range. Where the stated range includes one or both of the limits, ranges excluding either or both of those included limits are also included in the invention.

[0032] Unless defined otherwise, all technical and scientific terms used herein have the same meaning as commonly understood by one of ordinary skill in the art to which this invention belongs. Although any methods and materials similar or equivalent to those described herein can also be used in the practice or testing of the present invention, the preferred methods and materials are now described. All publications mentioned herein are incorporated herein by reference to disclose and describe the methods and/or materials in connection with which the publications are cited.

[0033] It must be noted that as used herein and in the appended claims, the singular forms "a," "an," and "the" include plural referents unless the context clearly dictates otherwise. Thus, for example, reference to "a target antigen" includes a plurality of such antigens and reference to "the system" includes reference to one or more systems and equivalents thereof known to those skilled in the art, and so forth. It is further noted that the claims may be drafted to exclude any optional element. As such, this statement is intended to serve as antecedent basis for use of such exclusive terminology as "solely," "only" and the like in connection with the recitation of claim elements, or use of a "negative" limitation.

[0034] It is appreciated that certain features of the invention, which are, for clarity, described in the context of separate embodiments, may also be provided in combination in a single embodiment. Conversely, various features of the invention, which are, for brevity, described in the context of a single

embodiment, may also be provided separately or in any suitable sub-combination. All combinations of the embodiments pertaining to the invention are specifically embraced by the present invention and are disclosed herein just as if each and every combination was individually and explicitly disclosed. In addition, all sub-combinations of the various embodiments and elements thereof are also specifically embraced by the present invention and are disclosed herein just as if each and every such sub-combination was individually and explicitly disclosed herein.

[0035] The publications discussed herein are provided solely for their disclosure prior to the filing date of the present application. Nothing herein is to be construed as an admission that the present invention is not entitled to antedate such publication by virtue of prior invention. Further, the dates of publication provided may be different from the actual publication dates which may need to be independently confirmed.

DETAILED DESCRIPTION

[0036] The present disclosure provides an immune cell genetically modified to produce two antigen-triggered polypeptides, each recognizing a different cell surface antigen. The present disclosure provides a system of two antigen-triggered polypeptides, each recognizing a different cell surface antigen. The present disclosure provides a method of killing a target cancer cell, using a genetically modified immune cell or a system of the present disclosure.

[0037] The present disclosure provides an *in vitro* genetically modified cytotoxic immune cell, where the cytotoxic immune cell is genetically modified to produce two different antigen-triggered polypeptides that recognize two different cell surface antigens, and where at least one of the two different cell surface antigens is present on the surface of a target cancer cell. In some cases, the two different antigen-triggered polypeptides comprise: a) a first antigen-triggered polypeptide that binds specifically to a first target cell surface antigen present on a target cancer cell; and b) a second antigen-triggered polypeptide that binds specifically to a second target cell surface antigen. In some cases, the genetically modified cytotoxic immune cell is a genetically modified cytotoxic T cell or a genetically modified natural killer cell. In some cases, the two different antigen-triggered polypeptides provide an AND gate; thus, for example, in some cases, the genetically modified cytotoxic immune cell is activated to kill a target cancer cell only when the target cancer cell expresses both of the two different cell surface antigens on its cell surface. In some cases, the two different antigen-triggered polypeptides provide an AND-NOT gate; thus, for example, in some cases, the genetically modified cytotoxic immune cell: a) is activated to kill a target cancer cell that expresses the first target cell surface antigen, but not the second target cell surface antigen, on its cell surface; and b) is inhibited from killing a non-cancerous cell if the non-cancerous cell expresses both the first target cell surface antigen and the second target cell surface antigen on its cell surface.

[0038] The present disclosure provides a system for killing a target cancer cell, the system comprising: a) a first antigen-triggered polypeptide that binds specifically to a first target antigen present on the target cancer cell, or a first nucleic acid comprising a nucleotide sequence encoding the first antigen-triggered polypeptide; and b) a second antigen-triggered polypeptide that binds specifically to a second target antigen, or a second nucleic acid comprising a nucleotide sequence encoding the second antigen-triggered polypeptide. In some cases, the system provides an AND gate; thus, for example, in some cases, the first target antigen and the second target antigen are both present on the surface of a target cancer cell. In some cases, the system provides an AND-NOT gate; thus, for example, in some cases: a) the first target antigen and the second target antigen are both present on the surface of a non-cancerous cell; and b) the first target antigen, but not the second target antigen, is present on the surface of a target cancer cell. A system of the present disclosure can be introduced *ex vivo* into an immune cell obtained from a patient, to generate a modified immune cell; and the modified immune cell can be introduced into the patient from whom the immune cell was obtained.

[0039] The present disclosure provides a method of killing a target cancer cell in an individual. In some cases, a method of the present disclosure for killing a target cell in an individual comprises: a) introducing a system of the present disclosure into an immune cell (e.g., a $CD8^+$ T cell; an NK cell) obtained from the individual, generating a modified immune cell; and b) administering the modified immune cell to the individual, where the modified immune cell kills the target cancer cell in the individual.

[0040] The present disclosure provides a method of killing a target cancer cell in an individual. In some cases, a method of the present disclosure for killing a target cell in an individual comprises administering a genetically modified cytotoxic immune cell (e.g., a genetically modified $CD8^+$ T cell; a genetically modified NK cell) of the present disclosure to the individual, where the genetically modified immune cell kills the target cancer cell in the individual.

[0041] As noted above, a genetically modified cytotoxic immune cell of the present disclosure, and a system of the present disclosure, involve at least two antigen-triggered polypeptides that recognize two different cell surface antigens. A pair of antigen-triggered polypeptides recognizes and binds to a pair of target antigens; antigen binding activates the antigen-triggered polypeptides. Thus, a first antigen-triggered polypeptide binds a first member of a target antigen pair; and a second antigen-triggered polypeptide binds a second member of the target antigen pair. Target antigen combinations (also referred to herein as "target antigen pairs") are provided in FIG. 1 and FIG. 9-14. At least one of the two antigens of a target antigen pair listed in FIG. 1 or FIG. 9-14 is present on the surface of a target cancer cell. In some cases, the second target antigen of a target antigen pair is present on the surface of the same target cancer cell as the first target antigen of the target antigen pair. In some cases, the first target antigen of the target antigen pair is present on the surface of a target cancer cell, and the second target antigen of a

target antigen pair is not present on the surface of the same target cancer cell; in these cases, both antigens of the target antigen pair are present on the surface of a non-cancerous cell. The target antigen combinations presented in FIG. 1 and FIG. 9-14 provide for an AND logic gate or an AND-NOT logic gate for a particular cancer cell type.

[0042] Where a target antigen pair provides for an AND logic gate, both antigens must be present on the surface of a target cancer cell in order for a genetically modified cytotoxic immune cell of the present disclosure to kill the target cancer cell, where in this case the genetically modified cytotoxic immune cell is genetically modified to express two antigen-triggered polypeptides, each recognizing one of the target antigens of the target antigen pair. For example, where a target antigen pair present in FIG. 1 and/or FIG. 9-14 is indicated as providing an AND logic gate, both target antigens of the target antigen pair must be present on the surface of a target cancer cell in order for a genetically modified cytotoxic immune cell of the present disclosure to kill the target cancer cell; and the genetically modified cytotoxic immune cell must express both a first antigen-triggered polypeptide that specifically binds the first target antigen of the target antigen pair and a second triggered polypeptide that specifically binds the second antigen of the target antigen pair. For example, in some cases, expression of the second antigen-triggered polypeptide is induced when the first antigen-triggered polypeptide binds to the first target antigen of the target antigen pair.

[0043] Where a target antigen pair provides an AND-NOT logic gate, a genetically modified cytotoxic immune cell of the present disclosure: a) is activated to kill a target cancer cell that expresses the first target cell surface antigen, but not the second target cell surface antigen, on its cell surface; and b) is inhibited from killing a non-cancerous cell if the non-cancerous cell expresses both the first target cell surface antigen and the second target cell surface antigen on its cell surface; in these cases, the genetically modified cytotoxic immune cell must express both a first antigen-triggered polypeptide that specifically binds the first target antigen of the target antigen pair and a second triggered polypeptide that specifically binds the second antigen of the target antigen pair. For example, in some cases, binding of the second antigen-triggered polypeptide to the second target cell surface antigen (expressed on a non-cancerous cell) inhibits T cell activation that would normally be induced by binding of the first antigen-triggered polypeptide to the first target antigen (present on the cancer cell surface and on the non-cancerous cell surface). In this manner, unintended/undesired killing of a non-cancerous cell is reduced, because the target cancer cell expressing the first target antigen and not the second target antigen will be preferentially killed over the non-cancerous cell expressing both the first target antigen and the second target antigen. Since the cancer cell does not express the second target cell surface antigen (expressed on a non-cancerous cell), binding of the first antigen-triggered polypeptide to the first target antigen (present on the cancer cell surface) results in activation of the genetically modified cytotoxic T cell and killing of the cancer cell.

[0044] In some cases, the first antigen-triggered polypeptide is a synNotch receptor and the second antigen-triggered polypeptide is a chimeric antigen receptor (CAR). In some cases, the first antigen-triggered polypeptide is a synNotch receptor and the second antigen-triggered polypeptide is a T cell receptor (TCR). In some cases, the first antigen-triggered polypeptide is a synNotch receptor, and the second antigen-triggered polypeptide is a split CAR (e.g., an ON-switch CAR). In some cases, the first antigen-triggered polypeptide is a synNotch receptor, and the second antigen-triggered polypeptide is one polypeptide chain of a split CAR (e.g., an ON-switch CAR). In some cases, the first antigen-triggered polypeptide is a synNotch receptor, and the second antigen-triggered polypeptide is another synNotch receptor. Any or either of the first and second antigen-triggered polypeptides of the subject systems may independently be a synNotch receptor, a CAR, a TCR or the like. In some cases, both the first and second antigen-triggered polypeptides of the subject systems may be a synNotch receptor, a CAR, a TCR or the like.

[0045] In some cases, the first antigen-triggered polypeptide is a CAR, and the second antigen-triggered polypeptide is an antigen-binding inhibitory polypeptide, such as e.g. an inhibitory CAR (iCAR). In some cases, the first antigen-triggered polypeptide is a TCR, and the second antigen-triggered polypeptide is an antigen-binding inhibitory polypeptide, such as e.g. an iCAR. In some cases, the first antigen-triggered polypeptide is an ON-switch CAR, and the second antigen-triggered polypeptide is an antigen-binding inhibitory polypeptide, such as e.g. an iCAR. In some cases, the first antigen-triggered polypeptide is a CAR, and the second antigen-triggered polypeptide is a synNotch receptor. In some cases, the first antigen-triggered polypeptide is a TCR, and the second antigen-triggered polypeptide is a synNotch receptor. In some cases, the first antigen-triggered polypeptide is an ON-switch CAR, and the second antigen-triggered polypeptide is a synNotch receptor.

[0046] In some cases, the target cancer cell is a liposarcoma, a glioblastoma, a breast cancer cell, a renal cancer cell, a pancreatic cancer cell, a melanoma, an anaplastic lymphoma, a leiomyosarcoma, an astrocytoma, an ovarian cancer cell, a neuroblastoma, a mantle cell lymphoma, a sarcoma, a non-small cell lung cancer cell, an AML cell, a stomach cancer cell, a B-cell cancer cell, a lung cancer cell, or an oligodendroglioma.

AND gate target antigen pairs

[0047] As noted above, in some cases, expression of the second antigen-triggered polypeptide in a genetically modified immune cell is induced only when the first antigen-triggered polypeptide binds to the first target antigen of the target antigen pair, where the binding to the first target antigen activates the first antigen-triggered polypeptide. Non-limiting examples of 2-input AND gates (AND gates based on 2 target antigens) and 3-input AND gates (AND gates based on 3 target antigens) are depicted schematically in FIG. 3A-3B.

[0048] For example, in some cases, the first antigen-triggered polypeptide is a synNotch receptor and activation of the synNotch receptor by binding to the first antigen (present on a target cancer cell) induces expression of the second antigen-triggered polypeptide. The second antigen-triggered polypeptide binds to the second antigen of the target antigen pair, where the second antigen is expressed on the surface of the target cancer cell. As an example, in some cases, the first antigen-triggered polypeptide is a synNotch receptor and the second antigen-triggered polypeptide is a single chain CAR. As another example, in some cases, the first antigen-triggered polypeptide is a synNotch receptor and the second antigen-triggered polypeptide is a TCR. For example, in some cases, the synNotch polypeptide comprises an intracellular domain comprising a transcriptional activator, and activation of the synNotch receptor by binding to the first antigen (present on a target cancer cell) induces release of the transcriptional activator; the released transcriptional activator activates transcription of the TCR or the single-chain CAR.

[0049] As another example, in some cases, the first antigen-triggered polypeptide is a synNotch receptor and activation of the synNotch receptor by binding to the first antigen (present on a target cancer cell) induces expression of the second antigen-triggered polypeptide, where the second antigen-triggered polypeptide is a heterodimeric ("two chain" or "split") CAR comprising a first polypeptide chain and a second polypeptide chain. The heterodimeric CAR binds to the second antigen of the target antigen pair, where the second antigen is expressed on the surface of the target cancer cell. For example, in some cases, the first antigen-triggered polypeptide is a synNotch receptor and the second antigen-triggered polypeptide is a split CAR (e.g., an ON-switch CAR). In some cases, activation of the synNotch receptor by binding to the first antigen (present on a target cancer cell) induces expression of only the first polypeptide chain of the heterodimeric CAR; expression of the second polypeptide chain of the heterodimeric CAR can be constitutive. For example, in some cases, the synNotch polypeptide comprises an intracellular domain comprising a transcriptional activator, and activation of the synNotch receptor by binding to the first antigen (present on a target cancer cell) induces release of the transcriptional activator; the released transcriptional activator activates transcription of the first polypeptide chain of the heterodimeric CAR. In some cases, activation of the synNotch receptor by binding to the first antigen (present on a target cancer cell) induces expression of only the second polypeptide chain of the heterodimeric CAR; expression of the first polypeptide chain of the heterodimeric CAR can be constitutive. Once the first polypeptide chain of the heterodimeric CAR is produced in the cell, it heterodimerizes with the second polypeptide chain of the heterodimeric CAR. As another example, in some cases, the synNotch polypeptide comprises an intracellular domain comprising a transcriptional activator, and activation of the synNotch receptor by binding to the first antigen (present on a target cancer cell) induces release of the transcriptional activator; the released transcriptional activator activates transcription of the second polypeptide chain of the heterodimeric CAR.

[0050] In AND gate systems, unintended/undesired killing of non-target cells is reduced; for example, a cell that expresses on its cell surface only one of the target antigen pair is not killed by a genetically modified cytotoxic immune cell of the present disclosure.

[0051] In some cases, a genetically modified immune cell or a system of the present disclosure provides for a 3-input AND gate, comprising: 1) a first target antigen of an AND-gate target antigen pair depicted in FIG. 1 or FIG. 9-14; 2) a second target antigen of the AND-gate target antigen pair; and 3) a third target antigen, where the third target antigen is a target antigen depicted in FIG. 4 or Table 3, where e.g., the third target antigen may be a target antigen for the same cancer cell type as the target antigen pair. As one non-limiting example, where the target AND-gate antigen pair is SYT11 AND TNFRSF17 (for a glioma cell) (as shown in FIG. 1), a suitable third target antigen is EphA2 (as shown in FIG. 4).

AND-NOT gate target antigen pairs

[0052] As noted above, in some cases, where a target antigen pair provides an AND-NOT logic gate, a genetically modified cytotoxic immune cell of the present disclosure: a) is activated to kill a target cancer cell that expresses the first target cell surface antigen, but not the second target cell surface antigen, on its cell surface; and b) is inhibited from killing a non-cancerous cell if the non-cancerous cell expresses both the first target cell surface antigen and the second target cell surface antigen on its cell surface; in these cases, the genetically modified cytotoxic immune cell must express both a first antigen-triggered polypeptide that specifically binds the first target antigen of the target antigen pair and a second triggered polypeptide that specifically binds the second antigen of the target antigen pair. Non-limiting examples of 2-input AND-NOT gates (AND-NOT gates based on 2 target antigens) and 3-input AND-NOT gates (AND-NOT gates based on 3 target antigens) are depicted schematically in FIG. 3C-3D.

[0053] As an example, in some cases, the first antigen-triggered polypeptide is a CAR, and the second antigen-triggered polypeptide is an iCAR. Binding of the iCAR to the second antigen (present on the surface of a non-cancerous cell, but not on the surface of a target cancer cell) of a target antigen pair inhibits T-cell activation mediated by activation of the CAR upon binding to the first antigen (present on the surface of the target cancer cell and on the surface of the non-cancerous cell) of the target antigen pair. As another example, in some cases, the first antigen-triggered polypeptide is a TCR, and the second antigen-triggered polypeptide is an iCAR. Binding of the iCAR to the second antigen (present on the surface of a non-cancerous cell, but not on the surface of a target cancer cell) of a target antigen pair blocks or reduces T-cell activation mediated by activation of the TCR upon binding to the first antigen (present on the surface of the target cancer cell and on the surface of the non-cancerous cell) of the target antigen pair.

[0054] The above provides examples of AND-NOT gates where the inhibitory component is an iCAR; however, as will be readily understood, the inhibitory components of combinatorial antigen gates having "NOT" functionality are not so limited and may generally include any polypeptide configured to inhibit

an activity, e.g., an activity induced by binding of a first activating antigen in an AND-NOT gate, including where such inhibition is conferred through the presence of an inhibitory domain. Inhibitory components of combinatorial antigen gates having "NOT" functionality may be specific for an antigen present on a non-target cell, including e.g., where such antigen is absent or present in low amounts on the surface of a target cell.

[0055] As another example, in some cases, the first antigen-triggered polypeptide is a CAR, and the second antigen-triggered polypeptide is a synNotch polypeptide comprising an intracellular domain that, when released upon activation of the synNotch polypeptide by binding to the second target antigen, induces expression of an intracellular inhibitor that inhibits T-cell activation mediated by activation of the CAR upon binding to the first antigen (present on the surface of the target cancer cell and on the surface of the non-cancerous cell) of the target antigen pair. As another example, in some cases, the first antigen-triggered polypeptide is a TCR, and the second antigen-triggered polypeptide is a synNotch polypeptide comprising an intracellular domain that, when released upon activation of the synNotch polypeptide by binding to the second target antigen, induces expression of an intracellular inhibitor that inhibits T-cell activation mediated by activation of the TCR upon binding to the first antigen (present on the surface of the target cancer cell and on the surface of the non-cancerous cell) of the target antigen pair.

[0056] As another example, in some cases, the first antigen-triggered polypeptide is a CAR, and the second antigen-triggered polypeptide is a synNotch polypeptide comprising an intracellular domain that, when released upon activation of the synNotch polypeptide by binding to the second target antigen, induces expression of an extracellular inhibitor that inhibits T-cell activation mediated by activation of the CAR upon binding to the first antigen (present on the surface of the target cancer cell and on the surface of the non-cancerous cell) of the target antigen pair. As another example, in some cases, the first antigen-triggered polypeptide is a TCR, and the second antigen-triggered polypeptide is a synNotch polypeptide comprising an intracellular domain that, when released upon activation of the synNotch polypeptide by binding to the second target antigen, induces expression of an extracellular inhibitor that inhibits T-cell activation mediated by activation of the TCR upon binding to the first antigen (present on the surface of the target cancer cell and on the surface of the non-cancerous cell) of the target antigen pair.

[0057] In some cases, a genetically modified immune cell or a system of the present disclosure provides for a 3-input AND, AND-NOT gate, comprising: 1) a first target antigen of an AND-NOT gate target antigen pair depicted in FIG. 1 or FIG. 9-14; 2) a second target antigen of the AND-NOT gate target antigen pair; and 3) a third target antigen, where the third target antigen is a target antigen depicted in FIG. 4 or Table 3, including e.g., where the third target antigen may be a target antigen for the same cancer cell type as the target antigen pair. In some cases, the 3-input AND, AND-NOT gate comprises:

1) the first target antigen of an AND-NOT gate target antigen pair depicted in FIG. 1 or FIG. 9-14; AND-NOT; 2) the second target antigen of the AND-NOT gate target antigen pair; AND 3) the third target antigen. In some cases, the 3-input AND, AND-NOT gate comprises: 1) the second target antigen of an AND-NOT gate target antigen pair depicted in FIG. 1 or FIG. 9-14; AND-NOT 2) the first target antigen of the AND-NOT gate target antigen pair; AND 3) the third target antigen.

[0058] As one non-limiting example, where the target AND-NOT gate antigen pair is NLGN1 AND-NOT TNFRSF17 (for a glioma cell) (as shown in FIG. 1), a suitable third target antigen is EphA2 (as shown in FIG. 4).

Antigen-triggered polypeptides

[0059] As noted above, an antigen-triggered polypeptide can be a synNotch polypeptide; a CAR; or a TCR. A CAR can be an ON-switch ("split") CAR, a single-chain CAR, an iCAR, etc. Schematic depictions of examples of antigen-triggered polypeptides are provided in FIG. 5 and FIG. 6A-6F.

synNotch polypeptides

[0060] As noted above, in some cases an antigen-triggered polypeptide produced in a genetically modified immune cell of the present disclosure, or present in a system of the present disclosure, or encoded by a nucleotide sequence in a nucleic acid present in a system of the present disclosure, is a synNotch receptor (a "synNotch polypeptide"). synNotch polypeptides are described in PCT/US16/19188, the disclosure of which is incorporated herein by reference in its entirety. Schematic depictions of exemplary synNotch polypeptides are provided in FIG. 5A-5G.

[0061] In some cases, a synNotch polypeptide is a chimeric Notch polypeptide comprising, from N-terminus to C-terminus and in covalent linkage: a) an extracellular domain comprising an antigen binding member that is not naturally present in a Notch receptor polypeptide and that specifically binds to an antigen; b) a Notch regulatory region comprising a Lin 12-Notch repeat, an S2 proteolytic cleavage site, and a transmembrane domain comprising an S3 proteolytic cleavage site; c) an intracellular domain comprising a transcriptional activator or a transcriptional repressor that is heterologous to the Notch regulatory region and replaces a naturally-occurring intracellular Notch domain, wherein binding of the antigen binding member to the antigen, present on a cell or other solid support, induces cleavage at the S2 and S3 proteolytic cleavage sites, thereby releasing the intracellular domain.

[0062] In some cases, a synNotch polypeptide is a chimeric Notch polypeptide comprising, from N-terminus to C-terminus and in covalent linkage: a) an extracellular domain comprising a single-chain Fv (scFv) or a nanobody that specifically binds to an antigen; b) a Notch regulatory region comprising a Lin 12-Notch repeat, a heterodimerization domain comprising an S2 proteolytic cleavage site and a transmembrane domain comprising an S3 proteolytic cleavage site; and c) an intracellular domain, heterologous to the Notch regulatory region, comprising a transcriptional activator comprising a DNA binding domain, wherein the transcriptional activator replaces a naturally-occurring intracellular notch

domain, and wherein binding of the scFv or the nanobody to the antigen in trans induces cleavage at the S2 and S3 proteolytic cleavage sites, thereby releasing the intracellular domain and wherein the chimeric Notch polypeptide does not bind its naturally-occurring ligand Delta.

[0063] In some cases, a synNotch polypeptide comprises, from N-terminal to C-terminal and in covalent linkage: a) an extracellular domain that specifically binds to an antigen (e.g., a single-chain Fv (scFv), a nanobody, and the like); b) a Notch receptor polypeptide comprising one or more proteolytic cleavage sites and having at least 50%, at least 55%, at least 60%, at least 65%, at least 70%, at least 75%, at least 80%, at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to a Notch receptor polypeptide as described below; and c) an intracellular domain comprising a transcriptional regulator (e.g., a transcriptional activator or a transcriptional repressor). When the synNotch polypeptide is present in a cell, binding of the extracellular domain (e.g., the scFv or the nanobody) to the antigen induces cleavage of the Notch receptor polypeptide at the one or more proteolytic cleavage sites, thereby releasing the intracellular domain. In some cases, the intracellular domain comprises a transcriptional activator. In some cases, the intracellular domain comprises a transcriptional repressor.

[0064] In some cases, a Notch receptor polypeptide comprises an amino acid sequence having at least 50%, at least 55%, at least 60%, at least 65%, at least 70%, at least 75%, at least 80%, at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence of a Notch receptor including e.g., any of SEQ ID NOs: 7240- 7245 and 7468-7475. In some instances, the Notch regulatory region of a Notch receptor polypeptide is a mammalian Notch regulatory region, including but not limited to e.g., a mouse Notch (e.g., mouse Notch1, mouse Notch2, mouse Notch3 or mouse Notch4) regulatory region, a rat Notch regulatory region (e.g., rat Notch1, rat Notch2 or rat Notch3), a human Notch regulatory region (e.g., human Notch1, human Notch2, human Notch3 or human Notch4), and the like or a Notch regulatory region derived from a mammalian Notch regulatory region and having at least 50%, at least 55%, at least 60%, at least 65%, at least 70%, at least 75%, at least 80%, at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence of a mammalian Notch regulatory region of a mammalian Notch receptor amino acid sequence, including e.g., SEQ ID NOs: 7240-7245 and 7468-7475.

[0065] Subject Notch regulatory regions may include or exclude various components (e.g., domains, cleavage sites, etc.) thereof. Examples of such components of Notch regulatory regions that may be present or absent in whole or in part, as appropriate, include e.g., one or more EGF-like repeat domains, one or more Lin12/Notch repeat domains, one or more heterodimerization domains (e.g., HD-N or HD-C), a transmembrane domain, one or more proteolytic cleavage sites (e.g., a furin-like protease site (e.g., an S1 site), an ADAM-family protease site (e.g., an S2 site) and/or a gamma-secretase protease site (e.g.,

an S3 site)), and the like. Notch receptor polypeptides may, in some instances, exclude all or a portion of one or more Notch extracellular domains, including e.g., Notch-ligand binding domains such as Delta-binding domains. Notch receptor polypeptides may, in some instances, include one or more non-functional versions of one or more Notch extracellular domains, including e.g., Notch-ligand binding domains such as Delta-binding domains. Notch receptor polypeptides may, in some instances, exclude all or a portion of one or more Notch intracellular domains, including e.g., Notch Rbp-associated molecule domains (i.e., RAM domains), Notch Ankyrin repeat domains, Notch transactivation domains, Notch PEST domains, and the like. Notch receptor polypeptides may, in some instances, include one or more non-functional versions of one or more Notch intracellular domains, including e.g., non-functional Notch Rbp-associated molecule domains (i.e., RAM domains), non-functional Notch Ankyrin repeat domains, non-functional Notch transactivation domains, non-functional Notch PEST domains, and the like.

[0066] In some cases, a Notch receptor polypeptide has a length of from about 310 amino acids (aa) to about 320 aa (e.g., 310 aa, 311 aa, 312 aa, 313 aa, 314 aa, 315 aa, 316 aa, 317 aa, 318 aa, 319 aa, or 320 aa).

[0067] In some cases, a Notch receptor polypeptide comprises an amino acid sequence having at least 50%, at least 55%, at least 60%, at least 65%, at least 70%, at least 75%, at least 80%, at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the following amino acid sequence:

PPQIEEACELPECQVDAGNKVCNLQCNNHACGWDGGDCSLNFNDPWKNCTQSLQCWKYFSDG HCDSQCNSAGCLFDGFDCQLTEGQCNPLYDQYCKDHFSDGHCDQGCNSAECEWDGLDCAEHV PERLAAGTLVLVVLLPPDQLRNNSFHFLRELSHVLHTNVVFKRDAQGQQMIFPYYGHEEELRKH PIKRSTVGWATSSLLPGTSGGRQRRELDPMDIRGSIVYLEIDNRQCVQSSSQCFQSATDVAAFLG ALASLGSLNIPYKIEAVKSEPVEPPLPSQLHLMYVAAAAFVLLFFVGCGVLLS (SEQ ID NO:7246); and has a length of from 300 amino acids to 310 amino acids (e.g., 300, 301, 302, 303, 304, 305, 306, 307, 308, 309, or 310 amino acids).

[0068] In some cases, a Notch receptor polypeptide comprises an amino acid sequence having at least 50%, at least 55%, at least 60%, at least 65%, at least 70%, at least 75%, at least 80%, at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the following amino acid sequence:

PCVGSNPCYNQGTCEPTSENPFYRCLCPAKFNGLLCHILDYSFTGGAGRDIPPPQIEEACELPECQ VDAGNKVCNLQCNNHACGWDGGDCSLNFNDPWKNCTQSLQCWKYFSDGHCDSQCNSAGCLF DGFDCQLTEGQCNPLYDQYCKDHFSDGHCDQGCNSAECEWDGLDCAEHVPERLAAGTLVLVV LLPPDQLRNNSFHFLRELSHVLHTNVVFKRDAQGQQMIFPYYGHEEELRKHPIKRSTVGWATSS LLPGTSGGRQRRELDPMDIRGSIVYLEIDNRQCVQSSSQCFQSATDVAAFLGALASLGSLNIPYKI EAVKSEPVEPPLPSQLHLMYVAAAAFVLLFFVGCGVLLS (SEQ ID NO:7247); and has a length of

from 350 amino acids to 370 amino acids (e.g., 350 351, 352, 353, 354, 355, 356, 357, 358, 359, 360, 361, 362, 363, 364, 365, 366, 367, 368, 369, or 370 amino acids).

[0069] In some cases, the intracellular domain comprises a transcriptional activator, and release of the intracellular domain causes the transcriptional activator to induce expression of an endogenous gene product in a cell. In some cases, the intracellular domain comprises a transcriptional activator, and release of the intracellular domain causes the transcriptional activator to induce expression of a heterologous gene product (e.g., a CAR; a TCR; a therapeutic antibody). For example, in some cases, a transcriptional control element, responsive to the transcriptional activator, is operably linked to a nucleotide sequence encoding a CAR or a TCR. As another example, a transcriptional control element, responsive to the transcriptional activator, is operably linked to a nucleotide sequence encoding a therapeutic antibody for the treatment of cancer.

[0070] In some cases, the intracellular domain comprises a transcriptional activator, and release of the intracellular domain causes the transcriptional activator to induce expression of an intracellular inhibitor that inhibits T-cell activation. In some cases, the intracellular domain comprises a transcriptional activator, and release of the intracellular domain causes the transcriptional activator to induce expression of an inhibitor that is secreted from the cell (an extracellular inhibitor), where the extracellular inhibitor inhibits T-cell activation.

[0071] In some cases, the Notch receptor polypeptide comprises, at its N-terminus, one or more epidermal growth factor (EGF) repeats. In some cases, the Notch receptor polypeptide comprises, at its N-terminus, 2 to 11 EGF repeats; e.g., in some cases, the Notch receptor polypeptide comprises, at its N-terminus, 2, 3, 4, 5, 6, 7, 8, 9, 10, or 11 epidermal growth factor (EGF) repeats. An EGF repeat can have a length of from 35 amino acids (aa) to 45 aa (e.g., 35, 36, 37, 38, 39, 40, 41, 42, 43, 44, or 45 aa). An EGF repeat can comprise an amino acid sequence having at least 75%, at least 80%, at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the following sequence: PCVGSNPCYNQGTCEPTSENPFYRCLCPAKFNGLLCH (SEQ ID NO:7248); and can have a length of from 35 amino acids to 40 amino acids (e.g., 35, 36, 37, 38, 39, or 40 amino acids. An EGF repeat can comprise an amino acid sequence having at least 75%, at least 80%, at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the following sequence: PCLGGNPCYNQGTCEPTSESPFYRCLCPAKFNGLLCH (SEQ ID NO:7249); and can have a length of from 37 amino acids to 40 amino acids (e.g., 37, 38, 39, or 40 amino acids.

[0072] In some cases, the Notch receptor polypeptide comprises a synthetic linker between the one or more EGF repeats and the one or more proteolytic cleavage sites.

[0073] In some cases, the Notch receptor polypeptide has a length from 50 amino acids to 1000 amino acids. For example, in some cases, the Notch receptor polypeptide has a length of from 50 amino acids (aa) to 75 aa, from 75 aa to 100 aa, from 100 aa to 150 aa, from 150 aa to 200 aa, from 200 aa to 250 aa,

from 250 aa to 300 aa, from 300 aa to 400 aa, from 400 aa to 500 aa, from 500 aa to 600 aa, from 600 aa to 700 aa, from 700 aa to 800 aa, from 800 aa to 900 aa, or from 900 aa to 1000 aa. In some cases, the Notch receptor polypeptide has a length from 300 amino acids to 400 amino acids.

[0074] In some cases, the one or more proteolytic cleavage sites comprises an S2 proteolytic cleavage site, an S3 proteolytic cleavage site or a combination thereof. In some cases, an S3 proteolytic cleavage site comprises the amino acid sequence VLLS (SEQ ID NO:7250). In some cases, an S3 proteolytic cleavage site comprises the amino acid sequence GVLLS (SEQ ID NO:7251). In some cases, the one or more proteolytic cleavage sites comprises an S2 proteolytic cleavage site that is an ADAM family type protease cleavage site, such as e.g., an ADAM-17-type protease cleavage site comprising an Ala-Val dipeptide sequence. In some cases, the one or more proteolytic cleavage sites comprises an S3 proteolytic cleavage site that is a gamma-secretase (γ-secretase) cleavage site comprising a Gly-Val dipeptide sequence. In some cases, the one or more proteolytic cleavage sites further comprises an S1 proteolytic cleavage site. In some cases, the S1 proteolytic cleavage site is a furin-like protease cleavage site comprising the amino acid sequence Arg-X-(Arg/Lys)-Arg, where X is any amino acid. In some cases, the Notch receptor polypeptide does not include an S1 proteolytic cleavage site.

CARs

[0075] As noted above, in some cases an antigen-triggered polypeptide produced in a genetically modified immune cell of the present disclosure, or present in a system of the present disclosure, or encoded by a nucleotide sequence in a nucleic acid present in a system of the present disclosure, is a chimeric antigen receptor. Schematic depictions of split CARs are provided in FIG. 6A-6F.

[0076] The terms "chimeric antigen receptor" and "CAR", used interchangeably herein, refer to artificial multi-module molecules capable of triggering or inhibiting the activation of an immune cell which generally but not exclusively comprise an extracellular domain (e.g., a ligand/antigen binding domain), a transmembrane domain and one or more intracellular signaling domains. The term CAR is not limited specifically to CAR molecules but also includes CAR variants. CAR variants include split CARs wherein the extracellular portion (e.g., the ligand binding portion) and the intracellular portion (e.g., the intracellular signaling portion) of a CAR are present on two separate molecules. CAR variants also include ON-switch CARs which are conditionally activatable CARs, e.g., comprising a split CAR wherein conditional hetero-dimerization of the two portions of the split CAR is pharmacologically controlled. CAR variants also include bispecific CARs, which include a secondary CAR binding domain that can either amplify or inhibit the activity of a primary CAR. CAR variants also include inhibitory chimeric antigen receptors (iCARs) which may, e.g., be used as a component of a bispecific CAR system, where binding of a secondary CAR binding domain results in inhibition of primary CAR activation. CAR molecules and derivatives thereof (i.e., CAR variants) are described, e.g., in PCT Application No. US2014/016527; Fedorov et al. Sci Transl Med (2013) ;5(215):215ra172; Glienke et al.

Front Pharmacol (2015) 6:21; Kakarla & Gottschalk 52 Cancer J (2014) 20(2):151-5; Riddell et al. Cancer J (2014) 20(2):141-4; Pegram et al. Cancer J (2014) 20(2):127-33; Cheadle et al. Immunol Rev (2014) 257(1):91-106; Barrett et al. Annu Rev Med (2014) 65:333-47; Sadelain et al. Cancer Discov (2013) 3(4):388-98; Cartellieri et al., J Biomed Biotechnol (2010) 956304; the disclosures of which are incorporated herein by reference in their entirety.

[0077] Spit CAR may be extracellularly split or intracellularly split and may or may not be conditionally heterodimerizable. For example, split CAR systems that are not conditionally heterodimerizable may contain a constitutive heterodimerization domain or other binding pair (e.g., a Fc binding pair or other orthogonal binding pair) that does not depend on the presence of one or more additional molecules for the heterodimerization that results in the formation of an active CAR from assembly of the split portions.

[0078] In some instances, an extracellularly split CAR may be split extracellularly at the antigen binding domain into two parts including e.g., where the first part of the split CAR contains an extracellular Fc binding domain that specifically binds to second part of the split CAR that contains the antigen recognition domain.

[0079] In some instances, an extracellularly split CAR may be split extracellularly at the antigen binding domain into two parts including e.g., where the first part of the split CAR contains an first part of an orthogonal protein binding pair that specifically binds to the second part of the orthogonal protein binding pair that is contained in the second part of the split CAR that contains the antigen recognition domain.

[0080] In some instances, an intracellularly split CAR may be split intracellularly proximal to the transmembrane domain into two parts including e.g., where the first part of the split CAR includes the antigen recognition domain, a transmembrane domain and an intracellular first portion of a constitutive heterodimerization domain and the second part of the split CAR includes a transmembrane domain, the second portion of the constitutive heterodimerization domain proximal to the transmembrane domain, one or more co-stimulatory domains and one or more signaling domains (e.g., ITAM domains).

[0081] In some instances, an intracellularly split CAR may be split into two parts intracellularly proximal to an intracellular domain or between two intracellular domains including e.g., where the first part of the split CAR includes the antigen recognition domain, a transmembrane domain, one or more co-stimulatory domains and an intracellular first portion of a constitutive heterodimerization domain and the second part of the split CAR includes a transmembrane domain, one or more co-stimulatory domains, one or more signaling domains (e.g., ITAM domains) and the second portion of the constitutive heterodimerization domain between the one or more co-stimulatory domains and the one or more signaling domains.

[0082] In some instances, an intracellularly split CAR may be split into two parts intracellularly between intracellular domains including e.g., where the first part of the split CAR includes the antigen recognition

domain, a transmembrane domain, one or more co-stimulatory domains and an intracellular first portion of a constitutive heterodimerization domain proximal to the intracellular terminus of the first part of the split CAR and the second part of the split CAR includes a transmembrane domain, one or more signaling domains (e.g., ITAM domains) and the second portion of the constitutive heterodimerization domain between the transmembrane domain and the one or more signaling domains.

[0083] An inhibitory CAR (iCAR) expressed on an immunoresponsive cell specifically binds to an antigen, whereupon binding its antigen the iCAR inhibits the immunoresponsive cell. By "inhibits an immunoresponsive cell" or "suppresses an immunoresponsive cell" is meant induction of signal transduction or changes in protein expression in the cell resulting in suppression of an immune response (e.g., decrease in cytokine production).

[0084] Generally, but not exclusively, an iCAR is employed as a component of a bispecific CAR system where the activity of an immunostimulatory CAR (e.g., a CAR or CAR variant) is repressed by the iCAR upon binding of the iCAR to its antigen. An iCAR will generally include an extracellular domain that binds an antigen; a transmembrane domain operably linked to the extracellular domain; and an intracellular domain that activates intracellular signaling to decrease an immune response, the intracellular domain operably linked to the transmembrane domain. In some embodiments, the intracellular signaling domain is selected from the group consisting of a CTLA-4 polypeptide, a PD-1 polypeptide, a LAG-3 polypeptide, a 2B4 polypeptide, and a BTLA polypeptide. In certain embodiments, the transmembrane domain is selected from the group consisting of a CD4 polypeptide, a CD8 polypeptide, a CTLA-4 polypeptide, a PD-1 polypeptide, a LAG-3 polypeptide, a 2B4 polypeptide, and a BTLA polypeptide. In some instances, an iCAR, as described herein, may be or may be derived from one or more of the iCARs described in U.S. Patent Application Publication No. 20150376296, the disclosure of which is incorporated herein by reference in its entirety.

[0085] Any convenient extracellular binding domain (i.e., antigen binding domain) may find use in an iCAR including but not limited to e.g., a Fab, scFv, a monovalent or polyvalent ligand, etc., provided the domain is sufficient for specific binding of the iCAR to its antigen. In the contexts of therapy, the antigen binding domain of an iCAR will generally bind a healthy cell antigen in order to repress an immune response that may be otherwise triggered by presentation of a target antigen on the surface of a healthy cell. For example, in the contexts of cancer therapy, the antigen binding domain of an iCAR will generally bind a non-tumor or healthy cell antigen. For example, the extracellular domain may be a binding domain that does not bind one or more tumor antigens, including but not limited to e.g., CD19, CAIX, CEA, CD5, CD7, CD10, CD20, CD22, CD30, CD33, CD34, CD38, CD41, CD44, CD49f, CD56, CD74, CD123, CD133, CD138, a cytomegalovirus (CMV) infected cell antigen, EGP-2, EGP-40, EpCAM, erb-B2,3,4, F8P, Fetal acetylcholine receptor, folate receptor-a, GD2, GD3, HER-2, hTERT, IL-13R-.alpha.2, K-light chain, KDR, LeY, L1 cell adhesion molecule, MAGE-A1, Mesothelin, Muc-1,

Muc-16, NKG2D ligands, NY-ESO-1, oncofetal antigen (h5T4), PSCA, PSMA, ROR1, TAG-72, VEGF-R2, WT-1, and the like. Antigens to which the extracellular domain of an iCAR does not bind are, however, not limited to cancer antigens and may likewise exclude any target antigen to which an immunostimulatory CAR is directed.

[0086] In some instances, the antigen binding domain of an iCAR will bind a non-tumor antigen including but not limited to e.g., CD33, CD38, a human leukocyte antigen (HLA), an organ specific antigen, a blood-brain barrier specific antigen, an Epithelial-mesenchymal transition (EMT) antigen, E-cadherin, cytokeratin, Opioid-binding protein/cell adhesion molecule (OPCML), HYLA2, Deleted in Colorectal Carcinoma (DCC), Scaffold/Matrix attachment region-binding protein 1 (SMAR1), cell surface carbohydrate, mucin type O-glycan, etc.

[0087] In certain instances, an antigen to which the extracellular domain of an iCAR binds may be an antigen listed in Table 2, provided as FIG. 2. In some cases, an antigen to which the extracellular domain of an iCAR binds may be an antigen listed in Table 1, provided as FIG. 1, or in FIG. 9-14, which antigen is described as the "NOT" portion of an antigen logic gate.

TCRs

[0088] As noted above, in some cases an antigen-triggered polypeptide produced in a genetically modified immune cell of the present disclosure, or present in a system of the present disclosure, or encoded by a nucleotide sequence in a nucleic acid present in a system of the present disclosure, is a T-cell receptor (TCR).

[0089] A TCR generally includes an alpha chain and a beta chain; and recognizes antigen when presented by a major histocompatibility complex. In some cases, the TCR is an engineered TCR. Any engineered TCR having immune cell activation function can be induced using a method of the present disclosure. Such TCRs include, e.g., antigen-specific TCRs, Monoclonal TCRs (MTCRs), Single chain MTCRs, High Affinity CDR2 Mutant TCRs, CD1-binding MTCRs, High Affinity NY-ESO TCRs, VYG HLA-A24 Telomerase TCRs, including e.g., those described in PCT Pub Nos. WO 2003/020763, WO 2004/033685, WO 2004/044004, WO 2005/114215, WO 2006/000830, WO 2008/038002, WO 2008/039818, WO 2004/074322, WO 2005/113595, WO 2006/125962; Strommes et al. Immunol Rev. 2014; 257(1):145-64; Schmitt et al. Blood. 2013; 122(3):348-56; Chapuls et al. Sci Transl Med. 2013; 5(174):174ra27; Thaxton et al. Hum Vaccin Immunother. 2014; 10(11):3313-21 (PMID:25483644); Gschweng et al. Immunol Rev. 2014; 257(1):237-49 (PMID:24329801); Hinrichs et al. Immunol Rev. 2014; 257(1):56-71 (PMID:24329789); Zoete et al. Front Immunol. 2013; 4:268 (PMID:24062738); Marr et al. Clin Exp Immunol. 2012; 167(2):216-25 (PMID:22235997); Zhang et al. Adv Drug Deliv Rev. 2012; 64(8):756-62 (PMID:22178904); Chhabra et al. Scientific World Journal. 2011; 11:121-9 (PMID:21218269); Boulter et al. Clin Exp Immunol. 2005; 142(3):454-60 (PMID:16297157); Sami et al. Protein Eng Des Sel. 2007; 20(8):397-403; Boulter et al. Protein Eng. 2003; 16(9):707-11; Ashfield et al.

IDrugs. 2006; 9(8):554-9; Li et al. Nat Biotechnol. 2005; 23(3):349-54; Dunn et al. Protein Sci. 2006; 15(4):710-21; Liddy et al. Mol Biotechnol. 2010; 45(2); Liddy et al. Nat Med. 2012; 18(6):980-7; Oates, et al. Oncoimmunology. 2013; 2(2):e22891; McCormack, et al. Cancer Immunol Immunother. 2013 Apr;62(4):773-85; Bossi et al. Cancer Immunol Immunother. 2014; 63(5):437-48 and Oates, et al. Mol Immunol. 2015 Oct;67(2 Pt A):67-74; the disclosures of which are incorporated herein by reference in their entirety.

Antigen-binding Inhibitory Polypeptides

[0090] In some cases an antigen-triggered polypeptide produced in a genetically modified immune cell of the present disclosure, or present in a system of the present disclosure, or encoded by a nucleotide sequence in a nucleic acid present in a system of the present disclosure, may be an inhibitory polypeptide. The term "antigen-binding inhibitory polypeptide", as used herein, will generally describe a polypeptide, specific for an antigen, that upon binding the antigen inhibits the activity of a second polypeptide (e.g., an activating antigen-specific polypeptide, such as a CAR or TCR or other synthetic stimulatory immune cell receptor) and/or an activity of a cell (e.g., immune activation). iCARs, as described above, are an example of an antigen-binding inhibitory polypeptide; however, the term antigen-binding inhibitory polypeptide is not so limited.

[0091] Antigen-binding inhibitory polypeptides will vary and will generally function to mediate repression of an activated or activatable immune cell, including e.g., an immune cell expressing a stimulatory receptor, such as a CAR or TCR. An antigen-binding inhibitory polypeptide will include an inhibitory domain that functions to repress immune cell activation, including e.g., immune cell activation attributed to a stimulatory receptor, such as a CAR or TCR. Domains useful as inhibitory domains will vary depending on the particular context of immune cell activation and repression, including e.g., the particular type of activated cell to be repressed and the desired degree of repression. Exemplary non-limited examples of inhibitory domains include but are not limited to domains and motifs thereof derived from immune receptors including, e.g., co-inhibitory molecules, immune checkpoint molecules, immune tolerance molecules, and the like.

[0092] Suitable intracellular inhibitory domains may be any functional unit of a polypeptide as short as a 3 amino acid linear motif and as long as an entire protein, where size of the inhibitory domain is restricted only in that the domain must be sufficiently large as to retain its function and sufficiently small so as to be compatible with the other components of the polypeptide. Accordingly, an inhibitory domain may range in size from 3 amino acids in length to 1000 amino acids or more and, in some instances, can have a length of from about 30 amino acids to about 70 amino acids (aa), e.g., an inhibitory domain can have a length of from about 30 aa to about 35 aa, from about 35 aa to about 40 aa, from about 40 aa to about 45 aa, from about 45 aa to about 50 aa, from about 50 aa to about 55 aa, from about 55 aa to about 60 aa, from about 60 aa to about 65 aa, or from about 65 aa to about 70 aa. In other cases, an inhibitory

domain can have a length of from about 70 aa to about 100 aa, from about 100 aa to about 200 aa, or greater than 200 aa.

[0093] In some instances, "co-inhibitory domains" find use in the subject polypeptides. Such co-inhibitory domains are generally polypeptides derived from receptors. Co-inhibition generally refers to the secondary inhibition of primary antigen-specific activation mechanisms which prevents co-stimulation. Co-inhibition, e.g., T cell co-inhibition, and the factors involved have been described in Chen & Flies. *Nat Rev Immunol* (2013) 13(4):227-42 and Thaventhiran et al. *J Clin Cell Immunol* (2012) S12, the disclosures of which are incorporated herein by reference in their entirety. In some embodiments, co-inhibitory domains homodimerize. In some instances, useful co-inhibitory domains have been modified to constitutively dimerize, including constitutively homodimerize. A subject co-inhibitory domain can be an intracellular portion of a transmembrane protein (i.e., the co-inhibitory domain can be derived from a transmembrane protein). Non-limiting examples of suitable co-inhibitory polypeptides include, but are not limited to, CTLA-4 and PD-1. In some instances, a co-inhibitory domain, e.g., as used in a subject polypeptide may include a co-inhibitory domain selected from PD-1, CTLA4, HPK1, SHP1, SHP2, Sts1 and Csk. In some instances, a co- inhibitory domain of subject polypeptide comprises an amino acid sequence having at least about 75%, at least about 80%, at least about 85%, at least about 90%, at least about 95%, at least about 98%, or 100% amino acid sequence identity to a co-inhibitory domain as described herein.

[0094] In some instances, an antigen-binding inhibitory polypeptide may include a domain of a dimerization pair, such as e.g., a synthetic immune cell receptor (ICR) repressor useful as a component of a heteromeric, conditionally repressible synthetic ICR. Components of a heteromeric, conditionally repressible synthetic ICR may include a synthetic stimulatory ICR and a synthetic ICR repressor, where e.g., the synthetic stimulatory ICR and the synthetic ICR repressor specifically bind the antigens of an antigen pair described herein. Heteromeric, conditionally repressible synthetic ICRs, and components thereof, are described in PCT Application No. PCT/US2016/062612; the disclosure of which is incorporated herein by reference in its entirety.

Genetically Modified Immune Cells

[0095] The present disclosure provides a cytotoxic immune cell genetically modified to produce two antigen-triggered polypeptides, each recognizing a different cell surface antigen.

[0096] To generate a genetically modified cytotoxic immune cell of the present disclosure, a parent cytotoxic immune cell is genetically modified to produce: a) a first antigen-triggered polypeptide that binds specifically to a first target cell surface antigen present on a target cancer cell; and b) a second antigen-triggered polypeptide that binds specifically to a second target cell surface antigen. Suitable parent cytotoxic immune cells include $CD8^+$ T cells, natural killer (NK) cells, and the like. Thus, in some cases, a genetically modified cytotoxic immune cell of the present disclosure is a genetically modified

$CD8^+$ T cell. In other cases, a genetically modified cytotoxic immune cell of the present disclosure is a genetically modified NK cell.

[0097] In some cases, the target cancer cell is a liposarcoma, a glioblastoma, a breast cancer cell, a renal cancer cell, a pancreatic cancer cell, a melanoma, an anaplastic lymphoma, a leiomyosarcoma, an astrocytoma, an ovarian cancer cell, a neuroblastoma, a mantle cell lymphoma, a sarcoma, a non-small cell lung cancer cell, an AML cell, a stomach cancer cell, a B-cell cancer cell, a lung cancer cell, or an oligodendroglioma.

[0098] In some cases, a genetically modified cytotoxic immune cell of the present disclosure is genetically modified to express a first antigen-triggered polypeptide and a second antigen-triggered polypeptide that bind to antigens of a 2-input AND-gate target antigen pair. In some cases, a genetically modified cytotoxic immune cell of the present disclosure is genetically modified to express a first antigen-triggered polypeptide and a second antigen-triggered polypeptide that bind to antigens of a 3-input AND-gate target antigen pair. Non-limiting examples of 2-input AND gates (AND gates based on 2 target antigens) and 3-input AND gates (AND gates based on 3 target antigens) are depicted schematically in FIG. 3A-3B.

[0099] For example, in some cases, the first antigen-triggered polypeptide is a synNotch receptor and activation of the synNotch receptor by binding to the first antigen (present on a target cancer cell) induces expression of the second antigen-triggered polypeptide. The second antigen-triggered polypeptide binds to the second antigen of the target antigen pair, where the second antigen is expressed on the surface of the target cancer cell. As an example, in some cases, the first antigen-triggered polypeptide is a synNotch receptor and the second antigen-triggered polypeptide is a single chain CAR. As another example, in some cases, the first antigen-triggered polypeptide is a synNotch receptor and the second antigen-triggered polypeptide is a TCR. For example, in some cases, the synNotch polypeptide comprises an intracellular domain comprising a transcriptional activator, and activation of the synNotch receptor by binding to the first antigen (present on a target cancer cell) induces release of the transcriptional activator; the released transcriptional activator activates transcription of the TCR or the single-chain CAR.

[00100] As another example, in some cases, the first antigen-triggered polypeptide is a synNotch receptor and activation of the synNotch receptor by binding to the first antigen (present on a target cancer cell) induces expression of the second antigen-triggered polypeptide, where the second antigen-triggered polypeptide is a heterodimeric ("two chain" or "split") CAR comprising a first polypeptide chain and a second polypeptide chain. The heterodimeric CAR binds to the second antigen of the target antigen pair, where the second antigen is expressed on the surface of the target cancer cell. For example, in some cases, the first antigen-triggered polypeptide is a synNotch receptor and the second antigen-triggered polypeptide is a split CAR (e.g., an ON-switch CAR). In some cases, activation of the synNotch receptor

by binding to the first antigen (present on a target cancer cell) induces expression of only the first polypeptide chain of the heterodimeric CAR; expression of the second polypeptide chain of the heterodimeric CAR can be constitutive. For example, in some cases, the synNotch polypeptide comprises an intracellular domain comprising a transcriptional activator, and activation of the synNotch receptor by binding to the first antigen (present on a target cancer cell) induces release of the transcriptional activator; the released transcriptional activator activates transcription of the first polypeptide chain of the heterodimeric CAR. In some cases, activation of the synNotch receptor by binding to the first antigen (present on a target cancer cell) induces expression of only the second polypeptide chain of the heterodimeric CAR; expression of the first polypeptide chain of the heterodimeric CAR can be constitutive. Once the first polypeptide chain of the heterodimeric CAR is produced in the cell, it heterodimerizes with the second polypeptide chain of the heterodimeric CAR. As another example, in some cases, the synNotch polypeptide comprises an intracellular domain comprising a transcriptional activator, and activation of the synNotch receptor by binding to the first antigen (present on a target cancer cell) induces release of the transcriptional activator; the released transcriptional activator activates transcription of the second polypeptide chain of the heterodimeric CAR.

[00101] In some cases, a genetically modified cytotoxic immune cell of the present disclosure is genetically modified to express a first antigen-triggered polypeptide and a second antigen-triggered polypeptide that bind to antigens of a 2-input AND-NOT-gate target antigen pair. In some cases, a genetically modified cytotoxic immune cell of the present disclosure is genetically modified to express a first antigen-triggered polypeptide and a second antigen-triggered polypeptide that bind to antigens of a 3-input AND-NOT-gate target antigen pair. Non-limiting examples of 2-input AND-NOT gates (AND-NOT gates based on 2 target antigens) and 3-input AND-NOT gates (AND-NOT gates based on 3 target antigens) are depicted schematically in FIG. 3C-3D.

[00102] As an example, in some cases, the first antigen-triggered polypeptide is a CAR, and the second antigen-triggered polypeptide is an iCAR. Binding of the iCAR to the second antigen (present on the surface of a non-cancerous cell, but not on the surface of a target cancer cell) of a target antigen pair inhibits T-cell activation mediated by activation of the CAR upon binding to the first antigen (present on the surface of the target cancer cell and on the surface of the non-cancerous cell) of the target antigen pair. As another example, in some cases, the first antigen-triggered polypeptide is a TCR, and the second antigen-triggered polypeptide is an iCAR. Binding of the iCAR to the second antigen (present on the surface of a non-cancerous cell, but not on the surface of a target cancer cell) of a target antigen pair blocks or reduces T-cell activation mediated by activation of the TCR upon binding to the first antigen (present on the surface of the target cancer cell and on the surface of the non-cancerous cell) of the target antigen pair.

[00103] As another example, in some cases, the first antigen-triggered polypeptide is a CAR, and the second antigen-triggered polypeptide is a synNotch polypeptide comprising an intracellular domain that, when released upon activation of the synNotch polypeptide by binding to the second target antigen, induces expression of an intracellular inhibitor that inhibits T-cell activation mediated by activation of the CAR upon binding to the first antigen (present on the surface of the target cancer cell and on the surface of the non-cancerous cell) of the target antigen pair. As another example, in some cases, the first antigen-triggered polypeptide is a TCR, and the second antigen-triggered polypeptide is a synNotch polypeptide comprising an intracellular domain that, when released upon activation of the synNotch polypeptide by binding to the second target antigen, induces expression of an intracellular inhibitor that inhibits T-cell activation mediated by activation of the TCR upon binding to the first antigen (present on the surface of the target cancer cell and on the surface of the non-cancerous cell) of the target antigen pair.

[00104] As another example, in some cases, the first antigen-triggered polypeptide is a CAR, and the second antigen-triggered polypeptide is a synNotch polypeptide comprising an intracellular domain that, when released upon activation of the synNotch polypeptide by binding to the second target antigen, induces expression of an extracellular inhibitor that inhibits T-cell activation mediated by activation of the CAR upon binding to the first antigen (present on the surface of the target cancer cell and on the surface of the non-cancerous cell) of the target antigen pair. As another example, in some cases, the first antigen-triggered polypeptide is a TCR, and the second antigen-triggered polypeptide is a synNotch polypeptide comprising an intracellular domain that, when released upon activation of the synNotch polypeptide by binding to the second target antigen, induces expression of an extracellular inhibitor that inhibits T-cell activation mediated by activation of the TCR upon binding to the first antigen (present on the surface of the target cancer cell and on the surface of the non-cancerous cell) of the target antigen pair.

[00105] In some cases, a genetically modified immune cell of the present disclosure provides for a 3-input AND, AND-NOT gate, comprising: 1) a first target antigen of an AND-NOT gate target antigen pair depicted in FIG. 1 or FIG. 9-14; 2) a second target antigen of the AND-NOT gate target antigen pair; and 3) a third target antigen, where the third target antigen is a target antigen depicted in FIG. 4 or Table 3, where the third target antigen is a target antigen for the same cancer cell type as the target antigen pair. In some cases, the 3-input AND, AND-NOT gate comprises: 1) the first target antigen of an AND-NOT gate target antigen pair depicted in FIG. 1 or FIG. 9-14; AND-NOT 2) the second target antigen of the AND-NOT gate target antigen pair; AND 3) the third target antigen. In some cases, the 3-input AND, AND-NOT gate comprises: 1) the second target antigen of an AND-NOT gate target antigen pair depicted in FIG. 1 or FIG. 9-14; AND-NOT 2) the first target antigen of the AND-NOT gate target antigen pair; AND 3) the third target antigen.

[00106] As one non-limiting example, where the target AND-NOT gate antigen pair is NLGN1 AND-NOT TNFRSF17 (for a glioma cell) (as shown in FIG. 1), a suitable third target antigen is EphA2 (as shown in FIG. 4).

Systems for Inhibiting Cancer Cells

[00107] The present disclosure provides a system for inhibiting or killing a target cancer cell. A system of the present disclosure comprises: a) a first antigen-triggered polypeptide that binds specifically to a first target antigen present on the target cancer cell, or a first nucleic acid comprising a nucleotide sequence encoding the first antigen-triggered polypeptide; and b) a second antigen-triggered polypeptide that binds specifically to a second target antigen, or a second nucleic acid comprising a nucleotide sequence encoding the second antigen-triggered polypeptide.

[00108] In some cases, the target cancer cell is a liposarcoma, a glioblastoma, a breast cancer cell, a renal cancer cell, a pancreatic cancer cell, a melanoma, an anaplastic lymphoma, a leiomyosarcoma, an astrocytoma, an ovarian cancer cell, a neuroblastoma, a mantle cell lymphoma, a sarcoma, a non-small cell lung cancer cell, an AML cell, a stomach cancer cell, a B-cell cancer cell, a lung cancer cell, or an oligodendroglioma.

[00109] In some cases, as noted above, a system of the present disclosure comprises: a) a first antigen-triggered polypeptide that binds specifically to a first target antigen present on a target cancer cell; and b) a second antigen-triggered polypeptide that binds specifically to a second target antigen. In these instances, the polypeptides per se are introduced into an immune cell (e.g., $CD8^+$ T cells and/or NK cells obtained from an individual). Methods of introducing polypeptides into a cell are known in the art; and any known method can be used. For example, in some cases, the first and the second antigen-triggered polypeptides comprise a protein transduction domain (PTD) at the N-terminus or the C-terminus of the polypeptides.

[00110] In some cases, as noted above, a system of the present disclosure comprises: a) a first nucleic acid comprising a nucleotide sequence encoding a first antigen-triggered polypeptide that binds specifically to a first target antigen present on a target cancer cell; and b) a second nucleic acid comprising a nucleotide sequence encoding a second antigen-triggered polypeptide that binds specifically to a second target antigen. In some cases, the first and the second antigen-triggered polypeptides are encoded by nucleotide sequences on separate nucleic acids. In other cases, the first and the second antigen-triggered polypeptides are encoded by nucleotide sequences present in the same nucleic acid. In some cases, the nucleic acid is a recombinant expression vector. In some cases, a system of the present disclosure comprises: a) a first recombinant expression vector comprising a nucleotide sequence encoding a first antigen-triggered polypeptide that binds specifically to a first target antigen present on a target cancer cell; and b) a second recombinant expression vector comprising a nucleotide sequence encoding a second antigen-triggered polypeptide that binds specifically to a second target antigen. In some cases, the

nucleotide sequences are operably linked to a constitutive promoter. In some cases, the nucleotide sequences are operably linked to a regulatable promoter (e.g., an inducible promoter, a reversible promoter, etc.). In some cases, the nucleotide sequences are operably linked to an immune cell promoter, e.g., a T-cell specific promoter. In some cases, a system of the present disclosure comprises a recombinant expression vector comprising nucleotide sequences encoding: a) a first antigen-triggered polypeptide that binds specifically to a first target antigen present on a target cancer cell; and b) a second antigen-triggered polypeptide that binds specifically to a second target antigen. In some cases, the nucleotide sequences are operably linked to a constitutive promoter. In some cases, the nucleotide sequences are operably linked to a regulatable promoter (e.g., an inducible promoter, a reversible promoter, etc.). In some cases, the nucleotide sequences are operably linked to an immune cell promoter, e.g., a T-cell specific promoter.

[00111] Suitable promoters include, but are not limited to; cytomegalovirus immediate early promoter; herpes simplex virus thymidine kinase promoter; early and late SV40 promoters; promoter present in long terminal repeats from a retrovirus; a metallothionein-I promoter; and various art-known promoters. Such reversible promoters, and systems based on such reversible promoters but also comprising additional control proteins, include, but are not limited to, alcohol regulated promoters (e.g., alcohol dehydrogenase I (alcA) gene promoter, promoters responsive to alcohol transactivator proteins (AlcR), etc.), tetracycline regulated promoters, (e.g., promoter systems including TetActivators, TetON, TetOFF, etc.), steroid regulated promoters (e.g., rat glucocorticoid receptor promoter systems, human estrogen receptor promoter systems, retinoid promoter systems, thyroid promoter systems, ecdysone promoter systems, mifepristone promoter systems, etc.), metal regulated promoters (e.g., metallothionein promoter systems, etc.), pathogenesis-related regulated promoters (e.g., salicylic acid regulated promoters, ethylene regulated promoters, benzothiadiazole regulated promoters, etc.), temperature regulated promoters (e.g., heat shock inducible promoters (e.g., HSP-70, HSP-90, soybean heat shock promoter, etc.), light regulated promoters, synthetic inducible promoters, and the like.

[00112] In some instances, nucleic acids present in a system of the present disclosure include immune cell specific promoters that are expressed in one or more immune cell types, including but not limited to lymphocytes, hematopoietic stem cells and/or progeny thereof (i.e., immune cell progenitors), etc. Any convenient and appropriate promoter of an immune cell specific gene may find use in nucleic acids of the present disclosure. In some instances, an immune cell specific promoter of a nucleic acid present in a system of the present disclosure may be a T cell specific promoter. In some instances, an immune cell specific promoter of a nucleic acid present in a system of the present disclosure may be a light and/or heavy chain immunoglobulin gene promoter and may or may not include one or more related enhancer elements.

[00113] In some instances, an immune cell specific promoter of a nucleic acid present in a system of the present disclosure may be a promoter of a B29 gene promoter, a CD14 gene promoter, a CD43 gene promoter, a CD45 gene promoter, a CD68 gene promoter, a IFN-β gene promoter, a WASP gene promoter, a T-cell receptor β -chain gene promoter, a V9 γ (TRGV9) gene promoter, a V2 δ (TRDV2) gene promoter, and the like.

[00114] In some instances, an immune cell specific promoter present in a system of a nucleic acid of the present disclosure may be a viral promoter expressed in immune cells. As such, in some instances, viral promoters useful in nucleic acids present in a system of the present disclosure include viral promoters derived from immune cells viruses, including but not limited to, e.g., lentivirus promoters (e.g., HIV, SIV, FIV, EIAV, or Visna promoters) including e.g., LTR promoter, etc., Retroviridae promoters including, e.g., HTLV-I promoter, HTLV-II promoter, etc., and the like.

[00115] In some cases, the promoter is a CD8 cell-specific promoter, a CD4 cell-specific promoter, a neutrophil-specific promoter, or an NK-specific promoter. For example, a CD4 gene promoter can be used; see, e.g., Salmon et al. (1993) *Proc. Natl. Acad. Sci. USA* 90:7739; and Marodon et al. (2003) *Blood* 101:3416. As another example, a CD8 gene promoter can be used. NK cell-specific expression can be achieved by use of an *Ncr1* (*p46*) promoter; see, e.g., Eckelhart et al. (2011) *Blood* 117:1565.

[00116] Expression vectors generally have convenient restriction sites located near the promoter sequence to provide for the insertion of nucleic acid sequences encoding heterologous proteins. A selectable marker operative in the expression host may be present. Suitable recombinant expression vectors include, but are not limited to, viral vectors (e.g. viral vectors based on vaccinia virus; poliovirus; adenovirus (see, e.g., Li et al., Invest Opthalmol Vis Sci 35:2543 2549, 1994; Borras et al., Gene Ther 6:515 524, 1999; Li and Davidson, PNAS 92:7700 7704, 1995; Sakamoto et al., H Gene Ther 5:1088 1097, 1999; WO 94/12649, WO 93/03769; WO 93/19191; WO 94/28938; WO 95/11984 and WO 95/00655); adeno-associated virus (see, e.g., Ali et al., Hum Gene Ther 9:81 86, 1998, Flannery et al., PNAS 94:6916 6921, 1997; Bennett et al., Invest Opthalmol Vis Sci 38:2857 2863, 1997; Jomary et al., Gene Ther 4:683 690, 1997, Rolling et al., Hum Gene Ther 10:641 648, 1999; Ali et al., Hum Mol Genet 5:591 594, 1996; Srivastava in WO 93/09239, Samulski et al., J. Vir. (1989) 63:3822-3828; Mendelson et al., Virol. (1988) 166:154-165; and Flotte et al., PNAS (1993) 90:10613-10617); SV40; herpes simplex virus; human immunodeficiency virus (see, e.g., Miyoshi et al., PNAS 94:10319 23, 1997; Takahashi et al., J Virol 73:7812 7816, 1999); a retroviral vector (e.g., Murine Leukemia Virus, spleen necrosis virus, and vectors derived from retroviruses such as Rous Sarcoma Virus, Harvey Sarcoma Virus, avian leukosis virus, human immunodeficiency virus, myeloproliferative sarcoma virus, and mammary tumor virus); and the like.

Antigen combinations

[00117] A system of the present disclosure targets antigen combinations, where the targeting provides for killing of a target cancer cell. A genetically modified immune cell of the present disclosure targets antigen combinations, where the targeting provides for killing of a target cancer cell. Antigen combinations may also reduce off-target effects and/or increase specificity for a target cancer cell, where e.g., an antigen combination includes one or more AND NOT combinations. Examples of target antigen combinations, and corresponding exemplary but non-limiting target cancer cells, are depicted in FIG. 1 and FIG. 9-14. The following antigen combinations are exemplary, and not meant to be limiting.

Liposarcoma antigen target combinations

[00118] In some cases, an antigen combination for targeting a liposarcoma cell comprises EVA1B AND NOT ITGA6. In some cases, the ITGA6 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 62 or 998. In some cases, the EVA1B polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 4400.

[00119] In some cases, an antigen combination for targeting a liposarcoma cell comprises ADAM12 AND NOT TACSTD2. In some cases, the TACSTD2 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 2631. In some cases, the ADAM12 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 2839 or 4815.

Glioblastoma antigen target combinations

[00120] In some cases, an antigen combination for targeting a glioblastoma cell comprises PTPRZ1 AND FOLR2. In some cases, the FOLR2 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 261, 1261, 1262 or 1263. In some cases, the PTPRZ1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 2726.

[00121] In some cases, an antigen combination for targeting a glioblastoma cell comprises PTPRZ1 AND IGF1R. In some cases, the IGF1R polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 308. In some cases, the PTPRZ1 polypeptide comprises an amino

acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 2726.

[00122] In some cases, an antigen combination for targeting a glioblastoma cell comprises PTPRZ1 AND NCAM1. In some cases, the NCAM1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 173, 949 or 6595. In some cases, the PTPRZ1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 2726.

[00123] In some cases, an antigen combination for targeting a glioblastoma cell comprises PTPRZ1 AND CD70. In some cases, the CD70 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 2331. In some cases, the PTPRZ1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 2726.

[00124] In some cases, an antigen combination for targeting a glioblastoma cell comprises PTPRZ1 AND AXL. In some cases, the AXL polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 2438 or 4847. In some cases, the PTPRZ1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 2726.

[00125] In some cases, an antigen combination for targeting a glioblastoma cell comprises PTPRZ1 AND NOT NAALAD2. In some cases, the NAALAD2 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3289. In some cases, the PTPRZ1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 2726.

[00126] In some cases, an antigen combination for targeting a glioblastoma cell comprises PTPRZ1 AND NOT TMEM170B. In some cases, the TMEM170B polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 1195. In some cases, the PTPRZ1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 2726.

[00127] In some cases, an antigen combination for targeting a glioblastoma cell comprises PTPRZ1 AND NOT PIGK. In some cases, the PIGK polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3293. In some cases, the PTPRZ1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 2726.

[00128] In some cases, an antigen combination for targeting a glioblastoma cell comprises PTPRZ1 AND CDH5. In some cases, the CDH5 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 2475. In some cases, the PTPRZ1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 2726.

[00129] In some cases, an antigen combination for targeting a glioblastoma cell comprises PTPRZ1 AND NOT 100507547?. In some cases, the 100507547? polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 7099. In some cases, the PTPRZ1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 2726.

[00130] Additional examples of target antigen pairs for targeting a glioblastoma cell are provided in FIG. 1 and FIG. 9-14.

Breast adenocarcinoma target antigen combinations

[00131] In some cases, an antigen combination for targeting a breast adenocarcinoma cell comprises CYB561 AND NOT SLC2A12. In some cases, the SLC2A12 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 5933. In some cases, the CYB561 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 646, 647 or 2500.

[00132] In some cases, an antigen combination for targeting a breast adenocarcinoma cell comprises RHBDF1 AND NOT TMEM256. In some cases, the TMEM256 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 6112. In some cases, the RHBDF1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least

98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 4919.

[00133] In some cases, an antigen combination for targeting a breast adenocarcinoma cell comprises HSD3B7 AND NOT TMEM256. In some cases, the TMEM256 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 6112. In some cases, the HSD3B7 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 1560, 1699, 1700, 4387 or 5132.

[00134] In some cases, an antigen combination for targeting a breast adenocarcinoma cell comprises ADAM12 AND NOT TMEM256. In some cases, the TMEM256 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 6112. In some cases, the ADAM12 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 2839 or 4815.

[00135] In some cases, an antigen combination for targeting a breast adenocarcinoma cell comprises SLC5A6 AND NOT TMEM256. In some cases, the TMEM256 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 6112. In some cases, the SLC5A6 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 4767.

[00136] In some cases, an antigen combination for targeting a breast adenocarcinoma cell comprises GGT5 AND NOT TSPAN8. In some cases, the TSPAN8 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3085. In some cases, the GGT5 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 1173, 1174 or 2968.

[00137] In some cases, an antigen combination for targeting a breast adenocarcinoma cell comprises ADAM12 AND NOT ADGRD1. In some cases, the ADGRD1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 6838. In some cases, the ADAM12 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least

95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 2839 or 4815.

[00138] In some cases, an antigen combination for targeting a breast adenocarcinoma cell comprises LTBP3 AND NOT HBD. In some cases, the HBD polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 148. In some cases, the LTBP3 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 1421, 2072 or 4761.

[00139] In some cases, an antigen combination for targeting a breast adenocarcinoma cell comprises MARVELD3 AND NOT TMEM220. In some cases, the TMEM220 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 429. In some cases, the MARVELD3 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 653 or 5518.

[00140] In some cases, an antigen combination for targeting a breast adenocarcinoma cell comprises RHBDF1 AND NOT SMIM5. In some cases, the SMIM5 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 2039. In some cases, the RHBDF1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 4919.

[00141] Additional examples of target antigen pairs for targeting a breast adenocarcinoma cell are provided in FIG. 1 and FIG. 9-14.

Glioma target antigen combinations

[00142] In some cases, an antigen combination for targeting a glioma cell comprises NLGN1 AND NOT TNFRSF17. In some cases, the TNFRSF17 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 2312. In some cases, the NLGN1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 4016.

[00143] In some cases, an antigen combination for targeting a glioma cell comprises ITGAV AND NOT TNFRSF17. In some cases, the TNFRSF17 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity

to the amino acid sequence set forth in SEQ ID NO: 2312. In some cases, the ITGAV polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 1802, 1803 or 2579.

[00144] In some cases, an antigen combination for targeting a glioma cell comprises SYT11 AND TNFRSF17. In some cases, the TNFRSF17 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 2312. In some cases, the SYT11 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 6023.

[00145] In some cases, an antigen combination for targeting a glioma cell comprises SCRG1 AND TNFRSF17. In some cases, the TNFRSF17 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 2312. In some cases, the SCRG1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3667.

[00146] In some cases, an antigen combination for targeting a glioma cell comprises AGPAT5 AND NOT TNFRSF17. In some cases, the TNFRSF17 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 2312. In some cases, the AGPAT5 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 4429.

[00147] In some cases, an antigen combination for targeting a glioma cell comprises PTPRZ1 AND NOT TNFRSF17. In some cases, the TNFRSF17 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 2312. In some cases, the PTPRZ1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 2726.

[00148] In some cases, an antigen combination for targeting a glioma cell comprises NRCAM AND NOT FOLH1. In some cases, the FOLH1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 631 or 3057. In some cases, the NRCAM polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 768, 769 or 3195.

[00149] In some cases, an antigen combination for targeting a glioma cell comprises PRAF2 AND NOT FOLH1. In some cases, the FOLH1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 631 or 3057. In some cases, the PRAF2 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3649.

[00150] In some cases, an antigen combination for targeting a glioma cell comprises XPR1 AND FOLH1. In some cases, the FOLH1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 631 or 3057. In some cases, the XPR1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 1546 or 3112.

[00151] In some cases, an antigen combination for targeting a glioma cell comprises SYT11 AND NOT FOLH1. In some cases, the FOLH1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 631 or 3057. In some cases, the SYT11 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 6023.

[00152] Additional examples of target antigen pairs for targeting a glioma cell are provided in FIG. 1 and FIG. 9-14.

Breast carcinoma (e.g., ductal breast carcinoma) target antigen combinations

[00153] In some cases, an antigen combination for targeting a breast carcinoma cell comprises CACFD1 AND NOT TMED6. In some cases, the TMED6 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 5889. In some cases, the CACFD1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 1566 or 4310.

[00154] In some cases, an antigen combination for targeting a breast carcinoma cell comprises CACFD1 AND NOT TMEM61. In some cases, the TMEM61 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 6652. In some cases, the CACFD1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 1566 or 4310.

[00155] In some cases, an antigen combination for targeting a breast carcinoma cell comprises CACFD1 AND NOT CDHR2. In some cases, the CDHR2 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 2306 or 4321. In some cases, the CACFD1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 1566 or 4310.

[00156] In some cases, an antigen combination for targeting a breast carcinoma cell comprises CACFD1 AND NOT TMEM246. In some cases, the TMEM246 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 5332. In some cases, the CACFD1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 1566 or 4310.

[00157] In some cases, an antigen combination for targeting a breast carcinoma cell comprises CACFD1 AND NOT SLC4A4. In some cases, the SLC4A4 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 1125, 1504 or 2885. In some cases, the CACFD1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 1566 or 4310.

[00158] In some cases, an antigen combination for targeting a breast carcinoma cell comprises C10orf35 AND NOT SLC4A4. In some cases, the SLC4A4 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 1125, 1504 or 2885. In some cases, the C10orf35 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 5959.

[00159] In some cases, an antigen combination for targeting a breast carcinoma cell comprises RNF121 AND NOT PTCH1. In some cases, the PTCH1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 77, 1071, 1072, 1073, 1074, 1075 or 1076. In some cases, the RNF121 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 4420.

Renal carcinoma target antigen combinations

[00160] In some cases, an antigen combination for targeting a renal carcinoma cell comprises CAV2 AND NOT SCARA5. In some cases, the SCARA5 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 6392. In some cases, the CAV2 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 2320 or 6759.

Pancreas carcinoma (e.g., pancreatic ductal carcinoma) target antigen combinations

[00161] In some cases, an antigen combination for targeting a pancreas carcinoma cell comprises CTRB2 AND NOT CDH18. In some cases, the CDH18 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 2192 or 3172. In some cases, the CTRB2 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 701.

[00162] In some cases, an antigen combination for targeting a pancreas carcinoma cell comprises TM4SF4 AND NOT ABCC2. In some cases, the ABCC2 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 117. In some cases, the TM4SF4 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3086.

[00163] In some cases, an antigen combination for targeting a pancreas carcinoma cell comprises TM4SF4 AND NOT TMEM56. In some cases, the TMEM56 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 6068. In some cases, the TM4SF4 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3086.

[00164] In some cases, an antigen combination for targeting a pancreas carcinoma cell comprises TM4SF4 AND NOT CYP3A7. In some cases, the CYP3A7 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 239. In some cases, the TM4SF4 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least

98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3086.

[00165] In some cases, an antigen combination for targeting a pancreas carcinoma cell comprises TM4SF4 AND NOT CYP2J2. In some cases, the CYP2J2 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 243. In some cases, the TM4SF4 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3086.

[00166] In some cases, an antigen combination for targeting a pancreas carcinoma cell comprises TM4SF4 AND NOT CLYBL. In some cases, the CLYBL polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 6965. In some cases, the TM4SF4 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3086.

[00167] In some cases, an antigen combination for targeting a pancreas carcinoma cell comprises TM4SF4 AND NOT SLC35D1. In some cases, the SLC35D1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 4040. In some cases, the TM4SF4 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3086.

[00168] In some cases, an antigen combination for targeting a pancreas carcinoma cell comprises TM4SF4 AND NOT G6PC. In some cases, the G6PC polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 45. In some cases, the TM4SF4 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3086.

[00169] In some cases, an antigen combination for targeting a pancreas carcinoma cell comprises TM4SF4 AND GJA1. In some cases, the GJA1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 51. In some cases, the TM4SF4 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3086.

[00170] In some cases, an antigen combination for targeting a pancreas carcinoma cell comprises CTRB2 AND NPTN. In some cases, the NPTN polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 2004, 2005, 3774 or 4295. In some cases, the CTRB2 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 701.

[00171] Additional examples of target antigen pairs for targeting a pancreas carcinoma cell are provided in FIG. 1 and FIG. 9-14.

Melanoma target antigen combinations

[00172] In some cases, an antigen combination for targeting a melanoma cell comprises DBI AND NOT FCN2. In some cases, the FCN2 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 2966 or 4122. In some cases, the DBI polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 1006, 1007 and 4684.

[00173] In some cases, an antigen combination for targeting a melanoma cell comprises KIAA1549L AND DBI. In some cases, the KIAA1549L polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3718. In some cases, the DBI polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 1006, 1007 and 4684.

[00174] In some cases, an antigen combination for targeting a melanoma cell comprises TMED3 AND DBI. In some cases, the TMED3 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3691. In some cases, the DBI polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 1006, 1007 and 4684.

[00175] In some cases, an antigen combination for targeting a melanoma cell comprises SFXN3 AND NOT DBI. In some cases, the SFXN3 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 5209. In some cases, the DBI polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 1006, 1007 and 4684.

[00176] In some cases, an antigen combination for targeting a melanoma cell comprises GPR19 AND DBI. In some cases, the GPR19 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3446. In some cases, the DBI polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 1006, 1007 and 4684.

[00177] In some cases, an antigen combination for targeting a melanoma cell comprises DBI AND NOT ITGA2B. In some cases, the ITGA2B polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 123. In some cases, the DBI polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 1006, 1007 and 4684.

[00178] In some cases, an antigen combination for targeting a melanoma cell comprises DBI AND NOT VSTM1. In some cases, the VSTM1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 6800. In some cases, the DBI polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 1006, 1007 and 4684.

[00179] In some cases, an antigen combination for targeting a melanoma cell comprises DBI AND BNIP1. In some cases, the BNIP1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 2317, 3842, 3843 or 3844. In some cases, the DBI polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 1006, 1007 and 4684.

[00180] In some cases, an antigen combination for targeting a melanoma cell comprises DBI AND NOT CD300LB. In some cases, the CD300LB polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 6404. In some cases, the DBI polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 1006, 1007 and 4684.

[00181] In some cases, an antigen combination for targeting a melanoma cell comprises MANBAL AND DBI. In some cases, the MANBAL polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 425 or 4873. In some cases, the DBI polypeptide

comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 1006, 1007 and 4684.

[00182] Additional examples of target antigen pairs for targeting a melanoma cell are provided in FIG. 1 and FIG. 9-14.

Anaplastic lymphoma (e.g., anaplastic large-cell lymphoma) target antigen combinations

[00183] In some cases, an antigen combination for targeting an anaplastic lymphoma cell comprises TNFRSF8 AND NOT GPC3. In some cases, the GPC3 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 2089, 2090, 2091 or 3062. In some cases, the TNFRSF8 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 2323 or 6140.

[00184] In some cases, an antigen combination for targeting an anaplastic lymphoma cell comprises TNFRSF8 AND NOT L1CAM. In some cases, the L1CAM polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 126, 1743 or 4990. In some cases, the TNFRSF8 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 2323 or 6140.

[00185] In some cases, an antigen combination for targeting an anaplastic lymphoma cell comprises TNFRSF8 AND NOT SDC1. In some cases, the SDC1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 496 or 2747. In some cases, the TNFRSF8 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 2323 or 6140.

[00186] In some cases, an antigen combination for targeting an anaplastic lymphoma cell comprises TNFRSF8 AND NOT TNFRSF10A. In some cases, the TNFRSF10A polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 2911. In some cases, the TNFRSF8 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 2323 or 6140.

[00187] In some cases, an antigen combination for targeting an anaplastic lymphoma cell comprises TNFRSF8 AND NOT ERBB2. In some cases, the ERBB2 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 474 or 3051. In some cases, the TNFRSF8 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 2323 or 6140.

[00188] In some cases, an antigen combination for targeting an anaplastic lymphoma cell comprises TNFRSF8 AND NOT IL11RA. In some cases, the IL11RA polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 1701, 3066 or 5993. In some cases, the TNFRSF8 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 2323 or 6140.

[00189] In some cases, an antigen combination for targeting an anaplastic lymphoma cell comprises TNFRSF8 AND NOT SYT15. In some cases, the SYT15 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 5268 or 6605. In some cases, the TNFRSF8 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 2323 or 6140.

[00190] In some cases, an antigen combination for targeting an anaplastic lymphoma cell comprises TNFRSF8 AND NOT CYP4F2. In some cases, the CYP4F2 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 988. In some cases, the TNFRSF8 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 2323 or 6140.

[00191] In some cases, an antigen combination for targeting an anaplastic lymphoma cell comprises TNFRSF8 AND NOT KCNK13. In some cases, the KCNK13 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 4869. In some cases, the TNFRSF8 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 2323 or 6140.

[00192] In some cases, an antigen combination for targeting an anaplastic lymphoma cell comprises TNFRSF8 AND NOT NTN1. In some cases, the NTN1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3138. In some cases, the TNFRSF8 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 2323 or 6140.

[00193] Additional examples of target antigen pairs for targeting an anaplastic lymphoma cell are provided in FIG. 1 and FIG. 9-14.

Leiomyosarcoma target antigen combinations

[00194] In some cases, an antigen combination for targeting a leiomyosarcoma cell comprises VANGL1 AND NOT SGPP2. In some cases, the SGPP2 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 6046. In some cases, the VANGL1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 5808.

[00195] In some cases, an antigen combination for targeting a leiomyosarcoma cell comprises VANGL1 AND NOT KRTCAP3. In some cases, the KRTCAP3 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 2225 or 6396. In some cases, the VANGL1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 5808.

[00196] In some cases, an antigen combination for targeting a leiomyosarcoma cell comprises NEMP1 AND NOT SMIM5. In some cases, the SMIM5 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 2039. In some cases, the NEMP1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 1451 or 4053.

[00197] In some cases, an antigen combination for targeting a leiomyosarcoma cell comprises ATP2A2 AND NOT CRB3. In some cases, the CRB3 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 5838 or 6403. In some cases, the ATP2A2 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least

98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 1561, 2432 or 6245.

[00198] In some cases, an antigen combination for targeting a leiomyosarcoma cell comprises VANGL1 AND NOT F11R. In some cases, the F11R polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 4278. In some cases, the VANGL1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 5808.

[00199] In some cases, an antigen combination for targeting a leiomyosarcoma cell comprises VANGL1 AND NOT RNF43. In some cases, the RNF43 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 4337. In some cases, the VANGL1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 5808.

[00200] In some cases, an antigen combination for targeting a leiomyosarcoma cell comprises VANGL1 AND NOT SCNN1A. In some cases, the SCNN1A polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 756, 1946 or 1947. In some cases, the VANGL1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 5808.

[00201] In some cases, an antigen combination for targeting a leiomyosarcoma cell comprises VANGL1 AND NOT SMIM5. In some cases, the SMIM5 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 2039. In some cases, the VANGL1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 5808.

[00202] In some cases, an antigen combination for targeting a leiomyosarcoma cell comprises VANGL1 AND NOT ERMP1. In some cases, the ERMP1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 5098. In some cases, the VANGL1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 5808.

[00203] In some cases, an antigen combination for targeting a leiomyosarcoma cell comprises VANGL1 AND NOT CRB3. In some cases, the CRB3 polypeptide comprises an amino acid sequence having at

least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 5838 or 6403. In some cases, the VANGL1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 5808.

[00204] Additional examples of target antigen pairs for targeting a leiomyocarcoma cell are provided in FIG. 1 and FIG. 9-14.

Astrocytoma target antigen combinations

[00205] In some cases, an antigen combination for targeting an astrocytoma cell comprises PTPRZ1 AND CLCN5. In some cases, the CLCN5 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 22, 1361 or 1362. In some cases, the PTPRZ1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 2726.

[00206] In some cases, an antigen combination for targeting an astrocytoma cell comprises PTPRZ1 AND ADGRE1. In some cases, the ADGRE1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 2514. In some cases, the PTPRZ1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 2726.

[00207] In some cases, an antigen combination for targeting an astrocytoma cell comprises PTPRZ1 AND F2R. In some cases, the F2R polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 2519. In some cases, the PTPRZ1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 2726.

[00208] In some cases, an antigen combination for targeting an astrocytoma cell comprises PTPRZ1 AND CYB561A3. In some cases, the CYB561A3 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 646, 647 or 2500. In some cases, the PTPRZ1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 2726.

[00209] In some cases, an antigen combination for targeting an astrocytoma cell comprises PTPRZ1 AND IL27. In some cases, the IL27 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 5969. In some cases, the PTPRZ1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 2726.

[00210] In some cases, an antigen combination for targeting an astrocytoma cell comprises PTPRZ1 AND NOT GHITM. In some cases, the GHITM polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3930. In some cases, the PTPRZ1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 2726.

[00211] In some cases, an antigen combination for targeting an astrocytoma cell comprises PTPRZ1 AND NOT PRRT3. In some cases, the PRRT3 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 7014. In some cases, the PTPRZ1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 2726.

[00212] In some cases, an antigen combination for targeting an astrocytoma cell comprises PTPRZ1 AND HAS2. In some cases, the HAS2 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3273. In some cases, the PTPRZ1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 2726.

[00213] In some cases, an antigen combination for targeting an astrocytoma cell comprises PTPRZ1 AND LCAT. In some cases, the LCAT polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 71. In some cases, the PTPRZ1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 2726.

[00214] In some cases, an antigen combination for targeting an astrocytoma cell comprises PTPRZ1 AND NOT EPCAM. In some cases, the EPCAM polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 2632. In some cases, the PTPRZ1 polypeptide

comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 2726.

[00215] Additional examples of target antigen pairs for targeting an astrocytoma are provided in FIG. 1 and FIG. 9-14.

Ovarian cancer (e.g., serous cystadenocarcinoma; papillary adenocarcinoma) target antigen combinations

[00216] In some cases, an antigen combination for targeting an ovarian cancer cell (e.g., a cystadenocarcinoma) comprises SVOPL AND NOT ARL6IP5. In some cases, the ARL6IP5 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3483. In some cases, the SVOPL polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 1618 or 6423.

[00217] In some cases, an antigen combination for targeting an ovarian cancer cell comprises FAAH2 AND NOT ARL6IP5. In some cases, the ARL6IP5 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3483. In some cases, the FAAH2 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 6408.

[00218] In some cases, an antigen combination for targeting an ovarian cancer cell comprises GAL3ST4 AND NOT ARL6IP5. In some cases, the ARL6IP5 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3483. In some cases, the GAL3ST4 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 5061.

[00219] In some cases, an antigen combination for targeting an ovarian cancer cell comprises SYNE4 AND NOT FAXDC2. In some cases, the FAXDC2 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 4206 or 5337. In some cases, the SYNE4 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 814.

[00220] In some cases, an antigen combination for targeting an ovarian cancer cell comprises SYNE4 AND NOT OS9. In some cases, the OS9 polypeptide comprises an amino acid sequence having at least 85%,

at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 650, 651, 652 or 3569. In some cases, the SYNE4 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 814.

[00221] In some cases, an antigen combination for targeting an ovarian cancer cell comprises SVOPL AND NOT ANTXR2. In some cases, the ANTXR2 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 1876 or 5582. In some cases, the SVOPL polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 1618 or 6423.

[00222] In some cases, an antigen combination for targeting an ovarian cancer cell comprises SVOPL AND NOT COMT. In some cases, the COMT polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 232, 1531, 1532 or 3683. In some cases, the SVOPL polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 1618 or 6423.

[00223] In some cases, an antigen combination for targeting an ovarian cancer cell comprises SVOPL AND NOT SLC31A2. In some cases, the SLC31A2 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 2490. In some cases, the SVOPL polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 1618 or 6423.

[00224] In some cases, an antigen combination for targeting an ovarian cancer cell comprises SVOPL AND NOT MFSD4. In some cases, the MFSD4 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 6617. In some cases, the SVOPL polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 1618 or 6423.

[00225] In some cases, an antigen combination for targeting an ovarian cancer cell comprises SVOPL AND NOT FMO5. In some cases, the FMO5 polypeptide comprises an amino acid sequence having at

least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 1775, 1776 or 2379. In some cases, the SVOPL polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 1618 or 6423.

[00226] Additional examples of target antigen pairs for targeting an ovarian cancer cell (e.g., a cystadenocarcinoma) are provided in FIG. 1 and FIG. 9-14.

Neuroblastoma target antigen combinations

[00227] In some cases, an antigen combination for targeting a neuroblastoma cell comprises MARCH11 AND SDC1. In some cases, the SDC1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 496 or 2747. In some cases, the MARCH11 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 1209.

[00228] In some cases, an antigen combination for targeting a neuroblastoma cell comprises SLC10A4 AND NOT SDC1. In some cases, the SDC1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 496 or 2747. In some cases, the SLC10A4 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 6095.

[00229] In some cases, an antigen combination for targeting a neuroblastoma cell comprises ST8SIA2 AND NOT SDC1. In some cases, the SDC1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 496 or 2747. In some cases, the ST8SIA2 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3413.

[00230] In some cases, an antigen combination for targeting a neuroblastoma cell comprises VANGL2 AND NOT SDC1. In some cases, the SDC1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 496 or 2747. In some cases, the VANGL2 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 4631.

[00231] In some cases, an antigen combination for targeting a neuroblastoma cell comprises DIABLO AND NOT SDC1. In some cases, the SDC1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 496 or 2747. In some cases, the DIABLO polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 4591 or 5805.

[00232] In some cases, an antigen combination for targeting a neuroblastoma cell comprises SLC10A4 AND NOT WT1. In some cases, the WT1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 111, 5031, 5032 or 5033. In some cases, the SLC10A4 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 6095.

[00233] In some cases, an antigen combination for targeting a neuroblastoma cell comprises ST8SIA2 AND NOT WT1. In some cases, the WT1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 111, 5031, 5032 or 5033. In some cases, the ST8SIA2 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3413.

[00234] In some cases, an antigen combination for targeting a neuroblastoma cell comprises ST8SIA2 AND NOT FAP. In some cases, the FAP polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3055. In some cases, the ST8SIA2 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3413.

[00235] In some cases, an antigen combination for targeting a neuroblastoma cell comprises SLC10A4 AND NOT MAGEA1. In some cases, the MAGEA1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3191. In some cases, the SLC10A4 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 6095.

[00236] In some cases, an antigen combination for targeting a neuroblastoma cell comprises MARCH11 AND NOT MET. In some cases, the MARCH11 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 1209. In some cases, the MET polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 76 or 1332.

[00237] Additional examples of target antigen pairs for targeting a neuroblastoma cell are provided in FIG. 1 and FIG. 9-14.

Mantle cell lymphoma target antigen combinations

[00238] In some cases, an antigen combination for targeting a mantle-cell lymphoma cell comprises CLECL1 AND NOT EPHA3. In some cases, the EPHA3 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 3237 or 6669. In some cases, the CLECL1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 6267.

[00239] In some cases, an antigen combination for targeting a mantle-cell lymphoma cell comprises CELSR1 AND NOT ERBB2. In some cases, the ERBB2 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 474 or 3051. In some cases, the CELSR1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3895.

[00240] In some cases, an antigen combination for targeting a mantle-cell lymphoma cell comprises QSOX2 AND NOT ERBB2. In some cases, the ERBB2 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 474 or 3051. In some cases, the QSOX2 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 6621.

[00241] In some cases, an antigen combination for targeting a mantle-cell lymphoma cell comprises CLECL1 AND NOT CD160. In some cases, the CD160 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3624. In some cases, the CLECL1

polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 6267.

[00242] In some cases, an antigen combination for targeting a mantle-cell lymphoma cell comprises CELSR1 AND NOT SDC1. In some cases, the SDC1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 496 or 2747. In some cases, the CELSR1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3895.

[00243] In some cases, an antigen combination for targeting a mantle-cell lymphoma cell comprises TNFRSF13C AND NOT SDC1. In some cases, the SDC1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 496 or 2747. In some cases, the TNFRSF13C polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 5547.

[00244] In some cases, an antigen combination for targeting a mantle-cell lymphoma cell comprises CELSR1 AND NOT ERBB3. In some cases, the ERBB3 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 476 or 2517. In some cases, the CELSR1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3895.

[00245] In some cases, an antigen combination for targeting a mantle-cell lymphoma cell comprises TNFRSF13C AND NOT FAP. In some cases, the FAP polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3055. In some cases, the TNFRSF13C polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 5547.

[00246] In some cases, an antigen combination for targeting a mantle-cell lymphoma cell comprises SNN AND NOT NCAM1. In some cases, the NCAM1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 173, 949 or 6595. In some cases, the SNN

polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 2843.

[00247] In some cases, an antigen combination for targeting a mantle-cell lymphoma cell comprises TNFRSF10A AND NOT NCAM1. In some cases, the NCAM1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 173, 949 or 6595. In some cases, the TNFRSF10A polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 2911.

[00248] Additional examples of target antigen pairs for targeting a mantle cell lymphoma are provided in FIG. 1 and FIG. 9-14.

Sarcoma target antigen combinations

[00249] In some cases, an antigen combination for targeting a sarcoma cell comprises ADAM12 AND NOT ERBB4. In some cases, the ERBB4 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 914 or 3238. In some cases, the ADAM12 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 2839 or 4815.

[00250] In some cases, an antigen combination for targeting a sarcoma cell comprises AXL AND NOT ST6GALNAC2. In some cases, the ST6GALNAC2 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3493. In some cases, the AXL polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 2438 or 4847.

[00251] In some cases, an antigen combination for targeting a sarcoma cell comprises ADAM12 AND NOT ERBB3. In some cases, the ERBB3 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 476 or 2517. In some cases, the ADAM12 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 2839 or 4815.

[00252] In some cases, an antigen combination for targeting a sarcoma cell comprises ADAM12 AND NOT NAALAD2. In some cases, the NAALAD2 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3289. In some cases, the ADAM12 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 2839 or 4815.

[00253] In some cases, an antigen combination for targeting a sarcoma cell comprises TRAM2 AND NOT TMEM265. In some cases, the TMEM265 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 7113. In some cases, the TRAM2 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3738.

[00254] In some cases, an antigen combination for targeting a sarcoma cell comprises ADAM12 AND NOT PDZK1IP1. In some cases, the PDZK1IP1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3359. In some cases, the ADAM12 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 2839 or 4815.

[00255] In some cases, an antigen combination for targeting a sarcoma cell comprises TRAM2 AND NOT LPCAT3. In some cases, the LPCAT3 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3362. In some cases, the TRAM2 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3738.

[00256] In some cases, an antigen combination for targeting a sarcoma cell comprises CD163L1 AND NOT PRSS16. In some cases, the PRSS16 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3389. In some cases, the CD163L1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 6416.

[00257] In some cases, an antigen combination for targeting a sarcoma cell comprises ADAM12 AND NOT PRSS16. In some cases, the PRSS16 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the

amino acid sequence set forth in SEQ ID NO: 3389. In some cases, the ADAM12 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 2839 or 4815.

[00258] In some cases, an antigen combination for targeting a sarcoma cell comprises TRAM2 AND NOT PRSS16. In some cases, the PRSS16 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3389. In some cases, the TRAM2 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3738.

[00259] Additional examples of target antigen pairs for targeting a sarcoma cell are provided in FIG. 1 and FIG. 9-14.

Non-small cell lung cancer target antigen combinations

[00260] In some cases, an antigen combination for targeting a non-small cell lung cancer cell comprises FAP AND NOT LRRN4CL. In some cases, the LRRN4CL polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 6935. In some cases, the FAP polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3055.

[00261] In some cases, an antigen combination for targeting a non-small-cell lung cancer cell comprises FAP AND NOT GALNT16. In some cases, the GALNT16 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 2226 or 4700. In some cases, the FAP polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3055.

[00262] In some cases, an antigen combination for targeting a non-small-cell lung cancer cell comprises FAP AND NOT PTGFR. In some cases, the PTGFR polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs 346 or 804. In some cases, the FAP polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3055.

[00263] In some cases, an antigen combination for targeting a non-small-cell lung cancer cell comprises FAP AND NOT LCAT. In some cases, the LCAT polypeptide comprises an amino acid sequence having

at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 71. In some cases, the FAP polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3055.

[00264] In some cases, an antigen combination for targeting a non-small-cell lung cancer cell comprises FAP AND NOT SFRP1. In some cases, the SFRP1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 2753. In some cases, the FAP polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3055.

[00265] In some cases, an antigen combination for targeting a non-small-cell lung cancer cell comprises FAP AND NOT ITGA7. In some cases, the ITGA7 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 1800, 1801 or 2576. In some cases, the FAP polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3055.

[00266] In some cases, an antigen combination for targeting a non-small-cell lung cancer cell comprises FAP AND NOT ABCB1. In some cases, the ABCB1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 331. In some cases, the FAP polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3055.

[00267] In some cases, an antigen combination for targeting a non-small-cell lung cancer cell comprises FAP AND NOT RECK. In some cases, the RECK polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 4774. In some cases, the FAP polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3055.

[00268] In some cases, an antigen combination for targeting a non-small-cell lung cancer cell comprises FAP AND NOT ENPP1. In some cases, the ENPP1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3456. In some cases, the FAP polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3055.

[00269] In some cases, an antigen combination for targeting a non-small-cell lung cancer cell comprises FAP AND NOT TGFBR3. In some cases, the TGFBR3 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 2803. In some cases, the FAP polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3055.

[00270] Additional examples of target antigen pairs for targeting a non-small-cell lung cancer cell are provided in FIG. 1 and FIG. 9-14.

Acute myeloid leukemia (AML) target antigen combinations

[00271] In some cases, an antigen combination for targeting an acute myeloid leukemia cell comprises SLC22A16 AND NOT SDC1. In some cases, the SDC1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 496 or 2747. In some cases, the SLC22A16 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 5452.

[00272] In some cases, an antigen combination for targeting an acute myeloid leukemia cell comprises FUT4 AND NOT SDC1. In some cases, the SDC1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 496 or 2747. In some cases, the FUT4 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 2534.

[00273] In some cases, an antigen combination for targeting an acute myeloid leukemia cell comprises ELANE AND NOT SDC1. In some cases, the SDC1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 496 or 2747. In some cases, the ELANE polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 2513.

[00274] In some cases, an antigen combination for targeting an acute myeloid leukemia cell comprises MS4A3 AND NOT SLAMF7. In some cases, the SLAMF7 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 4788. In some cases, the MS4A3 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least

98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 721, 735 or 3443.

[00275] In some cases, an antigen combination for targeting an acute myeloid leukemia cell comprises TMC8 AND NOT SLAMF7. In some cases, the SLAMF7 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 4788. In some cases, the TMC8 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 6065.

[00276] In some cases, an antigen combination for targeting an acute myeloid leukemia cell comprises NOT PFN2 AND NOT SLAMF7. In some cases, the SLAMF7 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 4788. In some cases, the PFN2 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 2695 or 5556.

[00277] In some cases, an antigen combination for targeting an acute myeloid leukemia cell comprises CD33 AND NOT SLAMF7. In some cases, the SLAMF7 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 4788. In some cases, the CD33 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 1062 or 2462.

[00278] In some cases, an antigen combination for targeting an acute myeloid leukemia cell comprises FUT4 AND NOT SLAMF7. In some cases, the SLAMF7 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 4788. In some cases, the FUT4 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 2534.

[00279] In some cases, an antigen combination for targeting an acute myeloid leukemia cell comprises NRROS AND NOT SLAMF7. In some cases, the SLAMF7 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 4788. In some cases, the NRROS polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least

98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 6817.

[00280] In some cases, an antigen combination for targeting an acute myeloid leukemia cell comprises FUT4 AND NOT SLAMF7. In some cases, the SLAMF7 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 4788. In some cases, the FUT4 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 2534.

[00281] Additional examples of target antigen pairs for targeting an AML cell are provided in FIG. 1 and FIG. 9-14.

Stomach cancer target antigen combinations

[00282] In some cases, an antigen combination for targeting a stomach cancer cell comprises MUC13 AND NOT SLC30A10. In some cases, the SLC30A10 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 4487. In some cases, the MUC13 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 5442.

[00283] In some cases, an antigen combination for targeting a stomach cancer cell comprises MUC13 AND NOT SCNN1B. In some cases, the SCNN1B polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 96. In some cases, the MUC13 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 5442.

[00284] In some cases, an antigen combination for targeting a stomach cancer cell comprises MUC13 AND APOLD1. In some cases, the APOLD1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 1423 or 5181. In some cases, the MUC13 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 5442.

[00285] In some cases, an antigen combination for targeting a stomach cancer cell comprises MUC13 AND GREM1. In some cases, the GREM1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3812. In some cases, the MUC13 polypeptide comprises

an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 5442.

[00286] In some cases, an antigen combination for targeting a stomach cancer cell comprises MUC13 AND NOT DHRS9. In some cases, the DHRS9 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 1628, 1629, 3364 or 6865. In some cases, the MUC13 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 5442.

[00287] In some cases, an antigen combination for targeting a stomach cancer cell comprises MUC13 AND NOT TRPM6. In some cases, the TRPM6 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 4319. In some cases, the MUC13 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 5442.

[00288] In some cases, an antigen combination for targeting a stomach cancer cell comprises MUC13 AND NOT CLDN8. In some cases, the CLDN8 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 6874. In some cases, the MUC13 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 5442.

[00289] In some cases, an antigen combination for targeting a stomach cancer cell comprises MUC13 AND NOT SLC26A2. In some cases, the SLC26A2 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 32. In some cases, the MUC13 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 5442.

[00290] In some cases, an antigen combination for targeting a stomach cancer cell comprises MUC13 AND NOT CWH43. In some cases, the CWH43 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 5119. In some cases, the MUC13 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 5442.

[00291] In some cases, an antigen combination for targeting a stomach cancer cell comprises GREM1 AND TNFRSF10B. In some cases, the TNFRSF10B polypeptide comprises an amino acid sequence

having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 2910 or 5994. In some cases, the GREM1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3812.

[00292] Additional examples of target antigen pairs for targeting a stomach cancer cell are provided in FIG. 1 and FIG. 9-14.

Diffuse large B-cell lymphoma target antigen combinations

[00293] In some cases, an antigen combination for targeting a diffuse large B-cell lymphoma cell comprises PLA2G2D AND NOT GABRD. In some cases, the GABRD polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 271. In some cases, the PLA2G2D polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3768.

[00294] In some cases, an antigen combination for targeting a diffuse large B-cell lymphoma cell comprises PLA2G2D AND NOT BMPR1B. In some cases, the BMPR1B polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 2315. In some cases, the PLA2G2D polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3768.

[00295] In some cases, an antigen combination for targeting a diffuse large B-cell lymphoma cell comprises PLA2G2D AND NOT CRYGC. In some cases, the CRYGC polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 4754. In some cases, the PLA2G2D polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3768/

[00296] In some cases, an antigen combination for targeting a diffuse large B-cell lymphoma cell comprises PLA2G2D AND NOT HCN4. In some cases, the HCN4 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3292. In some cases, the PLA2G2D polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least

95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3768.

[00297] In some cases, an antigen combination for targeting a diffuse large B-cell lymphoma cell comprises PLA2G2D AND NOT SLC17A4. In some cases, the SLC17A4 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3294. In some cases, the PLA2G2D polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3768.

[00298] In some cases, an antigen combination for targeting a diffuse large B-cell lymphoma cell comprises PLA2G2D AND NOT CDH7. In some cases, the CDH7 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 3031 or 5510. In some cases, the PLA2G2D polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3768.

[00299] In some cases, an antigen combination for targeting a diffuse large B-cell lymphoma cell comprises PLA2G2D AND NOT CDH12. In some cases, the CDH12 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 2955. In some cases, the PLA2G2D polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3768.

[00300] In some cases, an antigen combination for targeting a diffuse large B-cell lymphoma cell comprises PLA2G2D AND NOT CLEC3A. In some cases, the CLEC3A polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3356. In some cases, the PLA2G2D polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3768.

[00301] In some cases, an antigen combination for targeting a diffuse large B-cell lymphoma cell comprises PLA2G2D AND NOT PDZK1IP1. In some cases, the PDZK1IP1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3359. In some cases, the PLA2G2D polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least

95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3768.

[00302] In some cases, an antigen combination for targeting a diffuse large B-cell lymphoma cell comprises PLA2G2D AND NOT TENM1. In some cases, the TENM1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 2049, 2050 or 3896. In some cases, the PLA2G2D polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3768.

[00303] Additional examples of target antigen pairs for targeting a diffuse large B-cell lymphoma are provided in FIG. 1 and FIG. 9-14.

Lung cancer (e.g., lung adenocarcinoma) target antigen combinations

[00304] In some cases, an antigen combination for targeting a lung cancer cell comprises SFTPD AND NOT IL3RA. In some cases, the SFTPD polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 2755. In some cases, the IL3RA polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 2567.

[00305] In some cases, an antigen combination for targeting a lung cancer cell comprises SFTPD AND NOT CDH5. In some cases, the SFTPD polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 2755. In some cases, the CDH5 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 2475.

[00306] In some cases, an antigen combination for targeting a lung cancer cell comprises SFTPC AND NOT RAMP2. In some cases, the SFTPC polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 2754. In some cases, the RAMP2 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3385.

[00307] In some cases, an antigen combination for targeting a lung cancer cell comprises SFTPD AND NOT RAMP2. In some cases, the SFTPD polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 2755. In some cases, the RAMP2 polypeptide comprises

an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3385.

[00308] In some cases, an antigen combination for targeting a lung cancer cell comprises SFTPD AND NOT SIRPB1. In some cases, the SFTPD polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 2755. In some cases, the SIRPB1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 1082, 1573 or 3426.

[00309] In some cases, an antigen combination for targeting a lung cancer cell comprises SFTPD AND NOT ABCA8. In some cases, the SFTPD polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 2755. In some cases, the ABCA8 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3640.

[00310] In some cases, an antigen combination for targeting a lung cancer cell comprises SLC34A2 AND NOT DLL1. In some cases, the DLL1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3326. In some cases, the SLC34A2 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3490.

[00311] In some cases, an antigen combination for targeting a lung cancer cell comprises SFTPC AND NOT LYVE1. In some cases, the LYVE1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3547. In some cases, the SFTPC polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 2754.

[00312] In some cases, an antigen combination for targeting a lung cancer cell comprises SFTPD AND NOT LYVE1. In some cases, the SFTPD polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 2755. In some cases, the LYVE1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3547.

[00313] In some cases, an antigen combination for targeting a lung cancer cell comprises SFTPD AND NOT PNPLA6. In some cases, the SFTPD polypeptide comprises an amino acid sequence having at least

85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 2755. In some cases, the PNPLA6 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 2146, 2147, 2148, 2149 or 3551.

[00314] Additional examples of target antigen pairs for targeting a lung cancer cell are provided in FIG. 1 and FIG. 9-14.

Oligodendroglioma target antigen combinations

[00315] In some cases, an antigen combination for targeting an oligodendroglioma cell comprises NLGN3 AND NOT FOLR2. In some cases, the FOLR2 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 261, 1261, 1262 or 1263. In some cases, the NLGN3 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 2174, 4543 or 6589.

[00316] In some cases, an antigen combination for targeting an oligodendroglioma cell comprises NLGN3 AND NOT SLAMF7. In some cases, the SLAMF7 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 4788. In some cases, the NLGN3 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 2174, 4543 or 6589.

[00317] In some cases, an antigen combination for targeting an oligodendroglioma cell comprises NLGN3 AND NOT SDC1. In some cases, the SDC1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 496 or 2747. In some cases, the NLGN3 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 2174, 4543 or 6589.

[00318] In some cases, an antigen combination for targeting an oligodendroglioma cell comprises NLGN3 AND NOT TNFRSF8. In some cases, the TNFRSF8 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 2323 or 6140. In some cases, the NLGN3 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least

98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 2174, 4543 or 6589.

[00319] In some cases, an antigen combination for targeting an oligodendroglioma cell comprises NLGN3 AND NOT IL11RA. In some cases, the IL11RA polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 1701, 3066 or 5993. In some cases, the NLGN3 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 2174, 4543 or 6589.

[00320] In some cases, an antigen combination for targeting an oligodendroglioma cell comprises NLGN3 AND NOT MUC1. In some cases, the MUC1 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 661, 662, 922, 923, 924, 925 or 2657. In some cases, the NLGN3 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 2174, 4543 or 6589.

[00321] In some cases, an antigen combination for targeting an oligodendroglioma cell comprises NLGN3 AND NOT KDR. In some cases, the KDR polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 2607. In some cases, the NLGN3 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 2174, 4543 or 6589.

[00322] In some cases, an antigen combination for targeting an oligodendroglioma cell comprises NLGN3 AND NOT IGF1R. In some cases, the IGF1R polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 308. In some cases, the NLGN3 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 2174, 4543 or 6589.

[00323] In some cases, an antigen combination for targeting an oligodendroglioma cell comprises NLGN3 AND NOT ALK. In some cases, the ALK polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NO: 3013. In some cases, the NLGN3 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or

100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 2174, 4543 or 6589.

[00324] In some cases, an antigen combination for targeting an oligodendroglioma cell comprises NLGN3 AND NOT ERBB3. In some cases, the ERBB3 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 476 or 2517. In some cases, the NLGN3 polypeptide comprises an amino acid sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to the amino acid sequence set forth in SEQ ID NOs: 2174, 4543 or 6589.

[00325] Additional examples of target antigen pairs for targeting an oligodendroglioma cell are provided in FIG. 1 and FIG. 9-14.

Other Antigen Combinations

[00326] In some cases, an antigen combination may be selected for targeting a cancer where two antigens of the combination are "clinical antigens" (i.e., the antigen has been evaluated clinically (i.e., in an immunotherapy clinical trial) and found to be a marker for one or more cancers). An antigen combination having two or more clinical antigens may include any clinical antigen including those described herein.

[00327] Combinations of clinical antigens include combinations where an antigen is highly expressed (H) in a target cancer, where an antigen is lowly expressed (L) in a target cancer, where two antigens are highly expressed (HH) in a target cancer, where two antigens are lowly expressed (LL) in a target cancer, where a first antigen is highly expressed and the second antigen is lowly expressed (HL) in a target cancer, where a first antigen is lowly expressed and the second antigen is highly expressed (LH) in a target cancer, where two antigens are expressed (i.e., an "OR" gate) in a target cancer.

[00328] For example, in some instances, the following clinical-clinical antigen pairs may be employed in a subject antigen combination, alone or in combination with other antigens, where the pair is listed with one or more non-limiting exemplary cancers and an exemplary expression or gating structure as defined above: MLANA:PMEL (Melanoma; H), MLANA:B4GALNT1 (Melanoma; OR), MLANA:KDR (Melanoma; HL), EPCAM:MET (Breast; HL), MSLN:EPCAM (Breast; HL), IL3RA:CD33 (Leukemia, Myeloid, Acute; HH), EPCAM:TYR (Breast; HL), EPCAM:ROR1 (Breast; HH), CD19:MS4A1 (Lymphoma, Large B-Cell, Diffuse; OR), ERBB2:ROR1 (Breast; HL), ERBB2:TYR (Breast; HL), ERBB2:EPCAM (Breast; HH), MSLN:ERBB2 (Lung Adenocarcinoma; HL), MSLN:MUC1 (Lung Adenocarcinoma; H), ERBB2:MUC1 (Lung Adenocarcinoma; HL), ERBB2:MET (Breast; HL), MSLN:GPC3 (Colon; LH), EPHA2:ERBB2 (Glioma; LH), EPHA2:MUC1 (Glioma; LH), MUC1:ROR1 (Lung Carcinoma; LH), GPC3:MUC1 (Colon; LH), ERBB2:GPC3 (Colon; LH), CD19:CD22 (Lymphoma, Large B-Cell, Diffuse; HL), ROR1:TYR (Breast; LH), MS4A1:CD22 (Lymphoma, Large

B-Cell, Diffuse; HL), MSLN:ERBB2 (Carcinoma, Pancreatic Ductal; HL), MET:ROR1 (Breast; LH), MET:TYR (Breast; LH), MSLN:ROR1 (Breast; HL), MSLN:TYR (Breast; HL), MUC1:TNFRSF10A (Lung Carcinoma; OR), MSLN:MET (Breast; HL), ROR1:TNFRSF10A (Lung Carcinoma; LH), MSLN:ERBB2 (Lung Carcinoma; HL), MSLN:GPC3 (Lung Carcinoma; LH), MSLN:MUC1 (Lung Carcinoma; LH), MSLN:ROR1 (Lung Carcinoma; LH), MSLN:TYR (Lung Carcinoma; LH), MSLN:TNFRSF10A (Lung Carcinoma; LH), ERBB2:GPC3 (Lung Carcinoma; -), ERBB2:MUC1 (Lung Carcinoma; LH), ERBB2:ROR1 (Lung Carcinoma; LH), GPC3:MUC1 (Lung Carcinoma; LH), GPC3:ROR1 (Lung Carcinoma; LH), GPC3:TYR (Lung Carcinoma; LH), GPC3:TNFRSF10A (Lung Carcinoma; LH), MUC1:TYR (Lung Carcinoma; LH), ROR1:TYR (Lung Carcinoma; LH), TYR:TNFRSF10A (Lung Carcinoma; LH), MSLN:ERBB2 (Ovarian Neoplasms, Cystadenocarcinoma, Adenocarcinoma; HL), Flt3:CD123 (acute myeloid leukemia; HH), Flt3:CD33 (acute myeloid leukemia; HH) and GPC3:AFP (hepatocellular carcinoma; HH).

[00329] In some cases, useful clinical-clinical antigen pairs may include Flt3:CD123, for example as an AND-gate. In some instances, Flt3:CD123 may find use in targeting acute myeloid leukemia. In some cases, useful clinical-clinical antigen pairs may include Flt3:CD33, for example as an AND-gate. In some instances, Flt3:CD33 may find use in targeting acute myeloid leukemia. In some cases, useful clinical-clinical antigen pairs may include GPC3:AFP, for example as an AND-gate. In some instances, GPC3:AFP may find use in targeting hepatocellular carcinoma.

[00330] In some cases, an antigen combination may include one or more antigens of interest, where an "antigen of interest" may include clinical antigens as well as antigens that are of clinical interest for reasons other than having been identified as associated with a particular cancer type. Such other reasons include but are not limited to e.g., association with cancer stem cells (i.e., a cancer stem cell marker). Antigens of interest that may be included in antigen combinations include, e.g., those listed in Table 3 below.

[00331] Table 3.

ALK (SEQ ID NO:3013)	ULBP2 (SEQ ID NO:5133)	CLDN1 (SEQ ID NO:4771)
AXL (SEQ ID NO:2438)	ULBP3 (SEQ ID NO:7298)	CLDN10 (SEQ ID NO:1975)
CD276 (SEQ ID NO:685)	RAET1E (SEQ ID NO:5840)	CLDN11 (SEQ ID NO:3322)
NCR3LG1 (SEQ ID NO:7428)	RAET1G (SEQ ID NO:7369)	CLDN12 (SEQ ID NO:3711)
TNFRSF17 (SEQ ID NO:2312)	RAET1L (SEQ ID NO:7368)	CLDN14 (SEQ ID NO:1902)
CA9 (SEQ ID NO:2318)	CD34 (SEQ ID NO:697)	CLDN15 (SEQ ID NO:3920)
IL3RA (SEQ ID NO:2567)	ALDH1A1 (SEQ ID NO:7313)	CLDN16 (SEQ ID NO:3520)
SDC1 (SEQ ID NO:496)	PROM1 (SEQ ID NO:1878)	CLDN17 (SEQ ID NO:3713)
CD160 (SEQ ID NO:3624)	FLOT2 (SEQ ID NO:7359)	CLDN18 (SEQ ID NO:387)
L1CAM (SEQ ID NO:126)	CD24 (SEQ ID NO:7446)	CLDN19 (SEQ ID NO:1296)
CD19 (SEQ ID NO:2460)	CBX3 (SEQ ID NO:7304)	CLDN2 (SEQ ID NO:2267)
MS4A1 (SEQ ID NO:4853)	ABCA5 (SEQ ID NO:4483)	CLDN20 (SEQ ID NO:7334)
CD22 (SEQ ID NO:2461)	ABCB5 (SEQ ID NO:7412)	CLDN24 (SEQ ID NO:7422)
TNFRSF8 (SEQ ID NO:2323)	IGB1 (SEQ ID NO:7309)	CLDN22 (SEQ ID NO:7381)

CD33 (SEQ ID NO:1062)	LGR5 (SEQ ID NO:2866)	CLDN23 (SEQ ID NO:6713)
CD38 (SEQ ID NO:2465)	ALCAM (SEQ ID NO:2420)	CLDN3 (SEQ ID NO:2343)
CD44 (SEQ ID NO:172)	THY1 (SEQ ID NO:7454)	CLDN4 (SEQ ID NO:2342)
CD70 (SEQ ID NO:2331)	DNAJB8 (SEQ ID NO:7318)	CLDN5 (SEQ ID NO:1443)
SLAMF7 (SEQ ID NO:4788)	DDX3X (SEQ ID NO:7424)	CLDN6 (SEQ ID NO:4792)
EPCAM (SEQ ID NO:2632)	DLL3 (SEQ ID NO:4275)	CLDN7 (SEQ ID NO:2344)
EPHA2 (SEQ ID NO:3042)	BMPR1B (SEQ ID NO:2315)	CLDN8 (SEQ ID NO:6874)
EPHA3 (SEQ ID NO:3237)	SLC7A5 (SEQ ID NO:7352)	CLDN9 (SEQ ID NO:4752)
ERBB3 (SEQ ID NO:476)	STEAP1 (SEQ ID NO:3777)	CLEC14A (SEQ ID NO:6440)
ERBB4 (SEQ ID NO:914)	IL13RA1 (SEQ ID NO:2415)	CTAG1A (SEQ ID NO:7312)
FAP (SEQ ID NO:3055)	SLC34A2 (SEQ ID NO:3490)	CTAG2 (SEQ ID NO:7442)
FOLR1 (SEQ ID NO:260)	SEMA5B (SEQ ID NO:7357)	DKK1 (SEQ ID NO:7282)
FOLR2 (SEQ ID NO:261)	GPR37L1 (SEQ ID NO:3118)	DPEP1 (SEQ ID NO:7391)
B4GALNT1 (SEQ ID NO:2385)	RNF43 (SEQ ID NO:4337)	EDNRB (SEQ ID NO:33)
ST8SIA1 (SEQ ID NO:2758)	STEAP2 (SEQ ID NO:879)	EGFR (SEQ ID NO:3235)
GPA33 (SEQ ID NO:3374)	TRPM4 (SEQ ID NO:4315)	ENG (SEQ ID NO:35)
GPC3 (SEQ ID NO:2089)	TDGF1 (SEQ ID NO:7419)	ENPP3 (SEQ ID NO:3199)
ERBB2 (SEQ ID NO:474)	CR2 (SEQ ID NO:493)	GAGE1 (SEQ ID NO:7360)
CTAG1B (SEQ ID NO:7258)	CD79B (SEQ ID NO:178)	GPNMB (SEQ ID NO:454)
CSPG4 (SEQ ID NO:2497)	FCRL2 (SEQ ID NO:5166)	GUCY2C (SEQ ID NO:3179)
IGF1R (SEQ ID NO:308)	CEACAM6 (SEQ ID NO:7445)	HAVCR1 (SEQ ID NO:7417)
IL11RA (SEQ ID NO:1701)	IL20RA (SEQ ID NO:3936)	HHLA2 (SEQ ID NO:3626)
IL13RA2 (SEQ ID NO:185)	BCAN (SEQ ID NO:7332)	HSPA5 (SEQ ID NO:7305)
IGK (SEQ ID NO:7467)	EPHB2 (SEQ ID NO:3047)	IL2RA (SEQ ID NO:7271)
MAGEA1 (SEQ ID NO:3191)	TNFRSF13C (SEQ ID NO:5547)	ITGAV (SEQ ID NO:1802)
MAGEA3 (SEQ ID NO:7279)	CD74 (SEQ ID NO:7351)	ITGB6 (SEQ ID NO:313)
MLANA (SEQ ID NO:3302)	CD79A (SEQ ID NO:7257)	MAGEA11 (SEQ ID NO:7346)
MSLN (SEQ ID NO:3377)	CXCR5 (SEQ ID NO:7256)	MAGEA4 (SEQ ID NO:7341)
MET (SEQ ID NO:76)	HLA-DOB (SEQ ID NO:2555)	MOK (SEQ ID NO:7436)
MUC13 (SEQ ID NO:5442)	P2RX5 (SEQ ID NO:2680)	MST1R (SEQ ID NO:2656)
MUC17 (SEQ ID NO:835)	CD72 (SEQ ID NO:2470)	MUC4 (SEQ ID NO:3070)
MUC1 (SEQ ID NO:661)	CD180 (SEQ ID NO:3318)	OAS1 (SEQ ID NO:7354)
MUC16 (SEQ ID NO:5067)	FCRL1 (SEQ ID NO:1941)	PCYT1A (SEQ ID NO:7455)
NCAM1 (SEQ ID NO:173)	FCRL5 (SEQ ID NO:5213)	PMEL (SEQ ID NO:3597)
PSCA (SEQ ID NO:3340)	TMEFF2 (SEQ ID NO:4183)	PRR4 (SEQ ID NO:7429)
FOLH1 (SEQ ID NO:631)	SST (SEQ ID NO:7266)	PTK7 (SEQ ID NO:2709)
TNFSF11 (SEQ ID NO:2871)	SSTR2 (SEQ ID NO:928)	SLC39A6 (SEQ ID NO:1158)
ROR1 (SEQ ID NO:1070)	SSTR5 (SEQ ID NO:937)	SPON2 (SEQ ID NO:7392)
TNFRSF10A (SEQ ID NO:2911)	SSTR1 (SEQ ID NO:927)	SSTR4 (SEQ ID NO:936)
TNFRSF10B (SEQ ID NO:2910)	SSTR3 (SEQ ID NO:935)	SSX1 (SEQ ID NO:7438)
KDR (SEQ ID NO:2607)	AFP (SEQ ID NO:7252)	TDGF1P3 (SEQ ID NO:7466)
WT1 (SEQ ID NO:111)	AGER (SEQ ID NO:1470)	TNC (SEQ ID NO:2562)
ITGB1 (SEQ ID NO:7310)	ANXA1 (SEQ ID NO:7253)	TPBG (SEQ ID NO:2164)
ITGB3 (SEQ ID NO:64)	BAGE (SEQ ID NO:7269)	TYR (SEQ ID NO:110)
FUT3 (SEQ ID NO:43)	BIRC5 (SEQ ID NO:7342)	VCAM1 (SEQ ID NO:946)
CEACAM5 (SEQ ID NO:7444)	CD37 (SEQ ID NO:823)	VTCN1 (SEQ ID NO:5057)
ULBP1 (SEQ ID NO:7458)	CD52 (SEQ ID NO:2478)	

[00332] In some cases, useful antigen combinations may include an antigen combination that includes a clinical antigen in combination with a tissue antigen (e.g., an antigen expressed in normal brain tissue) in an AND NOT gate. For example, useful antigen combinations may include a GD2 (B4GALNT1; SEQ ID NO:2385) clinical antigen AND NOT a tissue antigen selected from: OPALIN (SEQ ID NO:833), TMEM235 (SEQ ID NO:1869), GABRA1 (SEQ ID NO:262), KCNJ9 (SEQ ID NO:3190), GRM3 (SEQ ID NO:290), SEZ6 (SEQ ID NO:1143), NTSR2 (SEQ ID NO:3753), KCNK4 (SEQ ID NO:5486), SLCO1A2 (SEQ ID NO:7290), SLC24A2 (SEQ ID NO:4632), MOG (SEQ ID NO:528), GABRG1 (SEQ ID NO:6354), GABRG2 (SEQ ID NO:272), CNTNAP4 (SEQ ID NO:5492), DSCAM (SEQ ID NO:2354), CACNG3 (SEQ ID NO:3510), CRB1 (SEQ ID NO:6891), CDH10 (SEQ ID NO:3552), HRH3 (SEQ ID NO:3657), GRIK1 (SEQ ID NO:280), SLC39A12 (SEQ ID NO:1826), GPR158 (SEQ ID NO:4716), CACNG2 (SEQ ID NO:3430), SYT3 (SEQ ID NO:7410), HTR5A (SEQ ID NO:4993), CACNG7 (SEQ ID NO:5264), GPR37L1 (SEQ ID NO:3118), LRRTM3 (SEQ ID NO:7362), GLRA2 (SEQ ID NO:1281), CHRNB2 (SEQ ID NO:228), KCNQ2 (SEQ ID NO:3067), JPH3 (SEQ ID NO:4690), GPR19 (SEQ ID NO:3446), ADCY8 (SEQ ID NO:1248), SPOCK3 (SEQ ID NO:845), SLC32A1 (SEQ ID NO:5606), OPCML (SEQ ID NO:594), GABRA3 (SEQ ID NO:264), GRM5 (SEQ ID NO:292), SCN1A (SEQ ID NO:2133), SLC5A11 (SEQ ID NO:5546), KCNC1 (SEQ ID NO:1249), SLC12A5 (SEQ ID NO:1506), GRM4 (SEQ ID NO:291), GRM1 (SEQ ID NO:288), GRIA4 (SEQ ID NO:279), MEGF11 (SEQ ID NO:5354), CACNA1B (SEQ ID NO:211), LYPD1 (SEQ ID NO:970), GRID2 (SEQ ID NO:2398), SCN2A (SEQ ID NO:842), NKAIN2 (SEQ ID NO:864), UNC5A (SEQ ID NO:5690), SLC4A10 (SEQ ID NO:7421), TMEFF2 (SEQ ID NO:4183), CSMD3 (SEQ ID NO:5532), PPAPDC1A (SEQ ID NO:715), HAPLN4 (SEQ ID NO:4968), GPR85 (SEQ ID NO:1920), ANTXR2 (SEQ ID NO:1876), CACNG4 (SEQ ID NO:3934), CSPG5 (SEQ ID NO:3516), KCNK10 (SEQ ID NO:4784), CHRNA4 (SEQ ID NO:225), CNTNAP2 (SEQ ID NO:3871), KCNJ10 (SEQ ID NO:2597), GABRB2 (SEQ ID NO:269), GRIN1 (SEQ ID NO:282), CRB2 (SEQ ID NO:6381), SHISA7 (SEQ ID NO:7406), NKAIN4 (SEQ ID NO:6122), HTR2C (SEQ ID NO:301), CACNG8 (SEQ ID NO:5263), NRG3 (SEQ ID NO:569), ABCG4 (SEQ ID NO:1668), CDH8 (SEQ ID NO:2476), GABRD (SEQ ID NO:271), KIRREL3 (SEQ ID NO:2022), GABRB1 (SEQ ID NO:268), KCNA2 (SEQ ID NO:3181), CDH20 (SEQ ID NO:5261), IGDCC3 (SEQ ID NO:3159), KCNJ6 (SEQ ID NO:2596), CNIH2 (SEQ ID NO:6658), KCNK12 (SEQ ID NO:7295), CDH18 (SEQ ID NO:2192), CSMD2 (SEQ ID NO:5530), SYT4 (SEQ ID NO:4722), OR2L13 (SEQ ID NO:6468), CDH9 (SEQ ID NO:4198), GABRA2 (SEQ ID NO:263), KCNF1 (SEQ ID NO:2592), MAG (SEQ ID NO:2634), CALN1 (SEQ ID NO:645), GRIN2B (SEQ ID NO:284), GRM7 (SEQ ID NO:294), VSTM2A (SEQ ID NO:7459), GPR61 (SEQ ID NO:5271), OMG (SEQ ID NO:2672), KCNA1 (SEQ ID NO:66), GPR83 (SEQ ID NO:4239), ATP8A2 (SEQ ID NO:4237), GABBR2 (SEQ ID NO:3288), GPR12 (SEQ ID NO:3255), TRPM3 (SEQ ID NO:519), SLC8A3 (SEQ ID NO:1424), KCND2 (SEQ ID NO:7288), GSG1L (SEQ ID NO:1236),

SLC30A10 (SEQ ID NO:4487), ASTN1 (SEQ ID NO:3016), GPR179 (SEQ ID NO:7447), LRFN2 (SEQ ID NO:4711), CACNA1E (SEQ ID NO:214), CALY (SEQ ID NO:4119), SLC6A15 (SEQ ID NO:1932), KIAA0319 (SEQ ID NO:2227), SYT6 (SEQ ID NO:6957), PTPRR (SEQ ID NO:2724), KCTD8 (SEQ ID NO:7331), GPR22 (SEQ ID NO:3261), SLC4A8 (SEQ ID NO:818), LAMP5 (SEQ ID NO:3729), MEGF10 (SEQ ID NO:5355), FXYD7 (SEQ ID NO:4864), KCNK9 (SEQ ID NO:4257), SLC1A6 (SEQ ID NO:3206), MLC1 (SEQ ID NO:4042), OPRK1 (SEQ ID NO:325), ATP2B2 (SEQ ID NO:353), ACSL6 (SEQ ID NO:560), THBD (SEQ ID NO:108), PTPRT (SEQ ID NO:3622), PCDHGC4 (SEQ ID NO:2688), CLDN10 (SEQ ID NO:1975), KCNV1 (SEQ ID NO:3925), LPPR3 (SEQ ID NO:5096), SLCO1C1 (SEQ ID NO:1885), PCDH8 (SEQ ID NO:2690), ANO4 (SEQ ID NO:6566), LRRTM4 (SEQ ID NO:1505), PCDH15 (SEQ ID NO:1688), CCKBR (SEQ ID NO:6481), GABRA5 (SEQ ID NO:266), SLC6A12 (SEQ ID NO:1290), GRIN2A (SEQ ID NO:283), SLC1A2 (SEQ ID NO:2977), SLC43A1 (SEQ ID NO:2856), KCNC2 (SEQ ID NO:5836), ELFN2 (SEQ ID NO:5534), ATP7A (SEQ ID NO:10), GRIK4 (SEQ ID NO:3965), LRRC55 (SEQ ID NO:448), HCN2 (SEQ ID NO:7377), NKAIN1 (SEQ ID NO:5039), DPP10 (SEQ ID NO:432), AJAP1 (SEQ ID NO:894), NPFFR1 (SEQ ID NO:4886), TRPC3 (SEQ ID NO:1434), TGFBI (SEQ ID NO:107), SLC6A7 (SEQ ID NO:7370), GABRA4 (SEQ ID NO:265), SLC13A5 (SEQ ID NO:1732), GRIN2C (SEQ ID NO:285), HCN1 (SEQ ID NO:4763), SLC26A8 (SEQ ID NO:5552), PPIC (SEQ ID NO:336), NETO1 (SEQ ID NO:7427), TNFRSF10B (SEQ ID NO:2910), CDH22 (SEQ ID NO:4803), SLC6A13 (SEQ ID NO:4264), DISP2 (SEQ ID NO:7320), SLC6A11 (SEQ ID NO:3890), CD93 (SEQ ID NO:3698), EPHA10 (SEQ ID NO:1160), PHLDB2 (SEQ ID NO:1490), OXTR (SEQ ID NO:7326), WNT7A (SEQ ID NO:3089), GYPC (SEQ ID NO:7432), KCNA4 (SEQ ID NO:2589), PCDHAC2 (SEQ ID NO:4498), HGFAC (SEQ ID NO:2405), DRD1 (SEQ ID NO:253), SHISA9 (SEQ ID NO:7418), SCN8A (SEQ ID NO:3879), ICAM1 (SEQ ID NO:7382), PIRT (SEQ ID NO:1201), A4GALT (SEQ ID NO:4291), MRGPRF (SEQ ID NO:1129), CD248 (SEQ ID NO:4649), CD58 (SEQ ID NO:7405), CD44 (SEQ ID NO:172), EPHA2 (SEQ ID NO:3042) or PROCR (SEQ ID NO:3480). In some instances, such antigen combinations may be useful in targeting melanoma.

[00333] Useful antigen combinations may include a MAGEA1 (SEQ ID NO:3191) clinical antigen AND NOT a tissue antigen (e.g., a brain tissue antigen) selected from: OPALIN (SEQ ID NO:833), TMEM235 (SEQ ID NO:1869), GABRA1 (SEQ ID NO:262), KCNJ9 (SEQ ID NO:3190), GRM3 (SEQ ID NO:290), SEZ6 (SEQ ID NO:1143), NTSR2 (SEQ ID NO:3753), KCNK4 (SEQ ID NO:5486), SLCO1A2 (SEQ ID NO:7290), SLC24A2 (SEQ ID NO:4632), MOG (SEQ ID NO:528), GABRG1 (SEQ ID NO:6354), GABRG2 (SEQ ID NO:272), CNTNAP4 (SEQ ID NO:5492), DSCAM (SEQ ID NO:2354), CACNG3 (SEQ ID NO:3510), CRB1 (SEQ ID NO:6891), CDH10 (SEQ ID NO:3552), HRH3 (SEQ ID NO:3657), GRIK1 (SEQ ID NO:280), SLC39A12 (SEQ ID NO:1826), GPR158 (SEQ ID NO:4716), CACNG2 (SEQ ID NO:3430), SYT3 (SEQ ID NO:7410), HTR5A (SEQ ID NO:4993),

CACNG7 (SEQ ID NO:5264), GPR37L1 (SEQ ID NO:3118), LRRTM3 (SEQ ID NO:7362), GLRA2 (SEQ ID NO:1281), CHRNB2 (SEQ ID NO:228), KCNQ2 (SEQ ID NO:3067), JPH3 (SEQ ID NO:4690), GPR19 (SEQ ID NO:3446), ADCY8 (SEQ ID NO:1248), SPOCK3 (SEQ ID NO:845), SLC32A1 (SEQ ID NO:5606), OPCML (SEQ ID NO:594), GABRA3 (SEQ ID NO:264), GRM5 (SEQ ID NO:292), SCN1A (SEQ ID NO:2133), SLC5A11 (SEQ ID NO:5546), KCNC1 (SEQ ID NO:1249), SLC12A5 (SEQ ID NO:1506), GRM4 (SEQ ID NO:291), GRM1 (SEQ ID NO:288), GRIA4 (SEQ ID NO:279), MEGF11 (SEQ ID NO:5354), CACNA1B (SEQ ID NO:211), LYPD1 (SEQ ID NO:970), GRID2 (SEQ ID NO:2398), SCN2A (SEQ ID NO:842), NKAIN2 (SEQ ID NO:864), UNC5A (SEQ ID NO:5690), SLC4A10 (SEQ ID NO:7421), TMEFF2 (SEQ ID NO:4183), CSMD3 (SEQ ID NO:5532), PPAPDC1A (SEQ ID NO:715), HAPLN4 (SEQ ID NO:4968), GPR85 (SEQ ID NO:1920), ANTXR2 (SEQ ID NO:1876), CACNG4 (SEQ ID NO:3934), CSPG5 (SEQ ID NO:3516), KCNK10 (SEQ ID NO:4784), CHRNA4 (SEQ ID NO:225), CNTNAP2 (SEQ ID NO:3871), KCNJ10 (SEQ ID NO:2597), GABRB2 (SEQ ID NO:269), GRIN1 (SEQ ID NO:282), CRB2 (SEQ ID NO:6381), SHISA7 (SEQ ID NO:7406), NKAIN4 (SEQ ID NO:6122), HTR2C (SEQ ID NO:301), CACNG8 (SEQ ID NO:5263), NRG3 (SEQ ID NO:569), ABCG4 (SEQ ID NO:1668), CDH8 (SEQ ID NO:2476), GABRD (SEQ ID NO:271), KIRREL3 (SEQ ID NO:2022), GABRB1 (SEQ ID NO:268), KCNA2 (SEQ ID NO:3181), CDH20 (SEQ ID NO:5261), IGDCC3 (SEQ ID NO:3159), KCNJ6 (SEQ ID NO:2596), CNIH2 (SEQ ID NO:6658), KCNK12 (SEQ ID NO:7295), CDH18 (SEQ ID NO:2192), CSMD2 (SEQ ID NO:5530), SYT4 (SEQ ID NO:4722), OR2L13 (SEQ ID NO:6468), CDH9 (SEQ ID NO:4198), GABRA2 (SEQ ID NO:263), KCNF1 (SEQ ID NO:2592), MAG (SEQ ID NO:2634), CALN1 (SEQ ID NO:645), GRIN2B (SEQ ID NO:284), GRM7 (SEQ ID NO:294), VSTM2A (SEQ ID NO:7459), GPR61 (SEQ ID NO:5271), OMG (SEQ ID NO:2672), KCNA1 (SEQ ID NO:66), GPR83 (SEQ ID NO:4239), ATP8A2 (SEQ ID NO:4237), GABBR2 (SEQ ID NO:3288), GPR12 (SEQ ID NO:3255), TRPM3 (SEQ ID NO:519), SLC8A3 (SEQ ID NO:1424), KCND2 (SEQ ID NO:7288), GSG1L (SEQ ID NO:1236), SLC30A10 (SEQ ID NO:4487), ASTN1 (SEQ ID NO:3016), GPR179 (SEQ ID NO:7447), LRFN2 (SEQ ID NO:4711), CACNA1E (SEQ ID NO:214), CALY (SEQ ID NO:4119), SLC6A15 (SEQ ID NO:1932), KIAA0319 (SEQ ID NO:2227), SYT6 (SEQ ID NO:6957), PTPRR (SEQ ID NO:2724), KCTD8 (SEQ ID NO:7331), GPR22 (SEQ ID NO:3261), SLC4A8 (SEQ ID NO:818), LAMP5 (SEQ ID NO:3729), MEGF10 (SEQ ID NO:5355), FXYD7 (SEQ ID NO:4864), KCNK9 (SEQ ID NO:4257), SLC1A6 (SEQ ID NO:3206), MLC1 (SEQ ID NO:4042), OPRK1 (SEQ ID NO:325), ATP2B2 (SEQ ID NO:353), ACSL6 (SEQ ID NO:560), THBD (SEQ ID NO:108), PTPRT (SEQ ID NO:3622), PCDHGC4 (SEQ ID NO:2688), CLDN10 (SEQ ID NO:1975), KCNV1 (SEQ ID NO:3925), LPPR3 (SEQ ID NO:5096), SLCO1C1 (SEQ ID NO:1885), PCDH8 (SEQ ID NO:2690), ANO4 (SEQ ID NO:6566), LRRTM4 (SEQ ID NO:1505), PCDH15 (SEQ ID NO:1688), CCKBR (SEQ ID NO:6481), GABRA5 (SEQ ID NO:266), SLC6A12 (SEQ ID NO:1290), GRIN2A (SEQ ID NO:283), SLC1A2 (SEQ ID

NO:2977), SLC43A1 (SEQ ID NO:2856), KCNC2 (SEQ ID NO:5836), ELFN2 (SEQ ID NO:5534), ATP7A (SEQ ID NO:10), GRIK4 (SEQ ID NO:3965), LRRC55 (SEQ ID NO:448), HCN2 (SEQ ID NO:7377), NKAIN1 (SEQ ID NO:5039), DPP10 (SEQ ID NO:432), AJAP1 (SEQ ID NO:894), NPFFR1 (SEQ ID NO:4886), TRPC3 (SEQ ID NO:1434), TGFBI (SEQ ID NO:107), SLC6A7 (SEQ ID NO:7370), GABRA4 (SEQ ID NO:265), SLC13A5 (SEQ ID NO:1732), GRIN2C (SEQ ID NO:285), HCN1 (SEQ ID NO:4763), SLC26A8 (SEQ ID NO:5552), PPIC (SEQ ID NO:336), NETO1 (SEQ ID NO:7427), TNFRSF10B (SEQ ID NO:2910), CDH22 (SEQ ID NO:4803), SLC6A13 (SEQ ID NO:4264), DISP2 (SEQ ID NO:7320), SLC6A11 (SEQ ID NO:3890), CD93 (SEQ ID NO:3698), EPHA10 (SEQ ID NO:1160), PHLDB2 (SEQ ID NO:1490), OXTR (SEQ ID NO:7326), WNT7A (SEQ ID NO:3089), GYPC (SEQ ID NO:7432), KCNA4 (SEQ ID NO:2589), PCDHAC2 (SEQ ID NO:4498), HGFAC (SEQ ID NO:2405), DRD1 (SEQ ID NO:253), SHISA9 (SEQ ID NO:7418), SCN8A (SEQ ID NO:3879), ICAM1 (SEQ ID NO:7382), PIRT (SEQ ID NO:1201), A4GALT (SEQ ID NO:4291), MRGPRF (SEQ ID NO:1129), CD248 (SEQ ID NO:4649), CD58 (SEQ ID NO:7405), CD44 (SEQ ID NO:172), EPHA2 (SEQ ID NO:3042) or PROCR (SEQ ID NO:3480). In some instances, such antigen combinations may be useful in targeting melanoma.

[00334] Useful antigen combinations may include a MAGEA3 (SEQ ID NO:7279) clinical antigen AND NOT a tissue antigen (e.g., a brain tissue antigen) selected from: OPALIN (SEQ ID NO:833), TMEM235 (SEQ ID NO:1869), GABRA1 (SEQ ID NO:262), KCNJ9 (SEQ ID NO:3190), GRM3 (SEQ ID NO:290), SEZ6 (SEQ ID NO:1143), NTSR2 (SEQ ID NO:3753), KCNK4 (SEQ ID NO:5486), SLCO1A2 (SEQ ID NO:7290), SLC24A2 (SEQ ID NO:4632), MOG (SEQ ID NO:528), GABRG1 (SEQ ID NO:6354), GABRG2 (SEQ ID NO:272), CNTNAP4 (SEQ ID NO:5492), DSCAM (SEQ ID NO:2354), CACNG3 (SEQ ID NO:3510), CRB1 (SEQ ID NO:6891), CDH10 (SEQ ID NO:3552), HRH3 (SEQ ID NO:3657), GRIK1 (SEQ ID NO:280), SLC39A12 (SEQ ID NO:1826), GPR158 (SEQ ID NO:4716), CACNG2 (SEQ ID NO:3430), SYT3 (SEQ ID NO:7410), HTR5A (SEQ ID NO:4993), CACNG7 (SEQ ID NO:5264), GPR37L1 (SEQ ID NO:3118), LRRTM3 (SEQ ID NO:7362), GLRA2 (SEQ ID NO:1281), CHRNB2 (SEQ ID NO:228), KCNQ2 (SEQ ID NO:3067), JPH3 (SEQ ID NO:4690), GPR19 (SEQ ID NO:3446), ADCY8 (SEQ ID NO:1248), SPOCK3 (SEQ ID NO:845), SLC32A1 (SEQ ID NO:5606), OPCML (SEQ ID NO:594), GABRA3 (SEQ ID NO:264), GRM5 (SEQ ID NO:292), SCN1A (SEQ ID NO:2133), SLC5A11 (SEQ ID NO:5546), KCNC1 (SEQ ID NO:1249), SLC12A5 (SEQ ID NO:1506), GRM4 (SEQ ID NO:291), GRM1 (SEQ ID NO:288), GRIA4 (SEQ ID NO:279), MEGF11 (SEQ ID NO:5354), CACNA1B (SEQ ID NO:211), LYPD1 (SEQ ID NO:970), GRID2 (SEQ ID NO:2398), SCN2A (SEQ ID NO:842), NKAIN2 (SEQ ID NO:864), UNC5A (SEQ ID NO:5690), SLC4A10 (SEQ ID NO:7421), TMEFF2 (SEQ ID NO:4183), CSMD3 (SEQ ID NO:5532), PPAPDC1A (SEQ ID NO:715), HAPLN4 (SEQ ID NO:4968), GPR85 (SEQ ID NO:1920), ANTXR2 (SEQ ID NO:1876), CACNG4 (SEQ ID NO:3934), CSPG5 (SEQ ID NO:3516), KCNK10 (SEQ ID

NO:4784), CHRNA4 (SEQ ID NO:225), CNTNAP2 (SEQ ID NO:3871), KCNJ10 (SEQ ID NO:2597), GABRB2 (SEQ ID NO:269), GRIN1 (SEQ ID NO:282), CRB2 (SEQ ID NO:6381), SHISA7 (SEQ ID NO:7406), NKAIN4 (SEQ ID NO:6122), HTR2C (SEQ ID NO:301), CACNG8 (SEQ ID NO:5263), NRG3 (SEQ ID NO:569), ABCG4 (SEQ ID NO:1668), CDH8 (SEQ ID NO:2476), GABRD (SEQ ID NO:271), KIRREL3 (SEQ ID NO:2022), GABRB1 (SEQ ID NO:268), KCNA2 (SEQ ID NO:3181), CDH20 (SEQ ID NO:5261), IGDCC3 (SEQ ID NO:3159), KCNJ6 (SEQ ID NO:2596), CNIH2 (SEQ ID NO:6658), KCNK12 (SEQ ID NO:7295), CDH18 (SEQ ID NO:2192), CSMD2 (SEQ ID NO:5530), SYT4 (SEQ ID NO:4722), OR2L13 (SEQ ID NO:6468), CDH9 (SEQ ID NO:4198), GABRA2 (SEQ ID NO:263), KCNF1 (SEQ ID NO:2592), MAG (SEQ ID NO:2634), CALN1 (SEQ ID NO:645), GRIN2B (SEQ ID NO:284), GRM7 (SEQ ID NO:294), VSTM2A (SEQ ID NO:7459), GPR61 (SEQ ID NO:5271), OMG (SEQ ID NO:2672), KCNA1 (SEQ ID NO:66), GPR83 (SEQ ID NO:4239), ATP8A2 (SEQ ID NO:4237), GABBR2 (SEQ ID NO:3288), GPR12 (SEQ ID NO:3255), TRPM3 (SEQ ID NO:519), SLC8A3 (SEQ ID NO:1424), KCND2 (SEQ ID NO:7288), GSG1L (SEQ ID NO:1236), SLC30A10 (SEQ ID NO:4487), ASTN1 (SEQ ID NO:3016), GPR179 (SEQ ID NO:7447), LRFN2 (SEQ ID NO:4711), CACNA1E (SEQ ID NO:214), CALY (SEQ ID NO:4119), SLC6A15 (SEQ ID NO:1932), KIAA0319 (SEQ ID NO:2227), SYT6 (SEQ ID NO:6957), PTPRR (SEQ ID NO:2724), KCTD8 (SEQ ID NO:7331), GPR22 (SEQ ID NO:3261), SLC4A8 (SEQ ID NO:818), LAMP5 (SEQ ID NO:3729), MEGF10 (SEQ ID NO:5355), FXYD7 (SEQ ID NO:4864), KCNK9 (SEQ ID NO:4257), SLC1A6 (SEQ ID NO:3206), MLC1 (SEQ ID NO:4042), OPRK1 (SEQ ID NO:325), ATP2B2 (SEQ ID NO:353), ACSL6 (SEQ ID NO:560), THBD (SEQ ID NO:108), PTPRT (SEQ ID NO:3622), PCDHGC4 (SEQ ID NO:2688), CLDN10 (SEQ ID NO:1975), KCNV1 (SEQ ID NO:3925), LPPR3 (SEQ ID NO:5096), SLCO1C1 (SEQ ID NO:1885), PCDH8 (SEQ ID NO:2690), ANO4 (SEQ ID NO:6566), LRRTM4 (SEQ ID NO:1505), PCDH15 (SEQ ID NO:1688), CCKBR (SEQ ID NO:6481), GABRA5 (SEQ ID NO:266), SLC6A12 (SEQ ID NO:1290), GRIN2A (SEQ ID NO:283), SLC1A2 (SEQ ID NO:2977), SLC43A1 (SEQ ID NO:2856), KCNC2 (SEQ ID NO:5836), ELFN2 (SEQ ID NO:5534), ATP7A (SEQ ID NO:10), GRIK4 (SEQ ID NO:3965), LRRC55 (SEQ ID NO:448), HCN2 (SEQ ID NO:7377), NKAIN1 (SEQ ID NO:5039), DPP10 (SEQ ID NO:432), AJAP1 (SEQ ID NO:894), NPFFR1 (SEQ ID NO:4886), TRPC3 (SEQ ID NO:1434), TGFBI (SEQ ID NO:107), SLC6A7 (SEQ ID NO:7370), GABRA4 (SEQ ID NO:265), SLC13A5 (SEQ ID NO:1732), GRIN2C (SEQ ID NO:285), HCN1 (SEQ ID NO:4763), SLC26A8 (SEQ ID NO:5552), PPIC (SEQ ID NO:336), NETO1 (SEQ ID NO:7427), TNFRSF10B (SEQ ID NO:2910), CDH22 (SEQ ID NO:4803), SLC6A13 (SEQ ID NO:4264), DISP2 (SEQ ID NO:7320), SLC6A11 (SEQ ID NO:3890), CD93 (SEQ ID NO:3698), EPHA10 (SEQ ID NO:1160), PHLDB2 (SEQ ID NO:1490), OXTR (SEQ ID NO:7326), WNT7A (SEQ ID NO:3089), GYPC (SEQ ID NO:7432), KCNA4 (SEQ ID NO:2589), PCDHAC2 (SEQ ID NO:4498), HGFAC (SEQ ID NO:2405), DRD1 (SEQ ID NO:253), SHISA9 (SEQ ID NO:7418), SCN8A (SEQ ID

NO:3879), ICAM1 (SEQ ID NO:7382), PIRT (SEQ ID NO:1201), A4GALT (SEQ ID NO:4291), MRGPRF (SEQ ID NO:1129), CD248 (SEQ ID NO:4649), CD58 (SEQ ID NO:7405), CD44 (SEQ ID NO:172), EPHA2 (SEQ ID NO:3042) or PROCR (SEQ ID NO:3480). In some instances, such antigen combinations may be useful in targeting melanoma.

[00335] Useful antigen combinations may include a MART1 (MLANA; SEQ ID NO:3302) clinical antigen AND NOT a tissue antigen (e.g., a brain tissue antigen) selected from: OPALIN (SEQ ID NO:833), TMEM235 (SEQ ID NO:1869), GABRA1 (SEQ ID NO:262), KCNJ9 (SEQ ID NO:3190), GRM3 (SEQ ID NO:290), SEZ6 (SEQ ID NO:1143), NTSR2 (SEQ ID NO:3753), KCNK4 (SEQ ID NO:5486), SLCO1A2 (SEQ ID NO:7290), SLC24A2 (SEQ ID NO:4632), MOG (SEQ ID NO:528), GABRG1 (SEQ ID NO:6354), GABRG2 (SEQ ID NO:272), CNTNAP4 (SEQ ID NO:5492), DSCAM (SEQ ID NO:2354), CACNG3 (SEQ ID NO:3510), CRB1 (SEQ ID NO:6891), CDH10 (SEQ ID NO:3552), HRH3 (SEQ ID NO:3657), GRIK1 (SEQ ID NO:280), SLC39A12 (SEQ ID NO:1826), GPR158 (SEQ ID NO:4716), CACNG2 (SEQ ID NO:3430), SYT3 (SEQ ID NO:7410), HTR5A (SEQ ID NO:4993), CACNG7 (SEQ ID NO:5264), GPR37L1 (SEQ ID NO:3118), LRRTM3 (SEQ ID NO:7362), GLRA2 (SEQ ID NO:1281), CHRNB2 (SEQ ID NO:228), KCNQ2 (SEQ ID NO:3067), JPH3 (SEQ ID NO:4690), GPR19 (SEQ ID NO:3446), ADCY8 (SEQ ID NO:1248), SPOCK3 (SEQ ID NO:845), SLC32A1 (SEQ ID NO:5606), OPCML (SEQ ID NO:594), GABRA3 (SEQ ID NO:264), GRM5 (SEQ ID NO:292), SCN1A (SEQ ID NO:2133), SLC5A11 (SEQ ID NO:5546), KCNC1 (SEQ ID NO:1249), SLC12A5 (SEQ ID NO:1506), GRM4 (SEQ ID NO:291), GRM1 (SEQ ID NO:288), GRIA4 (SEQ ID NO:279), MEGF11 (SEQ ID NO:5354), CACNA1B (SEQ ID NO:211), LYPD1 (SEQ ID NO:970), GRID2 (SEQ ID NO:2398), SCN2A (SEQ ID NO:842), NKAIN2 (SEQ ID NO:864), UNC5A (SEQ ID NO:5690), SLC4A10 (SEQ ID NO:7421), TMEFF2 (SEQ ID NO:4183), CSMD3 (SEQ ID NO:5532), PPAPDC1A (SEQ ID NO:715), HAPLN4 (SEQ ID NO:4968), GPR85 (SEQ ID NO:1920), ANTXR2 (SEQ ID NO:1876), CACNG4 (SEQ ID NO:3934), CSPG5 (SEQ ID NO:3516), KCNK10 (SEQ ID NO:4784), CHRNA4 (SEQ ID NO:225), CNTNAP2 (SEQ ID NO:3871), KCNJ10 (SEQ ID NO:2597), GABRB2 (SEQ ID NO:269), GRIN1 (SEQ ID NO:282), CRB2 (SEQ ID NO:6381), SHISA7 (SEQ ID NO:7406), NKAIN4 (SEQ ID NO:6122), HTR2C (SEQ ID NO:301), CACNG8 (SEQ ID NO:5263), NRG3 (SEQ ID NO:569), ABCG4 (SEQ ID NO:1668), CDH8 (SEQ ID NO:2476), GABRD (SEQ ID NO:271), KIRREL3 (SEQ ID NO:2022), GABRB1 (SEQ ID NO:268), KCNA2 (SEQ ID NO:3181), CDH20 (SEQ ID NO:5261), IGDCC3 (SEQ ID NO:3159), KCNJ6 (SEQ ID NO:2596), CNIH2 (SEQ ID NO:6658), KCNK12 (SEQ ID NO:7295), CDH18 (SEQ ID NO:2192), CSMD2 (SEQ ID NO:5530), SYT4 (SEQ ID NO:4722), OR2L13 (SEQ ID NO:6468), CDH9 (SEQ ID NO:4198), GABRA2 (SEQ ID NO:263), KCNF1 (SEQ ID NO:2592), MAG (SEQ ID NO:2634), CALN1 (SEQ ID NO:645), GRIN2B (SEQ ID NO:284), GRM7 (SEQ ID NO:294), VSTM2A (SEQ ID NO:7459), GPR61 (SEQ ID NO:5271), OMG (SEQ ID NO:2672), KCNA1 (SEQ ID NO:66), GPR83 (SEQ ID NO:4239), ATP8A2

(SEQ ID NO:4237), GABBR2 (SEQ ID NO:3288), GPR12 (SEQ ID NO:3255), TRPM3 (SEQ ID NO:519), SLC8A3 (SEQ ID NO:1424), KCND2 (SEQ ID NO:7288), GSG1L (SEQ ID NO:1236), SLC30A10 (SEQ ID NO:4487), ASTN1 (SEQ ID NO:3016), GPR179 (SEQ ID NO:7447), LRFN2 (SEQ ID NO:4711), CACNA1E (SEQ ID NO:214), CALY (SEQ ID NO:4119), SLC6A15 (SEQ ID NO:1932), KIAA0319 (SEQ ID NO:2227), SYT6 (SEQ ID NO:6957), PTPRR (SEQ ID NO:2724), KCTD8 (SEQ ID NO:7331), GPR22 (SEQ ID NO:3261), SLC4A8 (SEQ ID NO:818), LAMP5 (SEQ ID NO:3729), MEGF10 (SEQ ID NO:5355), FXYD7 (SEQ ID NO:4864), KCNK9 (SEQ ID NO:4257), SLC1A6 (SEQ ID NO:3206), MLC1 (SEQ ID NO:4042), OPRK1 (SEQ ID NO:325), ATP2B2 (SEQ ID NO:353), ACSL6 (SEQ ID NO:560), THBD (SEQ ID NO:108), PTPRT (SEQ ID NO:3622), PCDHGC4 (SEQ ID NO:2688), CLDN10 (SEQ ID NO:1975), KCNV1 (SEQ ID NO:3925), LPPR3 (SEQ ID NO:5096), SLCO1C1 (SEQ ID NO:1885), PCDH8 (SEQ ID NO:2690), ANO4 (SEQ ID NO:6566), LRRTM4 (SEQ ID NO:1505), PCDH15 (SEQ ID NO:1688), CCKBR (SEQ ID NO:6481), GABRA5 (SEQ ID NO:266), SLC6A12 (SEQ ID NO:1290), GRIN2A (SEQ ID NO:283), SLC1A2 (SEQ ID NO:2977), SLC43A1 (SEQ ID NO:2856), KCNC2 (SEQ ID NO:5836), ELFN2 (SEQ ID NO:5534), ATP7A (SEQ ID NO:10), GRIK4 (SEQ ID NO:3965), LRRC55 (SEQ ID NO:448), HCN2 (SEQ ID NO:7377), NKAIN1 (SEQ ID NO:5039), DPP10 (SEQ ID NO:432), AJAP1 (SEQ ID NO:894), NPFFR1 (SEQ ID NO:4886), TRPC3 (SEQ ID NO:1434), TGFBI (SEQ ID NO:107), SLC6A7 (SEQ ID NO:7370), GABRA4 (SEQ ID NO:265), SLC13A5 (SEQ ID NO:1732), GRIN2C (SEQ ID NO:285), HCN1 (SEQ ID NO:4763), SLC26A8 (SEQ ID NO:5552), PPIC (SEQ ID NO:336), NETO1 (SEQ ID NO:7427), TNFRSF10B (SEQ ID NO:2910), CDH22 (SEQ ID NO:4803), SLC6A13 (SEQ ID NO:4264), DISP2 (SEQ ID NO:7320), SLC6A11 (SEQ ID NO:3890), CD93 (SEQ ID NO:3698), EPHA10 (SEQ ID NO:1160), PHLDB2 (SEQ ID NO:1490), OXTR (SEQ ID NO:7326), WNT7A (SEQ ID NO:3089), GYPC (SEQ ID NO:7432), KCNA4 (SEQ ID NO:2589), PCDHAC2 (SEQ ID NO:4498), HGFAC (SEQ ID NO:2405), DRD1 (SEQ ID NO:253), SHISA9 (SEQ ID NO:7418), SCN8A (SEQ ID NO:3879), ICAM1 (SEQ ID NO:7382), PIRT (SEQ ID NO:1201), A4GALT (SEQ ID NO:4291), MRGPRF (SEQ ID NO:1129), CD248 (SEQ ID NO:4649), CD58 (SEQ ID NO:7405), CD44 (SEQ ID NO:172), EPHA2 (SEQ ID NO:3042) or PROCR (SEQ ID NO:3480). In some instances, such antigen combinations may be useful in targeting melanoma.

Third antigens

[00336] As noted above, in some cases, a genetically modified cytotoxic immune cell of the present disclosure, or a system of the present disclosure, can include an antigen-triggered polypeptide (or a nucleic acid comprising a nucleotide sequence encoding the same) that specifically binds a third target antigen present on the surface of a cancer cell. Examples of such third target antigens are depicted in FIG. 4 and Table 3. The third target antigen can be included except where contraindicated by a target antigen pair listed in FIG. 1 or FIG. 9-14. In some cases, the third antigen comprises an amino acid

sequence having at least 85%, at least 90%, at least 95%, at least 98%, at least 99%, or 100%, amino acid sequence identity to one of the amino acid sequences depicted in SEQ ID NOs: 7119-7467.

[00337] Where a third antigen (e.g., an antigen listed in FIG. 4 or Table 3) is included in the target antigen combination, the third antigen will be associated with the same cancer cell type as the target antigen pair (e.g., from FIG. 1 or FIG. 9-14). FIG. 1 and FIG. 9-14 provide exemplary target cancer cell types associate with a given target antigen pair. FIG. 4 also provides exemplary target cancer cell types associate with a given target antigen.

[00338] In some instances, a particular antigen (e.g., an antigen of an antigen pair as described in FIG. 1 or FIG. 9-14, a clinical antigen as described in FIG. 4, etc.) may be employed in an antigen combination, including combinations having two antigens, combinations having three antigens, etc., for treating a cancer other than the exemplary cancer with which it is identified. For example, although FIG. 4 identifies "exemplary cancers" for each antigen, use of the antigen in antigen combinations will not be limited to the specifically identified cancer(s) and the particular antigen may be employed in antigen combinations for the treatment of other cancers besides those specifically listed as exemplary.

[00339] Examples of hematological malignancy antigens include, e.g., CD19 (as expressed in e.g., B-cells), CD20 (as expressed in e.g., B-cells), CD22 (as expressed in e.g., B-cells), CD30 (as expressed in e.g., B-cells), CD33 (as expressed in e.g., Myeloid cells), CD70 (as expressed in e.g., B-cell/T-cells), CD123 (as expressed in e.g., Myeloid cells), Kappa (as expressed in e.g., B-cells), Lewis Y (as expressed in e.g., Myeloid cells), NKG2D ligands (as expressed in e.g., Myeloid cells), ROR1 (as expressed in e.g., B-cells), SLAMF7/CS1 (as expressed in e.g., myeloma cells, natural killer cells, T cells, and most B-cell types), CD138 (as expressed in e.g., malignant plasma cells in multiple myelomas), CD56 (as expressed in e.g., myeloma cells, neural cells, natural killer cells, T cells, and trabecular osteoblasts) CD38 (as expressed in e.g., B-cell/T-cells) and CD160 (as expressed in e.g., NK cells/T-cells), and the like. Examples of solid tumor antigens include, e.g., B7H3 (as expressed in e.g., Sarcoma, glioma), CAIX (as expressed in e.g., Kidney), CD44 v6/v7 (as expressed in e.g., Cervical cancer), CD171 (as expressed in e.g., Neuroblastoma), CEA (as expressed in e.g., Colon), EGFRvIII (as expressed in e.g., Glioma), EGP2 (as expressed in e.g., Carcinomas), EGP40 (as expressed in e.g., Colon), EphA2 (as expressed in e.g., Glioma, lung), ErbB2(HER2) (as expressed in e.g., Breast, lung, prostate, glioma), ErbB receptor family (as expressed in e.g., Breast, lung, prostate, glioma), ErbB3/4 (as expressed in e.g., Breast, ovarian), HLA-A1/MAGE1 (as expressed in e.g., Melanoma), HLA-A2/NY-ESO-1 (as expressed in e.g., Sarcoma, melanoma), FR-a (as expressed in e.g., Ovarian), FAP† (as expressed in e.g., Cancer associated fibroblasts), FAR (as expressed in e.g., Rhabdomyosarcoma), GD2 (as expressed in e.g., Neuroblastoma, sarcoma, melanoma), GD3 (as expressed in e.g., Melanoma, lung cancer), HMW-MAA (as expressed in e.g., Melanoma), IL11Ra (as expressed in e.g., Osteosarcoma), IL13Ra2 (as expressed in e.g., Glioma), Lewis Y (as expressed in e.g., Breast/ovarian/pancreatic), Mesothelin (as expressed in

e.g., Mesothelioma, breast, pancreas), Muc1 (as expressed in e.g., Ovarian, breast, prostate), NCAM (as expressed in e.g., Neuroblastoma, colorectal), NKG2D ligands (as expressed in e.g., Ovarian, sacoma), PSCA (as expressed in e.g., Prostate, pancreatic), PSMA (as expressed in e.g., Prostate), TAG72 (as expressed in e.g., Colon), VEGFR-2 (as expressed in e.g., Tumor vasculature), Axl (as expressed in e.g., Lung cancer), Met (as expressed in e.g., Lung cancer), α5β3 (as expressed in e.g., Tumor vasculature), α5β1 (as expressed in e.g., Tumor vasculature), TRAIL-R1/TRAIL-R2 (as expressed in e.g., Solid tumors (colon, lung, pancreas) and hematological malignancies), RANKL (as expressed in e.g., Prostate cancer and bone metastases), Tenacin (as expressed in e.g., Glioma, epithelial tumors (breast, prostate)), EpCAM (as expressed in e.g., Epithelial tumors (breast, colon, lung)), CEA (as expressed in e.g., Epithelial tumors (breast, colon, lung)), gpA33 (as expressed in e.g., Colorectal carcinoma), Mucins (as expressed in e.g., Epithelial tumors (breast, colon, lung, ovarian)), TAG-72 (as expressed in e.g., Epithelial tumors (breast, colon, lung)), EphA3 (as expressed in e.g., Lung, kidney, melanoma, glioma, hematological malignancies) and IGF1R (as expressed in e.g., Lung, breast, head and neck, prostate, thyroid, glioma). Examples of surface and intracellular antigens include, e.g., Her2 (gene symbol ERBB2), MAGE-A1 (gene symbol MAGEA1), MART-1 (gene symbol MLANA), NY-ESO (gene symbol CTAG1), WT1 (gene symbol WT1), MUC17 and MUC13. Examples of other antigens include, e.g., BCMA (gene symbol TNFRSF17), B7H6 (gene symbol NCR3LG1), CAIX (gene symbol CA9), CD123 (gene symbol IL3RA), CD138 (gene symbol SDC1), CD171 (gene symbol L1CAM), CD19 (gene symbol CD19), CD20 (gene symbol CD20), CD22 (gene symbol CD22), CD30 (gene symbol TNFRSF8), CD33 (gene symbol CD33), CD38 (gene symbol CD38), CD44, splice variants incl 7 and 8 (denoted vX in literature) (gene symbol CD44), CEA, CS1 (gene symbol SLAMF7), EGFRvIII (gene symbol EGFR, vIII deletion variant), EGP2, EGP40 (gene symbol EPCAM), Erb family member (gene symbol ERBB1, ERBB2, ERBB3, ERBB4), FAP (gene symbol FAP), fetal acetylcholine receptor (gene symbol AChR), Folate receptor alpha (gene symbol FOLR1), Folate receptor beta (gene symbol FOLR2), GD2, GD3, GPC3 (gene symbol GPC3), Her2/neu (gene symbol ERBB2), IL-13Ra2 (gene symbol IL13RA2), Kappa light chain (gene symbol IGK), Lewis-Y, Mesothelin (gene symbol MSLN), Mucin-1 (gene symbol MUC1), Mucin-16 (gene symbol MUC16), NKG2D ligands, prostate specific membrane antigen (PSMA) (gene symbol FOLH1), prostate stem cell antigen (PSCA) (gene symbol PSCA), receptor tyrosine kinase-like orphan receptor 1 (gene symbol ROR1), and Anaplastic Lymphoma Receptor Tyrosine Kinase (gene symbol ALK).

[00340] In some cases, useful third antigens may include an antigen selected from Table 3 above.

[00341] Useful three antigen combinations may or may not include a clinical antigen as an added "third antigen". For example, in some instances, a useful three antigen combination may include a clinical antigen and two or more antigens that provide AND NOT functionality, including e.g., the combinations including a clinical antigen and two tissue specific AND NOT antigens (e.g., brain tissue and cardiac

tissue) as described in more detail below. In some cases, useful three antigen combinations may not include a clinical antigen. Useful three antigen combinations may include various logic combinations including but not limited to e.g., antigen 1 AND antigen 2 AND antigen 3, antigen 1 AND antigen 2 AND NOT antigen 3, antigen 1 AND NOT antigen 2 AND NOT antigen 3, and the like. In some instances, the logic of a three antigen combination may be complex where "complex" logic, as used herein, refers to combinations having multiple positive prediction nodes in the associated tree or, e.g., where the logic contains one or more OR propositions. In some instances, useful three antigen combinations include those depicted in FIG. 15.

[00342] In some cases, useful three antigen combinations include an antigen combination that includes a clinical antigen in combination with two or more tissue antigens (e.g., a first antigen expressed in normal tissue, e.g., normal brain tissue, and a second antigen expressed in normal tissue, e.g., normal cardiac tissue) in a double AND NOT gate (i.e., clinical antigen AND NOT normal tissue antigen 1 AND NOT normal tissue antigen 2).

[00343] For example, useful antigen combinations may include a MAGEA1 (SEQ ID NO:3191) clinical antigen AND NOT a cardiac tissue specific antigen (such as GJA3 (SEQ ID NO:4854), HCN4 (SEQ ID NO:3292) or BMP10 (SEQ ID NO:3945)) AND NOT a brain tissue antigen selected from: OPALIN (SEQ ID NO:833), TMEM235 (SEQ ID NO:1869), GABRA1 (SEQ ID NO:262), KCNJ9 (SEQ ID NO:3190), GRM3 (SEQ ID NO:290), SEZ6 (SEQ ID NO:1143), NTSR2 (SEQ ID NO:3753), KCNK4 (SEQ ID NO:5486), SLCO1A2 (SEQ ID NO:7290), SLC24A2 (SEQ ID NO:4632), MOG (SEQ ID NO:528), GABRG1 (SEQ ID NO:6354), GABRG2 (SEQ ID NO:272), CNTNAP4 (SEQ ID NO:5492), DSCAM (SEQ ID NO:2354), CACNG3 (SEQ ID NO:3510), CRB1 (SEQ ID NO:6891), CDH10 (SEQ ID NO:3552), HRH3 (SEQ ID NO:3657), GRIK1 (SEQ ID NO:280), SLC39A12 (SEQ ID NO:1826), GPR158 (SEQ ID NO:4716), CACNG2 (SEQ ID NO:3430), SYT3 (SEQ ID NO:7410), HTR5A (SEQ ID NO:4993), CACNG7 (SEQ ID NO:5264), GPR37L1 (SEQ ID NO:3118), LRRTM3 (SEQ ID NO:7362), GLRA2 (SEQ ID NO:1281), CHRNB2 (SEQ ID NO:228), KCNQ2 (SEQ ID NO:3067), JPH3 (SEQ ID NO:4690), GPR19 (SEQ ID NO:3446), ADCY8 (SEQ ID NO:1248), SPOCK3 (SEQ ID NO:845), SLC32A1 (SEQ ID NO:5606), OPCML (SEQ ID NO:594), GABRA3 (SEQ ID NO:264), GRM5 (SEQ ID NO:292), SCN1A (SEQ ID NO:2133), SLC5A11 (SEQ ID NO:5546), KCNC1 (SEQ ID NO:1249), SLC12A5 (SEQ ID NO:1506), GRM4 (SEQ ID NO:291), GRM1 (SEQ ID NO:288), GRIA4 (SEQ ID NO:279), MEGF11 (SEQ ID NO:5354), CACNA1B (SEQ ID NO:211), LYPD1 (SEQ ID NO:970), GRID2 (SEQ ID NO:2398), SCN2A (SEQ ID NO:842), NKAIN2 (SEQ ID NO:864), UNC5A (SEQ ID NO:5690), SLC4A10 (SEQ ID NO:7421), TMEFF2 (SEQ ID NO:4183), CSMD3 (SEQ ID NO:5532), PPAPDC1A (SEQ ID NO:715), HAPLN4 (SEQ ID NO:4968), GPR85 (SEQ ID NO:1920), ANTXR2 (SEQ ID NO:1876), CACNG4 (SEQ ID NO:3934), CSPG5 (SEQ ID NO:3516), KCNK10 (SEQ ID NO:4784), CHRNA4 (SEQ ID NO:225), CNTNAP2 (SEQ ID NO:3871), KCNJ10

(SEQ ID NO:2597), GABRB2 (SEQ ID NO:269), GRIN1 (SEQ ID NO:282), CRB2 (SEQ ID NO:6381), SHISA7 (SEQ ID NO:7406), NKAIN4 (SEQ ID NO:6122), HTR2C (SEQ ID NO:301), CACNG8 (SEQ ID NO:5263), NRG3 (SEQ ID NO:569), ABCG4 (SEQ ID NO:1668), CDH8 (SEQ ID NO:2476), GABRD (SEQ ID NO:271), KIRREL3 (SEQ ID NO:2022), GABRB1 (SEQ ID NO:268), KCNA2 (SEQ ID NO:3181), CDH20 (SEQ ID NO:5261), IGDCC3 (SEQ ID NO:3159), KCNJ6 (SEQ ID NO:2596), CNIH2 (SEQ ID NO:6658), KCNK12 (SEQ ID NO:7295), CDH18 (SEQ ID NO:2192), CSMD2 (SEQ ID NO:5530), SYT4 (SEQ ID NO:4722), OR2L13 (SEQ ID NO:6468), CDH9 (SEQ ID NO:4198), GABRA2 (SEQ ID NO:263), KCNF1 (SEQ ID NO:2592), MAG (SEQ ID NO:2634), CALN1 (SEQ ID NO:645), GRIN2B (SEQ ID NO:284), GRM7 (SEQ ID NO:294), VSTM2A (SEQ ID NO:7459), GPR61 (SEQ ID NO:5271), OMG (SEQ ID NO:2672), KCNA1 (SEQ ID NO:66), GPR83 (SEQ ID NO:4239), ATP8A2 (SEQ ID NO:4237), GABBR2 (SEQ ID NO:3288), GPR12 (SEQ ID NO:3255), TRPM3 (SEQ ID NO:519), SLC8A3 (SEQ ID NO:1424), KCND2 (SEQ ID NO:7288), GSG1L (SEQ ID NO:1236), SLC30A10 (SEQ ID NO:4487), ASTN1 (SEQ ID NO:3016), GPR179 (SEQ ID NO:7447), LRFN2 (SEQ ID NO:4711), CACNA1E (SEQ ID NO:214), CALY (SEQ ID NO:4119), SLC6A15 (SEQ ID NO:1932), KIAA0319 (SEQ ID NO:2227), SYT6 (SEQ ID NO:6957), PTPRR (SEQ ID NO:2724), KCTD8 (SEQ ID NO:7331), GPR22 (SEQ ID NO:3261), SLC4A8 (SEQ ID NO:818), LAMP5 (SEQ ID NO:3729), MEGF10 (SEQ ID NO:5355), FXYD7 (SEQ ID NO:4864), KCNK9 (SEQ ID NO:4257), SLC1A6 (SEQ ID NO:3206), MLC1 (SEQ ID NO:4042), OPRK1 (SEQ ID NO:325), ATP2B2 (SEQ ID NO:353), ACSL6 (SEQ ID NO:560), THBD (SEQ ID NO:108), PTPRT (SEQ ID NO:3622), PCDHGC4 (SEQ ID NO:2688), CLDN10 (SEQ ID NO:1975), KCNV1 (SEQ ID NO:3925), LPPR3 (SEQ ID NO:5096), SLCO1C1 (SEQ ID NO:1885), PCDH8 (SEQ ID NO:2690), ANO4 (SEQ ID NO:6566), LRRTM4 (SEQ ID NO:1505), PCDH15 (SEQ ID NO:1688), CCKBR (SEQ ID NO:6481), GABRA5 (SEQ ID NO:266), SLC6A12 (SEQ ID NO:1290), GRIN2A (SEQ ID NO:283), SLC1A2 (SEQ ID NO:2977), SLC43A1 (SEQ ID NO:2856), KCNC2 (SEQ ID NO:5836), ELFN2 (SEQ ID NO:5534), ATP7A (SEQ ID NO:10), GRIK4 (SEQ ID NO:3965), LRRC55 (SEQ ID NO:448), HCN2 (SEQ ID NO:7377), NKAIN1 (SEQ ID NO:5039), DPP10 (SEQ ID NO:432), AJAP1 (SEQ ID NO:894), NPFFR1 (SEQ ID NO:4886), TRPC3 (SEQ ID NO:1434), TGFBI (SEQ ID NO:107), SLC6A7 (SEQ ID NO:7370), GABRA4 (SEQ ID NO:265), SLC13A5 (SEQ ID NO:1732), GRIN2C (SEQ ID NO:285), HCN1 (SEQ ID NO:4763), SLC26A8 (SEQ ID NO:5552), PPIC (SEQ ID NO:336), NETO1 (SEQ ID NO:7427), TNFRSF10B (SEQ ID NO:2910), CDH22 (SEQ ID NO:4803), SLC6A13 (SEQ ID NO:4264), DISP2 (SEQ ID NO:7320), SLC6A11 (SEQ ID NO:3890), CD93 (SEQ ID NO:3698), EPHA10 (SEQ ID NO:1160), PHLDB2 (SEQ ID NO:1490), OXTR (SEQ ID NO:7326), WNT7A (SEQ ID NO:3089), GYPC (SEQ ID NO:7432), KCNA4 (SEQ ID NO:2589), PCDHAC2 (SEQ ID NO:4498), HGFAC (SEQ ID NO:2405), DRD1 (SEQ ID NO:253), SHISA9 (SEQ ID NO:7418), SCN8A (SEQ ID NO:3879), ICAM1 (SEQ ID NO:7382), PIRT (SEQ ID NO:1201),

A4GALT (SEQ ID NO:4291), MRGPRF (SEQ ID NO:1129), CD248 (SEQ ID NO:4649), CD58 (SEQ ID NO:7405), CD44 (SEQ ID NO:172), EPHA2 (SEQ ID NO:3042) or PROCR (SEQ ID NO:3480). In some instances, such antigen combinations may be useful in targeting melanoma.

[00344] Useful antigen combinations may include a MAGEA3 (SEQ ID NO:7279) clinical antigen AND NOT a cardiac tissue specific antigen (such as GJA3 (SEQ ID NO:4854), HCN4 (SEQ ID NO:3292) or BMP10 (SEQ ID NO:3945)) AND NOT a brain tissue antigen selected from: OPALIN (SEQ ID NO:833), TMEM235 (SEQ ID NO:1869), GABRA1 (SEQ ID NO:262), KCNJ9 (SEQ ID NO:3190), GRM3 (SEQ ID NO:290), SEZ6 (SEQ ID NO:1143), NTSR2 (SEQ ID NO:3753), KCNK4 (SEQ ID NO:5486), SLCO1A2 (SEQ ID NO:7290), SLC24A2 (SEQ ID NO:4632), MOG (SEQ ID NO:528), GABRG1 (SEQ ID NO:6354), GABRG2 (SEQ ID NO:272), CNTNAP4 (SEQ ID NO:5492), DSCAM (SEQ ID NO:2354), CACNG3 (SEQ ID NO:3510), CRB1 (SEQ ID NO:6891), CDH10 (SEQ ID NO:3552), HRH3 (SEQ ID NO:3657), GRIK1 (SEQ ID NO:280), SLC39A12 (SEQ ID NO:1826), GPR158 (SEQ ID NO:4716), CACNG2 (SEQ ID NO:3430), SYT3 (SEQ ID NO:7410), HTR5A (SEQ ID NO:4993), CACNG7 (SEQ ID NO:5264), GPR37L1 (SEQ ID NO:3118), LRRTM3 (SEQ ID NO:7362), GLRA2 (SEQ ID NO:1281), CHRNB2 (SEQ ID NO:228), KCNQ2 (SEQ ID NO:3067), JPH3 (SEQ ID NO:4690), GPR19 (SEQ ID NO:3446), ADCY8 (SEQ ID NO:1248), SPOCK3 (SEQ ID NO:845), SLC32A1 (SEQ ID NO:5606), OPCML (SEQ ID NO:594), GABRA3 (SEQ ID NO:264), GRM5 (SEQ ID NO:292), SCN1A (SEQ ID NO:2133), SLC5A11 (SEQ ID NO:5546), KCNC1 (SEQ ID NO:1249), SLC12A5 (SEQ ID NO:1506), GRM4 (SEQ ID NO:291), GRM1 (SEQ ID NO:288), GRIA4 (SEQ ID NO:279), MEGF11 (SEQ ID NO:5354), CACNA1B (SEQ ID NO:211), LYPD1 (SEQ ID NO:970), GRID2 (SEQ ID NO:2398), SCN2A (SEQ ID NO:842), NKAIN2 (SEQ ID NO:864), UNC5A (SEQ ID NO:5690), SLC4A10 (SEQ ID NO:7421), TMEFF2 (SEQ ID NO:4183), CSMD3 (SEQ ID NO:5532), PPAPDC1A (SEQ ID NO:715), HAPLN4 (SEQ ID NO:4968), GPR85 (SEQ ID NO:1920), ANTXR2 (SEQ ID NO:1876), CACNG4 (SEQ ID NO:3934), CSPG5 (SEQ ID NO:3516), KCNK10 (SEQ ID NO:4784), CHRNA4 (SEQ ID NO:225), CNTNAP2 (SEQ ID NO:3871), KCNJ10 (SEQ ID NO:2597), GABRB2 (SEQ ID NO:269), GRIN1 (SEQ ID NO:282), CRB2 (SEQ ID NO:6381), SHISA7 (SEQ ID NO:7406), NKAIN4 (SEQ ID NO:6122), HTR2C (SEQ ID NO:301), CACNG8 (SEQ ID NO:5263), NRG3 (SEQ ID NO:569), ABCG4 (SEQ ID NO:1668), CDH8 (SEQ ID NO:2476), GABRD (SEQ ID NO:271), KIRREL3 (SEQ ID NO:2022), GABRB1 (SEQ ID NO:268), KCNA2 (SEQ ID NO:3181), CDH20 (SEQ ID NO:5261), IGDCC3 (SEQ ID NO:3159), KCNJ6 (SEQ ID NO:2596), CNIH2 (SEQ ID NO:6658), KCNK12 (SEQ ID NO:7295), CDH18 (SEQ ID NO:2192), CSMD2 (SEQ ID NO:5530), SYT4 (SEQ ID NO:4722), OR2L13 (SEQ ID NO:6468), CDH9 (SEQ ID NO:4198), GABRA2 (SEQ ID NO:263), KCNF1 (SEQ ID NO:2592), MAG (SEQ ID NO:2634), CALN1 (SEQ ID NO:645), GRIN2B (SEQ ID NO:284), GRM7 (SEQ ID NO:294), VSTM2A (SEQ ID NO:7459), GPR61 (SEQ ID NO:5271), OMG (SEQ ID NO:2672), KCNA1 (SEQ ID NO:66), GPR83

(SEQ ID NO:4239), ATP8A2 (SEQ ID NO:4237), GABBR2 (SEQ ID NO:3288), GPR12 (SEQ ID NO:3255), TRPM3 (SEQ ID NO:519), SLC8A3 (SEQ ID NO:1424), KCND2 (SEQ ID NO:7288), GSG1L (SEQ ID NO:1236), SLC30A10 (SEQ ID NO:4487), ASTN1 (SEQ ID NO:3016), GPR179 (SEQ ID NO:7447), LRFN2 (SEQ ID NO:4711), CACNA1E (SEQ ID NO:214), CALY (SEQ ID NO:4119), SLC6A15 (SEQ ID NO:1932), KIAA0319 (SEQ ID NO:2227), SYT6 (SEQ ID NO:6957), PTPRR (SEQ ID NO:2724), KCTD8 (SEQ ID NO:7331), GPR22 (SEQ ID NO:3261), SLC4A8 (SEQ ID NO:818), LAMP5 (SEQ ID NO:3729), MEGF10 (SEQ ID NO:5355), FXYD7 (SEQ ID NO:4864), KCNK9 (SEQ ID NO:4257), SLC1A6 (SEQ ID NO:3206), MLC1 (SEQ ID NO:4042), OPRK1 (SEQ ID NO:325), ATP2B2 (SEQ ID NO:353), ACSL6 (SEQ ID NO:560), THBD (SEQ ID NO:108), PTPRT (SEQ ID NO:3622), PCDHGC4 (SEQ ID NO:2688), CLDN10 (SEQ ID NO:1975), KCNV1 (SEQ ID NO:3925), LPPR3 (SEQ ID NO:5096), SLCO1C1 (SEQ ID NO:1885), PCDH8 (SEQ ID NO:2690), ANO4 (SEQ ID NO:6566), LRRTM4 (SEQ ID NO:1505), PCDH15 (SEQ ID NO:1688), CCKBR (SEQ ID NO:6481), GABRA5 (SEQ ID NO:266), SLC6A12 (SEQ ID NO:1290), GRIN2A (SEQ ID NO:283), SLC1A2 (SEQ ID NO:2977), SLC43A1 (SEQ ID NO:2856), KCNC2 (SEQ ID NO:5836), ELFN2 (SEQ ID NO:5534), ATP7A (SEQ ID NO:10), GRIK4 (SEQ ID NO:3965), LRRC55 (SEQ ID NO:448), HCN2 (SEQ ID NO:7377), NKAIN1 (SEQ ID NO:5039), DPP10 (SEQ ID NO:432), AJAP1 (SEQ ID NO:894), NPFFR1 (SEQ ID NO:4886), TRPC3 (SEQ ID NO:1434), TGFBI (SEQ ID NO:107), SLC6A7 (SEQ ID NO:7370), GABRA4 (SEQ ID NO:265), SLC13A5 (SEQ ID NO:1732), GRIN2C (SEQ ID NO:285), HCN1 (SEQ ID NO:4763), SLC26A8 (SEQ ID NO:5552), PPIC (SEQ ID NO:336), NETO1 (SEQ ID NO:7427), TNFRSF10B (SEQ ID NO:2910), CDH22 (SEQ ID NO:4803), SLC6A13 (SEQ ID NO:4264), DISP2 (SEQ ID NO:7320), SLC6A11 (SEQ ID NO:3890), CD93 (SEQ ID NO:3698), EPHA10 (SEQ ID NO:1160), PHLDB2 (SEQ ID NO:1490), OXTR (SEQ ID NO:7326), WNT7A (SEQ ID NO:3089), GYPC (SEQ ID NO:7432), KCNA4 (SEQ ID NO:2589), PCDHAC2 (SEQ ID NO:4498), HGFAC (SEQ ID NO:2405), DRD1 (SEQ ID NO:253), SHISA9 (SEQ ID NO:7418), SCN8A (SEQ ID NO:3879), ICAM1 (SEQ ID NO:7382), PIRT (SEQ ID NO:1201), A4GALT (SEQ ID NO:4291), MRGPRF (SEQ ID NO:1129), CD248 (SEQ ID NO:4649), CD58 (SEQ ID NO:7405), CD44 (SEQ ID NO:172), EPHA2 (SEQ ID NO:3042) or PROCR (SEQ ID NO:3480). In some instances, such antigen combinations may be useful in targeting melanoma.

Methods of Killing Target Cancer Cells

[00345] The present disclosure provides methods for killing a target cancer cell. The present disclosure provides a method of killing a target cancer cell in an individual. In some cases, a method of the present disclosure for killing a target cell in an individual comprises: a) introducing a system of the present disclosure into an immune cell (e.g., a $CD8^+$ T cell; an NK cell) obtained from the individual, generating a modified immune cell; and b) administering the modified immune cell to the individual, where the

modified immune cell kills the target cancer cell in the individual. In some cases, the modified cytotoxic T cell does not substantially kill non-target cells such as non-cancerous cells.

[00346] Example 1 provides an example of a method of killing a liposarcoma in an individual; and Example 2 provides an example of a method of killing a glioblastoma in an individual. This example is illustrative of how a $CD8^+$ cell (or other immune cell) genetically modified to express first and second antigen-triggered polypeptides targeting any of the target antigen pairs depicted in FIG. 1 or FIG. 9-14 to target the corresponding cancer cells listed in FIG. 1 or FIG. 9-14, can be used to kill a target cancer cell in an individual, and thereby to treat the cancer in the individual.

[00347] The present disclosure provides a method of killing a target cancer cell in an individual. In some cases, a method of the present disclosure for killing a target cell in an individual comprises administering a genetically modified cytotoxic immune cell (e.g., a genetically modified $CD8^+$ T cell; a genetically modified NK cell) of the present disclosure to the individual, where the genetically modified immune cell kills the target cancer cell in the individual. In some cases, the modified cytotoxic T cell does not substantially kill non-target cells such as non-cancerous cells.

[00348] Where the target antigen pair targeted by a method of the present disclosure is an AND-NOT gate target antigen pair, a method of the present disclosure provides for killing of a target cancer cell, but not a non-cancerous cell. For example, in some cases, a method of the present disclosure provides for a ratio of killing of cancer cells to non-cancerous cells of at least 10:1, at least 15:1, at least 20:1 at least 25:1, at least 50:1, at least 100:1, at least 500:1, at least 10^3:1, at least 10^4:1, or at least 10^5:1.

[00349] Where the target antigen pair targeted by a method of the present disclosure is an AND gate target antigen pair, a method of the present disclosure provides for highly specific killing of a target cancer cell, and not a non-target (e.g., non-cancerous cell). For example, in some cases, a method of the present disclosure provides for a ratio of killing of cancer cells to non-cancerous cells of at least 10:1, at least 15:1, at least 20:1 at least 25:1, at least 50:1, at least 100:1, at least 500:1, at least 10^3:1, at least 10^4:1, or at least 10^5:1.

Methods comprising use of a system of the present disclosure

[00350] As noted above, in some cases, a method of the present disclosure for killing a target cell in an individual comprises: a) introducing a system of the present disclosure into an immune cell (e.g., a $CD8^+$ T cell; an NK cell) obtained from the individual, generating a modified immune cell; and b) administering the modified immune cell to the individual, where the modified immune cell kills the target cancer cell in the individual. In some cases, the modified cytotoxic T cell does not substantially kill non-target cells such as non-cancerous cells.

[00351] T cells can be obtained from an individual (e.g., an individual having a cancer; an individual diagnosed as having a cancer; an individual being treated for a cancer with chemotherapy, radiation therapy, antibody therapy, surgery, etc.) using well-established methods. In some cases, a mixed

population of cells is obtained from an individual; and $CD8^+$ T cells and/or NK cells are isolated from the mixed population, such that a population of $CD8^+$ T cells and/or NK cells is obtained that is at least 75%, at least 80%, at least 85%, at least 90%, at least 95%, or more than 98% pure, i.e., the purified cell population includes less than 25%, less than 20%, less than 15%, less than 10%, less than 5%, or less than 2%, of cells other than $CD8^+$ T cells and or NK cells. A system of the present disclosure is then introduced into the purified $CD8^+$ T cells and/or NK cells, to generate modified $CD8^+$ T cells and/or modified NK cells that express the first antigen-triggered polypeptide and the second antigen-triggered polypeptide.

[00352] In some cases, as noted above, a system of the present disclosure comprises: a) a first antigen-triggered polypeptide that binds specifically to a first target antigen present on a target cancer cell; and b) a second antigen-triggered polypeptide that binds specifically to a second target antigen. In these instances, the polypeptides per se are introduced into an immune cell (e.g., $CD8^+$ T cells and/or NK cells obtained from an individual). Methods of introducing polypeptides into a cell are known in the art; and any known method can be used. For example, in some cases, the first and the second antigen-triggered polypeptides comprise a protein transduction domain (PTD) at the N-terminus or the C-terminus of the polypeptides.

[00353] In some cases, as noted above, a system of the present disclosure comprises: a) a first nucleic acid comprising a nucleotide sequence encoding a first antigen-triggered polypeptide that binds specifically to a first target antigen present on a target cancer cell; and b) a second nucleic acid comprising a nucleotide sequence encoding a second antigen-triggered polypeptide that binds specifically to a second target antigen. In some cases, the first and the second antigen-triggered polypeptides are encoded by nucleotide sequences on separate nucleic acids. In other cases, the first and the second antigen-triggered polypeptides are encoded by nucleotide sequences present in the same nucleic acid. In some cases, the nucleic acid is a recombinant expression vector. In some cases, a system of the present disclosure comprises: a) a first recombinant expression vector comprising a nucleotide sequence encoding a first antigen-triggered polypeptide that binds specifically to a first target antigen present on a target cancer cell; and b) a second recombinant expression vector comprising a nucleotide sequence encoding a second antigen-triggered polypeptide that binds specifically to a second target antigen. In some cases, the nucleotide sequences are operably linked to a constitutive promoter. In some cases, the nucleotide sequences are operably linked to a regulatable promoter (e.g., an inducible promoter, a reversible promoter, etc.). In some cases, the nucleotide sequences are operably linked to an immune cell promoter, e.g., a T-cell specific promoter. In some cases, a system of the present disclosure comprises a recombinant expression vector comprising nucleotide sequences encoding: a) a first antigen-triggered polypeptide that binds specifically to a first target antigen present on a target cancer cell; and b) a second antigen-triggered polypeptide that binds specifically to a second target antigen. In some cases, the

nucleotide sequences are operably linked to a constitutive promoter. In some cases, the nucleotide sequences are operably linked to a regulatable promoter (e.g., an inducible promoter, a reversible promoter, etc.). In some cases, the nucleotide sequences are operably linked to an immune cell promoter, e.g., a T-cell specific promoter. Suitable promoters include, but are not limited to; cytomegalovirus immediate early promoter; herpes simplex virus thymidine kinase promoter; early and late SV40 promoters; promoter present in long terminal repeats from a retrovirus; a metallothionein-I promoter; and various art-known promoters. Such reversible promoters, and systems based on such reversible promoters but also comprising additional control proteins, include, but are not limited to, alcohol regulated promoters (e.g., alcohol dehydrogenase I (alcA) gene promoter, promoters responsive to alcohol transactivator proteins (AlcR), etc.), tetracycline regulated promoters, (e.g., promoter systems including TetActivators, TetON, TetOFF, etc.), steroid regulated promoters (e.g., rat glucocorticoid receptor promoter systems, human estrogen receptor promoter systems, retinoid promoter systems, thyroid promoter systems, ecdysone promoter systems, mifepristone promoter systems, etc.), metal regulated promoters (e.g., metallothionein promoter systems, etc.), pathogenesis-related regulated promoters (e.g., salicylic acid regulated promoters, ethylene regulated promoters, benzothiadiazole regulated promoters, etc.), temperature regulated promoters (e.g., heat shock inducible promoters (e.g., HSP-70, HSP-90, soybean heat shock promoter, etc.), light regulated promoters, synthetic inducible promoters, and the like.

[00354] In some instances, nucleic acids present in a system of the present disclosure include immune cell specific promoters that are expressed in one or more immune cell types, including but not limited to lymphocytes, hematopoietic stem cells and/or progeny thereof (i.e., immune cell progenitors), etc. Any convenient and appropriate promoter of an immune cell specific gene may find use in nucleic acids of the present disclosure. In some instances, an immune cell specific promoter of a nucleic acid present in a system of the present disclosure may be a T cell specific promoter. In some instances, an immune cell specific promoter of a nucleic acid present in a system of the present disclosure may be a light and/or heavy chain immunoglobulin gene promoter and may or may not include one or more related enhancer elements.

[00355] In some instances, an immune cell specific promoter of a nucleic acid present in a system of the present disclosure may be a promoter of a B29 gene promoter, a CD14 gene promoter, a CD43 gene promoter, a CD45 gene promoter, a CD68 gene promoter, a IFN-β gene promoter, a WASP gene promoter, a T-cell receptor β -chain gene promoter, a V9 γ (TRGV9) gene promoter, a V2 δ (TRDV2) gene promoter, and the like.

[00356] In some instances, an immune cell specific promoter present in a system of a nucleic acid of the present disclosure may be a viral promoter expressed in immune cells. As such, in some instances, viral promoters useful in nucleic acids present in a system of the present disclosure include viral promoters

derived from immune cells viruses, including but not limited to, e.g., lentivirus promoters (e.g., human immunodeficiency virus (HIV), SIV, FIV, EIAV, or Visna promoters) including e.g., long terminal repeat (LTR) promoter, etc., Retroviridae promoters including, e.g., HTLV-I promoter, HTLV-II promoter, etc., and the like.

[00357] In some cases, the promoter is a CD8 cell-specific promoter, a CD4 cell-specific promoter, a neutrophil-specific promoter, or an NK-specific promoter. For example, a CD4 gene promoter can be used; see, e.g., Salmon et al. (1993) *Proc. Natl. Acad. Sci. USA* 90:7739; and Marodon et al. (2003) *Blood* 101:3416. As another example, a CD8 gene promoter can be used. NK cell-specific expression can be achieved by use of an *Ncr1* (*p46*) promoter; see, e.g., Eckelhart et al. (2011) *Blood* 117:1565.

[00358] Expression vectors generally have convenient restriction sites located near the promoter sequence to provide for the insertion of nucleic acid sequences encoding heterologous proteins. A selectable marker operative in the expression host may be present. Suitable recombinant expression vectors include, but are not limited to, viral vectors (e.g. viral vectors based on vaccinia virus; poliovirus; adenovirus (see, e.g., Li et al., Invest Opthalmol Vis Sci 35:2543 2549, 1994; Borras et al., Gene Ther 6:515 524, 1999; Li and Davidson, PNAS 92:7700 7704, 1995; Sakamoto et al., H Gene Ther 5:1088 1097, 1999; WO 94/12649, WO 93/03769; WO 93/19191; WO 94/28938; WO 95/11984 and WO 95/00655); adeno-associated virus (see, e.g., Ali et al., Hum Gene Ther 9:81 86, 1998, Flannery et al., PNAS 94:6916 6921, 1997; Bennett et al., Invest Opthalmol Vis Sci 38:2857 2863, 1997; Jomary et al., Gene Ther 4:683 690, 1997, Rolling et al., Hum Gene Ther 10:641 648, 1999; Ali et al., Hum Mol Genet 5:591 594, 1996; Srivastava in WO 93/09239, Samulski et al., J. Vir. (1989) 63:3822-3828; Mendelson et al., Virol. (1988) 166:154-165; and Flotte et al., PNAS (1993) 90:10613-10617); SV40; herpes simplex virus; human immunodeficiency virus (see, e.g., Miyoshi et al., PNAS 94:10319 23, 1997; Takahashi et al., J Virol 73:7812 7816, 1999); a retroviral vector (e.g., Murine Leukemia Virus, spleen necrosis virus, and vectors derived from retroviruses such as Rous Sarcoma Virus, Harvey Sarcoma Virus, avian leukosis virus, human immunodeficiency virus, myeloproliferative sarcoma virus, and mammary tumor virus); and the like.

[00359] A method of the present disclosure for killing a target cell in an individual comprising: a) introducing a system of the present disclosure into an immune cell (e.g., a $CD8^+$ T cell; an NK cell) obtained from the individual, generating a modified immune cell; and b) administering the modified immune cell to the individual, where the modified immune cell kills the target cancer cell in the individual, involves administering an effective amount of the modified immune cells to the individual.

[00360] In some cases, an effective amount (e.g., an effective number) of modified immune cells is an amount that, when administered in one or more doses to an individual having a cancer, decreases the number of cancer cells in the individual by at least about 10%, at least about 15%, at least about 20%, at least about 25%, at least about 30%, at least about 40%, at least about 50%, at least about 75%, at least

about 80%, at least about 90%, at least about 95%, or at least 98%, compared to the number of cancer cells in the individual before said administration.

[00361] In some cases, from about 10^2 to about 10^9 modified immune cells are administered to an individual in a single dose. In some cases, a single dose of modified immune cells disclosure contains from 10^2 to about 10^4, from about 10^4 to about 10^5, from about 10^5 to about 10^6, from about 10^6 to about 10^7, from about 10^7 to about 10^8, or from about 10^8 to about 10^9 modified immune cells. In some cases, a single dose of modified immune cells is administered. Multiple doses can also be administered, as needed and/or as determined by a medical professional. For example, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more than 10, doses can be administered. If multiple doses are administered, the multiple doses can be administered at various frequencies, including, e.g., once per week, twice per month, once per month, once every 2 months, once every 3 months, once every 4 months, once every 6 months, or once per year.

[00362] In some cases, the target cancer cell is a liposarcoma, a glioblastoma, a breast cancer cell, a renal cancer cell, a pancreatic cancer cell, a melanoma, an anaplastic lymphoma, a leiomyosarcoma, an astrocytoma, an ovarian cancer cell, a neuroblastoma, a mantle cell lymphoma, a sarcoma, a non-small cell lung cancer cell, an acute myeloid leukemia (AML) cell, a stomach cancer cell, a B-cell cancer cell, a lung cancer cell, or an oligodendroglioma.

Methods comprising use of a genetically modified cytotoxic T cell of the present disclosure

[00363] As noted above, in some cases, a method of the present disclosure for killing a target cell in an individual comprises administering a genetically modified cytotoxic immune cell (e.g., a genetically modified $CD8^+$ T cell; a genetically modified NK cell) of the present disclosure to the individual, where the genetically modified immune cell kills the target cancer cell in the individual. In some cases, the modified cytotoxic T cell does not substantially kill non-target cells such as non-cancerous cells.

[00364] T cells can be obtained from an individual (e.g., an individual having a cancer; an individual diagnosed as having a cancer; an individual being treated for a cancer with chemotherapy, radiation therapy, antibody therapy, surgery, etc.) using well-established methods. In some cases, a mixed population of cells is obtained from an individual; and $CD8^+$ T cells and/or NK cells are isolated from the mixed population, such that a population of $CD8^+$ T cells and/or NK cells is obtained that is at least 75%, at least 80%, at least 85%, at least 90%, at least 95%, or more than 98% pure, i.e., the purified cell population includes less than 25%, less than 20%, less than 15%, less than 10%, less than 5%, or less than 2%, of cells other than $CD8^+$ T cells and or NK cells. The purified $CD8^+$ T cells and/or NK cells are then genetically modified to express the first antigen-triggered polypeptide and the second antigen-triggered polypeptide.

[00365] A method of the present disclosure for killing a target cell in an individual comprising administering a genetically modified cytotoxic immune cell (e.g., a genetically modified $CD8^+$ T cell; a genetically modified NK cell) of the present disclosure to the individual involves administering an

effective amount of a genetically modified cytotoxic immune cell of the present disclosure to the individual.

[00366] In some cases, an effective amount (e.g., an effective number) of genetically modified cytotoxic immune cells of the present disclosure is an amount that, when administered in one or more doses to an individual having a cancer, decreases the number of cancer cells in the individual by at least about 10%, at least about 15%, at least about 20%, at least about 25%, at least about 30%, at least about 40%, at least about 50%, at least about 75%, at least about 80%, at least about 90%, at least about 95%, or at least 98%, compared to the number of cancer cells in the individual before said administration.

[00367] In some cases, from about 10^2 to about 10^9 genetically modified cytotoxic immune cells of the present disclosure are administered to an individual in a single dose. In some cases, a single dose of genetically modified cytotoxic immune cells of the present disclosure contains from 10^2 to about 10^4, from about 10^4 to about 10^5, from about 10^5 to about 10^6, from about 10^6 to about 10^7, from about 10^7 to about 10^8, or from about 10^8 to about 10^9 genetically modified cytotoxic immune cells of the present disclosure. In some cases, a single dose of genetically modified cytotoxic immune cells of the present disclosure is administered. Multiple doses can also be administered, as needed and/or as determined by a medical professional. For example, 2, 3, 4, 5, 6, 7, 8, 9, 10, or more than 10, doses can be administered. If multiple doses are administered, the multiple doses can be administered at various frequencies, including, e.g., once per week, twice per month, once per month, once every 2 months, once every 3 months, once every 4 months, once every 6 months, or once per year.

[00368] In some cases, the target cancer cell is a liposarcoma, a glioblastoma, a breast cancer cell, a renal cancer cell, a pancreatic cancer cell, a melanoma, an anaplastic lymphoma, a leiomyosarcoma, an astrocytoma, an ovarian cancer cell, a neuroblastoma, a mantle cell lymphoma, a sarcoma, a non-small cell lung cancer cell, an acute myeloid leukemia (AML) cell, a stomach cancer cell, a B-cell cancer cell, a lung cancer cell, or an oligodendroglioma.

Individuals suitable for treatment

[00369] Individuals suitable for treatment using a method of the present disclosure include an individual having a cancer; an individual diagnosed as having a cancer; an individual being treated for a cancer with chemotherapy, radiation therapy, antibody therapy, surgery, etc.); an individual who has been treated for a cancer (e.g., with one or more of chemotherapy, radiation therapy, antibody therapy, surgery, etc.), and who has failed to respond to the treatment; an individual who has been treated for a cancer (e.g., with one or more of chemotherapy, radiation therapy, antibody therapy, surgery, etc.), and who initially responded to the treatment but who subsequently relapsed, i.e., the cancer recurred.

[00370] Cancers that can be treated with a method of the present disclosure include liposarcoma, glioblastoma, breast cancer, renal cancer, pancreatic cancer, melanoma, anaplastic lymphoma, leiomyosarcoma, astrocytoma, ovarian cancer, neuroblastoma, mantle cell lymphoma, sarcoma, non-

small cell lung cancer, acute myeloid leukemia (AML), stomach cancer, B-cell cancer, lung cancer, and oligodendroglioma.

[00371] In some cases, an individual to which a treatment of the present disclosure is administered is an individual expressing one or more antigens relevant to the subject treatment, including e.g., one or more target (i.e., cancer) antigens and/or one or more non-target (i.e., non-cancer or normal) antigens. Antigen expression may be determined by any convenient means. For example, in some instances, a subject may be evaluated for expression (or lack thereof) of one or more antigens relevant to the subject treatment, including one or more or all of the antigens of a particular antigen combination utilized in the treatment. Such evaluations (i.e., antigen expression testing) may be performed at any convenient time before, during or after a particular treatment regimen and using any convenient sample obtained from a subject (e.g., a tissue sample, a biopsy sample, etc.). Evaluations of antigen expression may be employed predictively (e.g., to predict the efficacy of an antigen combination based therapy), concurrently (e.g., to confirm the expression of antigens of an antigen combination during therapy), retrospectively (e.g., to analyze the expression of antigens of an antigen combination after therapy, e.g., to correlate expression of treatment outcomes, e.g., as part of a clinical trial utilizing an antigen combination described herein), or the like.

Computational Methods

[00372] The instant disclosure includes computational methods of identifying antigen combinations for targeting cancer cells of various types. The computational methods will generally be based on the measured expression of a plurality of cell surface antigens and/or genes encoding cell surface antigens for a first specific population of cells relative to a second specific population of cells. For example, in some instances, the cell surface antigen expression within a population of cancer cells is measured relative to the cell surface antigen expression within a population of corresponding cells, where corresponding cells will generally be non-cancerous cells of the same type or derived from the same tissue as the cancer cells. In certain cases, gene or protein expression is compared between two tissues of the same type where one tissue contains a cancer and the other tissue is healthy or is otherwise suspected to not contain the cancer. In some cases, cell surface antigen expression within a population of cancer cells or a cancerous tissue is measured relative to the cell surface antigen expression within a population of corresponding cells or to the cell surface antigen expression within a corresponding tissue, where corresponding cells or tissue will generally be non-cancerous cells or tissue of the same type or derived from the same tissue as the cancer cells; in some cases, the non-cancerous cells (or normal tissue) are cells or tissue for which cell surface antigen expression data are publicly available. The antigen expression level(s) for a particular population of cells or tissue may be referred to as an expression dataset (e.g., a cancer tissue dataset, a normal tissue data set, a cancer cell dataset, a normal cell dataset, etc.).

[00373] Any convenient method may find use in comparing antigen expression of the cells and/or tissues, including methods for evaluating mRNA expression and methods for evaluating protein expression, including but not limited to e.g., RNA microarray methods, quantitative RNA sequencing (RNAseq) methods, quantitative polymerase chain reaction (qPCR) methods, protein microarray methods, quantitative mass-spec method, proteomics methods, and the like. In certain cases, existing data sets including but not limited to e.g., RNA microarray data sets, quantitative RNA sequencing (RNAseq) data sets, qPCR data sets, protein microarray data sets, quantitative mass-spec data sets, proteomics methods, etc., may be used in the computational methods as described herein. In some cases, a combination of two or three of these methods is used for comparing antigen expression of cells/tissues. In some instances, the data set may be a curated, including a human curated dataset or computationally annotated dataset. In some cases, single cell measurements, or micro dissected data, are used. In some cases, multiple (2, 3, 4, 5, 6, 7, 8, 9, 10, or more than 10) datasets are used. In some cases, e.g., where multiple data sets are used, each sample within each dataset is curated.

[00374] In some embodiments, an expression dataset comparing a cancerous biological sample vs. a corresponding normal biological sample may be utilized in the subject computational methods. A "biological sample" encompasses a variety of sample types obtained from an individual. The definition encompasses blood and other liquid samples of biological origin, solid tissue samples such as a biopsy specimen or tissue cultures or cells derived therefrom and the progeny thereof. The definition also includes samples that have been manipulated in any way after their procurement, such as by treatment with reagents, solubilization, microdissection, or enrichment for certain components, such as polynucleotides or polypeptides. A "biological sample" can be a single cell, or can be derived from a single cell. A "biological sample" can be a plurality of cells, or can be derived from a plurality of cells. The term "biological sample" encompasses a clinical sample, and also includes cells in culture, cell supernatants, cell lysates, serum, plasma, biological fluid, and tissue samples. The term "biological sample" includes urine, saliva, cerebrospinal fluid, interstitial fluid, ocular fluid, synovial fluid, bronchoalveolar lavage fluid, blood fractions such as plasma and serum, and the like, provided such samples contain the subject cells for which expression is to be measured. The term "biological sample" also includes solid tissue samples, tissue culture samples, and cellular samples.

[00375] Following procurement of expression datasets, whether measured de novo or obtained from a previously collected dataset or a combination thereof, the datasets will generally be normalized. In some cases, normalization is performed by computing the relative levels of expression to a reference gene or reference gene set. In some cases, normalization is performed by comparing all measurements of a dataset as a whole or globally, e.g., by computationally modeling the dataset. Useful normalization methods will vary and may include but are not limited to e.g., Single-channel array normalization (SCAN), Universal exPression Codes (UPC) normalization, and the like, as described in e.g., Piccolo et

al. (2012) Genomics, 100(6), pp. 337-344; Piccolo et al. (2013) PNAS, 110(44), pp. 17778-17783; the disclosures of which are incorporated herein by reference in their entirety. Normalization may allow not only for comparison of data points within a dataset but also between two or more datasets.

[00376] An expression dataset may be used to produce parameters for a target antigen and/or target antigen pairs. Useful target antigen parameters include, but are not limited to, the relative expression levels of a particular antigen and/or the relative expression levels of a pair of antigens compared to other antigen pairs derived from the data set. Useful target antigen parameters include, but are not limited to, relative expression levels of a particular antigen on a cancer cell or cancerous tissue compared to the expression levels of the antigen on a corresponding non-cancerous cell of the same cell type or corresponding non-cancerous tissues of the same tissue type and/or relative expression levels of a pair of antigens on a cancer cell or cancerous tissue compared to their levels on a corresponding non-cancerous cell of the same cell type or corresponding non-cancerous tissues of the same tissue type.

[00377] In some cases, a portion of the data (e.g., a portion of a larger dataset) is selected for training of a processor used in generating an algorithm for identifying antigen pairs. Such a selected portion of the data (e.g., a selected portion of a larger dataset) may be referred to as a "training data set" and may include data of various types including but not limited to e.g., any of those described herein. In using a training dataset, individual antigen expression levels may be compared pairwise to identify significant differences between the datasets and/or pairs of antigen expression levels may be compared pairwise to identify significant differences between the antigen pairs of the datasets. Such comparisons may be performed iteratively using the training set and/or may be expanded to include the entire dataset or datasets. From a training set of expression data, antigens and/or antigen pairs that discriminate between cell or tissue populations (e.g., cancer vs. non-cancer) may be selected and used in generating an algorithm for selecting and/or ranking antigens and/or antigen pairs of one or more larger datasets.

[00378] In some instances, the pairing algorithm may be constrained to select one or more antigens for each pairing that meets a specified criteria including e.g., where one of the antigens is a clinical antigen (i.e., has been evaluated clinically (i.e., in an immunotherapy clinical trial) and found to be a marker for one or more cancers), where two or more of the antigens are clinical antigens, where one or more of the antigens is not a clinical antigen, where two or more of the antigens are not clinical antigens, where one of the antigens is a clinical antigen and one of the antigens is not a clinical antigen, etc. In some instances, the pairing algorithm may be constrained to select only those antigens that contain a transmembrane domain, including or excluding putative transmembrane domains. In some cases, a clinical antigen that has been the target of immunotherapy for a first cancer cell type can be identified, using a computational method of the present disclosure, as a target antigen for a second cancer cell type (i.e., a cancer cell type that is different from the first cancer cell type).

[00379] An algorithm for identifying antigen combinations (e.g., antigen pairs) useful in antigen logic gates of the subject disclosure may include a decision-tree classifier, where the training dataset may be used to develop an algorithm that identifies a classification model that best fits the antigen parameters to the class labels (i.e., the populations) of the training dataset. Decision tree classifiers are organized as a series of test questions to separate antigens based on their different characteristics or expression parameters. The decision tree may iteratively test antigen pairs for the ability to differentiate a cancer cell from a non-cancer cell. A non-limiting examples of such decision trees are provided in FIG. 7, which schematically depicts a decision tree for identifying an AND-AND gate antigen pair, and FIG. 8, which schematically depicts a decision tree for identifying an AND-NOT gate antigen pair.

[00380] Various calculated performance measures or statistical values, and/or combinations thereof, may find use in an algorithm to identify useful antigen combinations. For example, in some instances, an algorithm, as described herein, may include a measure of the fraction of retrieved antigen combinations that are relevant (e.g., a precision statistic or a positive predictive value statistic) and/or a measure of the fraction of relevant antigen combinations that are retrieved (e.g., a recall statistic or a sensitivity statistic). In some instances, an algorithm, as described herein, may include a combined statistic of precision and recall including but not limited to e.g., the harmonic mean of precision and recall or, e.g., a F-score such as e.g., an F1 statistic, where the F1 statistic is calculated as ((the product of the precision and recall statistics) divided by (the sum or the precision and recall statistics)) multiplied by two. Precision, recall and F-score (e.g., F1 statistic), as well as various other statistical measures useful in an antigen combination determining algorithm that may be employed, include but are not limited to e.g., those described in Pizzuti et al. Evolutionary Computation, Machine Learning and Data Mining in Bioinformatics: 9th European Conference, EvoBIO (2011), Torino, Italy, April 27-29, 2011; Lee JK, Statistical Bioinformatics: For Biomedical and Life Science Researchers (2011), John Wiley & Sons; Mirkin B, Core Concepts in Data Analysis: Summarization, Correlation and Visualization (2011) Springer Science & Business Media; Kohane et al. Microarrays for an Integrative Genomics (2005) MIT Press; and the like.

[00381] The precision statistic, in this context, may be used to indicate the proportion of cancer samples of those samples from the test data that are predicted to trigger a cytotoxic response. Normal samples with expression patterns that correspond to the targeted logic may count as false positives and may impact the precision score. As normal samples come not only from multiple individuals, but from a wide variety of distinct body tissues, precision can be used to describe the level of risk of unwanted off-tumor reactivity associated with a particular targeting configuration.

[00382] The recall statistic, in this context, may be used to indicate the proportion of cancer samples in the test data with expression patterns that conform to the desired targeting logic. Cancer samples in the test data with expression patterns that do not match the targeting logic may count as false negatives and

impact the recall score. As tumor samples come from different tumors or patients, they reflect heterogeneity both of tumors and individuals. Therefore recall can be used to describe the efficacy or generalizability of a given targeting configuration.

[00383] An F-score (e.g., an F1 statistic) represents a combination of precision and recall, that, in this context, captures the tradeoff between safety and efficacy, and allows for a summary of the performance of different configurations along a single axis. Various constants for the F-score may be employed that will combine precision and recall in such a way as to emphasize one or the other, depending on the particular context.

[00384] Any convenient calculated statistic may be used independently or in combination as a cutoff value for computationally identifying useful antigen combinations as described herein. For example, in some instances, a precision statistic may be used as a precision cutoff in an algorithm for identifying a useful antigen combination. Useful precision cutoff values will vary depending on the particular computational method employed and may range from less than 0.7 to 1.0 including but not limited to e.g., 0.7 to 1.0, 0.75 to 1.0, 0.76 to 1, 0.77 to 1, 0.78 to 1, 0.79 to 1, 0.8 to 1, 0.81 to 1, 0.82 to 1, 0.83 to 1, 0.84 to 1, 0.85 to 1, 0.86 to 1, 0.87 to 1, 0.88 to 1, 0.89 to 1, 0.9 to 1, 0.91 to 1, 0.92 to 1, 0.93 to 1, 0.94 to 1, 0.95 to 1, 0.96 to 1, 0.97 to 1, 0.98 to 1, etc. In some cases, the precision cutoff values can range from 0.5 to 0.7 or 0.5 to 1.0; e.g., from 0.55 to 0.7, from 0.60 to 0.7, from 0.5 to 1.0, from 0.6 to 1.0, or from 0.65 to 0.7.

[00385] In some instances, a recall statistic may be used as a recall cutoff in an algorithm for identifying a useful antigen combination. Useful recall cutoff values will vary depending on the particular computational method employed and may range from less than 0.7 to 1.0 including but not limited to e.g., 0.7 to 1.0, 0.75 to 1.0, 0.76 to 1, 0.77 to 1, 0.78 to 1, 0.79 to 1, 0.8 to 1, 0.81 to 1, 0.82 to 1, 0.83 to 1, 0.84 to 1, 0.85 to 1, 0.86 to 1, 0.87 to 1, 0.88 to 1, 0.89 to 1, 0.9 to 1, 0.91 to 1, 0.92 to 1, 0.93 to 1, 0.94 to 1, 0.95 to 1, 0.96 to 1, 0.97 to 1, 0.98 to 1, etc.

[00386] In some instances, an F-score (such as an F1 statistic for example) may be used as a F1 cutoff in an algorithm for identifying a useful antigen combination. Useful F1 cutoff values will vary depending on the particular computational method employed and may range from less than 0.8 to 1.0 including but not limited to e.g., 0.8 to 1.0, 0.81 to 1, 0.82 to 1, 0.83 to 1, 0.84 to 1, 0.85 to 1, 0.86 to 1, 0.87 to 1, 0.88 to 1, 0.89 to 1, 0.9 to 1, 0.91 to 1, 0.92 to 1, 0.93 to 1, 0.94 to 1, 0.95 to 1, 0.96 to 1, 0.97 to 1, 0.98 to 1, 0.99 to 1, etc. In some cases, the F1 cutoff values can range from 0.5 to 1.0; e.g., from 0.55 to 0.7, from 0.60 to 0.7, from 0.5 to 1.0, from 0.6 to 1.0, from 0.65 to 0.8, or from 0.7 to 1.0.

[00387] In some instances, two or more statistics, including two or more different statistics may be used in combination including but not limited to e.g., 2 or more different statistics, 3 or more different statistics, 4 or more different statistics, 5 or more different statistics, 6 or more different statistics, 7 or more

different statistics, 8 or more different statistics, 9 or more different statistics, 10 or more different statistics, etc.

Examples of Non-Limiting Aspects of the Disclosure

[00388] Aspects, including embodiments, of the present subject matter described above may be beneficial alone or in combination, with one or more other aspects or embodiments. Without limiting the foregoing description, certain non-limiting aspects of the disclosure numbered as below are provided. As will be apparent to those of skill in the art upon reading this disclosure, each of the individually numbered aspects may be used or combined with any of the preceding or following individually numbered aspects. This is intended to provide support for all such combinations of aspects and is not limited to combinations of aspects explicitly provided below:

[00389] 1. An *in vitro* or *ex vivo* genetically modified cytotoxic immune cell, wherein the cytotoxic immune cell is genetically modified to produce two different polypeptides that recognize two different cell surface antigens, wherein at least one of the two different cell surface antigens is present on the surface of a target cancer cell.

[00390] 2. The genetically modified cytotoxic immune cell of aspect 1, wherein the two different polypeptides comprise: a) a first antigen-triggered polypeptide that binds specifically to a first target cell surface antigen present on a target cancer cell; and b) a second antigen-triggered polypeptide that binds specifically to a second target cell surface antigen.

[00391] 3. The genetically modified cytotoxic immune cell of aspect 1 or aspect 2, wherein the cytotoxic immune cell is a cytotoxic T cell or a natural killer cell.

[00392] 4. The genetically modified cytotoxic immune cell of any one of aspect 1-3, wherein the immune cell is activated to kill a target cancer cell only when the target cancer cell expresses both of the two different cell surface antigens on its cell surface.

[00393] 5. The genetically modified cytotoxic immune cell of aspect 2 or aspect 3, wherein the cytotoxic immune cell: a) is activated to kill a target cancer cell that expresses the first target cell surface antigen, but not the second target cell surface antigen, on its cell surface; and b) is inhibited from killing a non-cancerous cell if the non-cancerous cell expresses both the first target cell surface antigen and the second target cell surface antigen on its cell surface.

[00394] 6. The genetically modified cytotoxic immune cell any one of aspects 2-5, wherein the first antigen-triggered polypeptide is a synNotch receptor and wherein the second antigen-triggered polypeptide is a chimeric antigen receptor (CAR). In some cases, the CAR is an ON-switch CAR.

[00395] 7. The genetically modified cytotoxic immune cell any one of aspects 2-5, wherein the first antigen-triggered polypeptide is a synNotch receptor and wherein the second antigen-triggered polypeptide is a T cell receptor (TCR).

[00396] 8. The genetically modified cytotoxic immune cell any one of aspects 2-5, wherein the first antigen-triggered polypeptide is a CAR, and wherein the second antigen-triggered polypeptide is an iCAR.

[00397] 9. The genetically modified cytotoxic immune cell any one of aspects 2-5, wherein the first antigen-triggered polypeptide is a CAR, and wherein the second antigen-triggered polypeptide is a synNotch receptor.

[00398] 10. The genetically modified cytotoxic immune cell any one of aspects 2-5, wherein the first antigen-triggered polypeptide is a synNotch receptor and wherein activation of the synNotch receptor induces expression of the second antigen-triggered polypeptide.

[00399] 11. The genetically modified cytotoxic immune cell any one of aspects 2-5, wherein the first antigen-triggered polypeptide is a synNotch receptor and wherein activation of the synNotch receptor induces expression of an immune inhibitory polypeptide.

[00400] 12. The genetically modified cytotoxic immune cell any one of aspects 1-11, wherein the target cancer cell is a liposarcoma, a glioblastoma, a breast cancer cell, a renal cancer cell, a pancreatic cancer cell, a melanoma, an anaplastic lymphoma, a leiomyosarcoma, an astrocytoma, an ovarian cancer cell, a neuroblastoma, a mantle cell lymphoma, a sarcoma, a non-small cell lung cancer cell, an AML cell, a stomach cancer cell, a B-cell cancer cell, a lung cancer cell, or an oligodendroglioma.

[00401] 13. The genetically modified cytotoxic immune cell any one of aspects 1-12, wherein the two different cell surface antigens are a target antigen pair selected from a target antigen pair depicted in FIG. 1 or FIG. 9-14.

[00402] 14. The genetically modified cytotoxic immune cell of any one of aspects 1-13, wherein the genetically modified to produce a third antigen-triggered polypeptide that recognizes an additional cell surface antigen present on the target cancer cell.

[00403] 15. The genetically modified cytotoxic immune cell of aspect 14, wherein the additional cell surface antigen is selected from an antigen depicted in FIG. 4 or Table 3.

[00404] 16. The genetically modified cytotoxic immune cell of aspect 14, wherein the additional cell surface antigen is selected from: a) an α-folate receptor; b) carbonic anhydrase IX (CAIX); c) CD19; d) CD20; e) CD22; f) CD33; g) CD44v7/8; h) carcinoembryonic antigen; i) epithelial glycoprotein-2 (EGP-2); j) epithelial glycoprotein-40 (EGP-40); k) erb-B2; l) folate binding protein (FBP); m) fetal acetylcholine receptor; n) GD2 ganglioside; o) GD3 ganglioside; p) Her2/neu; q) IL-13R-a2; r) kinase insert domain receptor (KDR); s) immunoglobulin kappa light chain; t) LeY; u) melanoma antigen E-A1 (MAGE-A1); v) mesothelin; w) epithelial mucin MUC1; x) prostate stem cell antigen (PSCA); y) prostate-specific membrane antigen (PSMA); z) tumor-associated glycoprotein-72 (TAG-72); and aa) vascular endothelial growth factor receptor-2 (VEGF-R2).

[00405] 17. A method of killing a target cancer cell in an individual, the method comprising administering to the individual an effective number of the genetically modified cytotoxic immune cell of any one of aspects 1-16, wherein said genetically modified cytotoxic immune cell kills the target cancer cell in the individual.

[00406] 18. The method of aspect 17, wherein the target cancer cell is a liposarcoma, a glioblastoma, a breast cancer cell, a renal cancer cell, a pancreatic cancer cell, a melanoma, an anaplastic lymphoma, a leiomyosarcoma, an astrocytoma, an ovarian cancer cell, a neuroblastoma, a mantle cell lymphoma, a sarcoma, a non-small cell lung cancer cell, an AML cell, a stomach cancer cell, a B-cell cancer cell, a lung cancer cell, or an oligodendroglioma.

[00407] 19. A system for killing a target cancer cell, the system comprising: a) a first antigen-triggered polypeptide that binds specifically to a first target antigen present on the target cancer cell, or a first nucleic acid comprising a nucleotide sequence encoding the first antigen-triggered polypeptide; and b) a second antigen-triggered polypeptide that binds specifically to a second target antigen, or a second nucleic acid comprising a nucleotide sequence encoding the second antigen-triggered polypeptide. In some cases, the system comprises: a) a first nucleic acid comprising a nucleotide sequence encoding first antigen-triggered polypeptide that binds specifically to a first target antigen present on the target cancer cell; and b) a second nucleic acid comprising a nucleotide sequence encoding a second antigen-triggered polypeptide that binds specifically to a second target antigen. In some cases, the first and the second nucleic acids are recombinant expression vectors. In some cases, the first and the second nucleic acids are present in separate recombinant expression vectors. In some cases, the first and the second nucleic acids are present in a single recombinant expression vectors. In some cases, the nucleotide sequences are operably linked to promoters. In some cases, the promoters are T-cell-specific promoters. In some cases, the promoters are constitutive promoters. In some cases, the promoters are regulatable promoters.

[00408] 20. The system of aspect 19, wherein the first antigen-triggered polypeptide is a synNotch polypeptide, and the second antigen-triggered polypeptide is a chimeric antigen receptor (CAR).

[00409] 21. The system of aspect 19, wherein the first antigen-triggered polypeptide is a synNotch polypeptide, and the second antigen-triggered polypeptide is an inhibitory CAR (iCAR).

[00410] 22. The system of aspect 19, wherein the first antigen-triggered polypeptide is a synNotch polypeptide, and the second antigen-triggered polypeptide is a split CAR.

[00411] 23. The system of aspect 19, wherein the first target antigen and the second target antigen are both present on the surface of a target cancer cell.

[00412] 24. The system of aspect 19, wherein: a) the first target antigen and the second target antigen are both present on the surface of a non-cancerous cell; and b) wherein the first target antigen, but not the second target antigen, is present on the surface of a target cancer cell.

[00413] 25. The system of any one of aspects 19-24, wherein the first target antigen and the second target antigen are selected from a target antigen pair depicted in FIG. 1 or FIG. 9-14.

[00414] 26. The system of any one of aspects 19-24, wherein the target cancer cell is a liposarcoma, a glioblastoma, a breast cancer cell, a renal cancer cell, a pancreatic cancer cell, a melanoma, an anaplastic lymphoma, a leiomyosarcoma, an astrocytoma, an ovarian cancer cell, a neuroblastoma, a mantle cell lymphoma, a sarcoma, a non-small cell lung cancer cell, an AML cell, a stomach cancer cell, a B-cell cancer cell, a lung cancer cell, or an oligodendroglioma.

[00415] 27. The system of aspect 19, wherein the second target antigen is present on the target cancer cell.

[00416] 28. The system of aspect 19, wherein the second target antigen is not present on the target cancer cell.

[00417] 29. The system of aspect 19, wherein the first antigen-triggered polypeptide induces production of an antibody upon binding to the first target antigen.

[00418] 30. The system of aspect 19, wherein the first antigen-triggered polypeptide induces production of a cytokine upon binding to the first target antigen.

[00419] 31. A method of killing a target cancer cell in an individual, the method comprising: a) introducing the system of any one of aspects 19-30 into a cytotoxic T cell *in vitro* or *ex vivo*, generating a modified cytotoxic T cell; and b) administering the modified cytotoxic T cell to the individual.

[00420] 32. The method of aspect 31, wherein the target cancer cell is a liposarcoma, a glioblastoma, a breast cancer cell, a renal cancer cell, a pancreatic cancer cell, a melanoma, an anaplastic lymphoma, a leiomyosarcoma, an astrocytoma, an ovarian cancer cell, a neuroblastoma, a mantle cell lymphoma, a sarcoma, a non-small cell lung cancer cell, an AML cell, a stomach cancer cell, a B-cell cancer cell, a lung cancer cell, or an oligodendroglioma.

[00421] 33. The method of aspect 31 or 32, wherein the modified cytotoxic T cell is activated to kill the target cancer cell only when the target cancer cell expresses both the first target antigen and the second target antigen on its cell surface.

[00422] 34. The method of aspect 31 or 32, wherein the modified cytotoxic T cell: a) is activated to kill a target cancer cell that expresses the first target cell surface antigen, but not the second target cell surface antigen, on its cell surface; and b) is inhibited from killing a non-cancerous cell if the non-cancerous cell expresses both the first target cell surface antigen and the second target cell surface antigen on its cell surface.

[00423] 35. A method of identifying a combination of target antigens for targeting a cancer cell, the method comprising: a) generating a training set of target antigen parameters by determining, from a training set of expression data, pairs of antigens that discriminate between cancer cells and non-

cancerous cells, thereby generating an algorithm; b) applying the algorithm to a test set of expression data, to generate a combination of target antigens.

[00424] 36. The method of aspect 35, wherein step (a) comprises selecting pairs of target antigens that comprise: i) a first target antigen that is expressed at a high level on cancer cells of a selected cancer cell type; and ii) a second target antigen that is expressed at a high level on cancer cells of the selected cancer cell type; or i) a first target antigen that is expressed at a high level on cancer cells of a selected cancer cell type and on non-cancerous cells of the same cell type; and ii) a second target antigen that is expressed at a high level on non-cancerous cells of the same cell type but not on cancer cells of the same cell type.

[00425] 37. The method of aspects 35 or 36, wherein the algorithm comprises one or more of an F1 cutoff, a precision cutoff, a recall cutoff, or combination thereof.

[00426] 38. The method of any of aspects 35 to 37, wherein step (a) comprises training a computer software with the training set, and wherein step (b) comprises applying the algorithm using the trained computer software.

[00427] 39. The method of any of aspects 35 to 38, wherein the expression data comprises gene expression data, proteomics expression data or a combination thereof.

[00428] 40. The method of aspect 39, wherein the gene expression data comprises microarray data.

[00429] 41. The method of aspect 40, wherein the microarray data comprises mRNA expression data.

[00430] 42. The method of any of aspects 35 to 41, wherein the training comprises selecting target antigens that are transmembrane polypeptides.

[00431] 43. The method of any of aspects 35 to 42, further comprising comparing the output of the expression data to a reference dataset.

[00432] 44. The method of aspect 43, wherein the reference dataset comprises a RNAseq dataset.

[00433] 45. The method of any of aspects 35 to 44, wherein applying the algorithm is performed iteratively.

[00434] 46. An *in vitro* or *ex vivo* genetically modified cytotoxic immune cell, wherein the cytotoxic immune cell is genetically modified to produce a first polypeptide that recognizes a first cell surface antigen present on the surface of a target cancer cell and a second polypeptide that recognizes a second cell surface antigen present on the surface of a non-target cell.

[00435] 47. The genetically modified cytotoxic immune cell of aspect 46, wherein the non-target cell is a cell of a normal tissue.

[00436] 48. The genetically modified cytotoxic immune cell of aspect 47, wherein the normal tissue is brain tissue.

[00437] 49. The genetically modified cytotoxic immune cell of aspects 47 or 48, wherein the second cell surface antigen is selected from the group consisting of: OPALIN, TMEM235, GABRA1, KCNJ9, GRM3, SEZ6, NTSR2, KCNK4, SLCO1A2, SLC24A2, MOG, GABRG1, GABRG2, CNTNAP4, DSCAM, CACNG3, CRB1, CDH10, HRH3, GRIK1, SLC39A12, GPR158, CACNG2, SYT3, HTR5A, CACNG7, GPR37L1, LRRTM3, GLRA2, CHRNB2, KCNQ2, JPH3, GPR19, ADCY8, SPOCK3, SLC32A1, OPCML, GABRA3, GRM5, SCN1A, SLC5A11, KCNC1, SLC12A5, GRM4, GRM1, GRIA4, MEGF11, CACNA1B, LYPD1, GRID2, SCN2A, NKAIN2, UNC5A, SLC4A10, TMEFF2, CSMD3, PPAPDC1A, HAPLN4, GPR85, ANTXR2, CACNG4, CSPG5, KCNK10, CHRNA4, CNTNAP2, KCNJ10, GABRB2, GRIN1, CRB2, SHISA7, NKAIN4, HTR2C, CACNG8, NRG3, ABCG4, CDH8, GABRD, KIRREL3, GABRB1, KCNA2, CDH20, IGDCC3, KCNJ6, CNIH2, KCNK12, CDH18, CSMD2, SYT4, OR2L13, CDH9, GABRA2, KCNF1, MAG, CALN1, GRIN2B, GRM7, VSTM2A, GPR61, OMG, KCNA1, GPR83, ATP8A2, GABBR2, GPR12, TRPM3, SLC8A3, KCND2, GSG1L, SLC30A10, ASTN1, GPR179, LRFN2, CACNA1E, CALY, SLC6A15, KIAA0319, SYT6, PTPRR, KCTD8, GPR22, SLC4A8, LAMP5, MEGF10, FXYD7, KCNK9, SLC1A6, MLC1, OPRK1, ATP2B2, ACSL6, THBD, PTPRT, PCDHGC4, CLDN10, KCNV1, LPPR3, SLCO1C1, PCDH8, ANO4, LRRTM4, PCDH15, CCKBR, GABRA5, SLC6A12, GRIN2A, SLC1A2, SLC43A1, KCNC2, ELFN2, ATP7A, GRIK4, LRRC55, HCN2, NKAIN1, DPP10, AJAP1, NPFFR1, TRPC3, TGFBI, SLC6A7, GABRA4, SLC13A5, GRIN2C, HCN1, SLC26A8, PPIC, NETO1, TNFRSF10B, CDH22, SLC6A13, DISP2, SLC6A11, CD93, EPHA10, PHLDB2, OXTR, WNT7A, GYPC, KCNA4, PCDHAC2, HGFAC, DRD1, SHISA9, SCN8A, ICAM1, PIRT, A4GALT, MRGPRF, CD248, CD58, CD44, EPHA2 and PROCR.

[00438] 50. The genetically modified cytotoxic immune cell of any of aspects 46 to 49, wherein the cytotoxic immune cell is further genetically modified to produce a third polypeptides that recognizes a third cell surface antigen, wherein the third cell surface antigen is present on the surface of a second non-target cell.

[00439] 51. The genetically modified cytotoxic immune cell of aspect 50, wherein the second non-target cell is a cell of a second normal tissue.

[00440] 52. The genetically modified cytotoxic immune cell of aspect 51, wherein the second normal tissue is cardiac tissue.

[00441] 53. The genetically modified cytotoxic immune cell of aspect 52, wherein the third cell surface antigen is selected from the group consisting of: GJA3, HCN4 and BMP10.

[00442] 54. The genetically modified cytotoxic immune cell of any of aspects 46 to 53, wherein the first cell surface antigen is selected from the group consisting of: GD2 (B4GALNT1), MAGEA1, MAGEA3 and MART1 (MLANA).

[00443] 55. The genetically modified cytotoxic immune cell of any of aspects 46 to 54, wherein the first and second polypeptides each comprise an antigen-triggered polypeptide.

[00444] 56. The genetically modified cytotoxic immune cell of any of aspects 46 to 55, wherein the cytotoxic immune cell is a cytotoxic T cell or a natural killer cell.

[00445] 57. The genetically modified cytotoxic immune cell of any of aspects 46 to 55, wherein the target cell is a melanoma.

[00446] 58. A method of killing a target cancer cell in an individual, the method comprising administering to the individual an effective number of the genetically modified cytotoxic immune cell of any of aspects 46 to 57, wherein said genetically modified cytotoxic immune cell kills the target cancer cell in the individual.

[00447] 59. The method of aspect 58, wherein the genetically modified cytotoxic immune cell does not substantially kill the non-target cell.

EXAMPLES

[00448] The following examples are put forth so as to provide those of ordinary skill in the art with a complete disclosure and description of how to make and use the present invention, and are not intended to limit the scope of what the inventors regard as their invention nor are they intended to represent that the experiments below are all or the only experiments performed. Efforts have been made to ensure accuracy with respect to numbers used (e.g. amounts, temperature, etc.) but some experimental errors and deviations should be accounted for. Unless indicated otherwise, parts are parts by weight, molecular weight is weight average molecular weight, temperature is in degrees Celsius, and pressure is at or near atmospheric. Standard abbreviations may be used, e.g., bp, base pair(s); kb, kilobase(s); pl, picoliter(s); s or sec, second(s); min, minute(s); h or hr, hour(s); aa, amino acid(s); kb, kilobase(s); bp, base pair(s); nt, nucleotide(s); i.m., intramuscular(ly); i.p., intraperitoneal(ly); s.c., subcutaneous(ly); and the like.

Example 1: Use of an AND-NOT gate antigen pair as target antigens

[00449] $CD8^+$ T cells are isolated from a patient with a liposarcoma. The $CD8^+$ T cells are genetically modified with: a) a recombinant expression vector comprising a nucleotide sequence encoding a synNotch polypeptide that includes: i) a single-chain Fv (scFv) that is specific for an EVA1B polypeptide; and ii) an intracellular domain comprising a transcription activator; and b) a recombinant expression vector comprising a nucleotide sequence encoding an iCAR that comprises an antigen-binding portion (e.g., a scFv) specific for an ITGA6 polypeptide. From 10^6 to 10^9 genetically modified $CD8^+$ T cells are administered to the patient in a single dose intravenously.

Example 2: Use of an AND gate antigen pair as target antigens

[00450] $CD8^+$ T cells are isolated from a patient with a glioblastoma. The $CD8^+$ T cells are genetically modified with: a) a recombinant expression vector comprising a nucleotide sequence encoding a synNotch polypeptide that includes: i) an scFv that is specific for a PTPRZ1 polypeptide; and ii) an intracellular domain comprising a transcription activator; and b) a recombinant expression vector comprising a nucleotide sequence encoding a CAR that comprises an antigen-binding portion (e.g., a scFv) specific for a FOLR2 polypeptide. From 10^6 to 10^9 genetically modified $CD8^+$ T cells are administered to the patient in a single dose intravenously. Upon binding of the genetically modified $CD8^+$ T cells to a PTPRZ1 present on the surface of a glioblastoma in the patient, the transcription activator is released from the synNotch polypeptide. The CAR is encoded by a nucleotide sequence that is operably linked to a promoter element that is controlled by the released transcription activator. The released transcription activator induces expression of the CAR in the genetically modified $CD8^+$ T cells. The CAR binds to a FOLR2 polypeptide present on the surface of the glioblastoma in the patient.

[00451] While the present invention has been described with reference to the specific embodiments thereof, it should be understood by those skilled in the art that various changes may be made and equivalents may be substituted without departing from the true spirit and scope of the invention. In addition, many modifications may be made to adapt a particular situation, material, composition of matter, process, process step or steps, to the objective, spirit and scope of the present invention. All such modifications are intended to be within the scope of the claims appended hereto.

CLAIMS

What is claimed is:

1. An *in vitro* or *ex vivo* genetically modified cytotoxic immune cell, wherein the cytotoxic immune cell is genetically modified to produce two different polypeptides that recognize two different cell surface antigens, wherein at least one of the two different cell surface antigens is present on the surface of a target cancer cell.

2. The genetically modified cytotoxic immune cell of claim 1, wherein the two different polypeptides comprise:

a) a first antigen-triggered polypeptide that binds specifically to a first target cell surface antigen present on a target cancer cell; and

b) a second antigen-triggered polypeptide that binds specifically to a second target cell surface antigen.

3. The genetically modified cytotoxic immune cell of claim 1 or claim 2, wherein the cytotoxic immune cell is a cytotoxic T cell or a natural killer cell.

4. The genetically modified cytotoxic immune cell of any one of claims 1-3, wherein the immune cell is activated to kill a target cancer cell only when the target cancer cell expresses both of the two different cell surface antigens on its cell surface.

5. The genetically modified cytotoxic immune cell of claim 2 or claim 3, wherein the cytotoxic immune cell:

a) is activated to kill a target cancer cell that expresses the first target cell surface antigen, but not the second target cell surface antigen, on its cell surface; and

b) is inhibited from killing a non-cancerous cell if the non-cancerous cell expresses both the first target cell surface antigen and the second target cell surface antigen on its cell surface.

6. The genetically modified cytotoxic immune cell any one of claims 2-5, wherein the first antigen-triggered polypeptide is a synNotch receptor and wherein the second antigen-triggered polypeptide is a chimeric antigen receptor (CAR).

7. The genetically modified cytotoxic immune cell any one of claims 2-5, wherein the first antigen-triggered polypeptide is a synNotch receptor and wherein the second antigen-triggered polypeptide is a T cell receptor (TCR).

8. The genetically modified cytotoxic immune cell any one of claims 2-5, wherein the first antigen-triggered polypeptide is a CAR, and wherein the second antigen-triggered polypeptide is an iCAR.

9. The genetically modified cytotoxic immune cell any one of claims 2-5, wherein: a) the first antigen-triggered polypeptide is a CAR, and wherein the second antigen-triggered polypeptide is a synNotch receptor; or b) the first antigen-triggered polypeptide is a synNotch receptor, and wherein the second antigen-triggered polypeptide is a synNotch receptor.

10. The genetically modified cytotoxic immune cell any one of claims 2-5, wherein the first antigen-triggered polypeptide is a synNotch receptor and wherein activation of the synNotch receptor induces expression of the second antigen-triggered polypeptide.

11. The genetically modified cytotoxic immune cell any one of claims 2-5, wherein the first antigen-triggered polypeptide is a synNotch receptor and wherein activation of the synNotch receptor induces expression of an immune inhibitory polypeptide.

12. The genetically modified cytotoxic immune cell any one of claims 1-11, wherein the target cancer cell is an AML cell, an anaplastic lymphoma cell, an astrocytoma cell, a B-cell cancer cell, a breast cancer cell, a colon cancer cell, an ependymoma cell, an esophageal cancer cell, a glioblastoma cell, a glioma cell, a leiomyosarcoma cell, a liposarcoma cell, a liver cancer cell, a lung cancer cell, a mantle cell lymphoma cell, a melanoma cell, a neuroblastoma cell, a non-small cell lung cancer cell, an oligodendroglioma cell, an ovarian cancer cell, a pancreatic cancer cell, a peripheral T-Cell lymphoma cell, a renal cancer cell, a sarcoma cell or a stomach cancer cell.

13. The genetically modified cytotoxic immune cell any one of claims 1-12, wherein the two different cell surface antigens are a target antigen pair selected from a target antigen pair depicted in FIG. 1 or FIG. 9-14.

14. The genetically modified cytotoxic immune cell of any one of claims 1-13, wherein the cell is genetically modified to produce a third antigen-triggered polypeptide that recognizes an additional cell surface antigen present on the target cancer cell.

15. The genetically modified cytotoxic immune cell of claim 14, wherein the additional cell surface antigen is selected from an antigen depicted in FIG. 4 or Table 3.

16. The genetically modified cytotoxic immune cell of claim 14, wherein the additional cell surface antigen is selected from:

a) an α-folate receptor;

b) carbonic anhydrase IX (CAIX);

c) CD19;

d) CD20;

e) CD22;

f) CD33;

g) CD44v7/8;

h) carcinoembryonic antigen;

i) epithelial glycoprotein-2 (EGP-2);

j) epithelial glycoprotein-40 (EGP-40);

k) erb-B2;

l) folate binding protein (FBP);

m) fetal acetylcholine receptor;

n) GD2 ganglioside;

o) GD3 ganglioside;

p) Her2/neu;

q) IL-13R-a2;

r) kinase insert domain receptor (KDR);

s) immunoglobulin kappa light chain;

t) LeY;

u) melanoma antigen E-A1 (MAGE-A1);

v) mesothelin;

w) epithelial mucin MUC1;

x) prostate stem cell antigen (PSCA);

y) prostate-specific membrane antigen (PSMA);

z) tumor-associated glycoprotein-72 (TAG-72); and

aa) vascular endothelial growth factor receptor-2 (VEGF-R2).

17. A method of killing a target cancer cell in an individual, the method comprising administering to the individual an effective number of the genetically modified cytotoxic immune cell of any one of claims 1-16, wherein said genetically modified cytotoxic immune cell kills the target cancer cell in the individual.

18. The method of claim 17, wherein the target cancer cell is an AML cell, an anaplastic lymphoma cell, an astrocytoma cell, a B-cell cancer cell, a breast cancer cell, a colon cancer cell, an ependymoma cell, an esophageal cancer cell, a glioblastoma cell, a glioma cell, a leiomyosarcoma cell, a liposarcoma cell, a liver cancer cell, a lung cancer cell, a mantle cell lymphoma cell, a melanoma cell, a neuroblastoma cell, a non-small cell lung cancer cell, an oligodendroglioma cell, an ovarian cancer cell, a pancreatic cancer cell, a peripheral T-Cell lymphoma cell, a renal cancer cell, a sarcoma cell or a stomach cancer cell.

19. A system for killing a target cancer cell, the system comprising:

a) a first antigen-triggered polypeptide that binds specifically to a first target antigen present on the target cancer cell, or a first nucleic acid comprising a nucleotide sequence encoding the first antigen-triggered polypeptide; and

b) a second antigen-triggered polypeptide that binds specifically to a second target antigen, or a second nucleic acid comprising a nucleotide sequence encoding the second antigen-triggered polypeptide.

20. The system of claim 19, wherein the first antigen-triggered polypeptide is a synNotch polypeptide, and the second antigen-triggered polypeptide is a chimeric antigen receptor (CAR).

21. The system of claim 19, wherein the first antigen-triggered polypeptide is a synNotch polypeptide, and the second antigen-triggered polypeptide is an inhibitory CAR (iCAR).

22. The system of claim 19, wherein:

a) the first antigen-triggered polypeptide is a synNotch polypeptide, and the second antigen-triggered polypeptide is one polypeptide chain of a split CAR;

b) the first antigen-triggered polypeptide is a synNotch polypeptide, and the second antigen-triggered polypeptide is a T-cell receptor;

c) the first antigen-triggered polypeptide is a synNotch polypeptide, and the second antigen-triggered polypeptide is an immunoinhibitory polypeptide; or

d) the first antigen-triggered polypeptide is a first synNotch polypeptide, and the second antigen-triggered polypeptide is a second synNotch polypeptide

23. The system of claim 19, wherein the first target antigen and the second target antigen are both present on the surface of a target cancer cell.

24. The system of claim 19, wherein:

a) the first target antigen and the second target antigen are both present on the surface of a non-cancerous cell; and

b) wherein the first target antigen, but not the second target antigen, is present on the surface of a target cancer cell.

25. The system of any one of claims 19-24, wherein the first target antigen and the second target antigen are selected from a target antigen pair depicted in FIG. 1 or FIG. 9-14.

26. The system of any one of claims 19-24, wherein the target cancer cell is an AML cell, an anaplastic lymphoma cell, an astrocytoma cell, a B-cell cancer cell, a breast cancer cell, a colon cancer cell, an ependymoma cell, an esophageal cancer cell, a glioblastoma cell, a glioma cell, a leiomyosarcoma cell, a liposarcoma cell, a liver cancer cell, a lung cancer cell, a mantle cell lymphoma cell, a melanoma cell, a neuroblastoma cell, a non-small cell lung cancer cell, an oligodendroglioma cell, an ovarian cancer cell, a pancreatic cancer cell, a peripheral T-Cell lymphoma cell, a renal cancer cell, a sarcoma cell or a stomach cancer cell.

27. The system of claim 19, wherein the second target antigen is present on the target cancer cell.

28. The system of claim 19, wherein the second target antigen is not present on the target cancer cell.

29. The system of claim 19, wherein the first antigen-triggered polypeptide induces production of an antibody upon binding to the first target antigen.

30. The system of claim 19, wherein the first antigen-triggered polypeptide induces production of a cytokine upon binding to the first target antigen.

31. A method of killing a target cancer cell in an individual, the method comprising:

a) introducing the system of any one of claims 19-30 into a cytotoxic T cell *in vitro* or *ex vivo*, generating a modified cytotoxic T cell; and

b) administering the modified cytotoxic T cell to the individual.

32. The method of claim 31, wherein the target cancer cell is an AML cell, an anaplastic lymphoma cell, an astrocytoma cell, a B-cell cancer cell, a breast cancer cell, a colon cancer cell, an ependymoma cell, an esophageal cancer cell, a glioblastoma cell, a glioma cell, a leiomyosarcoma cell, a liposarcoma cell, a liver cancer cell, a lung cancer cell, a mantle cell lymphoma cell, a melanoma cell, a neuroblastoma cell, a non-small cell lung cancer cell, an oligodendroglioma cell, an ovarian cancer cell, a pancreatic cancer cell, a peripheral T-Cell lymphoma cell, a renal cancer cell, a sarcoma cell or a stomach cancer cell.

33. The method of claim 31, wherein the modified cytotoxic T cell is activated to kill the target cancer cell only when the target cancer cell expresses both the first target antigen and the second target antigen on its cell surface.

34. The method of claim 31, wherein the modified cytotoxic T cell:

a) is activated to kill a target cancer cell that expresses the first target cell surface antigen, but not the second target cell surface antigen, on its cell surface; and

b) is inhibited from killing a non-cancerous cell if the non-cancerous cell expresses both the first target cell surface antigen and the second target cell surface antigen on its cell surface.

35. A method of identifying a combination of target antigens for targeting a cancer cell, the method comprising:

a) generating a training set of target antigen parameters by determining, from a training set of expression data, pairs of antigens that discriminate between cancer cells and non-cancerous cells, thereby generating an algorithm;

b) applying the algorithm to a test set of expression data, to generate a combination of target antigens.

36. The method of claim 35, wherein step (a) comprises selecting pairs of target antigens that comprise:

i) a first target antigen that is expressed at a high level on cancer cells of a selected cancer cell type; and

ii) a second target antigen that is expressed at a high level on cancer cells of the selected cancer cell type; or

i) a first target antigen that is expressed at a high level on cancer cells of a selected cancer cell type and on non-cancerous cells of the same cell type; and

ii) a second target antigen that is expressed at a high level on non-cancerous cells of the same cell type but not on cancer cells of the same cell type.

37. The method of claims 35 or 36, wherein the algorithm comprises one or more of an F1 cutoff, a precision cutoff, a recall cutoff, or combination thereof.

38. The method of any of claims 35 to 37, wherein step (a) comprises training a computer software with the training set, and wherein step (b) comprises applying the algorithm using the trained computer software.

39. The method of any of claims 35 to 38, wherein the expression data comprises gene expression data, proteomics expression data or a combination thereof.

40. The method of claim 39, wherein the gene expression data comprises microarray data.

41. The method of claim 40, wherein the microarray data comprises mRNA expression data.

42. The method of any of claims 35 to 41, wherein the training comprises selecting target antigens that are transmembrane polypeptides.

43. The method of any of claims 35 to 42, further comprising comparing the output of the expression data to a reference dataset.

44. The method of claim 43, wherein the reference dataset comprises a RNAseq dataset.

45. The method of any of claims 35 to 44, wherein applying the algorithm is performed iteratively.

46. An *in vitro* or *ex vivo* genetically modified cytotoxic immune cell, wherein the cytotoxic immune cell is genetically modified to produce a first polypeptide that recognizes a first cell surface antigen present on the surface of a target cancer cell and a second polypeptide that recognizes a second cell surface antigen present on the surface of a non-target cell.

47. The genetically modified cytotoxic immune cell of claim 46, wherein the non-target cell is a cell of a normal tissue.

48. The genetically modified cytotoxic immune cell of claims 47, wherein the normal tissue is brain tissue.

49. The genetically modified cytotoxic immune cell of claims 47 or 48, wherein the second cell surface antigen is selected from the group consisting of: OPALIN, TMEM235, GABRA1, KCNJ9, GRM3, SEZ6, NTSR2, KCNK4, SLCO1A2, SLC24A2, MOG, GABRG1, GABRG2, CNTNAP4, DSCAM, CACNG3, CRB1, CDH10, HRH3, GRIK1, SLC39A12, GPR158, CACNG2, SYT3, HTR5A, CACNG7, GPR37L1, LRRTM3, GLRA2, CHRNB2, KCNQ2, JPH3, GPR19, ADCY8, SPOCK3, SLC32A1, OPCML, GABRA3, GRM5, SCN1A, SLC5A11, KCNC1, SLC12A5, GRM4, GRM1, GRIA4, MEGF11, CACNA1B, LYPD1, GRID2, SCN2A, NKAIN2, UNC5A, SLC4A10, TMEFF2, CSMD3, PPAPDC1A, HAPLN4, GPR85, ANTXR2, CACNG4, CSPG5, KCNK10, CHRNA4, CNTNAP2, KCNJ10, GABRB2, GRIN1, CRB2, SHISA7, NKAIN4, HTR2C, CACNG8, NRG3, ABCG4, CDH8, GABRD, KIRREL3, GABRB1, KCNA2, CDH20, IGDCC3, KCNJ6, CNIH2, KCNK12, CDH18, CSMD2, SYT4, OR2L13, CDH9, GABRA2, KCNF1, MAG, CALN1, GRIN2B, GRM7, VSTM2A, GPR61, OMG, KCNA1, GPR83, ATP8A2, GABBR2, GPR12, TRPM3, SLC8A3, KCND2, GSG1L, SLC30A10, ASTN1, GPR179, LRFN2, CACNA1E, CALY, SLC6A15, KIAA0319, SYT6, PTPRR, KCTD8, GPR22, SLC4A8, LAMP5, MEGF10, FXYD7, KCNK9, SLC1A6, MLC1, OPRK1, ATP2B2, ACSL6, THBD, PTPRT, PCDHGC4, CLDN10, KCNV1, LPPR3, SLCO1C1, PCDH8, ANO4, LRRTM4, PCDH15, CCKBR, GABRA5, SLC6A12, GRIN2A, SLC1A2, SLC43A1, KCNC2, ELFN2, ATP7A, GRIK4, LRRC55, HCN2, NKAIN1, DPP10, AJAP1, NPFFR1, TRPC3, TGFBI, SLC6A7, GABRA4, SLC13A5, GRIN2C, HCN1, SLC26A8, PPIC, NETO1, TNFRSF10B, CDH22, SLC6A13, DISP2, SLC6A11, CD93,

EPHA10, PHLDB2, OXTR, WNT7A, GYPC, KCNA4, PCDHAC2, HGFAC, DRD1, SHISA9, SCN8A, ICAM1, PIRT, A4GALT, MRGPRF, CD248, CD58, CD44, EPHA2 and PROCR.

50. The genetically modified cytotoxic immune cell of any of claims 46 to 49, wherein the cytotoxic immune cell is further genetically modified to produce a third polypeptides that recognizes a third cell surface antigen, wherein the third cell surface antigen is present on the surface of a second non-target cell.

51. The genetically modified cytotoxic immune cell of claim 50, wherein the second non-target cell is a cell of a second normal tissue.

52. The genetically modified cytotoxic immune cell of claim 51, wherein the second normal tissue is cardiac tissue.

53. The genetically modified cytotoxic immune cell of claim 52, wherein the third cell surface antigen is selected from the group consisting of: GJA3, HCN4 and BMP10.

54. The genetically modified cytotoxic immune cell of any of claims 46 to 53, wherein the first cell surface antigen is selected from the group consisting of: GD2 (B4GALNT1), MAGEA1, MAGEA3 and MART1 (MLANA).

55. The genetically modified cytotoxic immune cell of any of claims 46 to 54, wherein the first and second polypeptides each comprise an antigen-triggered polypeptide.

56. The genetically modified cytotoxic immune cell of any of claims 46 to 55, wherein the cytotoxic immune cell is a cytotoxic T cell or a natural killer cell.

57. The genetically modified cytotoxic immune cell of any of claims 46 to 55, wherein the target cell is a melanoma.

58. A method of killing a target cancer cell in an individual, the method comprising administering to the individual an effective number of the genetically modified cytotoxic immune cell of any of claims 46 to 57, wherein said genetically modified cytotoxic immune cell kills the target cancer cell in the individual.

59. The method of claim 58, wherein the genetically modified cytotoxic immune cell does not substantially kill the non-target cell.

FIG. 1

Antigen Pair Logic	F1	Precision	Recall
Liposarcoma (Liposarcoma)			
EVA1B AND NOT-ITGA6	0.904109589	0.89189189	0.916666667
ADAM12 AND NOT-TACSTD2	0.904109589	0.89189189	0.916666667
Glioblastoma (Glioblastoma)			
PTPRZ1	0.9375	1	0.882352941
PTPRZ1 AND FOLR2	0.903225806	1	0.823529412
PTPRZ1 AND IGF1R	0.9375	1	0.882352941
PTPRZ1 AND NCAM1	0.9375	1	0.882352941
PTPRZ1 AND CD70	0.9375	1	0.882352941
PTPRZ1 AND AXL	0.903225806	1	0.823529412
PTPRZ1 AND NOT-NAALAD2	0.9375	1	0.882352941
PTPRZ1 AND NOT-TMEM170B	0.903225806	1	0.823529412
PTPRZ1 AND NOT-PIGK	0.903225806	1	0.823529412
PTPRZ1 AND CDH5	0.9375	1	0.882352941
PTPRZ1 AND NOT-100507547?	0.9375	1	0.882352941
PTPRZ1 AND SCAMP2	0.903225806	1	0.823529412
PTPRZ1 AND NOT-CDH8	0.903225806	1	0.823529412
PTPRZ1 AND NOT-CDH12	0.903225806	1	0.823529412
PTPRZ1 AND NOT-LPAR6	0.9375	1	0.882352941
PTPRZ1 AND NOT-CDH18	0.903225806	1	0.823529412
PTPRZ1 AND TENM1	0.9375	1	0.882352941
PTPRZ1 AND LHFPL2	0.903225806	1	0.823529412
PTPRZ1 AND NOT-LHFP	0.903225806	1	0.823529412
PTPRZ1 AND NOT-TIMM17B	0.9375	1	0.882352941
PTPRZ1 AND NOT-PRSS16	0.9375	1	0.882352941
PTPRZ1 AND NOT-MARCH6	0.903225806	1	0.823529412
PTPRZ1 AND NOT-RTN3	0.9375	1	0.882352941
PTPRZ1 AND NOT-MRVI1	0.9375	1	0.882352941
PTPRZ1 AND NOT-PCGF3	0.9375	1	0.882352941
PTPRZ1 AND BTN3A3	0.9375	1	0.882352941
PTPRZ1 AND NOT-ATP8A1	0.9375	1	0.882352941
PTPRZ1 AND NOT-CDS1	0.903225806	1	0.823529412
PTPRZ1 AND NOT-CDIPT	0.9375	1	0.882352941
PTPRZ1 AND NOT-TMEM147	0.903225806	1	0.823529412
PTPRZ1 AND NOT-LYPLA1	0.903225806	1	0.823529412
PTPRZ1 AND NOT-CD52	0.9375	1	0.882352941
PTPRZ1 AND NOT-TIMM17A	0.9375	1	0.882352941
PTPRZ1 AND NOT-SLC25A17	0.9375	1	0.882352941
PTPRZ1 AND NOT-VTI1B	0.9375	1	0.882352941
PTPRZ1 AND CRTAP	0.9375	1	0.882352941
PTPRZ1 AND NOT-PROCR	0.9375	1	0.882352941
PTPRZ1 AND PRDX4	0.9375	1	0.882352941
PTPRZ1 AND NOT-AGPAT2	0.9375	1	0.882352941
PTPRZ1 AND NOT-SPTLC1	0.9375	1	0.882352941
PTPRZ1 AND IFITM2	0.903225806	1	0.823529412
PTPRZ1 AND NOT-ERLIN1	0.9375	1	0.882352941
PTPRZ1 AND TNFSF13B	0.9375	1	0.882352941
PTPRZ1 AND NOT-B3GNT2	0.9375	1	0.882352941
PTPRZ1 AND CERS1	0.9375	1	0.882352941
PTPRZ1 AND NOT-NUP50	0.9375	1	0.882352941
PTPRZ1 AND NOT-HBS1L	0.903225806	1	0.823529412
PTPRZ1 AND NOT-ZMYND11	0.903225806	1	0.823529412
PTPRZ1 AND VAMP5	0.9375	1	0.882352941
PTPRZ1 AND NOT-CFTR	0.9375	1	0.882352941
PTPRZ1 AND NOT-FAXDC2	0.903225806	1	0.823529412
PTPRZ1 AND NOT-TSPAN9	0.9375	1	0.882352941
PTPRZ1 AND USP19	0.903225806	1	0.823529412
PTPRZ1 AND NOT-GPR83	0.9375	1	0.882352941
PTPRZ1 AND NOT-ADCY2	0.9375	1	0.882352941
PTPRZ1 AND NOT-ERP29	0.903225806	1	0.823529412
PTPRZ1 AND NOT-UQCR11	0.9375	1	0.882352941
PTPRZ1 AND NOT-ADCY3	0.9375	1	0.882352941
PTPRZ1 AND NOT-SLC27A2	0.903225806	1	0.823529412
PTPRZ1 AND NOT-RER1	0.9375	1	0.882352941
PTPRZ1 AND ADAM29	0.903225806	1	0.823529412
PTPRZ1 AND BTN3A2	0.903225806	1	0.823529412
PTPRZ1 AND BTN3A1	0.903225806	1	0.823529412
PTPRZ1 AND NOT-BTN2A1	0.9375	1	0.882352941
PTPRZ1 AND NOT-PTPRT	0.903225806	1	0.823529412
PTPRZ1 AND NOT-IL1RAPL1	0.903225806	1	0.823529412
PTPRZ1 AND NOT-PLA2G16	0.903225806	1	0.823529412
PTPRZ1 AND NOT-C14orf1	0.9375	1	0.882352941
PTPRZ1 AND CHI3L2	0.903225806	1	0.823529412
PTPRZ1 AND NOT-FZD10	0.9375	1	0.882352941
PTPRZ1 AND NOT-GALNT6	0.9375	1	0.882352941
PTPRZ1 AND NOT-SEC63	0.9375	1	0.882352941

Antigen Pair Logic	F1	Precision	Recall
(cont.)			
CHRNA3 AND NOT-GOLGB1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-GP2	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-GP5	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-IFNL2	0.95	1	0.904761905
CHRNA3 AND NOT-OR51B5	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-OR5J2	0.956521739	1	0.916666667
CHRNA3 AND NOT-GPR3	0.95	1	0.904761905
CHRNA3 AND NOT-XCR1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-OR8D1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-OR8D2	0.956521739	1	0.916666667
CHRNA3 AND NOT-OR10A4	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-NPBWR2	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-B4GALNT3	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-CXCR3	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-DPY19L2	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-TMPRSS12	0.943396226	1	0.892857143
CHRNA3 AND NOT-ZDHHC22	0.936708861	1	0.880952381
CHRNA3 AND NOT-GPR12	0.95	1	0.904761905
CHRNA3 AND NOT-GPR15	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-C17orf78	0.956521739	1	0.916666667
CHRNA3 AND NOT-IZUMO1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-SIGLECL1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-GPR20	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-OR2L13	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-GPR25	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-CCDC141	0.936708861	1	0.880952381
CHRNA3 AND NOT-DNAJC5G	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-PRRT3	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-CYP4V2	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-GPR32	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-RNF180	0.943396226	1	0.892857143
CHRNA3 AND NOT-GPR37	0.956521739	1	0.916666667
CHRNA3 AND NOT-CRB2	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-GPR39	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-FFAR1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-ILDR1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-GRIA3	0.956521739	1	0.916666667
CHRNA3 AND NOT-SLC6A16	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-GRIK1	0.95	1	0.904761905
CHRNA3 AND NOT-TMPRSS11E	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-ABO	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-GRIN1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-GRIN2A	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-GRIN2B	0.956521739	1	0.916666667
CHRNA3 AND NOT-GRM2	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-GRM3	0.936708861	1	0.880952381
CHRNA3 AND NOT-GRM5	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-GRM6	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-GRPR	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-CYP2S1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-ZDHHC1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-GUCA2A	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-GUCY2C	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-TRPM5	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-GUCY2F	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-PCDHB1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-SLC39A3	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-SLC39A2	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-GUCY2D	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-HFE	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-PLA2G2E	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-ST6GALNAC6	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-ERVW-1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-KCNIP3	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-MR1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-HRH2	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-SERPINA9	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-NDST1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-HTR1A	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-HTR1F	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-HTR5A	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-HTR6	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-KRTAP19-3	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-IFNE	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-339166?	0.950617284	0.987179487	0.916666667

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
PTPRZ1 AND NOT-PRRT2	0.9375	1	0.882352941
PTPRZ1 AND NOT-KLHL2	0.9375	1	0.882352941
PTPRZ1 AND MGAT4A	0.9375	1	0.882352941
VSIG4 AND NOT-CYB5A	0.903225806	1	0.823529412
PTPRZ1 AND NOT-SMIM12	0.9375	1	0.882352941
PTPRZ1 AND SDSL	0.903225806	1	0.823529412
PTPRZ1 AND NOT-PIK3IP1	0.9375	1	0.882352941
PTPRZ1 AND NOT-SLC2A13	0.9375	1	0.882352941
PTPRZ1 AND NOT-MAL2	0.903225806	1	0.823529412
PTPRZ1 AND CSMD2	0.903225806	1	0.823529412
PTPRZ1 AND NOT-XKR4	0.9375	1	0.882352941
PTPRZ1 AND TMEM132B	0.903225806	1	0.823529412
PTPRZ1 AND MARCH3	0.903225806	1	0.823529412
PTPRZ1 AND NOT-GPR146	0.9375	1	0.882352941
PTPRZ1 AND NOT-LYSMD3	0.9375	1	0.882352941
PTPRZ1 AND NOT-LRRC3B	0.903225806	1	0.823529412
PTPRZ1 AND NOT-FAM210B	0.903225806	1	0.823529412
PTPRZ1 AND NOT-GINM1	0.903225806	1	0.823529412
PTPRZ1 AND TM4SF18	0.903225806	1	0.823529412
PTPRZ1 AND NOT-SLC18B1	0.9375	1	0.882352941
PTPRZ1 AND NOT-CLCN4	0.903225806	1	0.823529412
PTPRZ1 AND ANTXR2	0.903225806	1	0.823529412
PTPRZ1 AND NOT-ZFYVE27	0.9375	1	0.882352941
PTPRZ1 AND NOT-SLC36A4	0.903225806	1	0.823529412
PTPRZ1 AND FAT3	0.903225806	1	0.823529412
PTPRZ1 AND NOT-TMEM45B	0.9375	1	0.882352941
PTPRZ1 AND NOT-CLN5	0.9375	1	0.882352941
NOT-PTPRZ1 AND PTPRZ1	0.9375	1	0.882352941
PTPRZ1 AND NOT-SPPL3	0.9375	1	0.882352941
PTPRZ1 AND CCR1	0.903225806	1	0.823529412
PTPRZ1 AND NOT-C15orf27	0.9375	1	0.882352941
PTPRZ1 AND NOT-NIPA1	0.903225806	1	0.823529412
PTPRZ1 AND NOT-TMEM219	0.9375	1	0.882352941
PTPRZ1 AND NOT-CYB5D2	0.9375	1	0.882352941
PTPRZ1 AND NOT-SPNS2	0.9375	1	0.882352941
PTPRZ1 AND FAM69C	0.903225806	1	0.823529412
PTPRZ1 AND B3GALT6	0.903225806	1	0.823529412
PTPRZ1 AND CNR1	0.9375	1	0.882352941
PTPRZ1 AND SLC44A3	0.9375	1	0.882352941
PTPRZ1 AND NOT-CNTN1	0.9375	1	0.882352941
PTPRZ1 AND DRAM2	0.9375	1	0.882352941
PTPRZ1 AND C1orf162	0.9375	1	0.882352941
PTPRZ1 AND NOT-TBC1D20	0.9375	1	0.882352941
PTPRZ1 AND NOT-PIGU	0.9375	1	0.882352941
PTPRZ1 AND EMID1	0.903225806	1	0.823529412
PTPRZ1 AND NOT-MBOAT2	0.9375	1	0.882352941
PTPRZ1 AND NOT-TMEM18	0.903225806	1	0.823529412
PTPRZ1 AND NOT-SGPP2	0.903225806	1	0.823529412
PTPRZ1 AND NOT-ACVR1C	0.903225806	1	0.823529412
PTPRZ1 AND NOT-TMEM178A	0.903225806	1	0.823529412
PTPRZ1 AND NOT-COL17A1	0.9375	1	0.882352941
PTPRZ1 AND COMT	0.903225806	1	0.823529412
PTPRZ1 AND NOT-TMEM42	0.9375	1	0.882352941
PTPRZ1 AND NOT-SLC31A1	0.903225806	1	0.823529412
PTPRZ1 AND NOT-SLC31A2	0.903225806	1	0.823529412
PTPRZ1 AND NOT-SYNPR	0.903225806	1	0.823529412
PTPRZ1 AND NOT-SLC9B2	0.903225806	1	0.823529412
PTPRZ1 AND B3GAT2	0.903225806	1	0.823529412
PTPRZ1 AND NOT-TMEM68	0.9375	1	0.882352941
PTPRZ1 AND NOT-HGSNAT	0.9375	1	0.882352941
PTPRZ1 AND NOT-FAM69B	0.9375	1	0.882352941
PTPRZ1 AND NOT-SLITRK4	0.903225806	1	0.823529412
PTPRZ1 AND PTCHD1	0.9375	1	0.882352941
LTBP3 AND NOT-SELM	0.909090909	0.9375	0.882352941
PTPRZ1 AND NOT-SELM	0.9375	1	0.882352941
PTPRZ1 AND ROMO1	0.9375	1	0.882352941
PTPRZ1 AND NOT-CRY2	0.9375	1	0.882352941
TPST1 AND NOT-CRY2	0.903225806	1	0.823529412
PTPRZ1 AND ADORA3	0.903225806	1	0.823529412
PTPRZ1 AND SAMD8	0.9375	1	0.882352941
PTPRZ1 AND NOT-FAM76B	0.9375	1	0.882352941
PTPRZ1 AND CSF1R	0.903225806	1	0.823529412
PTPRZ1 AND NOT-ALG10B	0.9375	1	0.882352941
PTPRZ1 AND B3GLCT	0.903225806	1	0.823529412
PTPRZ1 AND NOT-CMTM4	0.9375	1	0.882352941
PTPRZ1 AND NOT-RBFOX3	0.903225806	1	0.823529412

Antigen Pair Logic	F1	Precision	Recall
CHRNA3 AND NOT-NAT8L	0.936708861	1	0.880952381
CHRNA3 AND NOT-SLC6A19	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-VSIG1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-GLDN	0.943396226	1	0.892857143
CHRNA3 AND NOT-AQP8	0.956521739	1	0.916666667
CHRNA3 AND NOT-ATP13A5	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-IFNB1	0.956521739	1	0.916666667
CHRNA3 AND NOT-MOGAT3	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-CATSPER3	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-FAM159A	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-HCN1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-IHH	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-FASLG	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-IL6R	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-IL12B	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-IL12RB2	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-AQP4	0.956521739	1	0.916666667
CHRNA3 AND NOT-AQP5	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-ITGA9	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-ITGAD	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-ITGB3	0.95	1	0.904761905
CHRNA3 AND NOT-ITGB6	0.943396226	1	0.892857143
CHRNA3 AND NOT-ITPR2	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-ITPR3	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-KCNA1	0.956521739	1	0.916666667
CHRNA3 AND NOT-KCNA2	0.956521739	1	0.916666667
CHRNA3 AND NOT-KCNA5	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-KCNA6	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-GALNT18	0.956521739	1	0.916666667
CHRNA3 AND NOT-KCNC1	0.95	1	0.904761905
CHRNA3 AND NOT-SPEM1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-KCNC2	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-KCNC3	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-C1orf95	0.936708861	1	0.880952381
CHRNA3 AND NOT-KCND1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-KCND3	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-KCNE1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-LHFPL3	0.956521739	1	0.916666667
CHRNA3 AND NOT-PNPLA7	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-KCNJ1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-KCNJ4	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-KCNJ6	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-KCNJ9	0.956521739	1	0.916666667
CHRNA3 AND NOT-KCNJ10	0.956521739	1	0.916666667
CHRNA3 AND NOT-KCNJ11	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-KCNJ13	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-KCNJ14	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-KCNK1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-KCNK2	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-KCNMA1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-KCNN3	0.95	1	0.904761905
CHRNA3 AND NOT-KCNS1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-KCNS2	0.95	1	0.904761905
CHRNA3 AND NOT-KEL	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-KIR2DL4	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-KRT5	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-C10orf99	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-CLEC12B	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-SCIMP	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-RPRML	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-RGS9BP	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-FAM132A	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-FAM209B	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-SPATA31D1	0.956521739	1	0.916666667
CHRNA3 AND NOT-OR51B6	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-OR51I1	0.943396226	1	0.892857143
CHRNA3 AND NOT-OR51I2	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-OR52D1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-OR7A10	0.936708861	1	0.880952381
CHRNA3 AND NOT-LCT	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-LETM1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-LHCGR	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-LIM2	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-FADS3	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-C11orf87	0.956521739	1	0.916666667
CHRNA3 AND NOT-C14orf180	0.950617284	0.987179487	0.916666667

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
PTPRZ1 AND NOT-ANKRD29	0.903225806	1	0.823529412
PTPRZ1 AND NOT-APCDD1	0.9375	1	0.882352941
PTPRZ1 AND NOT-SMIM17	0.903225806	1	0.823529412
PTPRZ1 AND NOT-MFSD4	0.903225806	1	0.823529412
PTPRZ1 AND NOT-SLC30A7	0.9375	1	0.882352941
PTPRZ1 AND CTNS	0.903225806	1	0.823529412
PTPRZ1 AND NOT-KANSL1L	0.9375	1	0.882352941
PTPRZ1 AND NOT-RMDN2	0.9375	1	0.882352941
PTPRZ1 AND NOT-GPR155	0.9375	1	0.882352941
PTPRZ1 AND NOT-XXYLT1	0.9375	1	0.882352941
PTPRZ1 AND NOT-CUX1	0.903225806	1	0.823529412
PTPRZ1 AND IGSF11	0.9375	1	0.882352941
PTPRZ1 AND PAQR3	0.903225806	1	0.823529412
PTPRZ1 AND NOT-CXADR	0.9375	1	0.882352941
PTPRZ1 AND NOT-CYB5A	0.903225806	1	0.823529412
PTPRZ1 AND NOT-ADRA2C	0.9375	1	0.882352941
PTPRZ1 AND NOT-CYB561	0.9375	1	0.882352941
PTPRZ1 AND CYBA	0.903225806	1	0.823529412
PTPRZ1 AND NOT-RNF145	0.9375	1	0.882352941
PTPRZ1 AND NOT-ADRB1	0.903225806	1	0.823529412
PTPRZ1 AND NOT-SLC2A12	0.9375	1	0.882352941
PTPRZ1 AND NOT-RNF217	0.9375	1	0.882352941
NRCAM AND NOT-CLEC2L	0.903225806	1	0.823529412
PTPRZ1 AND NOT-CLEC2L	0.9375	1	0.882352941
PTPRZ1 AND NOT-TMEM65	0.9375	1	0.882352941
PTPRZ1 AND RDH10	0.903225806	1	0.823529412
PTPRZ1 AND NOT-MOSPD2	0.903225806	1	0.823529412
PTPRZ1 AND NOT-FAM134C	0.9375	1	0.882352941
PTPRZ1 AND NOT-CNST	0.903225806	1	0.823529412
PTPRZ1 AND NOT-TMEM64	0.9375	1	0.882352941
PTPRZ1 AND NOT-SPTSSA	0.903225806	1	0.823529412
PTPRZ1 AND NOT-ABHD3	0.9375	1	0.882352941
PTPRZ1 AND NOT-DHCR7	0.903225806	1	0.823529412
PTPRZ1 AND DSCAM	0.903225806	1	0.823529412
PTPRZ1 AND DTNA	0.9375	1	0.882352941
PTPRZ1 AND NOT-AGTR1	0.9375	1	0.882352941
PTPRZ1 AND GPR183	0.903225806	1	0.823529412
PTPRZ1 AND NOT-LPAR1	0.903225806	1	0.823529412
PTPRZ1 AND EDNRB	0.9375	1	0.882352941
PTPRZ1 AND NOT-HIGD2A	0.9375	1	0.882352941
PTPRZ1 AND NOT-CELSR2	0.903225806	1	0.823529412
PTPRZ1 AND NOT-MPZL3	0.9375	1	0.882352941
PTPRZ1 AND ABCA1	0.903225806	1	0.823529412
PTPRZ1 AND EMP1	0.903225806	1	0.823529412
PTPRZ1 AND NOT-EMP2	0.9375	1	0.882352941
TPST1 AND NOT-EMP2	0.903225806	1	0.823529412
PTPRZ1 AND NOT-SMIM14	0.9375	1	0.882352941
NOT-PTPRZ1 AND PTPRZ1	0.9375	1	0.882352941
PTPRZ1 AND NOT-SLC25A43	0.9375	1	0.882352941
PTPRZ1 AND NOT-EPHA1	0.9375	1	0.882352941
PTPRZ1 AND NOT-CERS3	0.9375	1	0.882352941
PTPRZ1 AND NOT-EPHA7	0.903225806	1	0.823529412
PTPRZ1 AND NOT-EPHB6	0.903225806	1	0.823529412
PTPRZ1 AND NOT-CLN8	0.903225806	1	0.823529412
PTPRZ1 AND NOT-ETV6	0.9375	1	0.882352941
PTPRZ1 AND NOT-EVI2A	0.903225806	1	0.823529412
PTPRZ1 AND NOT-EXTL2	0.9375	1	0.882352941
PTPRZ1 AND NOT-F10	0.9375	1	0.882352941
PTPRZ1 AND NOT-FAAH	0.903225806	1	0.823529412
PTPRZ1 AND NOT-ACSL1	0.903225806	1	0.823529412
PTPRZ1 AND NOT-FAT1	0.9375	1	0.882352941
PTPRZ1 AND NOT-FAT2	0.9375	1	0.882352941
PTPRZ1 AND MPEG1	0.903225806	1	0.823529412
PTPRZ1 AND ABCA3	0.903225806	1	0.823529412
PTPRZ1 AND CYB561A3	0.9375	1	0.882352941
PTPRZ1 AND HEPACAM	0.9375	1	0.882352941
PTPRZ1 AND SLC16A9	0.9375	1	0.882352941
PTPRZ1 AND NOT-MARCH8	0.9375	1	0.882352941
PTPRZ1 AND NOT-SLC39A12	0.903225806	1	0.823529412
PTPRZ1 AND NOT-C6orf136	0.9375	1	0.882352941
PTPRZ1 AND FKTN	0.9375	1	0.882352941
PTPRZ1 AND NOT-THSD7A	0.9375	1	0.882352941
PTPRZ1 AND TMED4	0.903225806	1	0.823529412
GPM6A AND NOT-TMEM130	0.903225806	1	0.823529412
PTPRZ1 AND NOT-TMEM130	0.903225806	1	0.823529412
PTPRZ1 AND NOT-ALDH3A2	0.9375	1	0.882352941

Antigen Pair Logic	F1	Precision	Recall
CHRNA3 AND NOT-LMO7	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-HACD4	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-XKRX	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-IFITM10	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-LRP2	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-LRP4	0.95	1	0.904761905
CHRNA3 AND NOT-HAPLN4	0.956521739	1	0.916666667
CHRNA3 AND NOT-SPINK6	0.956521739	1	0.916666667
CHRNA3 AND NOT-LSAMP	0.95	1	0.904761905
CHRNA3 AND NOT-MAL	0.95	1	0.904761905
CHRNA3 AND NOT-STS	0.956521739	1	0.916666667
CHRNA3 AND NOT-LCN10	0.943396226	1	0.892857143
CHRNA3 AND NOT-MC5R	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-CD46	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-ADAM11	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-MEP1B	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-MGAT3	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-CTAGE5	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-MGST2	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-MIP	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-TRPM1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-MME	0.95	1	0.904761905
CHRNA3 AND NOT-MOG	0.956521739	1	0.916666667
CHRNA3 AND NOT-LRRC52	0.956521739	1	0.916666667
MARCH11 AND CHRNA3	0.952380952	0.952380952	0.952380952
CHRNA3 AND NOT-OR2B3	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-OR14J1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-OR10C1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-PTCHD4	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-MTHFD1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-MTNR1A	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-MUC1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-ATP1A2	0.956521739	1	0.916666667
CHRNA3 AND NOT-ATP12A	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-NINJ2	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-NMBR	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-NOTCH2	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-NPHS1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-NPY5R	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-NTRK2	0.956521739	1	0.916666667
CHRNA3 AND NOT-ATP2B2	0.956521739	1	0.916666667
CHRNA3 AND NOT-NTSR1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-ATP2B3	0.936708861	1	0.880952381
CHRNA3 AND NOT-GPR143	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-OMG	0.95	1	0.904761905
CHRNA3 AND NOT-OPCML	0.956521739	1	0.916666667
CHRNA3 AND NOT-OPRK1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-OR1D2	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-OR1F1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-OR2C1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-CLDN11	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-P2RY1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-P2RY2	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-GALNT9	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-IL22	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-ANO7	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-TAS2R3	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-TAS2R9	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-TAS2R8	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-TAS2R7	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-TAS2R13	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-NTM	0.956521739	1	0.916666667
CHRNA3 AND NOT-IMPG2	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-PCDH1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-SLC45A2	0.956521739	1	0.916666667
CHRNA3 AND NOT-CLDN18	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-CLEC1A	0.95	1	0.904761905
CHRNA3 AND NOT-RXFP3	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-GCNT4	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-KCNK9	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-PDCD1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-CHST15	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-WNT16	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-RHCG	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-ENPP3	0.943396226	1	0.892857143
CHRNA3 AND NOT-GHRL	0.950617284	0.987179487	0.916666667

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
PTPRZ1 AND NOT-FGFR2	0.9375	1	0.882352941
PTPRZ1 AND NLGN1	0.9375	1	0.882352941
NRCAM AND NOT-CLSTN1	0.903225806	1	0.823529412
PTPRZ1 AND NOT-CLSTN1	0.903225806	1	0.823529412
PTPRZ1 AND NOT-PCNX	0.9375	1	0.882352941
PTPRZ1 AND KLHDC10	0.9375	1	0.882352941
PTPRZ1 AND NOT-FAIM2	0.903225806	1	0.823529412
PTPRZ1 AND NFASC	0.903225806	1	0.823529412
PTPRZ1 AND ANKLE2	0.903225806	1	0.823529412
PTPRZ1 AND NOT-LRCH1	0.9375	1	0.882352941
PTPRZ1 AND GRAMD4	0.9375	1	0.882352941
PTPRZ1 AND STAB1	0.903225806	1	0.823529412
PTPRZ1 AND SYT11	0.9375	1	0.882352941
PTPRZ1 AND MLC1	0.903225806	1	0.823529412
PTPRZ1 AND ASTN2	0.9375	1	0.882352941
PTPRZ1 AND WSCD1	0.903225806	1	0.823529412
PTPRZ1 AND NOT-DNAJC16	0.9375	1	0.882352941
PTPRZ1 AND NOT-ESYT1	0.903225806	1	0.823529412
PTPRZ1 AND NOT-SLC35A3	0.9375	1	0.882352941
PTPRZ1 AND NOT-ABCB10	0.9375	1	0.882352941
PTPRZ1 AND NPTXR	0.903225806	1	0.823529412
PTPRZ1 AND TRAM1	0.9375	1	0.882352941
PTPRZ1 AND NOT-LEPROTL1	0.9375	1	0.882352941
PTPRZ1 AND NOT-PIGN	0.9375	1	0.882352941
PTPRZ1 AND FPR1	0.9375	1	0.882352941
PTPRZ1 AND SLC7A11	0.9375	1	0.882352941
PTPRZ1 AND CADM1	0.9375	1	0.882352941
PTPRZ1 AND NOT-TMEM245	0.9375	1	0.882352941
PTPRZ1 AND NOT-EML2	0.9375	1	0.882352941
PTPRZ1 AND PANX1	0.903225806	1	0.823529412
PTPRZ1 AND NOT-FUT8	0.9375	1	0.882352941
PTPRZ1 AND NOT-EPHX4	0.903225806	1	0.823529412
PTPRZ1 AND NOT-KDSR	0.9375	1	0.882352941
PTPRZ1 AND NOT-ACKR1	0.903225806	1	0.823529412
PTPRZ1 AND CADM2	0.903225806	1	0.823529412
NRCAM AND NOT-LPCAT4	0.903225806	1	0.823529412
PTPRZ1 AND NOT-LPCAT4	0.9375	1	0.882352941
PTPRZ1 AND NOT-TMEM256	0.903225806	1	0.823529412
PTPRZ1 AND NOT-GABRA4	0.903225806	1	0.823529412
NRCAM AND NOT-GABRB2	0.903225806	1	0.823529412
PTPRZ1 AND NOT-GABRB2	0.903225806	1	0.823529412
NRCAM AND NOT-GABRD	0.903225806	1	0.823529412
PTPRZ1 AND NOT-MFSD8	0.903225806	1	0.823529412
PTPRZ1 AND NOT-GABRG1	0.903225806	1	0.823529412
PTPRZ1 AND NOT-KIAA1549L	0.9375	1	0.882352941
PTPRZ1 AND NOT-SUN2	0.903225806	1	0.823529412
PTPRZ1 AND SLC39A6	0.9375	1	0.882352941
PTPRZ1 AND NOT-GALC	0.9375	1	0.882352941
PTPRZ1 AND NOT-TMEM184B	0.9375	1	0.882352941
PTPRZ1 AND NOT-YIPF3	0.9375	1	0.882352941
PTPRZ1 AND NOT-PARM1	0.903225806	1	0.823529412
PTPRZ1 AND NOT-ABHD14A	0.903225806	1	0.823529412
PTPRZ1 AND NOT-GALNS	0.9375	1	0.882352941
PTPRZ1 AND NOT-GALNT3	0.9375	1	0.882352941
PTPRZ1 AND NOT-ZDHHC5	0.9375	1	0.882352941
PTPRZ1 AND NOT-NALCN	0.903225806	1	0.823529412
PTPRZ1 AND NOT-PVRL3	0.903225806	1	0.823529412
PTPRZ1 AND NOT-PNKD	0.903225806	1	0.823529412
PTPRZ1 AND NOT-ADGRA2	0.9375	1	0.882352941
PTPRZ1 AND NOT-CYP4X1	0.903225806	1	0.823529412
PTPRZ1 AND NOT-SUSD5	0.903225806	1	0.823529412
PTPRZ1 AND NOT-GLCE	0.9375	1	0.882352941
PTPRZ1 AND LRRTM2	0.9375	1	0.882352941
PTPRZ1 AND NOT-CNTNAP2	0.903225806	1	0.823529412
PTPRZ1 AND NOT-SLITRK5	0.903225806	1	0.823529412
PTPRZ1 AND GIMAP2	0.903225806	1	0.823529412
PTPRZ1 AND NOT-NPHP4	0.903225806	1	0.823529412
PTPRZ1 AND TMEM251	0.903225806	1	0.823529412
PTPRZ1 AND NOT-FAM162A	0.9375	1	0.882352941
NRCAM AND NOT-SEZ6L2	0.903225806	1	0.823529412
PTPRZ1 AND NOT-CNNM4	0.9375	1	0.882352941
PTPRZ1 AND NOT-MYOF	0.9375	1	0.882352941
NRCAM AND NOT-HS6ST3	0.903225806	1	0.823529412
PTPRZ1 AND NOT-HS6ST3	0.903225806	1	0.823529412
PTPRZ1 AND AMFR	0.903225806	1	0.823529412
PTPRZ1 AND NOT-GJA1	0.9375	1	0.882352941

Antigen Pair Logic	F1	Precision	Recall
CHRNA3 AND NOT-CD244	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-ASIC5	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-ATP8B1	0.950617284	0.987179487	0.916666667
CHRNA3 AND PFN2	0.950617284	0.987179487	0.916666667
CHRNA3 AND SLC25A3	0.956521739	1	0.916666667
CHRNA3 AND NOT-PHEX	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-SERPINB13	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-IL17D	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-TM6SF2	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-LRP1B	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-TPCN1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-FXYD4	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-FGFRL1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-DUOX1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-SLCO1C1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-SMIM11	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-GPR88	0.950617284	0.987179487	0.916666667
CHRNA3 AND SLC38A2	0.936708861	1	0.880952381
CHRNA3 AND NOT-RNF186	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-DLL4	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-UGT1A8	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-VSIG10	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-DCHS2	0.956521739	1	0.916666667
CHRNA3 AND NOT-SIDT1	0.943396226	1	0.892857143
CHRNA3 AND NOT-GDPD2	0.936708861	1	0.880952381
CHRNA3 AND NOT-MS4A12	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-DNAJC25	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-RETSAT	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-PQLC2	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-C1orf159	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-SLC52A1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-PPP1R3A	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-KIRREL	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-TMEM39A	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-TMEM143	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-AVPR1A	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-TMEM144	0.95	1	0.904761905
CHRNA3 AND NOT-VNN3	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-LGR4	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-AVPR1B	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-ST6GALNAC1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-GABRQ	0.943396226	1	0.892857143
CHRNA3 AND NOT-GPRC5C	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-TMEM234	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-PCDHGC5	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-PCDHGC4	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-PCDHGA10	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-PCDHGA9	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-PCDHB9	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-PCDHAC2	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-PCDHAC1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-BTNL2	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-SLC7A10	0.956521739	1	0.916666667
CHRNA3 AND NOT-TMPRSS4	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-CLDND1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-KCNK13	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-PANX2	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-MUC13	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-AGPAT3	0.956521739	1	0.916666667
CHRNA3 AND NOT-OTOR	0.936708861	1	0.880952381
CHRNA3 AND NOT-C2orf83	0.936708861	1	0.880952381
CHRNA3 AND NOT-NMUR2	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-GPR108	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-PYY	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-ANO2	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-TRPC7	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-JPH2	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-GJC2	0.943396226	1	0.892857143
CHRNA3 AND NOT-SLC45A4	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-TAS2R38	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-PTGDR	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-DSCAML1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-HHATL	0.956521739	1	0.916666667
CHRNA3 AND NOT-SORCS2	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-PTCHD2	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-NCEH1	0.950617284	0.987179487	0.916666667

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
PTPRZ1 AND NOT-SEC22A	0.903225806	1	0.823529412
PTPRZ1 AND NOT-ATP2C1	0.9375	1	0.882352941
PTPRZ1 AND NOT-GHITM	0.903225806	1	0.823529412
PTPRZ1 AND NOT-GJB2	0.9375	1	0.882352941
PTPRZ1 AND NOT-NAAA	0.9375	1	0.882352941
PTPRZ1 AND PCDH17	0.903225806	1	0.823529412
PTPRZ1 AND NOT-GCLC	0.9375	1	0.882352941
PTPRZ1 AND GOLIM4	0.903225806	1	0.823529412
PTPRZ1 AND NOT-TMEM97	0.9375	1	0.882352941
PTPRZ1 AND NOT-TOR1B	0.903225806	1	0.823529412
PTPRZ1 AND NOT-TOR2A	0.9375	1	0.882352941
PTPRZ1 AND NOT-GLRB	0.903225806	1	0.823529412
PTPRZ1 AND NOT-GPD2	0.903225806	1	0.823529412
PTPRZ1 AND NOT-SLCO4A1	0.9375	1	0.882352941
PTPRZ1 AND SLCO1B3	0.903225806	1	0.823529412
PTPRZ1 AND GPM6B	0.903225806	1	0.823529412
GPM6B AND GAL3ST4	0.914285714	0.88888889	0.941176471
PTPRZ1 AND NOT-AQP11	0.9375	1	0.882352941
PTPRZ1 AND NOT-BRICD5	0.903225806	1	0.823529412
PTPRZ1 AND NOT-ANG	0.9375	1	0.882352941
PTPRZ1 AND NOT-PIGW	0.9375	1	0.882352941
PTPRZ1 AND NOT-GDPD1	0.903225806	1	0.823529412
PTPRZ1 AND NOT-GPR22	0.903225806	1	0.823529412
PTPRZ1 AND NOT-CYB561D1	0.9375	1	0.882352941
PTPRZ1 AND NOT-RNF149	0.9375	1	0.882352941
PTPRZ1 AND NOT-GPR26	0.903225806	1	0.823529412
PTPRZ1 AND NOT-EOGT	0.9375	1	0.882352941
PTPRZ1 AND NOT-CYP4V2	0.9375	1	0.882352941
PTPRZ1 AND GPR34	0.903225806	1	0.823529412
PTPRZ1 AND XKR6	0.903225806	1	0.823529412
PTPRZ1 AND NOT-MICU3	0.9375	1	0.882352941
PTPRZ1 AND GPR37	0.903225806	1	0.823529412
PTPRZ1 AND NOT-MLNR	0.9375	1	0.882352941
PTPRZ1 AND NOT-YIPF6	0.903225806	1	0.823529412
PTPRZ1 AND GRIA3	0.9375	1	0.882352941
PTPRZ1 AND NOT-DEXI	0.9375	1	0.882352941
PTPRZ1 AND TMEM176B	0.903225806	1	0.823529412
PTPRZ1 AND NOT-OSTM1	0.9375	1	0.882352941
PTPRZ1 AND NOT-FLVCR1	0.903225806	1	0.823529412
PTPRZ1 AND NOT-MAGEH1	0.903225806	1	0.823529412
PTPRZ1 AND GRIK2	0.903225806	1	0.823529412
PTPRZ1 AND NOT-GRIN2A	0.903225806	1	0.823529412
PTPRZ1 AND GRIN2C	0.903225806	1	0.823529412
PTPRZ1 AND NOT-ORMDL2	0.9375	1	0.882352941
PTPRZ1 AND NOT-DNAJC15	0.903225806	1	0.823529412
PTPRZ1 AND NOT-GRM3	0.903225806	1	0.823529412
PTPRZ1 AND NOT-GRM5	0.903225806	1	0.823529412
PTPRZ1 AND NOT-GRM7	0.9375	1	0.882352941
PTPRZ1 AND NOT-UBIAD1	0.903225806	1	0.823529412
PTPRZ1 AND NOT-TRHDE	0.903225806	1	0.823529412
PTPRZ1 AND CERS2	0.9375	1	0.882352941
PTPRZ1 AND NOT-SLC2A8	0.903225806	1	0.823529412
PTPRZ1 AND NOT-HAS1	0.9375	1	0.882352941
PTPRZ1 AND NOT-HDAC2	0.9375	1	0.882352941
PTPRZ1 AND NOT-ST6GALNAC6	0.903225806	1	0.823529412
PTPRZ1 AND HLA-B	0.9375	1	0.882352941
PTPRZ1 AND HLA-DMA	0.903225806	1	0.823529412
PTPRZ1 AND HLA-DMB	0.903225806	1	0.823529412
PTPRZ1 AND HLA-DRA	0.903225806	1	0.823529412
PTPRZ1 AND HLA-DRB1	0.903225806	1	0.823529412
PTPRZ1 AND HLA-F	0.903225806	1	0.823529412
PTPRZ1 AND NOT-HMOX2	0.9375	1	0.882352941
PTPRZ1 AND NOT-NDST1	0.9375	1	0.882352941
PTPRZ1 AND APLP2	0.9375	1	0.882352941
PTPRZ1 AND NOT-HTR2A	0.903225806	1	0.823529412
TNC AND NOT-AOC3	0.903225806	1	0.823529412
PTPRZ1 AND TMEM119	0.903225806	1	0.823529412
PTPRZ1 AND NOT-FAM19A2	0.903225806	1	0.823529412
PTPRZ1 AND NOT-NAT8L	0.9375	1	0.882352941
PTPRZ1 AND TSPAN33	0.903225806	1	0.823529412
PTPRZ1 AND APOC1	0.9375	1	0.882352941
PTPRZ1 AND NOT-FAM174A	0.9375	1	0.882352941
PTPRZ1 AND IGF1R	0.9375	1	0.882352941
PTPRZ1 AND NOT-HCN1	0.903225806	1	0.823529412
PTPRZ1 AND NOT-SLCO4C1	0.903225806	1	0.823529412
PTPRZ1 AND NOT-IL1R1	0.9375	1	0.882352941

Antigen Pair Logic	F1	Precision	Recall
CHRNA3 AND NOT-ADGRB1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-DPP10	0.936708861	1	0.880952381
CHRNA3 AND NOT-G6PC2	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-CATSPERG	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-SLC46A2	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-PVRL2	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-LY6G6D	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-ACE2	0.956521739	1	0.916666667
CHRNA3 AND NOT-CACNG8	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-CACNG7	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-CACNG6	0.943396226	1	0.892857143
CHRNA3 AND NOT-HRH4	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-EDA2R	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-CDH26	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-NYX	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-KCNK15	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-ROS1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-CTXN3	0.956521739	1	0.916666667
CHRNA3 AND NOT-RRBP1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-SLC22A23	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-SCN1A	0.956521739	1	0.916666667
CHRNA3 AND NOT-SCN1B	0.936708861	1	0.880952381
CHRNA3 AND NOT-SCN4A	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-SCN4B	0.936708861	1	0.880952381
CHRNA3 AND NOT-SCN8A	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-SCN10A	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-SCNN1A	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-SCNN1B	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-SCNN1D	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-SCNN1G	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-CEACAM1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-CX3CL1	0.956521739	1	0.916666667
CHRNA3 AND NOT-63914?	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-CDH23	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-TNMD	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-XYLT1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-ROBO3	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-MS4A5	0.95	1	0.904761905
CHRNA3 AND NOT-CCDC168	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-WBSCR17	0.956521739	1	0.916666667
CHRNA3 AND NOT-SFTPC	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-SFTPD	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-SGCA	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-SGCD	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-SMIM10	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-CSMD1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-NDST4	0.956521739	1	0.916666667
CHRNA3 AND NOT-GPR135	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-SYNDIG1L	0.95	1	0.904761905
CHRNA3 AND NOT-CTAGE1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-SHH	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-LMF1	0.956521739	1	0.916666667
CHRNA3 AND NOT-SLC13A3	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-ST3GAL4	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-LRRC19	0.956521739	1	0.916666667
CHRNA3 AND NOT-SLAMF1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-SLC1A3	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-SLC4A3	0.95	1	0.904761905
CHRNA3 AND NOT-SLC1A7	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-SLC2A4	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-PCDH15	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-SLC5A1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-SLC5A4	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-SLC6A1	0.956521739	1	0.916666667
CHRNA3 AND NOT-SLC6A4	0.95	1	0.904761905
CHRNA3 AND NOT-SLC6A12	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-SLC6A13	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-SLC8A2	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-LRTM2	0.956521739	1	0.916666667
CHRNA3 AND NOT-SLC8A3	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-SLC9A2	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-SLC9A5	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-SLC12A3	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-SLC13A1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-SLC15A2	0.956521739	1	0.916666667
CHRNA3 AND NOT-SLC34A1	0.950617284	0.987179487	0.916666667

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
PTPRZ1 AND NOT-IL4R	0.9375	1	0.882352941
PTPRZ1 AND IL6R	0.903225806	1	0.823529412
PTPRZ1 AND IL10RA	0.903225806	1	0.823529412
PTPRZ1 AND IL13RA1	0.903225806	1	0.823529412
PTPRZ1 AND NOT-IL15RA	0.9375	1	0.882352941
PTPRZ1 AND NOT-AQP3	0.903225806	1	0.823529412
PTPRZ1 AND AQP4	0.9375	1	0.882352941
PTPRZ1 AND NOT-INPP4A	0.9375	1	0.882352941
PTPRZ1 AND NOT-INSIG1	0.903225806	1	0.823529412
PTPRZ1 AND NOT-AQP7	0.9375	1	0.882352941
PTPRZ1 AND ITGA2	0.9375	1	0.882352941
PTPRZ1 AND ITGAM	0.903225806	1	0.823529412
PTPRZ1 AND NOT-JAG2	0.9375	1	0.882352941
PTPRZ1 AND NOT-KCNA1	0.903225806	1	0.823529412
PTPRZ1 AND TMEM179B	0.903225806	1	0.823529412
PTPRZ1 AND NOT-KCNB1	0.9375	1	0.882352941
PTPRZ1 AND NOT-KCNC1	0.9375	1	0.882352941
NOT-PTPRZ1 AND PTPRZ1	0.9375	1	0.882352941
PTPRZ1 AND NOT-TMEM205	0.9375	1	0.882352941
PTPRZ1 AND NOT-FAM73A	0.9375	1	0.882352941
PTPRZ1 AND NOT-MIA3	0.9375	1	0.882352941
PTPRZ1 AND NOT-C1orf95	0.9375	1	0.882352941
PTPRZ1 AND LHFPL4	0.9375	1	0.882352941
PTPRZ1 AND LHFPL3	0.903225806	1	0.823529412
PTPRZ1 AND NOT-ENHO	0.9375	1	0.882352941
NRCAM AND NOT-KCNJ3	0.903225806	1	0.823529412
PTPRZ1 AND NOT-KCNJ6	0.9375	1	0.882352941
NOT-PTPRZ1 AND PTPRZ1	0.9375	1	0.882352941
PTPRZ1 AND KCNJ8	0.903225806	1	0.823529412
PTPRZ1 AND KCNJ10	0.9375	1	0.882352941
PTPRZ1 AND NOT-KCNN2	0.903225806	1	0.823529412
PTPRZ1 AND NOT-GPR153	0.9375	1	0.882352941
PTPRZ1 AND C10orf99	0.9375	1	0.882352941
PTPRZ1 AND NOT-SHISA2	0.903225806	1	0.823529412
PTPRZ1 AND GLTPD2	0.903225806	1	0.823529412
PTPRZ1 AND NOT-USP27X	0.9375	1	0.882352941
PTPRZ1 AND OR51B6	0.903225806	1	0.823529412
PTPRZ1 AND NOT-LAMP1	0.9375	1	0.882352941
PTPRZ1 AND NOT-LDLR	0.9375	1	0.882352941
PTPRZ1 AND NOT-LEPR	0.9375	1	0.882352941
PTPRZ1 AND NOT-TMEM200B	0.9375	1	0.882352941
NRCAM AND NOT-C11orf87	0.903225806	1	0.823529412
PTPRZ1 AND NOT-C11orf87	0.9375	1	0.882352941
PTPRZ1 AND NOT-LMO7	0.903225806	1	0.823529412
PTPRZ1 AND NOT-C3orf80	0.903225806	1	0.823529412
PTPRZ1 AND NOT-C4orf3	0.9375	1	0.882352941
PTPRZ1 AND NOT-LRP3	0.9375	1	0.882352941
PTPRZ1 AND NOT-LRP4	0.903225806	1	0.823529412
PTPRZ1 AND LTBP3	0.903225806	1	0.823529412
LTBP3 AND NOT-DNAJC1	0.9375	1	0.882352941
PTPRZ1 AND NOT-LY75	0.9375	1	0.882352941
PTPRZ1 AND NOT-TM4SF1	0.903225806	1	0.823529412
PTPRZ1 AND NOT-EPCAM	0.903225806	1	0.823529412
PTPRZ1 AND NOT-FAM19A1	0.903225806	1	0.823529412
PTPRZ1 AND NOT-MAN1A1	0.9375	1	0.882352941
PTPRZ1 AND NOT-STS	0.9375	1	0.882352941
PTPRZ1 AND MARS	0.903225806	1	0.823529412
NRCAM AND NOT-ADAM11	0.903225806	1	0.823529412
PTPRZ1 AND NOT-DNAJB9	0.9375	1	0.882352941
PTPRZ1 AND MFAP3	0.903225806	1	0.823529412
PTPRZ1 AND NOT-MGAT5	0.9375	1	0.882352941
PTPRZ1 AND NOT-MGST1	0.9375	1	0.882352941
PTPRZ1 AND CD99	0.9375	1	0.882352941
PTPRZ1 AND NOT-MID1	0.903225806	1	0.823529412
PTPRZ1 AND MMP16	0.903225806	1	0.823529412
PTPRZ1 AND NOT-ALDH6A1	0.9375	1	0.882352941
PTPRZ1 AND NOT-MXRA7	0.9375	1	0.882352941
PTPRZ1 AND NOT-MINOS1	0.903225806	1	0.823529412
NRCAM AND NOT-TMEM151B	0.903225806	1	0.823529412
PTPRZ1 AND FAM26F	0.903225806	1	0.823529412
PTPRZ1 AND NOT-ASPH	0.9375	1	0.882352941
PTPRZ1 AND NOT-MSMB	0.9375	1	0.882352941
PTPRZ1 AND NOT-MST1R	0.9375	1	0.882352941
PTPRZ1 AND ASTN1	0.9375	1	0.882352941
PTPRZ1 AND NCAM1	0.9375	1	0.882352941
PTPRZ1 AND NOT-NDUFA1	0.9375	1	0.882352941

Antigen Pair Logic	F1	Precision	Recall
CHRNA3 AND NOT-SLC22A2	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-CYP4F12	0.943396226	1	0.892857143
CHRNA3 AND NOT-SOX1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-SPAM1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-SPINK2	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-SPN	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-SSTR3	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-SSTR5	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-BRS3	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-VAMP1	0.95	1	0.904761905
CHRNA3 AND NOT-SYPL1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-TACR2	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-TACR1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-TAZ	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-TGFA	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-ICAM5	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-TRHR	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-TRPC3	0.956521739	1	0.916666667
CHRNA3 AND NOT-TRPC4	0.95	1	0.904761905
CHRNA3 AND NOT-TRPC5	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-TRPC6	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-TLCD2	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-TYR	0.956521739	1	0.916666667
CHRNA3 AND NOT-TYRO3	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-UPK1B	0.95	1	0.904761905
CHRNA3 AND NOT-UGT8	0.936708861	1	0.880952381
CHRNA3 AND NOT-UMOD	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-UPK3A	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-VIPR2	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-BEST1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-MYRF	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-WNT1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-WNT7B	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-WNT11	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-XG	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-CA12	0.95	1	0.904761905
CHRNA3 AND NOT-CACNA1A	0.95	1	0.904761905
CHRNA3 AND NOT-CACNA1C	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-SLC30A2	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-CACNA1F	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-CACNA1S	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-BSND	0.936708861	1	0.880952381
CHRNA3 AND NOT-NPHS2	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-RNF103	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-CACNG1	0.936708861	1	0.880952381
CHRNA3 AND NOT-SLC25A20	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-SMIM2	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-FA2H	0.956521739	1	0.916666667
CHRNA3 AND NOT-OR51B4	0.95	1	0.904761905
CHRNA3 AND NOT-OR51B2	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-NOX5	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-ACSS3	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-ZDHHC14	0.95	1	0.904761905
CHRNA3 AND NOT-STEAP4	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-MCTP1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-HHIPL2	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-NIPAL2	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-ST7	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-TMC5	0.956521739	1	0.916666667
CHRNA3 AND NOT-ZDHHC11	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-TM4SF20	0.95	1	0.904761905
CHRNA3 AND NOT-THSD4	0.943396226	1	0.892857143
CHRNA3 AND NOT-TMC7	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-BTNL8	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-PAQR6	0.956521739	1	0.916666667
CHRNA3 AND NOT-WLS	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-CPED1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-GLRA3	0.956521739	1	0.916666667
CHRNA3 AND NOT-LRRC8E	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-CWH43	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-TMEM254	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-ALPK1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-SLC35F5	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-DNAJC5	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-BPIFB2	0.95	1	0.904761905
CHRNA3 AND NOT-PDCD1LG2	0.950617284	0.987179487	0.916666667

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
PTPRZ1 AND NOT-NDUFB1	0.9375	1	0.882352941
PTPRZ1 AND NOT-NDUFB8	0.9375	1	0.882352941
PTPRZ1 AND NOT-NEO1	0.9375	1	0.882352941
PTPRZ1 AND ATP1A2	0.9375	1	0.882352941
PTPRZ1 AND NMB	0.903225806	1	0.823529412
PTPRZ1 AND ATP1B2	0.9375	1	0.882352941
PTPRZ1 AND ATP2A2	0.9375	1	0.882352941
NRCAM AND NOT-CALY	0.903225806	1	0.823529412
NRCAM AND NOT-TMEM63C	0.903225806	1	0.823529412
NRCAM AND NOT-SLC9A7	0.903225806	1	0.823529412
NRCAM AND NOT-GABBR2	0.903225806	1	0.823529412
PTPRZ1 AND NOT-ATP2B1	0.903225806	1	0.823529412
PTPRZ1 AND NOT-ATP2B2	0.903225806	1	0.823529412
PTPRZ1 AND NOT-ATP2B3	0.9375	1	0.882352941
PTPRZ1 AND NOT-SLC22A18	0.903225806	1	0.823529412
PTPRZ1 AND NOT-CLDN11	0.9375	1	0.882352941
PTPRZ1 AND P2RY1	0.903225806	1	0.823529412
PTPRZ1 AND NOT-CALY	0.903225806	1	0.823529412
PTPRZ1 AND NOT-F11R	0.9375	1	0.882352941
PTPRZ1 AND CLEC4A	0.9375	1	0.882352941
PTPRZ1 AND TMED5	0.9375	1	0.882352941
PTPRZ1 AND NOT-SLC35B3	0.9375	1	0.882352941
PTPRZ1 AND SLC35C2	0.903225806	1	0.823529412
PTPRZ1 AND PCDH8	0.9375	1	0.882352941
PTPRZ1 AND NOT-PCDH9	0.903225806	1	0.823529412
PTPRZ1 AND NOT-GOLT1B	0.903225806	1	0.823529412
PTPRZ1 AND NOT-ST8SIA3	0.903225806	1	0.823529412
PTPRZ1 AND NDUFA13	0.9375	1	0.882352941
PTPRZ1 AND GAL	0.903225806	1	0.823529412
PTPRZ1 AND NOT-SIDT2	0.903225806	1	0.823529412
PTPRZ1 AND NOT-ADIPOR1	0.9375	1	0.882352941
PTPRZ1 AND NOT-TUBD1	0.9375	1	0.882352941
PTPRZ1 AND NOT-ZDHHC2	0.9375	1	0.882352941
PTPRZ1 AND NOT-PIGP	0.9375	1	0.882352941
PTPRZ1 AND NOT-CRIM1	0.9375	1	0.882352941
PTPRZ1 AND NOT-TMEM216	0.903225806	1	0.823529412
PTPRZ1 AND PIPOX	0.903225806	1	0.823529412
PTPRZ1 AND NOT-CEND1	0.903225806	1	0.823529412
PTPRZ1 AND NOT-TIMMDC1	0.9375	1	0.882352941
PTPRZ1 AND NOT-CYP39A1	0.9375	1	0.882352941
PTPRZ1 AND NOT-FKBP11	0.9375	1	0.882352941
PTPRZ1 AND NOT-ZDHHC3	0.903225806	1	0.823529412
PTPRZ1 AND SLC22A17	0.903225806	1	0.823529412
PTPRZ1 AND MBTPS2	0.9375	1	0.882352941
PTPRZ1 AND CHST15	0.903225806	1	0.823529412
PTPRZ1 AND NOT-TEX264	0.9375	1	0.882352941
PTPRZ1 AND NOT-ATRAID	0.903225806	1	0.823529412
PTPRZ1 AND NOT-GULP1	0.9375	1	0.882352941
PTPRZ1 AND RHCG	0.9375	1	0.882352941
PTPRZ1 AND NOT-UBE2J1	0.903225806	1	0.823529412
PTPRZ1 AND TMEM138	0.9375	1	0.882352941
PTPRZ1 AND NOT-GDE1	0.9375	1	0.882352941
PTPRZ1 AND NOT-LSR	0.9375	1	0.882352941
PTPRZ1 AND ERGIC3	0.9375	1	0.882352941
PTPRZ1 AND NOT-HMP19	0.9375	1	0.882352941
PTPRZ1 AND NOT-MPC1	0.903225806	1	0.823529412
PTPRZ1 AND NOT-CYB5R2	0.903225806	1	0.823529412
PTPRZ1 AND NOT-EMCN	0.9375	1	0.882352941
PTPRZ1 AND NOT-CYB5R1	0.9375	1	0.882352941
PTPRZ1 AND NOT-SELT	0.903225806	1	0.823529412
PTPRZ1 AND DNAJB11	0.9375	1	0.882352941
PTPRZ1 AND PECAM1	0.903225806	1	0.823529412
PTPRZ1 AND NOT-ATP8A2	0.903225806	1	0.823529412
PTPRZ1 AND CECR1	0.903225806	1	0.823529412
PTPRZ1 AND NOT-ATP8B1	0.9375	1	0.882352941
PTPRZ1 AND NOT-ATP6V0C	0.903225806	1	0.823529412
PTPRZ1 AND NOT-PIGF	0.9375	1	0.882352941
PTPRZ1 AND NOT-PIGH	0.9375	1	0.882352941
PTPRZ1 AND PLP1	0.903225806	1	0.823529412
PTPRZ1 AND PLXNA1	0.903225806	1	0.823529412
PTPRZ1 AND NOT-S1PR5	0.903225806	1	0.823529412
PTPRZ1 AND NOT-IL20RA	0.9375	1	0.882352941
PTPRZ1 AND SLCO1C1	0.9375	1	0.882352941
PTPRZ1 AND NOT-FTHL17	0.903225806	1	0.823529412
PTPRZ1 AND NOT-SLC37A1	0.9375	1	0.882352941
PTPRZ1 AND NOT-PODXL	0.9375	1	0.882352941

Antigen Pair Logic	F1	Precision	Recall
CHRNA3 AND NOT-SLC44A4	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-LY6G5C	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-UPK3B	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-TAS1R1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-TMPRSS5	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-KCNH6	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-OR2C3	0.95	1	0.904761905
CHRNA3 AND NOT-ACSBG2	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-DEFB126	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-OR5V1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-OR2B2	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-SLC14A2	0.943396226	1	0.892857143
CHRNA3 AND NOT-OR12D3	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-SPACA1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-SLC10A3	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-FZD9	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-ELOVL3	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-FCRL4	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-SLC25A31	0.95	1	0.904761905
CHRNA3 AND NOT-CHST9	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-GPR101	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-CD99L2	0.956521739	1	0.916666667
CHRNA3 AND NOT-SLC25A18	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-SYT15	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-SPATA9	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-OR1G1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-SLC4A11	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-KREMEN1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-PLA2G10	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-USP48	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-ATP13A4	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-ACSS1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-FUT10	0.956521739	1	0.916666667
CHRNA3 AND NOT-IL17RC	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-TMEM25	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-PTPN5	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-FIBCD1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-DISP1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-ITGA10	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-ABCC11	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-KIAA1644	0.956521739	1	0.916666667
CHRNA3 AND NOT-SLC45A3	0.956521739	1	0.916666667
CHRNA3 AND NOT-CDC14B	0.936708861	1	0.880952381
CHRNA3 AND NOT-SLC4A4	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-DGAT1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-B3GALT2	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-ABCC3	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-ADAM23	0.956521739	1	0.916666667
CHRNA3 AND NOT-ADAM18	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-ADAM7	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-TNFRSF10C	0.956521739	1	0.916666667
CHRNA3 AND NOT-CACNA1I	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-LMLN	0.943396226	1	0.892857143
CHRNA3 AND NOT-CEACAM21	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-ANGPTL1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-CLDN2	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-CLDN1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-CLDN9	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-TMEM67	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-SLC16A5	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-P2RX6	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-SLC28A1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-SYT12	0.943396226	1	0.892857143
CHRNA3 AND NOT-KIAA1919	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-HTR3B	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-SLC33A1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-GGTLC1	0.95	1	0.904761905
CHRNA3 AND NOT-VAPB	0.943396226	1	0.892857143
CHRNA3 AND NOT-DAPL1	0.95	1	0.904761905
CHRNA3 AND NOT-TMEM129	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-REEP6	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-TAAR2	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-GPR55	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-GPR52	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-PEX11G	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-KCNB2	0.950617284	0.987179487	0.916666667

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
PTPRZ1 AND NOT-GDAP1	0.903225806	1	0.823529412
PTPRZ1 AND PON2	0.9375	1	0.882352941
PTPRZ1 AND NOT-FAM134B	0.9375	1	0.882352941
PTPRZ1 AND NOT-PON3	0.9375	1	0.882352941
PTPRZ1 AND NOT-TMX3	0.9375	1	0.882352941
PTPRZ1 AND NOT-TMCO1	0.903225806	1	0.823529412
PTPRZ1 AND NOT-TOMM7	0.9375	1	0.882352941
PTPRZ1 AND NOT-NDFIP2	0.9375	1	0.882352941
PTPRZ1 AND NOT-VSIG10	0.9375	1	0.882352941
PTPRZ1 AND LRRN3	0.9375	1	0.882352941
PTPRZ1 AND NOT-MANSC1	0.9375	1	0.882352941
PTPRZ1 AND NOT-MARCH5	0.9375	1	0.882352941
PTPRZ1 AND SLC35F2	0.903225806	1	0.823529412
PTPRZ1 AND NOT-LEPROT	0.9375	1	0.882352941
PTPRZ1 AND NOT-GRAMD1C	0.903225806	1	0.823529412
PTPRZ1 AND NOT-CTSA	0.903225806	1	0.823529412
PTPRZ1 AND NOT-DNAJB12	0.9375	1	0.882352941
PTPRZ1 AND NOT-TRPM4	0.9375	1	0.882352941
PTPRZ1 AND PPIB	0.9375	1	0.882352941
PTPRZ1 AND NOT-TRPM7	0.9375	1	0.882352941
PTPRZ1 AND NOT-CDHR2	0.9375	1	0.882352941
PTPRZ1 AND NOT-WBP1L	0.9375	1	0.882352941
PTPRZ1 AND NOT-MFSD6	0.903225806	1	0.823529412
PTPRZ1 AND NOT-PAQR5	0.9375	1	0.882352941
PTPRZ1 AND MS4A12	0.903225806	1	0.823529412
PTPRZ1 AND NOT-DPP8	0.9375	1	0.882352941
PTPRZ1 AND LPPR1	0.903225806	1	0.823529412
PTPRZ1 AND NOT-RNF43	0.9375	1	0.882352941
PTPRZ1 AND CMTM6	0.9375	1	0.882352941
PTPRZ1 AND NOT-RHBDL2	0.903225806	1	0.823529412
PTPRZ1 AND NOT-SLC41A3	0.9375	1	0.882352941
PTPRZ1 AND NOT-C1orf27	0.9375	1	0.882352941
PTPRZ1 AND NOT-TMEM160	0.903225806	1	0.823529412
PTPRZ1 AND NOT-PIGX	0.9375	1	0.882352941
PTPRZ1 AND NOT-MARC2	0.9375	1	0.882352941
PTPRZ1 AND NOT-TMCO3	0.9375	1	0.882352941
PTPRZ1 AND NOT-STX17	0.9375	1	0.882352941
PTPRZ1 AND TMEM248	0.903225806	1	0.823529412
PTPRZ1 AND TMEM45A	0.9375	1	0.882352941
PTPRZ1 AND NOT-SLC6A15	0.903225806	1	0.823529412
PTPRZ1 AND NOT-ANO10	0.9375	1	0.882352941
PTPRZ1 AND NOT-TMEM38B	0.903225806	1	0.823529412
PTPRZ1 AND TMEM33	0.903225806	1	0.823529412
PTPRZ1 AND SEC61A2	0.903225806	1	0.823529412
PTPRZ1 AND NOT-RMDN3	0.903225806	1	0.823529412
PTPRZ1 AND NOT-TMEM57	0.9375	1	0.882352941
PTPRZ1 AND NAT10	0.903225806	1	0.823529412
PTPRZ1 AND NOT-TMEM19	0.903225806	1	0.823529412
PTPRZ1 AND TMEM100	0.9375	1	0.882352941
PTPRZ1 AND NOT-FBXW7	0.903225806	1	0.823529412
PTPRZ1 AND NOT-RNF121	0.9375	1	0.882352941
PTPRZ1 AND NOT-TMEM74B	0.9375	1	0.882352941
PTPRZ1 AND NOT-SLC39A9	0.9375	1	0.882352941
PTPRZ1 AND NOT-WDR33	0.9375	1	0.882352941
PTPRZ1 AND NOT-STYK1	0.903225806	1	0.823529412
PTPRZ1 AND NOT-LGR4	0.9375	1	0.882352941
PTPRZ1 AND LRRC59	0.9375	1	0.882352941
PTPRZ1 AND NOT-ETNK1	0.9375	1	0.882352941
PTPRZ1 AND CHST12	0.903225806	1	0.823529412
PTPRZ1 AND NOT-SMPD3	0.9375	1	0.882352941
PTPRZ1 AND NOT-SVOP	0.903225806	1	0.823529412
PTPRZ1 AND IL17RB	0.9375	1	0.882352941
PTPRZ1 AND NOT-SLC39A4	0.9375	1	0.882352941
PTPRZ1 AND NOT-TMEM127	0.9375	1	0.882352941
PTPRZ1 AND SLC30A6	0.903225806	1	0.823529412
PTPRZ1 AND NDC1	0.9375	1	0.882352941
PTPRZ1 AND SAYSD1	0.903225806	1	0.823529412
PTPRZ1 AND NOT-CSGALNACT1	0.9375	1	0.882352941
PTPRZ1 AND NOT-SCN3B	0.903225806	1	0.823529412
PTPRZ1 AND PAG1	0.9375	1	0.882352941
PTPRZ1 AND NOT-GLT8D1	0.9375	1	0.882352941
PTPRZ1 AND NOT-TEX2	0.903225806	1	0.823529412
PTPRZ1 AND TMEM165	0.9375	1	0.882352941
PTPRZ1 AND NOT-TMEM126B	0.903225806	1	0.823529412
PTPRZ1 AND NOT-PKN2	0.9375	1	0.882352941
PTPRZ1 AND AXL	0.903225806	1	0.823529412

Antigen Pair Logic	F1	Precision	Recall
CHRNA3 AND NOT-MMP20	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-OPALIN	0.956521739	1	0.916666667
CHRNA3 AND NOT-GLP2R	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-SLC22A6	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-ACPT	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-NRXN3	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-TTYH2	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-MUC16	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-TMPRSS11D	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-SFXN5	0.95	1	0.904761905
CHRNA3 AND NOT-ABCC12	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-CD80	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-CYP7B1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-OPN4	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-NCR2	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-CHST3	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-SIGLEC6	0.95	1	0.904761905
CHRNA3 AND NOT-PIGL	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-ENTPD2	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-CELSR1	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-CLCA2	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-TMEM63A	0.956521739	1	0.916666667
CHRNA3 AND NOT-KIAA0319	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-SV2B	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-ATP2C2	0.950617284	0.987179487	0.916666667
CHRNA3 AND NOT-KCNE2	0.950617284	0.987179487	0.916666667
FAM163A AND NOT-SLC2A13	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-SLC2A13	0.959537572	0.93258427	0.988095238
MARCH11 AND NOT-SLC2A13	0.946745562	0.941176471	0.952380952
SLC10A4 AND NOT-CHRNB2	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-CHRNB3	0.93258427	0.882978723	0.988095238
SLC10A4 AND NOT-CSMD3	0.922222222	0.864583333	0.988095238
MARCH11 AND NOT-CSMD3	0.946745562	0.941176471	0.952380952
FAM163A AND NOT-ELFN2	0.922222222	0.864583333	0.988095238
ST8SIA2 AND NOT-ELFN2	0.943396226	1	0.892857143
SLC10A4 AND NOT-TMEM132B	0.927374302	0.873684211	0.988095238
MARCH11 AND NOT-TMEM132B	0.946745562	0.941176471	0.952380952
SLC10A4 AND NOT-C1QTNF1	0.922222222	0.864583333	0.988095238
ST8SIA2 AND NOT-SLC26A9	0.95	1	0.904761905
SLC10A4 AND NOT-OMA1	0.922222222	0.864583333	0.988095238
FAM163A AND NOT-SLC5A11	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-SLC5A11	0.970760234	0.954022989	0.988095238
ST8SIA2 AND NOT-SLC5A11	0.943396226	1	0.892857143
SLC10A4 AND NOT-CMTM5	0.965116279	0.943181818	0.988095238
ST8SIA2 AND NOT-CMTM5	0.943396226	1	0.892857143
MARCH11 AND NOT-SLC26A8	0.963855422	0.975609756	0.952380952
FAM163A AND NOT-GRIN3B	0.93258427	0.882978723	0.988095238
SLC10A4 AND NOT-GRIN3B	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-SLC18B1	0.954022989	0.922222222	0.988095238
GPR19 AND NOT-SLC18B1	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-SLC18B1	0.946745562	0.941176471	0.952380952
FAM163A AND NOT-MRGPRX2	0.948571429	0.912087912	0.988095238
SLC10A4 AND NOT-MRGPRX2	0.948571429	0.912087912	0.988095238
ST8SIA2 AND NOT-MRGPRX2	0.956521739	1	0.916666667
FAM163A AND NOT-MRGPRX3	0.965116279	0.943181818	0.988095238
SLC10A4 AND NOT-MRGPRX3	0.959537572	0.93258427	0.988095238
ST8SIA2 AND NOT-MRGPRX3	0.956521739	1	0.916666667
SLC10A4 AND NOT-GALNT15	0.976470588	0.965116279	0.988095238
ST8SIA2 AND NOT-GALNT15	0.943396226	1	0.892857143
SLC10A4 AND NOT-ADCYAP1R1	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-GPR62	0.937853107	0.892473118	0.988095238
GPR19 AND NOT-GPR62	0.951219512	0.975	0.928571429
MARCH11 AND NOT-GPR62	0.946745562	0.941176471	0.952380952
ST8SIA2 AND NOT-GPR62	0.943396226	1	0.892857143
SLC10A4 AND NOT-CLCN5	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-CLCN7	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-SFXN2	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-PDZD8	0.927374302	0.873684211	0.988095238
ST8SIA2 AND NOT-OR5P2	0.950617284	0.987179487	0.916666667
ST8SIA2 AND NOT-OR5P3	0.950617284	0.987179487	0.916666667
FAM163A AND NOT-TMEM52B	0.943181818	0.902173913	0.988095238
SLC10A4 AND NOT-TMEM52B	0.93258427	0.882978723	0.988095238
MARCH11 AND NOT-TMEM52B	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-TEX29	0.963855422	0.975609756	0.952380952
FAM163A AND NOT-SLC24A4	0.976470588	0.965116279	0.988095238
SLC10A4 AND NOT-SLC24A4	0.982248521	0.976470588	0.988095238

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
PTPRZ1 AND NOT-ACSS2	0.903225806	1	0.823529412
PTPRZ1 AND NOT-KMT2E	0.9375	1	0.882352941
PTPRZ1 AND NOT-BCAP29	0.9375	1	0.882352941
PTPRZ1 AND PCDHB9	0.9375	1	0.882352941
PTPRZ1 AND NOT-ANKH	0.903225806	1	0.823529412
PTPRZ1 AND NOT-MOSPD1	0.903225806	1	0.823529412
PTPRZ1 AND NOT-PRNP	0.903225806	1	0.823529412
PTPRZ1 AND PROS1	0.903225806	1	0.823529412
PTPRZ1 AND NOT-CYP26B1	0.9375	1	0.882352941
PTPRZ1 AND NOT-PSEN1	0.9375	1	0.882352941
PTPRZ1 AND NOT-PSEN2	0.9375	1	0.882352941
PTPRZ1 AND NOT-CLDND1	0.9375	1	0.882352941
PTPRZ1 AND NOT-TMEM9B	0.9375	1	0.882352941
PTPRZ1 AND NOT-JPH1	0.903225806	1	0.823529412
PTPRZ1 AND NOT-CELF4	0.903225806	1	0.823529412
PTPRZ1 AND NOT-TM9SF3	0.9375	1	0.882352941
TPST1 AND NOT-TM9SF3	0.903225806	1	0.823529412
NOT-PTPRZ1 AND PTPRZ1	0.9375	1	0.882352941
PTPRZ1 AND PMEPA1	0.903225806	1	0.823529412
PTPRZ1 AND NOT-MFF	0.9375	1	0.882352941
PTPRZ1 AND ACKR3	0.903225806	1	0.823529412
PTPRZ1 AND PNPLA2	0.9375	1	0.882352941
PTPRZ1 AND NOT-RHBG	0.9375	1	0.882352941
PTPRZ1 AND NOT-RTN4	0.9375	1	0.882352941
PTPRZ1 AND NOT-SMIM8	0.903225806	1	0.823529412
PTPRZ1 AND NOT-TMEM63C	0.9375	1	0.882352941
PTPRZ1 AND NOT-SNX14	0.9375	1	0.882352941
PTPRZ1 AND PTCH1	0.9375	1	0.882352941
TPST1 AND NOT-PTGER2	0.903225806	1	0.823529412
PTPRZ1 AND NOT-ZNF304	0.9375	1	0.882352941
PTPRZ1 AND NOT-MRS2	0.903225806	1	0.823529412
PTPRZ1 AND NOT-ABHD6	0.903225806	1	0.823529412
PTPRZ1 AND NOT-SLC24A3	0.9375	1	0.882352941
PTPRZ1 AND PTGS1	0.903225806	1	0.823529412
PTPRZ1 AND NOT-TMCC3	0.903225806	1	0.823529412
PTPRZ1 AND NOT-PTH1R	0.9375	1	0.882352941
PTPRZ1 AND NOT-SLC12A5	0.903225806	1	0.823529412
PTPRZ1 AND GRAMD1B	0.9375	1	0.882352941
PTPRZ1 AND HEG1	0.9375	1	0.882352941
PTPRZ1 AND GPR158	0.9375	1	0.882352941
PTPRZ1 AND SERINC1	0.9375	1	0.882352941
PTPRZ1 AND NOT-KIAA1324	0.9375	1	0.882352941
PTPRZ1 AND NOT-NCEH1	0.9375	1	0.882352941
PTPRZ1 AND NOT-SYT13	0.903225806	1	0.823529412
PTPRZ1 AND NOT-KIAA1467	0.903225806	1	0.823529412
PTPRZ1 AND LRFN1	0.903225806	1	0.823529412
PTPRZ1 AND NOT-LRRC4C	0.903225806	1	0.823529412
PTPRZ1 AND PCDHB16	0.903225806	1	0.823529412
PTPRZ1 AND NOT-PTPN2	0.903225806	1	0.823529412
PTPRZ1 AND PTPRA	0.9375	1	0.882352941
PTPRZ1 AND PTPRC	0.903225806	1	0.823529412
PTPRZ1 AND NOT-PTPRF	0.9375	1	0.882352941
PTPRZ1 AND PTPRO	0.903225806	1	0.823529412
PTPRZ1 AND NOT-PTPRR	0.9375	1	0.882352941
PTPRZ1 AND NOT-PTPRS	0.9375	1	0.882352941
PTPRZ1 AND MS4A7	0.9375	1	0.882352941
PTPRZ1 AND NOT-OSTC	0.9375	1	0.882352941
PTPRZ1 AND NOT-IL22RA1	0.9375	1	0.882352941
PTPRZ1 AND RARRES3	0.903225806	1	0.823529412
PTPRZ1 AND NOT-SIGIRR	0.9375	1	0.882352941
PTPRZ1 AND NOT-TMEM35	0.9375	1	0.882352941
PTPRZ1 AND NOT-BCL2	0.9375	1	0.882352941
PTPRZ1 AND NOT-BCL2L2	0.903225806	1	0.823529412
PTPRZ1 AND NOT-FAM3A	0.9375	1	0.882352941
PTPRZ1 AND RNASE4	0.903225806	1	0.823529412
PTPRZ1 AND RNASE6	0.903225806	1	0.823529412
PTPRZ1 AND KCNK15	0.903225806	1	0.823529412
PTPRZ1 AND NOT-ROBO2	0.903225806	1	0.823529412
PTPRZ1 AND NOT-RPL37A	0.9375	1	0.882352941
PTPRZ1 AND RPN2	0.903225806	1	0.823529412
PTPRZ1 AND NOT-RPS23	0.903225806	1	0.823529412
PTPRZ1 AND RYK	0.903225806	1	0.823529412
PTPRZ1 AND NOT-RYR2	0.903225806	1	0.823529412
PTPRZ1 AND SORT1	0.903225806	1	0.823529412
PTPRZ1 AND NOT-SLC22A23	0.9375	1	0.882352941
PTPRZ1 AND NOT-SC5D	0.9375	1	0.882352941

Antigen Pair Logic	F1	Precision	Recall
GPR19 AND NOT-SLC24A4	0.96969697	0.987654321	0.952380952
MARCH11 AND NOT-SLC24A4	0.946745562	0.941176471	0.952380952
ST8SIA2 AND NOT-SLC24A4	0.956521739	1	0.916666667
SLC10A4 AND NOT-SLC51B	0.93258427	0.882978723	0.988095238
ST8SIA2 AND NOT-SLC51B	0.950617284	0.987179487	0.916666667
SLC10A4 AND NOT-NIPA1	0.927374302	0.873684211	0.988095238
RTN1 AND CANT1	0.950617284	0.987179487	0.916666667
ST8SIA2 AND NOT-SPNS2	0.95	1	0.904761905
SLC10A4 AND NOT-FAM69C	0.963414634	0.9875	0.94047619
MARCH11 AND NOT-FAM69C	0.951807229	0.963414634	0.94047619
ST8SIA2 AND NOT-FAM69C	0.936708861	1	0.880952381
ST8SIA2 AND NOT-IZUMO2	0.950617284	0.987179487	0.916666667
SLC10A4 AND NOT-CNGA4	0.93258427	0.882978723	0.988095238
ST8SIA2 AND NOT-CNGA4	0.950617284	0.987179487	0.916666667
FAM163A AND NOT-TMEM125	0.988095238	0.988095238	0.988095238
SLC10A4 AND NOT-TMEM125	0.988095238	0.988095238	0.988095238
GABRB3 AND NOT-TMEM125	0.950617284	0.987179487	0.916666667
GPR19 AND NOT-TMEM125	0.96969697	0.987654321	0.952380952
MARCH11 AND NOT-TMEM125	0.952380952	0.952380952	0.952380952
ST8SIA3 AND NOT-TMEM125	0.946745562	0.941176471	0.952380952
SCN3B AND NOT-TMEM125	0.97005988	0.975903614	0.964285714
ST8SIA2 AND NOT-TMEM125	0.956521739	1	0.916666667
SLC10A4 AND NOT-DRAM2	0.922222222	0.864583333	0.988095238
FAM163A AND MBOAT2	0.931818182	0.891304348	0.976190476
SLC10A4 AND NOT-CNTNAP5	0.941860465	0.920454545	0.964285714
ST8SIA2 AND NOT-CNTNAP5	0.95	1	0.904761905
SLC10A4 AND NOT-KCNH8	0.927374302	0.873684211	0.988095238
MARCH11 AND NOT-SYNPR	0.946745562	0.941176471	0.952380952
ST8SIA2 AND NOT-TMPRSS11B	0.956521739	1	0.916666667
FAM163A AND COX4I1	0.943181818	0.902173913	0.988095238
ST8SIA2 AND COX4I1	0.936708861	1	0.880952381
SLC10A4 AND NOT-SLC9B2	0.922222222	0.864583333	0.988095238
ST8SIA2 AND NOT-IL31RA	0.936708861	1	0.880952381
FAM163A AND NOT-TAAR9	0.959537572	0.93258427	0.988095238
SLC10A4 AND NOT-TAAR9	0.948571429	0.912087912	0.988095238
FAM163A AND NOT-ADORA1	0.941860465	0.920454545	0.964285714
SLC10A4 AND NOT-ADORA1	0.964285714	0.964285714	0.964285714
MARCH11 AND NOT-ADORA1	0.957575758	0.975308642	0.94047619
ST8SIA2 AND NOT-ADORA1	0.943396226	1	0.892857143
MARCH11 AND NOT-B3GAT2	0.946745562	0.941176471	0.952380952
ST8SIA2 AND NOT-B3GAT2	0.950617284	0.987179487	0.916666667
ST8SIA2 AND NOT-TMEM139	0.956521739	1	0.916666667
MARCH11 AND NOT-ADORA2B	0.946745562	0.941176471	0.952380952
SLC10A4 AND NOT-TMEM71	0.922222222	0.864583333	0.988095238
FAM163A AND AGPAT6	0.927374302	0.873684211	0.988095238
ST8SIA2 AND NOT-RNF183	0.950617284	0.987179487	0.916666667
SLC10A4 AND NOT-CRHR2	0.948571429	0.912087912	0.988095238
ST8SIA2 AND NOT-CRHR2	0.943396226	1	0.892857143
FAM163A AND NOT-ASB5	0.947976879	0.921348315	0.976190476
SLC10A4 AND NOT-ASB5	0.982035928	0.987951807	0.976190476
ST8SIA2 AND NOT-ASB5	0.950617284	0.987179487	0.916666667
NRSN1 AND NOT-SLC2A12	0.95	1	0.904761905
NRSN1 AND NOT-MLC1	0.95	1	0.904761905
NRSN1 AND NOT-SLC7A11	0.936708861	1	0.880952381
NRSN1 AND NOT-GPR37	0.936708861	1	0.880952381
NRSN1 AND NOT-GRM3	0.95	1	0.904761905
NRSN1 AND NOT-AQP4	0.936708861	1	0.880952381
NRSN1 AND NOT-KCNA2	0.943396226	1	0.892857143
NRSN1 AND NOT-KCNJ10	0.95	1	0.904761905
NRSN1 AND NOT-KCNJ16	0.95	1	0.904761905
NRSN1 AND NOT-LRP4	0.95	1	0.904761905
NRSN1 AND NOT-MOG	0.95	1	0.904761905
NRSN1 AND NOT-HHATL	0.95	1	0.904761905
NRSN1 AND NOT-SLC6A1	0.95	1	0.904761905
NRSN1 AND NOT-SLC15A2	0.95	1	0.904761905
NRSN1 AND NOT-FA2H	0.95	1	0.904761905
NRSN1 AND NOT-PAQR6	0.936708861	1	0.880952381
NRSN1 AND NOT-SLC25A18	0.95	1	0.904761905
NRSN1 AND NOT-SLITRK2	0.943396226	1	0.892857143
NRSN1 AND NOT-SLC4A4	0.95	1	0.904761905
NRSN1 AND NOT-OPALIN	0.95	1	0.904761905
ST8SIA2 AND NOT-TRPM6	0.956521739	1	0.916666667
SLC10A4 AND NOT-SIRPA	0.937853107	0.892473118	0.988095238
MARCH11 AND NOT-SIRPA	0.946745562	0.941176471	0.952380952
SLC10A4 AND NOT-SAMD8	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-VTI1A	0.931818182	0.891304348	0.976190476

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
PTPRZ1 AND SCD	0.903225806	1	0.823529412
PTPRZ1 AND SCN1A	0.903225806	1	0.823529412
PTPRZ1 AND NOT-SCN1B	0.903225806	1	0.823529412
PTPRZ1 AND NOT-SCN2A	0.903225806	1	0.823529412
PTPRZ1 AND NOT-SCN2B	0.9375	1	0.882352941
PTPRZ1 AND NOT-SCN8A	0.903225806	1	0.823529412
PTPRZ1 AND NOT-SCNN1G	0.9375	1	0.882352941
PTPRZ1 AND CCL4	0.903225806	1	0.823529412
PTPRZ1 AND NOT-SDC2	0.903225806	1	0.823529412
PTPRZ1 AND THADA	0.903225806	1	0.823529412
PTPRZ1 AND NOT-SDHC	0.9375	1	0.882352941
PTPRZ1 AND NOT-SEL1L	0.9375	1	0.882352941
PTPRZ1 AND NOT-PERP	0.9375	1	0.882352941
PTPRZ1 AND NOT-POPDC2	0.9375	1	0.882352941
PTPRZ1 AND NOT-SMOC2	0.903225806	1	0.823529412
PTPRZ1 AND NOT-LRRC4	0.9375	1	0.882352941
PTPRZ1 AND NOT-ADGRL4	0.9375	1	0.882352941
PTPRZ1 AND NOT-SEPP1	0.9375	1	0.882352941
PTPRZ1 AND NOT-TMEM91	0.9375	1	0.882352941
PTPRZ1 AND NOT-DNAJC1	0.9375	1	0.882352941
PTPRZ1 AND NOT-SEMA4A	0.9375	1	0.882352941
PTPRZ1 AND NOT-ATL2	0.9375	1	0.882352941
PTPRZ1 AND MS4A6A	0.903225806	1	0.823529412
PTPRZ1 AND NOT-C1orf233	0.9375	1	0.882352941
PTPRZ1 AND NOT-CDH22	0.9375	1	0.882352941
PTPRZ1 AND NOT-C5orf28	0.9375	1	0.882352941
PTPRZ1 AND NOT-TMEM168	0.9375	1	0.882352941
PTPRZ1 AND FNDC3B	0.903225806	1	0.823529412
PTPRZ1 AND NOT-ELOVL1	0.9375	1	0.882352941
PTPRZ1 AND FNDC4	0.903225806	1	0.823529412
PTPRZ1 AND NOT-FBXL17	0.9375	1	0.882352941
PTPRZ1 AND NOT-NDRG4	0.9375	1	0.882352941
PTPRZ1 AND NOT-TMEM237	0.9375	1	0.882352941
PTPRZ1 AND BMP2	0.903225806	1	0.823529412
PTPRZ1 AND NOT-SLC1A6	0.9375	1	0.882352941
PTPRZ1 AND SLC2A3	0.9375	1	0.882352941
PTPRZ1 AND NOT-SLC3A1	0.9375	1	0.882352941
PTPRZ1 AND SLC3A2	0.9375	1	0.882352941
PTPRZ1 AND SLC4A2	0.9375	1	0.882352941
PTPRZ1 AND NOT-SLC6A1	0.903225806	1	0.823529412
PTPRZ1 AND NOT-SLC6A6	0.9375	1	0.882352941
PTPRZ1 AND NOT-SLC6A12	0.903225806	1	0.823529412
PTPRZ1 AND NOT-SLC12A2	0.9375	1	0.882352941
PTPRZ1 AND NOT-SLC18A2	0.9375	1	0.882352941
PTPRZ1 AND NOT-SLC20A1	0.903225806	1	0.823529412
PTPRZ1 AND BMPR1A	0.903225806	1	0.823529412
PTPRZ1 AND SLC22A4	0.903225806	1	0.823529412
PTPRZ1 AND BMPR1B	0.9375	1	0.882352941
PTPRZ1 AND NOT-DLK2	0.9375	1	0.882352941
PTPRZ1 AND SOAT1	0.9375	1	0.882352941
PTPRZ1 AND NOT-BNIP3	0.9375	1	0.882352941
PTPRZ1 AND SPAM1	0.903225806	1	0.823529412
PTPRZ1 AND NOT-DST	0.9375	1	0.882352941
PTPRZ1 AND NOT-SPAST	0.9375	1	0.882352941
PTPRZ1 AND NOT-SPG7	0.9375	1	0.882352941
PTPRZ1 AND NOT-SQLE	0.9375	1	0.882352941
PTPRZ1 AND BRCA1	0.903225806	1	0.823529412
PTPRZ1 AND NOT-SSR1	0.9375	1	0.882352941
PTPRZ1 AND NOT-SSTR1	0.9375	1	0.882352941
PTPRZ1 AND HSPA13	0.9375	1	0.882352941
PTPRZ1 AND NOT-ELOVL4	0.903225806	1	0.823529412
PTPRZ1 AND NOT-STIM1	0.9375	1	0.882352941
PTPRZ1 AND NOT-STX3	0.903225806	1	0.823529412
PTPRZ1 AND NOT-VAMP1	0.903225806	1	0.823529412
PTPRZ1 AND BST2	0.9375	1	0.882352941
PTPRZ1 AND NOT-SYT4	0.9375	1	0.882352941
PTPRZ1 AND TEK	0.9375	1	0.882352941
PTPRZ1 AND TGFBR2	0.903225806	1	0.823529412
PTPRZ1 AND NOT-TSPO	0.903225806	1	0.823529412
PTPRZ1 AND NOT-ICAM5	0.903225806	1	0.823529412
PTPRZ1 AND NOT-SEC62	0.903225806	1	0.823529412
PTPRZ1 AND TLR2	0.903225806	1	0.823529412
PTPRZ1 AND TSPAN6	0.903225806	1	0.823529412
PTPRZ1 AND C1QB	0.903225806	1	0.823529412
PTPRZ1 AND NOT-TPBG	0.903225806	1	0.823529412
PTPRZ1 AND C3AR1	0.9375	1	0.882352941

Antigen Pair Logic	F1	Precision	Recall
SLC10A4 AND NOT-SYT9	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-CSF1	0.922222222	0.864583333	0.988095238
FAM163A AND ALG10B	0.947368421	0.931034483	0.964285714
MARCH11 AND NOT-B3GLCT	0.952380952	0.952380952	0.952380952
SLC10A4 AND NOT-C14orf37	0.93258427	0.882978723	0.988095238
MARCH11 AND NOT-C14orf37	0.946745562	0.941176471	0.952380952
ST8SIA2 AND NOT-C14orf37	0.943396226	1	0.892857143
SLC10A4 AND NOT-LRFN5	0.927374302	0.873684211	0.988095238
MARCH11 AND NOT-LRFN5	0.946745562	0.941176471	0.952380952
ST8SIA2 AND NOT-LRFN5	0.950617284	0.987179487	0.916666667
MARCH11 AND NOT-LYSMD4	0.946745562	0.941176471	0.952380952
ST8SIA2 AND NOT-TMCO5A	0.956521739	1	0.916666667
ST8SIA2 AND NOT-NRG4	0.95	1	0.904761905
ST8SIA2 AND NOT-GSG1L	0.936708861	1	0.880952381
MARCH11 AND NOT-RBFOX3	0.946745562	0.941176471	0.952380952
SLC10A4 AND NOT-CD300LF	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-ANKRD29	0.927374302	0.873684211	0.988095238
MARCH11 AND NOT-ANKRD29	0.946745562	0.941176471	0.952380952
SLC10A4 AND NOT-APCDD1	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-VSIG10L	0.922222222	0.864583333	0.988095238
SMIM17 AND NOT-FGFR3	0.95	1	0.904761905
FAM163A AND NOT-C19orf18	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-C19orf18	0.93258427	0.882978723	0.988095238
SLC10A4 AND NOT-TMEM190	0.922222222	0.864583333	0.988095238
ST8SIA2 AND NOT-TMC4	0.943396226	1	0.892857143
FAM163A AND NOT-DCST1	0.954022989	0.922222222	0.988095238
FAM163A AND NOT-SLC9B1	0.93258427	0.882978723	0.988095238
FAM163A AND NOT-GPBAR1	0.941860465	0.920454545	0.964285714
FAM163A AND NOT-SLC25A48	0.965116279	0.943181818	0.988095238
FAM163A AND NOT-MARVELD2	0.93258427	0.882978723	0.988095238
FAM163A AND CCDC112	0.942528736	0.911111111	0.976190476
FAM163A AND NOT-NKAIN2	0.937853107	0.892473118	0.988095238
FAM163A AND CCDC167	0.947368421	0.931034483	0.964285714
FAM163A AND NOT-CLEC2L	0.93258427	0.882978723	0.988095238
FAM163A AND NOT-PEBP4	0.964285714	0.964285714	0.964285714
FAM163A AND NOT-KCNU1	0.947976879	0.921348315	0.976190476
FAM163A AND NOT-CYP4B1	0.954022989	0.922222222	0.988095238
FAM163A AND NOT-OR1Q1	0.982248521	0.976470588	0.988095238
FAM163A AND NOT-FAAH2	0.931818182	0.891304348	0.976190476
FAM163A AND NOT-TMEM229B	0.922222222	0.864583333	0.988095238
FAM163A AND NOT-HTR3C	0.936416185	0.91011236	0.964285714
FAM163A AND SPTSSA	0.931818182	0.891304348	0.976190476
FAM163A AND NOT-DIO2	0.959064327	0.942528736	0.976190476
FAM163A AND NOT-EGFR	0.922222222	0.864583333	0.988095238
FAM163A AND NOT-TMEM61	0.927374302	0.873684211	0.988095238
FAM163A AND NOT-CERS3	0.937142857	0.901098901	0.976190476
FAM163A AND NOT-FAAH	0.954022989	0.922222222	0.988095238
FAM163A AND NOT-OPN5	0.954022989	0.922222222	0.988095238
FAM163A AND NOT-GPRC6A	0.953488372	0.931818182	0.976190476
FAM163A AND NOT-MLC1	0.948571429	0.912087912	0.988095238
FAM163A AND NOT-SLC7A8	0.936416185	0.91011236	0.964285714
FAM163A AND ABCB10	0.937142857	0.901098901	0.976190476
FAM163A AND SEC11A	0.952380952	0.952380952	0.952380952
FAM163A AND SEC61G	0.947976879	0.921348315	0.976190476
FAM163A AND MMD	0.970760234	0.954022989	0.988095238
FAM163A AND LCLAT1	0.93258427	0.882978723	0.988095238
FAM163A AND NOT-GABRA6	0.937142857	0.901098901	0.976190476
FAM163A AND NOT-GABRD	0.937142857	0.901098901	0.976190476
FAM163A AND NOT-SLC24A2	0.927374302	0.873684211	0.988095238
FAM163A AND LETMD1	0.93258427	0.882978723	0.988095238
FAM163A AND NOT-TAS2R41	0.954022989	0.922222222	0.988095238
FAM163A AND NOT-OR7A17	0.93258427	0.882978723	0.988095238
FAM163A AND NOT-OR8G1	0.943181818	0.902173913	0.988095238
FAM163A AND NOT-OR10A3	0.93258427	0.882978723	0.988095238
FAM163A AND NOT-OR10H3	0.982248521	0.976470588	0.988095238
FAM163A AND NOT-OR4D1	0.958579882	0.952941176	0.964285714
FAM163A AND NOT-ADGRF1	0.954022989	0.922222222	0.988095238
FAM163A AND NOT-PCDH11X	0.952941176	0.941860465	0.964285714
FAM163A AND NOT-GLP1R	0.947976879	0.921348315	0.976190476
FAM163A AND NOT-GNRHR	0.948571429	0.912087912	0.988095238
FAM163A AND NOT-XCR1	0.93258427	0.882978723	0.988095238
FAM163A AND NOT-GPR6	0.922222222	0.864583333	0.988095238
FAM163A AND NOT-IZUMO1	0.922222222	0.864583333	0.988095238
FAM163A AND NOT-GPR37	0.947976879	0.921348315	0.976190476
FAM163A AND NOT-SLC6A16	0.922222222	0.864583333	0.988095238
FAM163A AND TMEM14A	0.937142857	0.901098901	0.976190476

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
PTPRZ1 AND NOT-TRPC1	0.9375	1	0.882352941
PTPRZ1 AND NOT-TLCD2	0.9375	1	0.882352941
PTPRZ1 AND NOT-C12orf73	0.9375	1	0.882352941
PTPRZ1 AND NOT-TMEM242	0.9375	1	0.882352941
PTPRZ1 AND TYROBP	0.9375	1	0.882352941
PTPRZ1 AND VCAM1	0.9375	1	0.882352941
PTPRZ1 AND NOT-VIPR1	0.9375	1	0.882352941
PTPRZ1 AND NOT-TRPV1	0.9375	1	0.882352941
PTPRZ1 AND NOT-WFS1	0.903225806	1	0.823529412
PTPRZ1 AND NOT-TMEM258	0.903225806	1	0.823529412
PTPRZ1 AND NOT-WNT3	0.9375	1	0.882352941
PTPRZ1 AND WNT11	0.9375	1	0.882352941
PTPRZ1 AND NOT-XK	0.903225806	1	0.823529412
PTPRZ1 AND NOT-ZNF7	0.9375	1	0.882352941
PTPRZ1 AND NOT-TMEM50B	0.9375	1	0.882352941
PTPRZ1 AND RELL1	0.9375	1	0.882352941
PTPRZ1 AND NOT-ZNF138	0.9375	1	0.882352941
PTPRZ1 AND NOT-CACNA1A	0.9375	1	0.882352941
PTPRZ1 AND NOT-CACNA1B	0.903225806	1	0.823529412
PTPRZ1 AND NOT-CACNA1E	0.903225806	1	0.823529412
PTPRZ1 AND LAPTM5	0.9375	1	0.882352941
PTPRZ1 AND DDR1	0.903225806	1	0.823529412
PTPRZ1 AND PXDN	0.903225806	1	0.823529412
PTPRZ1 AND RNF103	0.9375	1	0.882352941
PTPRZ1 AND SCG2	0.903225806	1	0.823529412
PTPRZ1 AND NOT-SLMAP	0.9375	1	0.882352941
PTPRZ1 AND NOT-SLC25A20	0.903225806	1	0.823529412
PTPRZ1 AND NOT-CRELD1	0.9375	1	0.882352941
PTPRZ1 AND NOT-VKORC1	0.903225806	1	0.823529412
PTPRZ1 AND ALG8	0.903225806	1	0.823529412
PTPRZ1 AND NOT-REEP5	0.9375	1	0.882352941
PTPRZ1 AND SMIM7	0.903225806	1	0.823529412
PTPRZ1 AND NOT-APOO	0.9375	1	0.882352941
PTPRZ1 AND NOT-DHRS11	0.9375	1	0.882352941
PTPRZ1 AND NOT-MFSD11	0.9375	1	0.882352941
PTPRZ1 AND NOT-GNPTAB	0.903225806	1	0.823529412
PTPRZ1 AND NOT-TMEM243	0.9375	1	0.882352941
PTPRZ1 AND NOT-TMEM43	0.9375	1	0.882352941
PTPRZ1 AND NOT-LRFN3	0.9375	1	0.882352941
PTPRZ1 AND MUL1	0.9375	1	0.882352941
PTPRZ1 AND NOT-GALNT14	0.903225806	1	0.823529412
PTPRZ1 AND NOT-SRD5A3	0.9375	1	0.882352941
PTPRZ1 AND GAL3ST4	0.9375	1	0.882352941
PTPRZ1 AND MANEA	0.9375	1	0.882352941
PTPRZ1 AND NOT-GALNT12	0.903225806	1	0.823529412
PTPRZ1 AND NOT-C1orf115	0.903225806	1	0.823529412
PTPRZ1 AND NOT-NIPAL2	0.9375	1	0.882352941
PTPRZ1 AND NOT-CAMKMT	0.9375	1	0.882352941
PTPRZ1 AND NOT-CCDC134	0.9375	1	0.882352941
PTPRZ1 AND CYBRD1	0.903225806	1	0.823529412
PTPRZ1 AND NOT-SLC35E1	0.903225806	1	0.823529412
PTPRZ1 AND SYNDIG1	0.903225806	1	0.823529412
PTPRZ1 AND NOT-ERMP1	0.903225806	1	0.823529412
PTPRZ1 AND NOT-PCNXL2	0.903225806	1	0.823529412
PTPRZ1 AND SEMA6D	0.903225806	1	0.823529412
PTPRZ1 AND NOT-PGAP1	0.9375	1	0.882352941
PTPRZ1 AND LRRTM4	0.9375	1	0.882352941
PTPRZ1 AND NOT-MYCT1	0.9375	1	0.882352941
PTPRZ1 AND NOT-TMEM254	0.9375	1	0.882352941
PTPRZ1 AND NOT-PIGZ	0.9375	1	0.882352941
PTPRZ1 AND NOT-MLLT10	0.9375	1	0.882352941
PTPRZ1 AND NOT-DNAJC5	0.9375	1	0.882352941
PTPRZ1 AND NOT-REEP4	0.9375	1	0.882352941
PTPRZ1 AND NOT-CYB5B	0.9375	1	0.882352941
PTPRZ1 AND NOT-PRRT1	0.9375	1	0.882352941
PTPRZ1 AND SLC2A10	0.903225806	1	0.823529412
PTPRZ1 AND TMX1	0.903225806	1	0.823529412
PTPRZ1 AND NOT-GDPD5	0.9375	1	0.882352941
PTPRZ1 AND NOT-YIPF5	0.9375	1	0.882352941
PTPRZ1 AND VWA9	0.9375	1	0.882352941
PTPRZ1 AND LMAN2L	0.903225806	1	0.823529412
PTPRZ1 AND APOLD1	0.9375	1	0.882352941
PTPRZ1 AND TMEM163	0.903225806	1	0.823529412
PTPRZ1 AND NOT-TSPAN14	0.9375	1	0.882352941
PTPRZ1 AND NOT-AKAP1	0.9375	1	0.882352941
PTPRZ1 AND NOT-RNF170	0.9375	1	0.882352941

Antigen Pair Logic	F1	Precision	Recall
FAM163A AND NOT-GRIK1	0.931818182	0.891304348	0.976190476
FAM163A AND CNIH4	0.927374302	0.873684211	0.988095238
FAM163A AND ALG5	0.927374302	0.873684211	0.988095238
FAM163A AND GUSB	0.947976879	0.921348315	0.976190476
FAM163A AND NOT-VN1R2	0.957575758	0.975308642	0.94047619
FAM163A AND NOT-HTR1B	0.927374302	0.873684211	0.988095238
FAM163A AND NOT-HTR6	0.922222222	0.864583333	0.988095238
FAM163A AND NOT-CLEC4D	0.948571429	0.912087912	0.988095238
FAM163A AND NOT-339166?	0.93258427	0.882978723	0.988095238
FAM163A AND NOT-FREM2	0.947368421	0.931034483	0.964285714
FAM163A AND NOT-COX8C	0.959537572	0.93258427	0.988095238
FAM163A AND NOT-IL12B	0.954022989	0.922222222	0.988095238
FAM163A AND NOT-IL12RB2	0.93258427	0.882978723	0.988095238
FAM163A AND NOT-ITGB6	0.952941176	0.941860465	0.964285714
FAM163A AND NOT-KCNA1	0.982248521	0.976470588	0.988095238
FAM163A AND NOT-SPEM1	0.922222222	0.864583333	0.988095238
FAM163A AND NOT-KCND3	0.922222222	0.864583333	0.988095238
FAM163A AND NOT-KCNJ10	0.937853107	0.892473118	0.988095238
FAM163A AND NOT-CLEC12B	0.948571429	0.912087912	0.988095238
FAM163A AND NOT-LHCGR	0.922222222	0.864583333	0.988095238
FAM163A AND PPAPDC2	0.942528736	0.911111111	0.976190476
FAM163A AND NOT-LRP4	0.937142857	0.901098901	0.976190476
FAM163A AND NOT-HAPLN4	0.931818182	0.891304348	0.976190476
FAM163A AND NOT-ERVFRD-1	0.927374302	0.873684211	0.988095238
FAM163A AND NOT-LCN10	0.931818182	0.891304348	0.976190476
FAM163A AND NOT-MME	0.953488372	0.931818182	0.976190476
FAM163A AND NOT-MOG	0.943181818	0.902173913	0.988095238
FAM163A AND MARCH11	0.931818182	0.891304348	0.976190476
FAM163A AND NOT-ATP1A2	0.927374302	0.873684211	0.988095238
FAM163A AND NOT-ATP1B2	0.958579882	0.952941176	0.964285714
FAM163A AND NOT-NPY5R	0.970760234	0.954022989	0.988095238
FAM163A AND NOT-NTRK2	0.927374302	0.873684211	0.988095238
FAM163A AND NOT-ATP2B2	0.93258427	0.882978723	0.988095238
FAM163A AND NOT-CD207	0.937142857	0.901098901	0.976190476
FAM163A AND IER3IP1	0.948571429	0.912087912	0.988095238
FAM163A AND FAM8A1	0.964705882	0.953488372	0.976190476
FAM163A AND TMEM14C	0.942528736	0.911111111	0.976190476
FAM163A AND ATP5F1	0.954022989	0.922222222	0.988095238
FAM163A AND PFN2	0.959537572	0.93258427	0.988095238
FAM163A AND SLC25A3	0.965116279	0.943181818	0.988095238
FAM163A AND NOT-PI3	0.947976879	0.921348315	0.976190476
FAM163A AND SLC38A2	0.96969697	0.987654321	0.952380952
FAM163A AND NDUFB11	0.954022989	0.922222222	0.988095238
FAM163A AND NOT-EQTN	0.937142857	0.901098901	0.976190476
FAM163A AND CRLS1	0.922222222	0.864583333	0.988095238
FAM163A AND NOT-DCHS2	0.976470588	0.965116279	0.988095238
FAM163A AND ZDHHC4	0.93258427	0.882978723	0.988095238
FAM163A AND NOT-TMEM144	0.937142857	0.901098901	0.976190476
FAM163A AND NOT-STYK1	0.975903614	0.987804878	0.964285714
FAM163A AND LRRC59	0.93258427	0.882978723	0.988095238
FAM163A AND PLGRKT	0.948571429	0.912087912	0.988095238
FAM163A AND NOT-SLC7A10	0.948571429	0.912087912	0.988095238
FAM163A AND DIABLO	0.957575758	0.975308642	0.94047619
FAM163A AND NOT-C2orf83	0.959064327	0.942528736	0.976190476
FAM163A AND C5orf15	0.937142857	0.901098901	0.976190476
FAM163A AND NOT-SLC17A7	0.927374302	0.873684211	0.988095238
FAM163A AND VANGL2	0.942528736	0.911111111	0.976190476
FAM163A AND NOT-HHATL	0.937853107	0.892473118	0.988095238
FAM163A AND TXNDC16	0.952941176	0.941860465	0.964285714
FAM163A AND SELK	0.936416185	0.91011236	0.964285714
FAM163A AND NOT-RAD51B	0.953488372	0.931818182	0.976190476
FAM163A AND NOT-CACNG6	0.954022989	0.922222222	0.988095238
FAM163A AND NOT-LGR6	0.958083832	0.963855422	0.952380952
FAM163A AND NOT-CDH26	0.922222222	0.864583333	0.988095238
FAM163A AND NOT-ROS1	0.93258427	0.882978723	0.988095238
FAM163A AND NOT-CTXN3	0.965116279	0.943181818	0.988095238
FAM163A AND RTN1	0.937142857	0.901098901	0.976190476
FAM163A AND NOT-SLC22A23	0.927374302	0.873684211	0.988095238
FAM163A AND NOT-SCN1A	0.93258427	0.882978723	0.988095238
FAM163A AND NOT-SCN1B	0.941860465	0.920454545	0.964285714
FAM163A AND NOT-SCN10A	0.93258427	0.882978723	0.988095238
FAM163A AND SMIM15	0.922222222	0.864583333	0.988095238
FAM163A AND NOT-SMIM10	0.93258427	0.882978723	0.988095238
FAM163A AND MFSD1	0.941860465	0.920454545	0.964285714
FAM163A AND NOT-LMF1	0.937853107	0.892473118	0.988095238
FAM163A AND NOT-LRRC19	0.959537572	0.93258427	0.988095238

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
PTPRZ1 AND NETO2	0.9375	1	0.882352941
PTPRZ1 AND NOT-SFXN3	0.9375	1	0.882352941
PTPRZ1 AND NOT-PNPLA4	0.903225806	1	0.823529412
PTPRZ1 AND NOT-TMEM187	0.903225806	1	0.823529412
PTPRZ1 AND KLRC4	0.903225806	1	0.823529412
PTPRZ1 AND NOT-FZD6	0.9375	1	0.882352941
PTPRZ1 AND FZD7	0.9375	1	0.882352941
PTPRZ1 AND FZD8	0.9375	1	0.882352941
PTPRZ1 AND EMC6	0.9375	1	0.882352941
PTPRZ1 AND NOT-TMEM47	0.9375	1	0.882352941
PTPRZ1 AND NOT-CRISPLD2	0.903225806	1	0.823529412
PTPRZ1 AND L3MBTL2	0.9375	1	0.882352941
PTPRZ1 AND NOT-ARMC10	0.9375	1	0.882352941
PTPRZ1 AND OR1A1	0.9375	1	0.882352941
PTPRZ1 AND NOT-ESYT3	0.9375	1	0.882352941
PTPRZ1 AND NOT-TMTC1	0.903225806	1	0.823529412
PTPRZ1 AND TMEM120A	0.9375	1	0.882352941
PTPRZ1 AND NOT-TM2D2	0.903225806	1	0.823529412
PTPRZ1 AND NOT-OR3A3	0.9375	1	0.882352941
PTPRZ1 AND NOT-STARD3NL	0.9375	1	0.882352941
PTPRZ1 AND TMEM133	0.903225806	1	0.823529412
PTPRZ1 AND NOT-TM2D1	0.9375	1	0.882352941
PTPRZ1 AND B3GNT5	0.903225806	1	0.823529412
PTPRZ1 AND NOT-ZNRF3	0.9375	1	0.882352941
PTPRZ1 AND NOT-SLITRK6	0.9375	1	0.882352941
PTPRZ1 AND NOT-USP48	0.903225806	1	0.823529412
PTPRZ1 AND NOT-TMEM117	0.9375	1	0.882352941
PTPRZ1 AND NOT-RHBDD1	0.9375	1	0.882352941
PTPRZ1 AND NOT-SLC37A3	0.9375	1	0.882352941
PTPRZ1 AND NOT-FAM213A	0.903225806	1	0.823529412
PTPRZ1 AND NOT-TMEM246	0.903225806	1	0.823529412
PTPRZ1 AND NOT-PPAPDC1B	0.9375	1	0.882352941
PTPRZ1 AND ZNF559	0.9375	1	0.882352941
PTPRZ1 AND NOT-ATRN	0.9375	1	0.882352941
PTPRZ1 AND NOT-SLC12A8	0.9375	1	0.882352941
PTPRZ1 AND NOT-TPST2	0.903225806	1	0.823529412
PTPRZ1 AND TPST1	0.903225806	1	0.823529412
PTPRZ1 AND NOT-KIRREL3	0.903225806	1	0.823529412
PTPRZ1 AND NOT-HIATL1	0.9375	1	0.882352941
PTPRZ1 AND NOT-DGAT2	0.9375	1	0.882352941
PTPRZ1 AND MGARP	0.903225806	1	0.823529412
PTPRZ1 AND NOT-USP30	0.9375	1	0.882352941
PTPRZ1 AND NOT-PPAPDC3	0.9375	1	0.882352941
PTPRZ1 AND NOT-LINGO1	0.9375	1	0.882352941
PTPRZ1 AND NOT-TMTC4	0.903225806	1	0.823529412
PTPRZ1 AND NOT-TMEM87B	0.9375	1	0.882352941
PTPRZ1 AND NOT-LRP11	0.903225806	1	0.823529412
PTPRZ1 AND DISP1	0.903225806	1	0.823529412
PTPRZ1 AND FRMD5	0.9375	1	0.882352941
PTPRZ1 AND NOT-COX14	0.903225806	1	0.823529412
PTPRZ1 AND NOT-TMEM128	0.9375	1	0.882352941
PTPRZ1 AND NOT-TMEM141	0.903225806	1	0.823529412
PTPRZ1 AND NOT-NDST2	0.9375	1	0.882352941
PTPRZ1 AND NOT-IFITM1	0.903225806	1	0.823529412
PTPRZ1 AND NOT-DGKE	0.9375	1	0.882352941
PTPRZ1 AND NOT-KIAA1644	0.903225806	1	0.823529412
PTPRZ1 AND NOT-LGR5	0.903225806	1	0.823529412
PTPRZ1 AND NOT-CDC14B	0.9375	1	0.882352941
PTPRZ1 AND NOT-CAV1	0.903225806	1	0.823529412
PTPRZ1 AND NOT-LY6D	0.9375	1	0.882352941
PTPRZ1 AND NOT-CAV2	0.9375	1	0.882352941
PTPRZ1 AND NOT-PPAP2A	0.9375	1	0.882352941
PTPRZ1 AND VAMP8	0.903225806	1	0.823529412
PTPRZ1 AND NOT-DGAT1	0.9375	1	0.882352941
PTPRZ1 AND NOT-B3GALT4	0.903225806	1	0.823529412
PTPRZ1 AND NOT-B3GALNT1	0.9375	1	0.882352941
PTPRZ1 AND NOT-TNFSF10	0.9375	1	0.882352941
PTPRZ1 AND ADAM23	0.903225806	1	0.823529412
PTPRZ1 AND NOT-CDS2	0.9375	1	0.882352941
PTPRZ1 AND NOT-PEX11A	0.9375	1	0.882352941
PTPRZ1 AND NOT-PROM1	0.9375	1	0.882352941
PTPRZ1 AND NOT-SGPL1	0.9375	1	0.882352941
PTPRZ1 AND NOT-CCKBR	0.9375	1	0.882352941
PTPRZ1 AND NOT-PPP1R3F	0.9375	1	0.882352941
PTPRZ1 AND NOT-ABCC10	0.9375	1	0.882352941
PTPRZ1 AND NOT-RHOT2	0.9375	1	0.882352941

Antigen Pair Logic	F1	Precision	Recall
FAM163A AND NOT-SLC1A3	0.93258427	0.882978723	0.988095238
FAM163A AND NOT-SLC5A1	0.954022989	0.922222222	0.988095238
FAM163A AND NOT-SLC13A1	0.982248521	0.976470588	0.988095238
FAM163A AND NOT-SLC15A2	0.959537572	0.93258427	0.988095238
FAM163A AND SLC20A1	0.93258427	0.882978723	0.988095238
FAM163A AND NOT-CYP4F12	0.941860465	0.920454545	0.964285714
FAM163A AND SSR2	0.922222222	0.864583333	0.988095238
FAM163A AND NOT-SSTR1	0.927374302	0.873684211	0.988095238
FAM163A AND NOT-BRS3	0.937853107	0.892473118	0.988095238
FAM163A AND NOT-BTC	0.93258427	0.882978723	0.988095238
FAM163A AND NOT-TRPC3	0.987951807	1	0.976190476
FAM163A AND NOT-C1orf234	0.937142857	0.901098901	0.976190476
FAM163A AND NOT-TYR	0.937853107	0.892473118	0.988095238
FAM163A AND NOT-UPK1B	0.964705882	0.953488372	0.976190476
FAM163A AND TMEM258	0.943181818	0.902173913	0.988095238
FAM163A AND NOT-CACNA1A	0.931818182	0.891304348	0.976190476
FAM163A AND NOT-BSND	0.946745562	0.941176471	0.952380952
FAM163A AND NOT-YIPF2	0.943181818	0.902173913	0.988095238
FAM163A AND GDAP1L1	0.927374302	0.873684211	0.988095238
FAM163A AND NOT-SMIM2	0.943181818	0.902173913	0.988095238
FAM163A AND DERL1	0.922222222	0.864583333	0.988095238
FAM163A AND NOT-C1orf115	0.931818182	0.891304348	0.976190476
FAM163A AND NOT-TMC5	0.948571429	0.912087912	0.988095238
FAM163A AND TUSC3	0.93258427	0.882978723	0.988095238
FAM163A AND NOT-PAQR6	0.937853107	0.892473118	0.988095238
FAM163A AND NOT-GLRA3	0.988095238	0.988095238	0.988095238
FAM163A AND NOT-PRRT1	0.942528736	0.911111111	0.976190476
FAM163A AND NIPA2	0.947976879	0.921348315	0.976190476
FAM163A AND EMC6	0.93258427	0.882978723	0.988095238
FAM163A AND NOT-ESYT3	0.964705882	0.953488372	0.976190476
FAM163A AND NOT-SLC25A2	0.931818182	0.891304348	0.976190476
FAM163A AND NOT-SPATA9	0.937853107	0.892473118	0.988095238
FAM163A AND NOT-C2orf40	0.927374302	0.873684211	0.988095238
FAM163A AND JAGN1	0.937853107	0.892473118	0.988095238
FAM163A AND NOT-FUT10	0.970760234	0.954022989	0.988095238
FAM163A AND NOT-SLC22A16	0.931818182	0.891304348	0.976190476
FAM163A AND NOT-SLC4A4	0.93258427	0.882978723	0.988095238
FAM163A AND NOT-ADAM7	0.93258427	0.882978723	0.988095238
FAM163A AND NOT-CACNA1I	0.937853107	0.892473118	0.988095238
FAM163A AND NOT-LMLN	0.927374302	0.873684211	0.988095238
FAM163A AND NOT-OPALIN	0.943181818	0.902173913	0.988095238
FAM163A AND NOT-SFXN5	0.922222222	0.864583333	0.988095238
FAM163A AND NOT-SIGLEC6	0.976190476	0.976190476	0.976190476
FAM163A AND NOT-TMEM63A	0.937853107	0.892473118	0.988095238
SLC10A4 AND NOT-MFSD4	0.927374302	0.873684211	0.988095238
MARCH11 AND NOT-MFSD4	0.946745562	0.941176471	0.952380952
SLC10A4 AND NOT-DCST1	0.954022989	0.922222222	0.988095238
ST8SIA2 AND NOT-DCST1	0.950617284	0.987179487	0.916666667
SLC10A4 AND NOT-SLC9B1	0.948571429	0.912087912	0.988095238
ST8SIA2 AND NOT-SLC9B1	0.950617284	0.987179487	0.916666667
MARCH11 AND NOT-SLC38A11	0.946745562	0.941176471	0.952380952
ST8SIA2 AND NOT-SLC38A11	0.936708861	1	0.880952381
ST8SIA2 AND NOT-SLC23A3	0.936708861	1	0.880952381
SLC10A4 AND NOT-PPM1L	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-IGSF11	0.982035928	0.987951807	0.976190476
ST8SIA2 AND NOT-IGSF11	0.95	1	0.904761905
SLC10A4 AND NOT-SLC38A9	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-SLC25A48	0.976470588	0.965116279	0.988095238
ST8SIA2 AND NOT-SLC25A48	0.956521739	1	0.916666667
SLC10A4 AND NOT-MARVELD2	0.927374302	0.873684211	0.988095238
SLC10A4 AND CCDC112	0.953488372	0.931818182	0.976190476
SLC10A4 AND NOT-ADRB1	0.982035928	0.987951807	0.976190476
GPR19 AND NOT-ADRB1	0.957575758	0.975308642	0.94047619
MARCH11 AND NOT-ADRB1	0.958083832	0.963855422	0.952380952
SLC10A4 AND NOT-SLC2A12	0.959537572	0.93258427	0.988095238
GPR19 AND NOT-SLC2A12	0.96969697	0.987654321	0.952380952
MARCH11 AND NOT-SLC2A12	0.958083832	0.963855422	0.952380952
ADGRB3 AND NOT-SLC2A12	0.950617284	0.987179487	0.916666667
ST8SIA2 AND NOT-SLC2A12	0.956521739	1	0.916666667
SLC10A4 AND NOT-MBOAT1	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-NKAIN2	0.988095238	0.988095238	0.988095238
ST8SIA2 AND NOT-CYP1A1	0.956521739	1	0.916666667
SLC10A4 AND CCDC167	0.947368421	0.931034483	0.964285714
MARCH11 AND NOT-CLEC2L	0.946745562	0.941176471	0.952380952
SLC10A4 AND NOT-PEBP4	0.975903614	0.987804878	0.964285714
MARCH11 AND NOT-PEBP4	0.969325153	1	0.94047619

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
PTPRZ1 AND NOT-C9orf69	0.9375	1	0.882352941
PTPRZ1 AND PKD2L1	0.9375	1	0.882352941
PTPRZ1 AND NOT-MCU	0.9375	1	0.882352941
PTPRZ1 AND SLC7A7	0.903225806	1	0.823529412
PTPRZ1 AND SLC7A6	0.9375	1	0.882352941
PTPRZ1 AND NOT-ANGPTL1	0.9375	1	0.882352941
PTPRZ1 AND NOT-CCDC126	0.9375	1	0.882352941
PTPRZ1 AND NOT-CLDN12	0.9375	1	0.882352941
PTPRZ1 AND NOT-CLDN8	0.9375	1	0.882352941
PTPRZ1 AND NOT-TMEM261	0.9375	1	0.882352941
PTPRZ1 AND NOT-ACVR1	0.9375	1	0.882352941
PTPRZ1 AND AIFM1	0.903225806	1	0.823529412
PTPRZ1 AND NOT-SYNGR3	0.903225806	1	0.823529412
PTPRZ1 AND NOT-SYNGR2	0.9375	1	0.882352941
PTPRZ1 AND NOT-SYNGR1	0.9375	1	0.882352941
PTPRZ1 AND NOT-PKDCC	0.9375	1	0.882352941
PTPRZ1 AND NOT-BOC	0.9375	1	0.882352941
PTPRZ1 AND NOT-MYADM	0.9375	1	0.882352941
PTPRZ1 AND NOT-RFT1	0.9375	1	0.882352941
PTPRZ1 AND NOT-SLC33A1	0.9375	1	0.882352941
PTPRZ1 AND NOT-ACVR1B	0.9375	1	0.882352941
PTPRZ1 AND NOT-LARGE	0.903225806	1	0.823529412
PTPRZ1 AND NOT-VAPA	0.903225806	1	0.823529412
PTPRZ1 AND NOT-DAPL1	0.903225806	1	0.823529412
PTPRZ1 AND NOT-CCPG1	0.903225806	1	0.823529412
PTPRZ1 AND PXYLP1	0.903225806	1	0.823529412
PTPRZ1 AND NOT-DHRS3	0.9375	1	0.882352941
PTPRZ1 AND TIMM50	0.903225806	1	0.823529412
PTPRZ1 AND NOT-DTD1	0.9375	1	0.882352941
PTPRZ1 AND NOT-TMEM183A	0.9375	1	0.882352941
PTPRZ1 AND NOT-BPIFB1	0.9375	1	0.882352941
PTPRZ1 AND NOT-REEP6	0.9375	1	0.882352941
PTPRZ1 AND CD83	0.903225806	1	0.823529412
PTPRZ1 AND PIGM	0.903225806	1	0.823529412
PTPRZ1 AND NOT-IGSF8	0.9375	1	0.882352941
PTPRZ1 AND NOT-PGAP3	0.9375	1	0.882352941
PTPRZ1 AND NOT-GTF3C3	0.903225806	1	0.823529412
PTPRZ1 AND B4GALT5	0.903225806	1	0.823529412
PTPRZ1 AND NOT-OPALIN	0.903225806	1	0.823529412
PTPRZ1 AND NOT-SDR42E1	0.9375	1	0.882352941
PTPRZ1 AND NOT-TM9SF2	0.9375	1	0.882352941
PTPRZ1 AND NOT-NRXN1	0.9375	1	0.882352941
PTPRZ1 AND NOT-NRXN2	0.9375	1	0.882352941
PTPRZ1 AND NOT-STOML1	0.903225806	1	0.823529412
PTPRZ1 AND TTYH2	0.903225806	1	0.823529412
PTPRZ1 AND NOT-SFXN1	0.9375	1	0.882352941
PTPRZ1 AND SFXN5	0.903225806	1	0.823529412
PTPRZ1 AND NOT-TMEM203	0.9375	1	0.882352941
PTPRZ1 AND ABCG2	0.9375	1	0.882352941
PTPRZ1 AND EIF2AK3	0.9375	1	0.882352941
PTPRZ1 AND NOT-ITM2A	0.9375	1	0.882352941
PTPRZ1 AND NOT-AKAP6	0.903225806	1	0.823529412
PTPRZ1 AND NOT-STX8	0.903225806	1	0.823529412
PTPRZ1 AND SLC4A8	0.9375	1	0.882352941
PTPRZ1 AND NOT-SCAMP1	0.903225806	1	0.823529412
PTPRZ1 AND NOT-GOSR1	0.9375	1	0.882352941
PTPRZ1 AND ENTPD1	0.903225806	1	0.823529412
PTPRZ1 AND NOT-GABBR2	0.903225806	1	0.823529412
PTPRZ1 AND SLC25A44	0.903225806	1	0.823529412
PTPRZ1 AND CD69	0.903225806	1	0.823529412
PTPRZ1 AND CD70	0.9375	1	0.882352941
PTPRZ1 AND NOT-LAPTM4A	0.9375	1	0.882352941
PTPRZ1 AND NOT-CLSTN3	0.9375	1	0.882352941
PTPRZ1 AND MLEC	0.903225806	1	0.823529412
PTPRZ1 AND SUSD6	0.903225806	1	0.823529412
PTPRZ1 AND NOT-TOMM20	0.9375	1	0.882352941
PTPRZ1 AND NOT-C2CD5	0.903225806	1	0.823529412
PTPRZ1 AND NOT-MFAP3L	0.9375	1	0.882352941
PTPRZ1 AND NOT-C2CD2L	0.9375	1	0.882352941
PTPRZ1 AND NOT-KIAA0319	0.903225806	1	0.823529412
PTPRZ1 AND NOT-TOMM70A	0.9375	1	0.882352941
PTPRZ1 AND NOT-LPPR4	0.9375	1	0.882352941
PTPRZ1 AND NOT-SV2B	0.903225806	1	0.823529412
PTPRZ1 AND SV2A	0.9375	1	0.882352941
PTPRZ1 AND NOT-ATP2C2	0.9375	1	0.882352941
PTPRZ1 AND NOT-FAM20B	0.9375	1	0.882352941

Antigen Pair Logic	F1	Precision	Recall
SLC10A4 AND NOT-KCNU1	0.947976879	0.921348315	0.976190476
SLC10A4 AND NOT-CYP4B1	0.943181818	0.902173913	0.988095238
SLC10A4 AND NOT-OR1Q1	0.976470588	0.965116279	0.988095238
MARCH11 AND NOT-FAAH2	0.946745562	0.941176471	0.952380952
ST8SIA2 AND NOT-MOSPD2	0.956521739	1	0.916666667
SLC10A4 AND NOT-CLEC12A	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-TMTC3	0.922222222	0.864583333	0.988095238
ST8SIA2 AND NOT-TMEM229B	0.95	1	0.904761905
MARCH11 AND NOT-RHBDL3	0.946745562	0.941176471	0.952380952
SLC10A4 AND NOT-CNST	0.927374302	0.873684211	0.988095238
ST8SIA2 AND NOT-CABP7	0.95	1	0.904761905
SLC10A4 AND NOT-FAM171B	0.927374302	0.873684211	0.988095238
MARCH11 AND NOT-SPTSSB	0.946745562	0.941176471	0.952380952
ST8SIA2 AND NOT-SPTSSB	0.936708861	1	0.880952381
MARCH11 AND NOT-GALNTL5	0.957575758	0.975308642	0.94047619
ST8SIA2 AND NOT-PKD1L1	0.956521739	1	0.916666667
SLC10A4 AND NOT-HTR3C	0.941860465	0.920454545	0.964285714
SLC10A4 AND NOT-KCNG3	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-CLYBL	0.927374302	0.873684211	0.988095238
MARCH11 AND NOT-CLYBL	0.951219512	0.975	0.928571429
GPR19 AND NOT-DIO2	0.963414634	0.9875	0.94047619
MARCH11 AND NOT-DIO2	0.957575758	0.975308642	0.94047619
ST8SIA2 AND NOT-DIO2	0.95	1	0.904761905
ST8SIA2 AND NOT-SLC26A3	0.956521739	1	0.916666667
ST8SIA2 AND NOT-DRD1	0.950617284	0.987179487	0.916666667
MARCH11 AND NOT-DSCAM	0.946745562	0.941176471	0.952380952
SLC10A4 AND NOT-DTNA	0.937853107	0.892473118	0.988095238
SLC10A4 AND NOT-HBEGF	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-APLNR	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-S1PR1	0.931818182	0.891304348	0.976190476
SLC10A4 AND NOT-EDNRB	0.931818182	0.891304348	0.976190476
GPR19 AND NOT-EGF	0.950617284	0.987179487	0.916666667
MARCH11 AND NOT-EGF	0.958083832	0.963855422	0.952380952
ST8SIA2 AND NOT-EGF	0.950617284	0.987179487	0.916666667
SLC10A4 AND NOT-MEGF8	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-MEGF9	0.943181818	0.902173913	0.988095238
SLC10A4 AND NOT-EGFR	0.948571429	0.912087912	0.988095238
GPR19 AND NOT-EGFR	0.951807229	0.963414634	0.94047619
MARCH11 AND NOT-EGFR	0.963414634	0.9875	0.94047619
ST8SIA2 AND NOT-EGFR	0.956521739	1	0.916666667
GPR19 AND NOT-GRAMD2	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-GRAMD2	0.946745562	0.941176471	0.952380952
ST8SIA2 AND NOT-GRAMD2	0.950617284	0.987179487	0.916666667
SLC10A4 AND NOT-MCEMP1	0.922222222	0.864583333	0.988095238
ST8SIA2 AND NOT-APLF	0.943396226	1	0.892857143
SLC10A4 AND NOT-SPNS3	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-GAPT	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-C9orf91	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-VMA21	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-CERS3	0.942528736	0.911111111	0.976190476
SLC10A4 AND NOT-ERBB4	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-ABCA2	0.959537572	0.93258427	0.988095238
SLC10A4 AND NOT-ETV6	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-F3	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-FAAH	0.959537572	0.93258427	0.988095238
SLC10A4 AND NOT-LRRC55	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-MS4A15	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-FCER1A	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-SLC39A12	0.948571429	0.912087912	0.988095238
SLC10A4 AND NOT-OPN5	0.954022989	0.922222222	0.988095238
SLC10A4 AND NOT-MMD2	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-GPRC6A	0.947976879	0.921348315	0.976190476
SLC10A4 AND NOT-LHFPL5	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-CLCA4	0.937853107	0.892473118	0.988095238
SLC10A4 AND NOT-SLITRK3	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-ENPP4	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-KLHDC10	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-FAIM2	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-NFASC	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-CLCC1	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-MLC1	0.959537572	0.93258427	0.988095238
SLC10A4 AND NOT-FLT3	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-KIAA1024	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-ACSL6	0.948571429	0.912087912	0.988095238
SLC10A4 AND NOT-FAM189A1	0.952941176	0.941860465	0.964285714
SLC10A4 AND NOT-SLC7A8	0.936416185	0.91011236	0.964285714

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
PTPRZ1 AND NOT-MFN2	0.9375	1	0.882352941
PTPRZ1 AND NOT-CLEC2B	0.903225806	1	0.823529412
PTPRZ1 AND NOT-SLC12A6	0.9375	1	0.882352941
PTPRZ1 AND DGCR2	0.9375	1	0.882352941
TPST1 AND NOT-SIGIRR	0.903225806	1	0.823529412
TPST1 AND NOT-PERP	0.903225806	1	0.823529412
TPST1 AND NOT-VIPR1	0.903225806	1	0.823529412
TPST1 AND NOT-SLMAP	0.903225806	1	0.823529412
NOT-TNFSF10 AND GAL3ST4	0.9375	1	0.882352941
TPST1 AND NOT-TNFSF10	0.903225806	1	0.823529412
TPST1 AND NOT-TM9SF2	0.903225806	1	0.823529412
Adenocarcinoma|Breast Neoplasms (Breast A)			
CYB561 AND NOT-SLC2A12	0.923076923	0.92307692	0.923076923
RHBDF1 AND NOT-TMEM256	0.916666667	1	0.846153846
HSD3B7 AND NOT-TMEM256	0.916666667	1	0.846153846
ADAM12 AND NOT-TMEM256	0.916666667	1	0.846153846
SLC5A6 AND NOT-TMEM256	0.916666667	1	0.846153846
GGT5 AND NOT-TSPAN8	0.916666667	1	0.846153846
ADAM12 AND NOT-ADGRD1	0.916666667	1	0.846153846
LTBP3 AND NOT-HBD	0.916666667	1	0.846153846
MARVELD3 AND NOT-TMEM220	0.916666667	1	0.846153846
RHBDF1 AND NOT-SMIM5	0.96	1	0.923076923
LMLN AND NOT-XK	0.916666667	1	0.846153846
Glioma (Glioma)			
PTPRZ1	0.985507246	0.98550725	0.985507246
NLGN1	0.917293233	0.953125	0.884057971
SCRG1	0.909090909	0.87837838	0.942028986
PRAF2	0.905109489	0.91176471	0.898550725
SYT11	0.951048951	0.91891892	0.985507246
AGPAT5	0.905109489	0.91176471	0.898550725
CHST10	0.915492958	0.89041096	0.942028986
TPST1	0.917293233	0.953125	0.884057971
NLGN1 AND NOT-TNFRSF17	0.938461538	1	0.884057971
ITGAV AND NOT-TNFRSF17	0.928571429	0.91549296	0.942028986
PRAF2 AND PRAF2	0.905109489	0.91176471	0.898550725
SYT11 AND TNFRSF17	0.943661972	0.91780822	0.971014493
SCRG1 AND TNFRSF17	0.907801418	0.88888889	0.927536232
TPST1 AND TPST1	0.917293233	0.953125	0.884057971
AGPAT5 AND NOT-TNFRSF17	0.946564885	1	0.898550725
PTPRZ1 AND NOT-TNFRSF17	0.97810219	0.98529412	0.971014493
NRCAM AND NOT-FOLH1	0.909090909	0.95238095	0.869565217
NLGN1 AND NLGN1	0.917293233	0.953125	0.884057971
PRAF2 AND NOT-FOLH1	0.909090909	0.95238095	0.869565217
XPR1 AND FOLH1	0.909090909	0.95238095	0.869565217
SYT11 AND NOT-FOLH1	0.956521739	0.95652174	0.956521739
SCRG1 AND NOT-FOLH1	0.919708029	0.92647059	0.913043478
TPST1 AND TPST1	0.917293233	0.953125	0.884057971
AGPAT5 AND FOLH1	0.905109489	0.91176471	0.898550725
CHST10 AND NOT-FOLH1	0.935251799	0.92857143	0.942028986
PTPRZ1 AND NOT-FOLH1	0.970588235	0.98507463	0.956521739
PRAF2 AND PRAF2	0.905109489	0.91176471	0.898550725
SYT11 AND CD44	0.928571429	0.91549296	0.942028986
SCRG1 AND CD44	0.909090909	0.87837838	0.942028986
TPST1 AND TPST1	0.917293233	0.953125	0.884057971
CHST10 AND CD44	0.925373134	0.95384615	0.898550725
PTPRZ1 AND CD44	0.985507246	0.98550725	0.985507246
NLGN1 AND NLGN1	0.917293233	0.953125	0.884057971
ITGAV AND NOT-WT1	0.921985816	0.90277778	0.942028986
PRAF2 AND PRAF2	0.905109489	0.91176471	0.898550725
SYT11 AND WT1	0.920863309	0.91428571	0.927536232
SCRG1 AND SCRG1	0.909090909	0.87837838	0.942028986
TPST1 AND TPST1	0.917293233	0.953125	0.884057971
AGPAT5 AND WT1	0.905109489	0.91176471	0.898550725
PTPRZ1 AND WT1	0.985507246	0.98550725	0.985507246
NLGN1 AND NLGN1	0.917293233	0.953125	0.884057971
ITGAV AND CD33	0.929577465	0.90410959	0.956521739
PRAF2 AND PRAF2	0.905109489	0.91176471	0.898550725
SYT11 AND CD33	0.951048951	0.91891892	0.985507246
SCRG1 AND CD33	0.909090909	0.87837838	0.942028986
TPST1 AND TPST1	0.917293233	0.953125	0.884057971
AGPAT5 AND NOT-CD33	0.939393939	0.98412698	0.898550725
CHST10 AND CD33	0.9	0.88732394	0.913043478
PTPRZ1 AND CD33	0.985507246	0.98550725	0.985507246
NLGN1 AND NLGN1	0.917293233	0.953125	0.884057971
NOT-PRAF2 AND ST8SIA1	0.90647482	0.9	0.913043478
SYT11 AND ST8SIA1	0.951048951	0.91891892	0.985507246

Antigen Pair Logic	F1	Precision	Recall
SLC10A4 AND NOT-SLC44A1	0.970760234	0.954022989	0.988095238
SLC10A4 AND NOT-FOLH1	0.954022989	0.922222222	0.988095238
SLC10A4 AND SEC11A	0.946745562	0.941176471	0.952380952
SLC10A4 AND SEC61G	0.947976879	0.921348315	0.976190476
SLC10A4 AND MMD	0.954022989	0.922222222	0.988095238
SLC10A4 AND NOT-SEZ6L	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-HYAL4	0.937142857	0.901098901	0.976190476
SLC10A4 AND NOT-PIGN	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-BACE1	0.948571429	0.912087912	0.988095238
SLC10A4 AND NOT-CA14	0.93258427	0.882978723	0.988095238
SLC10A4 AND NOT-RABGAP1	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-SLC7A11	0.954022989	0.922222222	0.988095238
SLC10A4 AND NOT-FRRS1L	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-APOL2	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-EML2	0.958579882	0.952941176	0.964285714
SLC10A4 AND NOT-ABCA4	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-FUT2	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-FUT4	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-FUT7	0.941860465	0.920454545	0.964285714
SLC10A4 AND NOT-CADM2	0.957055215	0.987341772	0.928571429
SLC10A4 AND NOT-ZDHHC20	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-MYADML2	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-RNF144B	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-GABRA1	0.948571429	0.912087912	0.988095238
SLC10A4 AND NOT-GABRA3	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-GABRA4	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-GABRA6	0.959064327	0.942528736	0.976190476
SLC10A4 AND NOT-GABRB2	0.937853107	0.892473118	0.988095238
SLC10A4 AND NOT-SLC24A2	0.976470588	0.965116279	0.988095238
SLC10A4 AND NOT-METTL7A	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-SUMF2	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-LY6G6F	0.937853107	0.892473118	0.988095238
SLC10A4 AND NOT-SGMS1	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-NALCN	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-TAS2R41	0.954022989	0.922222222	0.988095238
SLC10A4 AND NOT-OR2L2	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-IL1RAPL2	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-OR7A17	0.948571429	0.912087912	0.988095238
SLC10A4 AND NOT-OR8G1	0.943181818	0.902173913	0.988095238
SLC10A4 AND NOT-OR10A3	0.943181818	0.902173913	0.988095238
SLC10A4 AND NOT-GCNT1	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-OR10H3	0.948571429	0.912087912	0.988095238
SLC10A4 AND NOT-OR7C2	0.951807229	0.963414634	0.94047619
SLC10A4 AND NOT-OR7A5	0.93258427	0.882978723	0.988095238
SLC10A4 AND NOT-OR4D1	0.941860465	0.920454545	0.964285714
SLC10A4 AND NOT-ADGRF1	0.954022989	0.922222222	0.988095238
SLC10A4 AND NOT-B3GAT1	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-C5AR2	0.937853107	0.892473118	0.988095238
SLC10A4 AND NOT-COQ2	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-PCDH11X	0.952941176	0.941860465	0.964285714
SLC10A4 AND NOT-GLP1R	0.947976879	0.921348315	0.976190476
SLC10A4 AND NOT-GNRHR	0.948571429	0.912087912	0.988095238
SLC10A4 AND NOT-SLCO4A1	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-XCR1	0.954022989	0.922222222	0.988095238
SLC10A4 AND NOT-GPR6	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-C16orf54	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-TMEM235	0.93258427	0.882978723	0.988095238
SLC10A4 AND NOT-IZUMO1	0.93258427	0.882978723	0.988095238
SLC10A4 AND NOT-SMIM24	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-OR2L13	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-GPR26	0.93258427	0.882978723	0.988095238
SLC10A4 AND NOT-PRRT3	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-ARL10	0.976190476	0.976190476	0.976190476
SLC10A4 AND NOT-RELL2	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-RNF180	0.937853107	0.892473118	0.988095238
SLC10A4 AND NOT-GPR37	0.982248521	0.976470588	0.988095238
SLC10A4 AND NOT-GPR39	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-YIPF6	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-ILDR1	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-ANK1	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-GRIA3	0.976470588	0.965116279	0.988095238
SLC10A4 AND NOT-GRID2	0.959064327	0.942528736	0.976190476
SLC10A4 AND NOT-SLC6A16	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-GRIK1	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-GRIN2A	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-GRM3	0.965116279	0.943181818	0.988095238

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
SCRG1 AND ST8SIA1	0.909090909	0.87837838	0.942028986
PTPRZ1 AND ST8SIA1	0.985507246	0.98550725	0.985507246
NCAM1 AND NOT-CYP4F12	0.904761905	1	0.826086957
NCAM1 AND TMEM14C	0.900763359	0.9516129	0.855072464
NRCAM AND NOT-NCAM1	0.909090909	0.95238095	0.869565217
NCAM1 AND WDR83OS	0.903703704	0.92424242	0.884057971
NCAM1 AND NOT-OTOR	0.916030534	0.96774194	0.869565217
NLGN1 AND NLGN1	0.917293233	0.953125	0.884057971
NCAM1 AND NOT-SIGLEC6	0.918518519	0.93939394	0.898550725
NCAM1 AND PTTG1IP	0.907692308	0.96721311	0.855072464
NCAM1 AND ITGAV	0.914728682	0.98333333	0.855072464
NCAM1 AND TMX1	0.903703704	0.92424242	0.884057971
NCAM1 AND SERP1	0.903703704	0.92424242	0.884057971
NCAM1 AND NOT-CYP3A4	0.924242424	0.96825397	0.884057971
NOT-PRAF2 AND NCAM1	0.905109489	0.91176471	0.898550725
NCAM1 AND NOT-TMEM26	0.911764706	0.92537313	0.898550725
NCAM1 AND UBXN8	0.905109489	0.91176471	0.898550725
NCAM1 AND NOT-AQP8	0.918518519	0.93939394	0.898550725
NCAM1 AND SLC38A9	0.903703704	0.92424242	0.884057971
NCAM1 AND NOT-CHST4	0.914728682	0.98333333	0.855072464
PON2 AND NCAM1	0.918518519	0.93939394	0.898550725
XPR1 AND NCAM1	0.909090909	0.95238095	0.869565217
NCAM1 AND GOLM1	0.916030534	0.96774194	0.869565217
NCAM1 AND WRB	0.917293233	0.953125	0.884057971
NCAM1 AND SELK	0.916030534	0.96774194	0.869565217
NCAM1 AND GALNT1	0.911764706	0.92537313	0.898550725
NCAM1 AND ATRAID	0.903703704	0.92424242	0.884057971
NCAM1 AND ATP5F1	0.903703704	0.92424242	0.884057971
NCAM1 AND SLC30A5	0.903703704	0.92424242	0.884057971
NCAM1 AND NOT-CD1B	0.930232558	1	0.869565217
NCAM1 AND NOT-AADAC	0.921875	1	0.855072464
FJX1 AND NCAM1	0.905109489	0.91176471	0.898550725
SYT11 AND SYT11	0.951048951	0.91891892	0.985507246
SCRG1 AND SCRG1	0.909090909	0.87837838	0.942028986
NCAM1 AND SPCS1	0.907692308	0.96721311	0.855072464
NCAM1 AND JTB	0.907692308	0.96721311	0.855072464
NCAM1 AND UBA2	0.932330827	0.96875	0.898550725
NCAM1 AND NOT-UGT2B15	0.914728682	0.98333333	0.855072464
NCAM1 AND E2F5	0.903703704	0.92424242	0.884057971
TPST1 AND NCAM1	0.917293233	0.953125	0.884057971
AGPAT5 AND NCAM1	0.918518519	0.93939394	0.898550725
NCAM1 AND SEC11A	0.902255639	0.9375	0.869565217
NCAM1 AND NOT-TMPRSS15	0.925373134	0.95384615	0.898550725
NCAM1 AND TMEM69	0.911764706	0.92537313	0.898550725
CHST10 AND NCAM1	0.915492958	0.89041096	0.942028986
NCAM1 AND IER3IP1	0.909090909	0.95238095	0.869565217
NCAM1 AND PIGF	0.903703704	0.92424242	0.884057971
PTPRZ1 AND NCAM1	0.985507246	0.98550725	0.985507246
NCAM1 AND CDYL	0.900763359	0.9516129	0.855072464
NCAM1 AND NDUFA4	0.910447761	0.93846154	0.884057971
NCAM1 AND NOT-SIGLECL1	0.909090909	0.95238095	0.869565217
NCAM1 AND TMEM203	0.903703704	0.92424242	0.884057971
NLGN1 AND NOT-CD19	0.938461538	1	0.884057971
PRAF2 AND PRAF2	0.905109489	0.91176471	0.898550725
SYT11 AND CD19	0.928571429	0.91549296	0.942028986
SCRG1 AND CD19	0.907801418	0.88888889	0.927536232
TPST1 AND TPST1	0.917293233	0.953125	0.884057971
AGPAT5 AND NOT-CD19	0.946564885	1	0.898550725
PTPRZ1 AND NOT-CD19	0.985507246	0.98550725	0.985507246
NLGN1 AND NLGN1	0.917293233	0.953125	0.884057971
ITGAV AND NOT-GPC3	0.913385827	1	0.84057971
PRAF2 AND PRAF2	0.905109489	0.91176471	0.898550725
FJX1 AND NOT-GPC3	0.909090909	0.95238095	0.869565217
SYT11 AND GPC3	0.928571429	0.91549296	0.942028986
SCRG1 AND GPC3	0.901408451	0.87671233	0.927536232
TPST1 AND TPST1	0.917293233	0.953125	0.884057971
CHST10 AND GPC3	0.907801418	0.88888889	0.927536232
PTPRZ1 AND NOT-GPC3	0.985507246	0.98550725	0.985507246
NLGN1 AND NLGN1	0.917293233	0.953125	0.884057971
PRAF2 AND PRAF2	0.905109489	0.91176471	0.898550725
SYT11 AND SYT11	0.951048951	0.91891892	0.985507246
SCRG1 AND SCRG1	0.909090909	0.87837838	0.942028986
TPST1 AND NOT-MET	0.917293233	0.953125	0.884057971
CHST10 AND NOT-MET	0.915492958	0.89041096	0.942028986
PTPRZ1 AND NOT-MET	0.985507246	0.98550725	0.985507246
NRCAM AND NOT-B4GALNT1	0.921875	1	0.855072464

Antigen Pair Logic	F1	Precision	Recall
SLC10A4 AND NOT-GRM5	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-SLC25A4	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-CYP2S1	0.959537572	0.93258427	0.988095238
SLC10A4 AND NOT-ALG6	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-HFE	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-ADGRE2	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-SLC29A2	0.937142857	0.901098901	0.976190476
SLC10A4 AND NOT-HRH2	0.93258427	0.882978723	0.988095238
SLC10A4 AND NOT-HSD11B1	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-APLP1	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-HTR1B	0.954022989	0.922222222	0.988095238
SLC10A4 AND NOT-HTR1F	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-HTR6	0.943181818	0.902173913	0.988095238
SLC10A4 AND NOT-CLEC4D	0.943181818	0.902173913	0.988095238
SLC10A4 AND NOT-ICAM4	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-339166?	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-NAT8L	0.943181818	0.902173913	0.988095238
SLC10A4 AND NOT-TREML1	0.959064327	0.942528736	0.976190476
SLC10A4 AND NOT-COX8C	0.937853107	0.892473118	0.988095238
SLC10A4 AND NOT-GLDN	0.976190476	0.976190476	0.976190476
SLC10A4 AND NOT-SLC35B2	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-IL6ST	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-AQP1	0.937853107	0.892473118	0.988095238
SLC10A4 AND NOT-IL12B	0.948571429	0.912087912	0.988095238
SLC10A4 AND NOT-IL12RB2	0.943181818	0.902173913	0.988095238
SLC10A4 AND NOT-AQP4	0.982248521	0.976470588	0.988095238
SLC10A4 AND NOT-INPP4A	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-INSIG1	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-ITGA2B	0.947976879	0.921348315	0.976190476
SLC10A4 AND NOT-ITGA4	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-ITGA9	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-ITGB6	0.941860465	0.920454545	0.964285714
SLC10A4 AND NOT-KCNA2	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-KCNA3	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-KCNA6	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-KCNA7	0.941860465	0.920454545	0.964285714
SLC10A4 AND NOT-KCNA10	0.931818182	0.891304348	0.976190476
SLC10A4 AND NOT-SPEM1	0.93258427	0.882978723	0.988095238
SLC10A4 AND NOT-SFT2D2	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-KCND3	0.937853107	0.892473118	0.988095238
SLC10A4 AND NOT-LHFPL3	0.982248521	0.976470588	0.988095238
SLC10A4 AND NOT-ENHO	0.943181818	0.902173913	0.988095238
SLC10A4 AND NOT-KCNJ2	0.93258427	0.882978723	0.988095238
SLC10A4 AND NOT-KCNJ4	0.943181818	0.902173913	0.988095238
SLC10A4 AND NOT-KCNJ6	0.948571429	0.912087912	0.988095238
SLC10A4 AND NOT-KCNJ9	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-KCNJ10	0.954022989	0.922222222	0.988095238
SLC10A4 AND NOT-KCNJ11	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-KCNJ16	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-KCNK1	0.954022989	0.922222222	0.988095238
SLC10A4 AND NOT-KCNN3	0.982035928	0.987951807	0.976190476
SLC10A4 AND NOT-KCNQ3	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-CLEC12B	0.943181818	0.902173913	0.988095238
SLC10A4 AND NOT-USP27X	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-LAIR1	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-LHCGR	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-LIFR	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-FADS3	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-C14orf180	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-LNPEP	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-HACD4	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-LRP2	0.943181818	0.902173913	0.988095238
SLC10A4 AND NOT-LRP4	0.988095238	0.988095238	0.988095238
SLC10A4 AND NOT-LSAMP	0.965116279	0.943181818	0.988095238
SLC10A4 AND NOT-ERVFRD-1	0.954022989	0.922222222	0.988095238
SLC10A4 AND NOT-M6PR	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-MAG	0.976470588	0.965116279	0.988095238
SLC10A4 AND NOT-MAL	0.970414201	0.964705882	0.976190476
SLC10A4 AND NOT-MAN2A2	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-STS	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-LCN10	0.942528736	0.911111111	0.976190476
SLC10A4 AND NOT-MGST1	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-MGST2	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-MME	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-ALDH6A1	0.948571429	0.912087912	0.988095238
SLC10A4 AND NOT-MOG	0.976470588	0.965116279	0.988095238

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
NLGN1 AND NOT-B4GALNT1	0.930232558	1	0.869565217
PRAF2 AND NOT-B4GALNT1	0.909090909	0.95238095	0.869565217
SYT11 AND NOT-B4GALNT1	0.956521739	0.95652174	0.956521739
SCRG1 AND NOT-B4GALNT1	0.913043478	0.91304348	0.913043478
AGPAT5 AND B4GALNT1	0.905109489	0.91176471	0.898550725
CHST10 AND NOT-B4GALNT1	0.940298507	0.96923077	0.913043478
PTPRZ1 AND NOT-B4GALNT1	0.970588235	0.98507463	0.956521739
NRCAM AND NOT-CD160	0.900763359	0.9516129	0.855072464
FKTN AND NOT-CD160	0.904761905	1	0.826086957
NLGN1 AND NOT-CD160	0.921875	1	0.855072464
LRRC37A3 AND NOT-CD160	0.939393939	0.98412698	0.898550725
ITGAV AND NOT-CD160	0.954545455	1	0.913043478
MARS AND NOT-CD160	0.911764706	0.92537313	0.898550725
PRAF2 AND NOT-CD160	0.923076923	0.98360656	0.869565217
XPR1 AND NOT-CD160	0.904761905	1	0.826086957
GOLM1 AND NOT-CD160	0.925373134	0.95384615	0.898550725
WRB AND NOT-CD160	0.90625	0.98305085	0.84057971
FJX1 AND NOT-CD160	0.918518519	0.93939394	0.898550725
SYT11 AND NOT-CD160	0.96350365	0.97058824	0.956521739
SCRG1 AND NOT-CD160	0.933333333	0.95454545	0.913043478
B3GALT6 AND NOT-CD160	0.923076923	0.98360656	0.869565217
TPST1 AND NOT-CD160	0.913385827	1	0.84057971
AGPAT5 AND NOT-CD160	0.93129771	0.98387097	0.884057971
CHST10 AND CD160	0.915492958	0.89041096	0.942028986
JAM2 AND NOT-CD160	0.900763359	0.9516129	0.855072464
PTPRZ1 AND NOT-CD160	0.970588235	0.98507463	0.956521739
FKTN AND NOT-ERBB2	0.90625	0.98305085	0.84057971
NLGN1 AND ERBB2	0.917293233	0.953125	0.884057971
LRRC37A3 AND NOT-ERBB2	0.90647482	0.9	0.913043478
ITGAV AND NOT-ERBB2	0.948148148	0.96969697	0.927536232
PRAF2 AND PRAF2	0.905109489	0.91176471	0.898550725
FJX1 AND NOT-ERBB2	0.90647482	0.9	0.913043478
SYT11 AND ERBB2	0.951048951	0.91891892	0.985507246
SCRG1 AND ERBB2	0.909090909	0.87837838	0.942028986
TPST1 AND NOT-ERBB2	0.930232558	1	0.869565217
AGPAT5 AND ERBB2	0.905109489	0.91176471	0.898550725
CHST10 AND ERBB2	0.915492958	0.89041096	0.942028986
PTPRZ1 AND NOT-ERBB2	0.97810219	0.98529412	0.971014493
NLGN1 AND NLGN1	0.917293233	0.953125	0.884057971
ITGAV AND NOT-IL11RA	0.924242424	0.96825397	0.884057971
PRAF2 AND PRAF2	0.905109489	0.91176471	0.898550725
FJX1 AND NOT-IL11RA	0.913043478	0.91304348	0.913043478
SYT11 AND SYT11	0.951048951	0.91891892	0.985507246
SCRG1 AND SCRG1	0.909090909	0.87837838	0.942028986
TPST1 AND TPST1	0.917293233	0.953125	0.884057971
AGPAT5 AND AGPAT5	0.905109489	0.91176471	0.898550725
CHST10 AND IL11RA	0.915492958	0.89041096	0.942028986
PTPRZ1 AND NOT-IL11RA	0.97810219	0.98529412	0.971014493
NLGN1 AND NOT-PSCA	0.921875	1	0.855072464
LRRC37A3 AND NOT-PSCA	0.9	0.88732394	0.913043478
ITGAV AND NOT-PSCA	0.948148148	0.96969697	0.927536232
PRAF2 AND PRAF2	0.905109489	0.91176471	0.898550725
FJX1 AND NOT-PSCA	0.905109489	0.91176471	0.898550725
SYT11 AND PSCA	0.943661972	0.91780822	0.971014493
SCRG1 AND PSCA	0.907801418	0.88888889	0.927536232
TPST1 AND NOT-PSCA	0.938461538	1	0.884057971
AGPAT5 AND PSCA	0.905109489	0.91176471	0.898550725
CHST10 AND PSCA	0.907801418	0.88888889	0.927536232
PCDHB10 AND NOT-PSCA	0.904761905	1	0.826086957
PTPRZ1 AND NOT-PSCA	0.970588235	0.98507463	0.956521739
NLGN1 AND NLGN1	0.917293233	0.953125	0.884057971
LRRC37A3 AND NOT-MS4A1	0.90647482	0.9	0.913043478
ITGAV AND NOT-MS4A1	0.941176471	0.95522388	0.927536232
PRAF2 AND PRAF2	0.905109489	0.91176471	0.898550725
XPR1 AND NOT-MS4A1	0.913385827	1	0.84057971
TMEM255A AND NOT-MS4A1	0.901408451	0.87671233	0.927536232
SYT11 AND MS4A1	0.943661972	0.91780822	0.971014493
SCRG1 AND MS4A1	0.901408451	0.87671233	0.927536232
TPST1 AND NOT-MS4A1	0.913385827	1	0.84057971
AGPAT5 AND NOT-MS4A1	0.938461538	1	0.884057971
CHST10 AND MS4A1	0.907801418	0.88888889	0.927536232
PTPRZ1 AND NOT-MS4A1	0.977777778	1	0.956521739
KCNH2 AND NOT-MS4A1	0.907801418	0.88888889	0.927536232
BCHE AND NOT-MS4A1	0.924242424	0.96825397	0.884057971
NOT-RNF180 AND IL13RA2	0.910344828	0.86842105	0.956521739
NLGN1 AND NLGN1	0.917293233	0.953125	0.884057971

Antigen Pair Logic	F1	Precision	Recall
SLC10A4 AND NOT-C1orf186	0.922222222	0.864583333	0.988095238
SLC10A4 AND MARCH11	0.953488372	0.931818182	0.976190476
SLC10A4 AND NOT-NCAM1	0.937142857	0.901098901	0.976190476
SLC10A4 AND NOT-ATP1A2	0.976470588	0.965116279	0.988095238
SLC10A4 AND NOT-ATP1A3	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-NINJ2	0.937853107	0.892473118	0.988095238
SLC10A4 AND NNAT	0.947368421	0.931034483	0.964285714
SLC10A4 AND NOT-NMBR	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-ATP1B2	0.97005988	0.975903614	0.964285714
SLC10A4 AND NOT-NPC1	0.941860465	0.920454545	0.964285714
SLC10A4 AND NOT-NPY5R	0.954022989	0.922222222	0.988095238
SLC10A4 AND NOT-NTRK2	0.943181818	0.902173913	0.988095238
SLC10A4 AND NOT-ATP2B2	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-GPR143	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-OLR1	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-OMG	0.982035928	0.987951807	0.976190476
SLC10A4 AND NOT-P2RX1	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-P2RX5	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-P2RY2	0.93258427	0.882978723	0.988095238
SLC10A4 AND NOT-CD207	0.942528736	0.911111111	0.976190476
SLC10A4 AND NOT-DUOX2	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-CHST11	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-TAS2R3	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-TAS2R1	0.946745562	0.941176471	0.952380952
SLC10A4 AND NOT-SPOCK3	0.937853107	0.892473118	0.988095238
SLC10A4 AND NOT-NTM	0.948571429	0.912087912	0.988095238
SLC10A4 AND NOT-ST8SIA3	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-PLLP	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-ZDHHC9	0.937853107	0.892473118	0.988095238
SLC10A4 AND IER3IP1	0.937853107	0.892473118	0.988095238
SLC10A4 AND NOT-ZDHHC2	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-CRIM1	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-RXFP3	0.922222222	0.864583333	0.988095238
SLC10A4 AND FAM8A1	0.942528736	0.911111111	0.976190476
SLC10A4 AND CUTA	0.931818182	0.891304348	0.976190476
SLC10A4 AND ATP5F1	0.93258427	0.882978723	0.988095238
SLC10A4 AND NOT-ACSL5	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-GPRC5B	0.959537572	0.93258427	0.988095238
SLC10A4 AND NOT-ATP8A2	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-TM7SF3	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-GALNT7	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-PEX10	0.937853107	0.892473118	0.988095238
SLC10A4 AND PFN2	0.937853107	0.892473118	0.988095238
SLC10A4 AND NOT-PHEX	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-IL17D	0.965116279	0.943181818	0.988095238
SLC10A4 AND NOT-STX18	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-FXYD1	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-ADAM22	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-SMIM11	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-PON2	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-DCHS2	0.959537572	0.93258427	0.988095238
SLC10A4 AND NOT-DPP8	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-LPCAT2	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-STX17	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-SUSD4	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-SLC52A1	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-AUP1	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-NAT10	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-FBXW7	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-SPTLC3	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-TMEM144	0.982035928	0.987951807	0.976190476
SLC10A4 AND NOT-ACER3	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-SLC39A9	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-STYK1	0.975903614	0.987804878	0.964285714
SLC10A4 AND NOT-AVPR2	0.931818182	0.891304348	0.976190476
SLC10A4 AND NOT-CDCA7L	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-GALNT10	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-GPATCH2L	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-RNF130	0.922222222	0.864583333	0.988095238
SLC10A4 AND PCDHB10	0.946745562	0.941176471	0.952380952
SLC10A4 AND NOT-PCDHAC2	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-PCDHAC1	0.948571429	0.912087912	0.988095238
SLC10A4 AND NOT-SLC7A10	0.988095238	0.988095238	0.988095238
SLC10A4 AND NOT-PRRG1	0.976470588	0.965116279	0.988095238
SLC10A4 AND DIABLO	0.951807229	0.963414634	0.94047619
SLC10A4 AND NOT-AGPAT3	0.954022989	0.922222222	0.988095238

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
PRAF2 AND PRAF2	0.905109489	0.91176471	0.898550725
SYT11 AND IL13RA2	0.951048951	0.91891892	0.985507246
SCRG1 AND IL13RA2	0.909090909	0.87837838	0.942028986
TPST1 AND TPST1	0.917293233	0.953125	0.884057971
AGPAT5 AND IL13RA2	0.918518519	0.93939394	0.898550725
CHST10 AND IL13RA2	0.915492958	0.89041096	0.942028986
PTPRZ1 AND IL13RA2	0.985507246	0.98550725	0.985507246
NOT-CRB1 AND IL13RA2	0.905109489	0.91176471	0.898550725
NLGN1 AND NLGN1	0.917293233	0.953125	0.884057971
PRAF2 AND PRAF2	0.905109489	0.91176471	0.898550725
SYT11 AND SYT11	0.951048951	0.91891892	0.985507246
SCRG1 AND SCRG1	0.909090909	0.87837838	0.942028986
TPST1 AND NOT-IGF1R	0.938461538	1	0.884057971
AGPAT5 AND AGPAT5	0.905109489	0.91176471	0.898550725
PTPRZ1 AND NOT-IGF1R	0.97810219	0.98529412	0.971014493
NLGN1 AND NOT-MAGEA1	0.904761905	1	0.826086957
ITGAV AND MAGEA1	0.914285714	0.90140845	0.927536232
PRAF2 AND PRAF2	0.905109489	0.91176471	0.898550725
SYT11 AND MAGEA1	0.936170213	0.91666667	0.956521739
SCRG1 AND MAGEA1	0.907801418	0.88888889	0.927536232
TPST1 AND MAGEA1	0.909090909	0.95238095	0.869565217
CHST10 AND MAGEA1	0.9	0.88732394	0.913043478
PTPRZ1 AND NOT-MAGEA1	0.955223881	0.98461538	0.927536232
SRR AND NOT-MLANA	0.904761905	1	0.826086957
NLGN1 AND MLANA	0.917293233	0.953125	0.884057971
LRRC37A3 AND NOT-MLANA	0.915492958	0.89041096	0.942028986
ITGAV AND NOT-MLANA	0.96350365	0.97058824	0.956521739
PRAF2 AND NOT-MLANA	0.925373134	0.95384615	0.898550725
FJX1 AND NOT-MLANA	0.920863309	0.91428571	0.927536232
SYT11 AND MLANA	0.928571429	0.91549296	0.942028986
TPST1 AND TPST1	0.917293233	0.953125	0.884057971
PTPRZ1 AND NOT-MLANA	0.985507246	0.98550725	0.985507246
NLGN1 AND NLGN1	0.917293233	0.953125	0.884057971
PRAF2 AND PRAF2	0.905109489	0.91176471	0.898550725
SYT11 AND SYT11	0.951048951	0.91891892	0.985507246
SCRG1 AND SCRG1	0.909090909	0.87837838	0.942028986
TPST1 AND TPST1	0.917293233	0.953125	0.884057971
CHST10 AND ERBB3	0.915492958	0.89041096	0.942028986
PTPRZ1 AND NOT-ERBB3	0.985507246	0.98550725	0.985507246
NLGN1 AND NLGN1	0.917293233	0.953125	0.884057971
PRAF2 AND PRAF2	0.905109489	0.91176471	0.898550725
SYT11 AND NOT-ERBB4	0.964028777	0.95714286	0.971014493
SCRG1 AND NOT-ERBB4	0.909090909	0.87837838	0.942028986
TPST1 AND ERBB4	0.900763359	0.9516129	0.855072464
AGPAT5 AND ERBB4	0.910447761	0.93846154	0.884057971
CHST10 AND NOT-ERBB4	0.927536232	0.92753623	0.927536232
PTPRZ1 AND NOT-ERBB4	0.985507246	0.98550725	0.985507246
NLGN1 AND NOT-GPA33	0.938461538	1	0.884057971
ITGAV AND NOT-GPA33	0.927536232	0.92753623	0.927536232
GPR19 AND NOT-GPA33	0.914728682	0.98333333	0.855072464
PRAF2 AND PRAF2	0.905109489	0.91176471	0.898550725
XPR1 AND NOT-GPA33	0.921875	1	0.855072464
TMEM255A AND NOT-GPA33	0.901408451	0.87671233	0.927536232
SYT11 AND GPA33	0.943661972	0.91780822	0.971014493
SCRG1 AND GPA33	0.901408451	0.87671233	0.927536232
HSPA13 AND NOT-GPA33	0.904761905	1	0.826086957
PCDH17 AND NOT-GPA33	0.900763359	0.9516129	0.855072464
TPST1 AND TPST1	0.917293233	0.953125	0.884057971
AGPAT5 AND NOT-GPA33	0.930232558	1	0.869565217
GPR137C AND NOT-GPA33	0.907692308	0.96721311	0.855072464
CHST10 AND GPA33	0.907801418	0.88888889	0.927536232
CDH2 AND NOT-GPA33	0.916666667	0.88	0.956521739
PCDHB10 AND NOT-GPA33	0.907692308	0.96721311	0.855072464
PTPRZ1 AND NOT-GPA33	0.985507246	0.98550725	0.985507246
TMEM206 AND NOT-GPA33	0.916030534	0.96774194	0.869565217
BCHE AND NOT-GPA33	0.911764706	0.92537313	0.898550725
NRCAM AND CD276	0.910447761	0.93846154	0.884057971
NLGN1 AND NLGN1	0.917293233	0.953125	0.884057971
ITGAV AND CD276	0.942857143	0.92957746	0.956521739
NOT-C1orf186 AND CD276	0.962962963	0.98484848	0.942028986
NOT-C1orf210 AND CD276	0.955882353	0.97014925	0.942028986
GPM6A AND CD276	0.936170213	0.91666667	0.956521739
XPR1 AND CD276	0.900763359	0.9516129	0.855072464
NOT-AADAC AND CD276	0.940298507	0.96923077	0.913043478
SYT11 AND CD276	0.943661972	0.91780822	0.971014493
SCRG1 AND CD276	0.914285714	0.90140845	0.927536232

Antigen Pair Logic	F1	Precision	Recall
SLC10A4 AND NOT-C2orf83	0.937142857	0.901098901	0.976190476
SLC10A4 AND NOT-SLC17A7	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-MAN1C1	0.937853107	0.892473118	0.988095238
SLC10A4 AND NOT-NIPAL3	0.969325153	1	0.94047619
SLC10A4 AND NOT-PTGER2	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-TTYH1	0.970760234	0.954022989	0.988095238
SLC10A4 AND NOT-NDRG2	0.982248521	0.976470588	0.988095238
SLC10A4 AND NOT-TMCC3	0.942528736	0.911111111	0.976190476
SLC10A4 AND NOT-KIAA1161	0.931818182	0.891304348	0.976190476
SLC10A4 AND NOT-HHATL	0.976470588	0.965116279	0.988095238
SLC10A4 AND NOT-SLC12A5	0.964705882	0.953488372	0.976190476
SLC10A4 AND NOT-NCEH1	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-PCDH10	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-SLC7A14	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-CADM3	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-SLC46A2	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-SRPRB	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-BBS4	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-RAD51B	0.953488372	0.931818182	0.976190476
SLC10A4 AND NOT-ENPP5	0.97005988	0.975903614	0.964285714
SLC10A4 AND NOT-CACNG6	0.948571429	0.912087912	0.988095238
SLC10A4 AND NOT-LGR6	0.963855422	0.975609756	0.952380952
SLC10A4 AND NOT-RET	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-BCL2L2	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-CDH26	0.937853107	0.892473118	0.988095238
SLC10A4 AND NOT-ROS1	0.937853107	0.892473118	0.988095238
SLC10A4 AND NOT-CTXN3	0.988095238	0.988095238	0.988095238
SLC10A4 AND RTN2	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-SORT1	0.948571429	0.912087912	0.988095238
SLC10A4 AND NOT-BDNF	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-SC5D	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-SCD	0.954022989	0.922222222	0.988095238
SLC10A4 AND NOT-SCN1B	0.964285714	0.964285714	0.964285714
SLC10A4 AND NOT-SCN2B	0.963414634	0.9875	0.94047619
SLC10A4 AND NOT-SCN4B	0.958083832	0.963855422	0.952380952
SLC10A4 AND NOT-SCN8A	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-SCN10A	0.937853107	0.892473118	0.988095238
SLC10A4 AND NOT-SCNN1G	0.959537572	0.93258427	0.988095238
SLC10A4 AND NOT-CX3CL1	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-SDC1	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-SDHC	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-SEL1L	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-SMIM10	0.948571429	0.912087912	0.988095238
SLC10A4 AND NOT-TMEM200C	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-LMF1	0.976470588	0.965116279	0.988095238
SLC10A4 AND NOT-P2RY12	0.957575758	0.975308642	0.94047619
SLC10A4 AND NOT-ELOVL1	0.948571429	0.912087912	0.988095238
SLC10A4 AND NOT-LRRC19	0.943181818	0.902173913	0.988095238
SLC10A4 AND NOT-SLC1A2	0.93258427	0.882978723	0.988095238
SLC10A4 AND NOT-SLC2A5	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-SLC3A1	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-SLC5A1	0.954022989	0.922222222	0.988095238
SLC10A4 AND NOT-SLC5A3	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-SLC6A1	0.976470588	0.965116279	0.988095238
SLC10A4 AND NOT-SLC6A3	0.937142857	0.901098901	0.976190476
SLC10A4 AND NOT-SLC6A4	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-SLC6A9	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-SLC8A1	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-SLC12A1	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-SLC12A3	0.93258427	0.882978723	0.988095238
SLC10A4 AND NOT-SLC13A1	0.959537572	0.93258427	0.988095238
SLC10A4 AND NOT-SLC14A1	0.963855422	0.975609756	0.952380952
SLC10A4 AND NOT-SLC15A2	0.988095238	0.988095238	0.988095238
SLC10A4 AND NOT-SLC20A2	0.937853107	0.892473118	0.988095238
SLC10A4 AND NOT-GRAMD3	0.953488372	0.931818182	0.976190476
SLC10A4 AND NOT-BNIP2	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-SPINK2	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-SRD5A1	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-SSR3	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-SSTR1	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-BRS3	0.937853107	0.892473118	0.988095238
SLC10A4 AND NOT-STX4	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-BST1	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-VAMP1	0.93258427	0.882978723	0.988095238
SLC10A4 AND NOT-SYP	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-SYPL1	0.937853107	0.892473118	0.988095238

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
TPST1 AND TPST1	0.917293233	0.953125	0.884057971
GRIA2 AND CD276	0.90647482	0.9	0.913043478
AGPAT5 AND CD276	0.918518519	0.93939394	0.898550725
CHST10 AND CD276	0.934306569	0.94117647	0.927536232
PTPRZ1 AND CD276	0.985294118	1	0.971014493
NRCAM AND NOT-CD22	0.917293233	0.953125	0.884057971
NLGN1 AND NOT-CD22	0.930232558	1	0.869565217
ITGAV AND NOT-CD22	0.948905109	0.95588235	0.942028986
PRAF2 AND NOT-CD22	0.938461538	1	0.884057971
XPR1 AND NOT-CD22	0.909090909	0.95238095	0.869565217
SYT11 AND NOT-CD22	0.971428571	0.95774648	0.985507246
SCRG1 AND NOT-CD22	0.942028986	0.94202899	0.942028986
TPST1 AND TPST1	0.917293233	0.953125	0.884057971
AGPAT5 AND NOT-CD22	0.939393939	0.98412698	0.898550725
CHST10 AND CD22	0.907801418	0.88888889	0.927536232
PTPRZ1 AND NOT-CD22	0.985294118	1	0.971014493
TMEM206 AND NOT-CD22	0.932330827	0.96875	0.898550725
NLGN1 AND NLGN1	0.917293233	0.953125	0.884057971
ITGAV AND NOT-ROR1	0.916030534	0.96774194	0.869565217
PRAF2 AND PRAF2	0.905109489	0.91176471	0.898550725
SYT11 AND ROR1	0.951048951	0.91891892	0.985507246
SCRG1 AND ROR1	0.909090909	0.87837838	0.942028986
TPST1 AND TPST1	0.917293233	0.953125	0.884057971
AGPAT5 AND ROR1	0.905109489	0.91176471	0.898550725
CHST10 AND ROR1	0.915492958	0.89041096	0.942028986
PTPRZ1 AND NOT-ROR1	0.985507246	0.98550725	0.985507246
NLGN1 AND NLGN1	0.917293233	0.953125	0.884057971
PRAF2 AND PRAF2	0.905109489	0.91176471	0.898550725
SYT11 AND CA9	0.943661972	0.91780822	0.971014493
SCRG1 AND CA9	0.907801418	0.88888889	0.927536232
TPST1 AND TPST1	0.917293233	0.953125	0.884057971
AGPAT5 AND CA9	0.905109489	0.91176471	0.898550725
CHST10 AND CA9	0.9	0.88732394	0.913043478
PTPRZ1 AND CA9	0.985507246	0.98550725	0.985507246
NLGN1 AND NLGN1	0.917293233	0.953125	0.884057971
ITGAV AND NOT-EPHA2	0.90625	0.98305085	0.84057971
PRAF2 AND PRAF2	0.905109489	0.91176471	0.898550725
SYT11 AND EPHA2	0.943661972	0.91780822	0.971014493
SCRG1 AND EPHA2	0.909090909	0.87837838	0.942028986
TPST1 AND NOT-EPHA2	0.904761905	1	0.826086957
CHST10 AND EPHA2	0.907801418	0.88888889	0.927536232
PTPRZ1 AND NOT-EPHA2	0.970588235	0.98507463	0.956521739
NLGN1 AND NOT-TNFSF11	0.930232558	1	0.869565217
ITGAV AND NOT-TNFSF11	0.928571429	0.91549296	0.942028986
PRAF2 AND PRAF2	0.905109489	0.91176471	0.898550725
SYT11 AND TNFSF11	0.936170213	0.91666667	0.956521739
SCRG1 AND TNFSF11	0.9	0.88732394	0.913043478
TPST1 AND TPST1	0.917293233	0.953125	0.884057971
CHST10 AND TNFSF11	0.9	0.88732394	0.913043478
PTPRZ1 AND NOT-TNFSF11	0.97810219	0.98529412	0.971014493
FKTN AND NOT-SDC1	0.904761905	1	0.826086957
ITGAV AND NOT-SDC1	0.955882353	0.97014925	0.942028986
PRAF2 AND NOT-SDC1	0.917293233	0.953125	0.884057971
FJX1 AND NOT-SDC1	0.905109489	0.91176471	0.898550725
SYT11 AND SDC1	0.928571429	0.91549296	0.942028986
JAM2 AND NOT-SDC1	0.909090909	0.95238095	0.869565217
PTPRZ1 AND NOT-SDC1	0.985507246	0.98550725	0.985507246
NLGN1 AND NOT-TNFRSF8	0.930232558	1	0.869565217
LRRC37A3 AND NOT-TNFRSF8	0.909090909	0.87837838	0.942028986
ITGAV AND NOT-TNFRSF8	0.940298507	0.96923077	0.913043478
PRAF2 AND PRAF2	0.905109489	0.91176471	0.898550725
XPR1 AND NOT-TNFRSF8	0.914728682	0.98333333	0.855072464
FJX1 AND NOT-TNFRSF8	0.913043478	0.91304348	0.913043478
SYT11 AND SYT11	0.951048951	0.91891892	0.985507246
SCRG1 AND SCRG1	0.909090909	0.87837838	0.942028986
TPST1 AND TPST1	0.917293233	0.953125	0.884057971
AGPAT5 AND NOT-TNFRSF8	0.930232558	1	0.869565217
CHST10 AND TNFRSF8	0.9	0.88732394	0.913043478
PCDHB10 AND NOT-TNFRSF8	0.904761905	1	0.826086957
PTPRZ1 AND NOT-TNFRSF8	0.97810219	0.98529412	0.971014493
NLGN1 AND NLGN1	0.917293233	0.953125	0.884057971
PTPRZ1 AND CSPG4	0.985507246	0.98550725	0.985507246
NLGN1 AND NLGN1	0.917293233	0.953125	0.884057971
PRAF2 AND PRAF2	0.905109489	0.91176471	0.898550725
SYT11 AND CD70	0.936170213	0.91666667	0.956521739
TPST1 AND TPST1	0.917293233	0.953125	0.884057971

Antigen Pair Logic	F1	Precision	Recall
SLC10A4 AND NOT-BTC	0.93258427	0.882978723	0.988095238
SLC10A4 AND NOT-TRPC4	0.953488372	0.931818182	0.976190476
SLC10A4 AND NOT-TRPC5	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-TRPC6	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-C1orf234	0.931818182	0.891304348	0.976190476
SLC10A4 AND NOT-TNFSF4	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-TYR	0.948571429	0.912087912	0.988095238
SLC10A4 AND NOT-UPK1B	0.959064327	0.942528736	0.976190476
SLC10A4 AND NOT-UGT8	0.976190476	0.976190476	0.976190476
SLC10A4 AND NOT-BEST1	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-TRPV1	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-VRK2	0.922222222	0.864583333	0.988095238
SLC10A4 AND TMEM258	0.954022989	0.922222222	0.988095238
SLC10A4 AND NOT-LDLRAD4	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-CACNA1A	0.970414201	0.964705882	0.976190476
SLC10A4 AND NOT-CACNA1E	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-BSND	0.946745562	0.941176471	0.952380952
SLC10A4 AND NOT-DDR1	0.937853107	0.892473118	0.988095238
SLC10A4 AND NOT-RNF103	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-SEMA3B	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-SLC25A20	0.922222222	0.864583333	0.988095238
SLC10A4 AND GDAP1L1	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-SMIM2	0.959537572	0.93258427	0.988095238
SLC10A4 AND NOT-TMEM109	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-FA2H	0.937853107	0.892473118	0.988095238
SLC10A4 AND NOT-NIPAL2	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-TMC5	0.954022989	0.922222222	0.988095238
SLC10A4 AND NOT-ATP8B4	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-ERMP1	0.965116279	0.943181818	0.988095238
SLC10A4 AND NOT-PAQR6	0.988095238	0.988095238	0.988095238
SLC10A4 AND NOT-WLS	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-ELOVL7	0.937142857	0.901098901	0.976190476
SLC10A4 AND NOT-GLRA3	0.965116279	0.943181818	0.988095238
SLC10A4 AND NOT-LRRC8E	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-ALPK1	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-PIGZ	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-PRRT1	0.982035928	0.987951807	0.976190476
SLC10A4 AND NOT-ELOVL3	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-TMEM47	0.953488372	0.931818182	0.976190476
SLC10A4 AND NOT-CD99L2	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-SLC4A9	0.937142857	0.901098901	0.976190476
SLC10A4 AND NOT-CALN1	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-SLC25A18	0.943181818	0.902173913	0.988095238
SLC10A4 AND NOT-ESYT3	0.959064327	0.942528736	0.976190476
SLC10A4 AND NOT-SPATA9	0.93258427	0.882978723	0.988095238
SLC10A4 AND NOT-SLC41A2	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-KIAA1109	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-USP48	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-DCUN1D5	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-MEGF10	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-ACSS1	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-CASR	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-FUT10	0.976470588	0.965116279	0.988095238
SLC10A4 AND NOT-MFSD2A	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-PLXDC2	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-DGKE	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-PAQR8	0.93258427	0.882978723	0.988095238
SLC10A4 AND NOT-SLC22A16	0.947976879	0.921348315	0.976190476
SLC10A4 AND NOT-EPT1	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-SLC4A4	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-S1PR4	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-B3GALT1	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-TNFSF9	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-ADAM23	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-ADAM7	0.948571429	0.912087912	0.988095238
SLC10A4 AND NOT-CDS2	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-PROM1	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-APLN	0.959064327	0.942528736	0.976190476
SLC10A4 AND NOT-CCKAR	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-SLC5A6	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-CACNA1I	0.93258427	0.882978723	0.988095238
SLC10A4 AND NOT-LMLN	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-DUOXA1	0.941860465	0.920454545	0.964285714
SLC10A4 AND NOT-GPRC5A	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-MCU	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-CLDN10	0.922222222	0.864583333	0.988095238

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
CHST10 AND CD70	0.907801418	0.88888889	0.927536232
PTPRZ1 AND CD70	0.962962963	0.98484848	0.942028986
NLGN1 AND NLGN1	0.917293233	0.953125	0.884057971
PRAF2 AND NOT-AXL	0.932330827	0.96875	0.898550725
NOT-AADAC AND AXL	0.907692308	0.96721311	0.855072464
SYT11 AND AXL	0.943661972	0.91780822	0.971014493
SCRG1 AND SCRG1	0.909090909	0.87837838	0.942028986
TPST1 AND TPST1	0.917293233	0.953125	0.884057971
AGPAT5 AND AXL	0.910447761	0.93846154	0.884057971
CHST10 AND AXL	0.919708029	0.92647059	0.913043478
PTPRZ1 AND AXL	0.985507246	0.98550725	0.985507246
NLGN1 AND NOT-MUC13	0.938461538	1	0.884057971
PRAF2 AND NOT-MUC13	0.946564885	1	0.898550725
GOLM1 AND NOT-MUC13	0.934306569	0.94117647	0.927536232
SYT11 AND MUC13	0.951048951	0.91891892	0.985507246
SCRG1 AND MUC13	0.909090909	0.87837838	0.942028986
TPST1 AND NOT-MUC13	0.938461538	1	0.884057971
AGPAT5 AND MUC13	0.905109489	0.91176471	0.898550725
CHST10 AND NOT-MUC13	0.935251799	0.92857143	0.942028986
PTPRZ1 AND NOT-MUC13	0.97810219	0.98529412	0.971014493
NLGN1 AND NLGN1	0.917293233	0.953125	0.884057971
PRAF2 AND PRAF2	0.905109489	0.91176471	0.898550725
SYT11 AND SYT11	0.951048951	0.91891892	0.985507246
SCRG1 AND SCRG1	0.909090909	0.87837838	0.942028986
TPST1 AND ALK	0.900763359	0.9516129	0.855072464
CHST10 AND ALK	0.915492958	0.89041096	0.942028986
PTPRZ1 AND ALK	0.985507246	0.98550725	0.985507246
SYT11 AND TNFRSF10B	0.928571429	0.91549296	0.942028986
TPST1 AND TPST1	0.917293233	0.953125	0.884057971
AGPAT5 AND AGPAT5	0.905109489	0.91176471	0.898550725
PTPRZ1 AND TNFRSF10B	0.985507246	0.98550725	0.985507246
NLGN1 AND NLGN1	0.917293233	0.953125	0.884057971
PRAF2 AND PRAF2	0.905109489	0.91176471	0.898550725
SYT11 AND SYT11	0.951048951	0.91891892	0.985507246
SCRG1 AND SCRG1	0.909090909	0.87837838	0.942028986
PTPRZ1 AND NOT-FAP	0.985507246	0.98550725	0.985507246
NLGN1 AND NLGN1	0.917293233	0.953125	0.884057971
PRAF2 AND PRAF2	0.905109489	0.91176471	0.898550725
SYT11 AND FOLR2	0.928571429	0.91549296	0.942028986
SCRG1 AND FOLR2	0.909090909	0.87837838	0.942028986
AGPAT5 AND AGPAT5	0.905109489	0.91176471	0.898550725
PTPRZ1 AND FOLR2	0.985507246	0.98550725	0.985507246
NLGN1 AND KDR	0.917293233	0.953125	0.884057971
PRAF2 AND PRAF2	0.905109489	0.91176471	0.898550725
SYT11 AND KDR	0.951048951	0.91891892	0.985507246
SCRG1 AND KDR	0.909090909	0.87837838	0.942028986
TPST1 AND TPST1	0.917293233	0.953125	0.884057971
AGPAT5 AND KDR	0.902255639	0.9375	0.869565217
CHST10 AND KDR	0.907801418	0.88888889	0.927536232
PTPRZ1 AND KDR	0.977777778	1	0.956521739
NLGN1 AND NLGN1	0.917293233	0.953125	0.884057971
PRAF2 AND PRAF2	0.905109489	0.91176471	0.898550725
SYT11 AND CD38	0.951048951	0.91891892	0.985507246
SCRG1 AND CD38	0.909090909	0.87837838	0.942028986
TPST1 AND NOT-CD38	0.921875	1	0.855072464
CHST10 AND CD38	0.915492958	0.89041096	0.942028986
PTPRZ1 AND NOT-CD38	0.970588235	0.98507463	0.956521739
NRCAM AND NOT-ITGB3	0.900763359	0.9516129	0.855072464
NLGN1 AND NOT-ITGB3	0.904761905	1	0.826086957
PRAF2 AND PRAF2	0.905109489	0.91176471	0.898550725
SYT11 AND NOT-ITGB3	0.913385827	1	0.84057971
TPST1 AND TPST1	0.917293233	0.953125	0.884057971
AGPAT5 AND NOT-ITGB3	0.913385827	1	0.84057971
PTPRZ1 AND NOT-ITGB3	0.985507246	0.98550725	0.985507246
NLGN1 AND NLGN1	0.917293233	0.953125	0.884057971
ITGAV AND NOT-MUC16	0.948148148	0.96969697	0.927536232
PRAF2 AND PRAF2	0.905109489	0.91176471	0.898550725
FJX1 AND NOT-MUC16	0.902255639	0.9375	0.869565217
SYT11 AND MUC16	0.951048951	0.91891892	0.985507246
SCRG1 AND MUC16	0.915492958	0.89041096	0.942028986
TPST1 AND NOT-MUC16	0.938461538	1	0.884057971
AGPAT5 AND MUC16	0.905109489	0.91176471	0.898550725
CHST10 AND MUC16	0.915492958	0.89041096	0.942028986
PTPRZ1 AND NOT-MUC16	0.962962963	0.98484848	0.942028986
FKTN AND NOT-IL3RA	0.913385827	1	0.84057971
NLGN1 AND NOT-IL3RA	0.938461538	1	0.884057971

Antigen Pair Logic	F1	Precision	Recall
SLC10A4 AND NOT-ACBD5	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-SLC6A5	0.936416185	0.91011236	0.964285714
SLC10A4 AND NOT-SLC28A1	0.937853107	0.892473118	0.988095238
SLC10A4 AND NOT-IL1RL1	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-VAPB	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-DAPL1	0.943181818	0.902173913	0.988095238
SLC10A4 AND NOT-LMBRD2	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-OPALIN	0.982248521	0.976470588	0.988095238
SLC10A4 AND NOT-NRXN3	0.943181818	0.902173913	0.988095238
SLC10A4 AND NOT-SLC22A8	0.941860465	0.920454545	0.964285714
SLC10A4 AND NOT-TTYH2	0.937853107	0.892473118	0.988095238
SLC10A4 AND NOT-SFXN5	0.982248521	0.976470588	0.988095238
SLC10A4 AND NOT-FAM189A2	0.941860465	0.920454545	0.964285714
SLC10A4 AND NOT-CYP7B1	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-ABCG2	0.976190476	0.976190476	0.976190476
SLC10A4 AND NOT-CD33	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-CHST3	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-SIGLEC6	0.953488372	0.931818182	0.976190476
SLC10A4 AND NOT-AKAP6	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-CD34	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-PIGB	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-GABBR2	0.964285714	0.964285714	0.964285714
SLC10A4 AND NOT-ABCG1	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-CD47	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-EDEM1	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-TMEM63A	0.976470588	0.965116279	0.988095238
SLC10A4 AND NOT-CD79B	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-EFCAB14	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-MFAP3L	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-KIAA0319	0.927374302	0.873684211	0.988095238
SLC10A4 AND NOT-P2RY14	0.922222222	0.864583333	0.988095238
SLC10A4 AND NOT-SLC12A6	0.927374302	0.873684211	0.988095238
MARCH11 AND NOT-CCDC107	0.951807229	0.963414634	0.94047619
MARCH11 AND NOT-CERS3	0.969325153	1	0.94047619
MARCH11 AND NOT-EPHA4	0.963414634	0.9875	0.94047619
MARCH11 AND NOT-ERBB4	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-F3	0.952380952	0.952380952	0.952380952
ST8SIA2 AND NOT-F3	0.95	1	0.904761905
MARCH11 AND NOT-FAAH	0.963855422	0.975609756	0.952380952
ST8SIA2 AND NOT-FAT2	0.956521739	1	0.916666667
RTN1 AND NOT-HEPACAM	0.95	1	0.904761905
ST8SIA2 AND NOT-HEPACAM	0.95	1	0.904761905
GPR19 AND NOT-SLC39A12	0.963855422	0.975609756	0.952380952
MARCH11 AND NOT-SLC39A12	0.946745562	0.941176471	0.952380952
ST8SIA2 AND NOT-SLC39A12	0.956521739	1	0.916666667
ST8SIA2 AND NOT-OPN5	0.956521739	1	0.916666667
MARCH11 AND NOT-MMD2	0.952380952	0.952380952	0.952380952
ST8SIA2 AND NOT-GPRC6A	0.95	1	0.904761905
GABRB3 AND NOT-FGFR3	0.956521739	1	0.916666667
PCSK1N AND NOT-FGFR3	0.946745562	0.941176471	0.952380952
GPR19 AND NOT-FGFR3	0.96969697	0.987654321	0.952380952
MARCH11 AND NOT-FGFR3	0.946745562	0.941176471	0.952380952
ST8SIA3 AND NOT-FGFR3	0.975609756	1	0.952380952
HMP19 AND NOT-FGFR3	0.962962963	1	0.928571429
SCN3B AND NOT-FGFR3	0.97005988	0.975903614	0.964285714
SCN2A AND NOT-FGFR3	0.936708861	1	0.880952381
ST8SIA2 AND NOT-FGFR3	0.956521739	1	0.916666667
MARCH11 AND NOT-CLCA4	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-SLITRK3	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-PLA2R1	0.946745562	0.941176471	0.952380952
FAIM2 AND NOT-SLC15A2	0.952941176	0.941860465	0.964285714
FAIM2 AND NOT-SLC4A4	0.943396226	1	0.892857143
ST8SIA2 AND NOT-ATP10B	0.943396226	1	0.892857143
SYT11 AND NOT-NIPAL3	0.936708861	1	0.880952381
GPR19 AND NOT-MLC1	0.958083832	0.963855422	0.952380952
MARCH11 AND NOT-MLC1	0.952380952	0.952380952	0.952380952
HMP19 AND NOT-MLC1	0.962962963	1	0.928571429
ST8SIA2 AND NOT-MLC1	0.956521739	1	0.916666667
MARCH11 AND NOT-FLT3	0.952380952	0.952380952	0.952380952
MARCH11 AND NOT-ACSL6	0.946745562	0.941176471	0.952380952
ST8SIA2 AND NOT-FAM189A1	0.943396226	1	0.892857143
MARCH11 AND CTDNEP1	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-SLC44A1	0.946745562	0.941176471	0.952380952
ST8SIA2 AND NOT-SLC44A1	0.956521739	1	0.916666667
MARCH11 AND MMD	0.943396226	1	0.892857143
ST8SIA2 AND NOT-CA14	0.95	1	0.904761905

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
ITGAV AND NOT-IL3RA	0.924242424	0.96825397	0.884057971
PRAF2 AND PRAF2	0.905109489	0.91176471	0.898550725
XPR1 AND NOT-IL3RA	0.930232558	1	0.869565217
FJX1 AND NOT-IL3RA	0.920863309	0.91428571	0.927536232
SYT11 AND IL3RA	0.913043478	0.91304348	0.913043478
SCRG1 AND IL3RA	0.901408451	0.87671233	0.927536232
HSPA13 AND NOT-IL3RA	0.904761905	1	0.826086957
TPST1 AND IL3RA	0.909090909	0.95238095	0.869565217
AGPAT5 AND NOT-IL3RA	0.946564885	1	0.898550725
CHST10 AND IL3RA	0.907801418	0.88888889	0.927536232
PCDHB10 AND NOT-IL3RA	0.921875	1	0.855072464
PTPRZ1 AND NOT-IL3RA	0.985507246	0.98550725	0.985507246
NRCAM AND NOT-MUC17	0.938461538	1	0.884057971
SRR AND NOT-MUC17	0.904761905	1	0.826086957
FKTN AND NOT-MUC17	0.913385827	1	0.84057971
NLGN1 AND NOT-MUC17	0.938461538	1	0.884057971
LRRC37A3 AND NOT-MUC17	0.955882353	0.97014925	0.942028986
ITGAV AND NOT-MUC17	0.96350365	0.97058824	0.956521739
PRAF2 AND PRAF2	0.905109489	0.91176471	0.898550725
FJX1 AND NOT-MUC17	0.927536232	0.92753623	0.927536232
SYT11 AND MUC17	0.951048951	0.91891892	0.985507246
SCRG1 AND NOT-MUC17	0.935251799	0.92857143	0.942028986
TPST1 AND NOT-MUC17	0.938461538	1	0.884057971
AGPAT5 AND NOT-MUC17	0.932330827	0.96875	0.898550725
CHST10 AND MUC17	0.915492958	0.89041096	0.942028986
PCDHB10 AND NOT-MUC17	0.921875	1	0.855072464
PTPRZ1 AND NOT-MUC17	0.985507246	0.98550725	0.985507246
NLGN1 AND NLGN1	0.917293233	0.953125	0.884057971
ITGAV AND L1CAM	0.920863309	0.91428571	0.927536232
PRAF2 AND PRAF2	0.905109489	0.91176471	0.898550725
SYT11 AND SYT11	0.951048951	0.91891892	0.985507246
SCRG1 AND L1CAM	0.909090909	0.87837838	0.942028986
TPST1 AND L1CAM	0.900763359	0.9516129	0.855072464
CHST10 AND NOT-L1CAM	0.915492958	0.89041096	0.942028986
PTPRZ1 AND NOT-L1CAM	0.985507246	0.98550725	0.985507246
NLGN1 AND NLGN1	0.917293233	0.953125	0.884057971
ITGAV AND EPHA3	0.914285714	0.90140845	0.927536232
PRAF2 AND EPHA3	0.910447761	0.93846154	0.884057971
SYT11 AND SYT11	0.951048951	0.91891892	0.985507246
SCRG1 AND EPHA3	0.901408451	0.87671233	0.927536232
TPST1 AND EPHA3	0.917293233	0.953125	0.884057971
CHST10 AND EPHA3	0.907801418	0.88888889	0.927536232
PTPRZ1 AND EPHA3	0.97810219	0.98529412	0.971014493
NLGN1 AND NOT-MSLN	0.930232558	1	0.869565217
ITGAV AND MSLN	0.936170213	0.91666667	0.956521739
NOT-C1orf210 AND MSLN	0.905405405	0.84810127	0.971014493
PRAF2 AND PRAF2	0.905109489	0.91176471	0.898550725
SYT11 AND MSLN	0.951048951	0.91891892	0.985507246
SCRG1 AND MSLN	0.909090909	0.87837838	0.942028986
TPST1 AND NOT-MSLN	0.930232558	1	0.869565217
AGPAT5 AND MSLN	0.905109489	0.91176471	0.898550725
CHST10 AND MSLN	0.915492958	0.89041096	0.942028986
PTPRZ1 AND NOT-MSLN	0.970588235	0.98507463	0.956521739
NOT-SLAMF7 AND TNFSF13B	0.900763359	0.9516129	0.855072464
NOT-SLAMF7 AND C5orf28	0.921985816	0.90277778	0.942028986
NOT-SLAMF7 AND ATP2A2	0.905405405	0.84810127	0.971014493
NOT-SLAMF7 AND LPCAT1	0.925373134	0.95384615	0.898550725
TM2D2 AND NOT-SLAMF7	0.926470588	0.94029851	0.913043478
NOT-SLAMF7 AND ERGIC2	0.903703704	0.92424242	0.884057971
NOT-SLAMF7 AND WDR83OS	0.927536232	0.92753623	0.927536232
NOT-SLAMF7 AND TSPAN6	0.90647482	0.9	0.913043478
NOT-SLAMF7 AND TMEM209	0.909090909	0.95238095	0.869565217
NLGN1 AND NLGN1	0.917293233	0.953125	0.884057971
NOT-SLAMF7 AND C17orf62	0.933333333	0.95454545	0.913043478
LRRC37A3 AND NOT-SLAMF7	0.914285714	0.90140845	0.927536232
NOT-SLAMF7 AND TMEM230	0.913043478	0.91304348	0.913043478
NOT-SLAMF7 AND ALG8	0.919708029	0.92647059	0.913043478
NOT-SLAMF7 AND TMEM167A	0.923076923	0.98360656	0.869565217
ITGAV AND NOT-SLAMF7	0.962962963	0.98484848	0.942028986
NOT-SLAMF7 AND TMX1	0.96350365	0.97058824	0.956521739
NOT-SLAMF7 AND MANBAL	0.913043478	0.91304348	0.913043478
NOT-SLAMF7 AND MARS	0.941176471	0.95522388	0.927536232
NOT-SLAMF7 AND COX8A	0.921985816	0.90277778	0.942028986
NOT-SLAMF7 AND TSPAN3	0.910344828	0.86842105	0.956521739
NOT-SLAMF7 AND TMEM68	0.909090909	0.95238095	0.869565217
PRAF2 AND NOT-SLAMF7	0.938461538	1	0.884057971

Antigen Pair Logic	F1	Precision	Recall
ST8SIA2 AND NOT-SLC7A11	0.95	1	0.904761905
MARCH11 AND NOT-FLRT2	0.946745562	0.941176471	0.952380952
ST8SIA2 AND NOT-CADM2	0.936708861	1	0.880952381
MARCH11 AND NOT-LPCAT4	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-ZDHHC23	0.958083832	0.963855422	0.952380952
ST8SIA2 AND NOT-CALHM1	0.956521739	1	0.916666667
MARCH11 AND NOT-GABRA1	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-GABRA2	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-COL24A1	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-GABRA4	0.952380952	0.952380952	0.952380952
MARCH11 AND NOT-GABRB2	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-GABRD	0.946745562	0.941176471	0.952380952
ST8SIA2 AND NOT-GABRD	0.95	1	0.904761905
MARCH11 AND NOT-C1orf101	0.963855422	0.975609756	0.952380952
MARCH11 AND NOT-SLC24A2	0.946745562	0.941176471	0.952380952
ST8SIA2 AND NOT-SLC24A2	0.936708861	1	0.880952381
ST8SIA2 AND NOT-MRGPRX1	0.95	1	0.904761905
MARCH11 AND NOT-HIGD1A	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-GLCE	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-STEAP2	0.946745562	0.941176471	0.952380952
ST8SIA2 AND NOT-OR1C1	0.956521739	1	0.916666667
MARCH11 AND NOT-OR2F1	0.951807229	0.963414634	0.94047619
ST8SIA2 AND NOT-OR1J4	0.943396226	1	0.892857143
ST8SIA2 AND NOT-OR5K1	0.956521739	1	0.916666667
ST8SIA2 AND NOT-OR10A3	0.956521739	1	0.916666667
ST8SIA2 AND NOT-TSPAN16	0.95	1	0.904761905
ST8SIA2 AND NOT-OR12D2	0.95	1	0.904761905
ST8SIA2 AND NOT-OR10H1	0.95	1	0.904761905
ST8SIA2 AND NOT-OR8B2	0.956521739	1	0.916666667
ST8SIA2 AND NOT-OR7C2	0.936708861	1	0.880952381
MARCH11 AND NOT-OR7A5	0.95	1	0.904761905
ST8SIA2 AND NOT-OR4D1	0.95	1	0.904761905
ST8SIA2 AND NOT-ADGRF1	0.956521739	1	0.916666667
ST8SIA2 AND NOT-OR2H1	0.956521739	1	0.916666667
ST8SIA2 AND NOT-OR1J2	0.936708861	1	0.880952381
MARCH11 AND NOT-KCNV1	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-ATP2C1	0.946745562	0.941176471	0.952380952
ST8SIA2 AND NOT-GJB1	0.956521739	1	0.916666667
MARCH11 AND NOT-B3GAT1	0.946745562	0.941176471	0.952380952
DIABLO AND NOT-SERP1	0.936708861	1	0.880952381
PCSK1N AND TMEM258	0.957575758	0.975308642	0.94047619
ST8SIA2 AND NOT-GLP1R	0.95	1	0.904761905
CECR6 AND NOT-SLC15A2	0.953488372	0.931818182	0.976190476
MARCH11 AND NOT-GLRB	0.946745562	0.941176471	0.952380952
ST8SIA2 AND NOT-OR51B5	0.95	1	0.904761905
MARCH11 AND NOT-GPR6	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-SLC46A3	0.946745562	0.941176471	0.952380952
ST8SIA2 AND NOT-C17orf78	0.956521739	1	0.916666667
GPR19 AND NOT-GRM3	0.952380952	0.952380952	0.952380952
GPR19 AND NOT-KCNA1	0.958083832	0.963855422	0.952380952
GPR19 AND NOT-PNPLA7	0.951219512	0.975	0.928571429
GPR19 AND NOT-KCNJ16	0.963855422	0.975609756	0.952380952
GPR19 AND NOT-LRP4	0.951807229	0.963414634	0.94047619
GPR19 AND NOT-MOG	0.963855422	0.975609756	0.952380952
GPR19 AND NOT-SLCO1C1	0.951219512	0.975	0.928571429
GPR19 AND NOT-GRAMD1C	0.969325153	1	0.94047619
GPR19 AND NOT-STYK1	0.95	1	0.904761905
GPR19 AND NOT-HHATL	0.952380952	0.952380952	0.952380952
GPR19 AND NOT-CTXN3	0.952380952	0.952380952	0.952380952
GPR19 AND NOT-SLC22A23	0.951807229	0.963414634	0.94047619
GPR19 AND NOT-SLC1A3	0.952380952	0.952380952	0.952380952
GPR19 AND NOT-SLC15A2	0.975609756	1	0.952380952
GPR19 AND NOT-TRPC3	0.957055215	0.987341772	0.928571429
GPR19 AND TMEM258	0.951219512	0.975	0.928571429
GPR19 AND NOT-CA12	0.936708861	1	0.880952381
GPR19 AND NOT-CACNA1A	0.957055215	0.987341772	0.928571429
GPR19 AND NOT-FA2H	0.96969697	0.987654321	0.952380952
GPR19 AND NOT-PRRT1	0.943396226	1	0.892857143
GPR19 AND NOT-SLC25A18	0.952380952	0.952380952	0.952380952
GPR19 AND NOT-ATP13A4	0.963855422	0.975609756	0.952380952
GPR19 AND NOT-SLC4A4	0.943396226	1	0.892857143
GPR19 AND NOT-CACNA1I	0.952380952	0.952380952	0.952380952
GPR19 AND NOT-SYT12	0.957055215	0.987341772	0.928571429
GPR19 AND NOT-OPALIN	0.963855422	0.975609756	0.952380952
GPR19 AND NOT-SFXN5	0.951807229	0.963414634	0.94047619
TMEM145 AND NOT-MOG	0.937142857	0.901098901	0.976190476

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
TMX2 AND NOT-SLAMF7	0.917293233	0.953125	0.884057971
COA1 AND NOT-SLAMF7	0.914285714	0.90140845	0.927536232
NOT-SLAMF7 AND ITGAE	0.927536232	0.92753623	0.927536232
PON2 AND NOT-SLAMF7	0.954545455	1	0.913043478
XPR1 AND NOT-SLAMF7	0.921875	1	0.855072464
GOLM1 AND NOT-SLAMF7	0.941176471	0.95522388	0.927536232
NOT-SLAMF7 AND ETNK1	0.921985816	0.90277778	0.942028986
WRB AND NOT-SLAMF7	0.924242424	0.96825397	0.884057971
NOT-SLAMF7 AND METTL21A	0.901408451	0.87671233	0.927536232
NOT-SLAMF7 AND ARL6IP1	0.909090909	0.87837838	0.942028986
NOT-SLAMF7 AND CNEP1R1	0.925373134	0.95384615	0.898550725
CANX AND NOT-SLAMF7	0.924242424	0.96825397	0.884057971
VMA21 AND NOT-SLAMF7	0.900763359	0.9516129	0.855072464
NOT-SLAMF7 AND CHST11	0.955882353	0.97014925	0.942028986
NOT-SLAMF7 AND DNAJC19	0.920863309	0.91428571	0.927536232
NOT-SLAMF7 AND CCDC127	0.911764706	0.92537313	0.898550725
NOT-SLAMF7 AND MCUR1	0.925373134	0.95384615	0.898550725
SPNS1 AND NOT-SLAMF7	0.913385827	1	0.84057971
NOT-SLAMF7 AND DPY19L4	0.920863309	0.91428571	0.927536232
NOT-SLAMF7 AND TMX3	0.911764706	0.92537313	0.898550725
NOT-SLAMF7 AND ORMDL1	0.934306569	0.94117647	0.927536232
NOT-SLAMF7 AND BET1	0.910447761	0.93846154	0.884057971
NOT-SLAMF7 AND STX18	0.914285714	0.90140845	0.927536232
FJX1 AND NOT-SLAMF7	0.913043478	0.91304348	0.913043478
NOT-SLAMF7 AND BRCA1	0.905109489	0.91176471	0.898550725
NOT-SLAMF7 AND TOR1B	0.948905109	0.95588235	0.942028986
SYT11 AND SLAMF7	0.951048951	0.91891892	0.985507246
RNF145 AND NOT-SLAMF7	0.909090909	0.95238095	0.869565217
SCRG1 AND SLAMF7	0.909090909	0.87837838	0.942028986
NOT-SLAMF7 AND STIM2	0.902255639	0.9375	0.869565217
NOT-SLAMF7 AND TMEM126B	0.913043478	0.91304348	0.913043478
NOT-SLAMF7 AND EEF1E1	0.936170213	0.91666667	0.956521739
NOT-SLAMF7 AND RNASE6	0.900763359	0.9516129	0.855072464
NOT-SLAMF7 AND TMCO1	0.932330827	0.96875	0.898550725
NOT-SLAMF7 AND SLC20A1	0.911564626	0.85897436	0.971014493
B3GALT6 AND NOT-SLAMF7	0.946564885	1	0.898550725
TPST1 AND NOT-SLAMF7	0.930232558	1	0.869565217
NOT-SLAMF7 AND NIPA2	0.911764706	0.92537313	0.898550725
ZDHHC4 AND NOT-SLAMF7	0.914285714	0.90140845	0.927536232
NOT-SLAMF7 AND UBE2J1	0.9	0.88732394	0.913043478
AGPAT5 AND NOT-SLAMF7	0.946564885	1	0.898550725
NOT-SLAMF7 AND ATP13A1	0.914728682	0.98333333	0.855072464
CHST10 AND SLAMF7	0.915492958	0.89041096	0.942028986
NOT-SLAMF7 AND PAG1	0.919708029	0.92647059	0.913043478
NOT-SLAMF7 AND TMED5	0.909090909	0.95238095	0.869565217
JAM2 AND NOT-SLAMF7	0.903703704	0.92424242	0.884057971
NOT-SLAMF7 AND TMEM9B	0.933333333	0.95454545	0.913043478
NOT-SLAMF7 AND TMBIM6	0.916030534	0.96774194	0.869565217
NOT-SLAMF7 AND PEX2	0.905109489	0.91176471	0.898550725
PTPRZ1 AND NOT-SLAMF7	0.985294118	1	0.971014493
NOT-SLAMF7 AND LRRC59	0.934306569	0.94117647	0.927536232
NOT-SLAMF7 AND NEMP1	0.948905109	0.95588235	0.942028986
HTRA2 AND NOT-SLAMF7	0.916030534	0.96774194	0.869565217
NDUFA4 AND NOT-SLAMF7	0.918518519	0.93939394	0.898550725
NOT-SLAMF7 AND C12orf73	0.910344828	0.86842105	0.956521739
NOT-SLAMF7 AND GAL3ST4	0.920863309	0.91428571	0.927536232
NOT-SLAMF7 AND LEPROTL1	0.934306569	0.94117647	0.927536232
NOT-SLAMF7 AND MANEA	0.947368421	0.984375	0.913043478
BCHE AND NOT-SLAMF7	0.916030534	0.96774194	0.869565217
NOT-SLAMF7 AND SSR2	0.948148148	0.96969697	0.927536232
NRCAM AND NOT-TNFRSF10A	0.900763359	0.9516129	0.855072464
FKTN AND NOT-TNFRSF10A	0.913385827	1	0.84057971
NLGN1 AND NOT-TNFRSF10A	0.930232558	1	0.869565217
LRRC37A3 AND NOT-TNFRSF10A	0.905109489	0.91176471	0.898550725
ITGAV AND NOT-TNFRSF10A	0.948148148	0.96969697	0.927536232
MARS AND NOT-TNFRSF10A	0.913043478	0.91304348	0.913043478
PRAF2 AND PRAF2	0.905109489	0.91176471	0.898550725
XPR1 AND NOT-TNFRSF10A	0.913385827	1	0.84057971
GOLM1 AND NOT-TNFRSF10A	0.913043478	0.91304348	0.913043478
FJX1 AND NOT-TNFRSF10A	0.932330827	0.96875	0.898550725
SYT11 AND TNFRSF10A	0.936170213	0.91666667	0.956521739
SCRG1 AND NOT-TNFRSF10A	0.926470588	0.94029851	0.913043478
CERS5 AND NOT-TNFRSF10A	0.904761905	1	0.826086957
B3GALT6 AND NOT-TNFRSF10A	0.910447761	0.93846154	0.884057971
TPST1 AND NOT-TNFRSF10A	0.921875	1	0.855072464
AGPAT5 AND NOT-TNFRSF10A	0.930232558	1	0.869565217

Antigen Pair Logic	F1	Precision	Recall
TMEM145 AND SLC38A2	0.951219512	0.975	0.928571429
TMEM145 AND TMEM258	0.957575758	0.975308642	0.94047619
MARCH11 AND NOT-GPR21	0.951807229	0.963414634	0.94047619
MARCH11 AND NOT-OR2L13	0.946745562	0.941176471	0.952380952
ST8SIA2 AND NOT-OR2L13	0.936708861	1	0.880952381
MARCH11 AND NOT-GPR26	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-RNF180	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-GPR37	0.958083832	0.963855422	0.952380952
ST8SIA2 AND NOT-GPR37	0.95	1	0.904761905
ST8SIA2 AND NOT-GRIA3	0.95	1	0.904761905
MARCH11 AND NOT-SLC6A16	0.963855422	0.975609756	0.952380952
MARCH11 AND NOT-GRIK1	0.946745562	0.941176471	0.952380952
ST8SIA2 AND NOT-GRIK1	0.956521739	1	0.916666667
MARCH11 AND NOT-GRIN1	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-GRIN2A	0.946745562	0.941176471	0.952380952
ST8SIA2 AND NOT-GRIN2B	0.956521739	1	0.916666667
MARCH11 AND NOT-GRIN2C	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-GRM1	0.952380952	0.952380952	0.952380952
MARCH11 AND NOT-GRM2	0.958083832	0.963855422	0.952380952
ST8SIA2 AND NOT-GRM2	0.956521739	1	0.916666667
MARCH11 AND NOT-GRM3	0.946745562	0.941176471	0.952380952
HMP19 AND NOT-GRM3	0.95	1	0.904761905
ST8SIA2 AND NOT-GRM3	0.956521739	1	0.916666667
MARCH11 AND NOT-GRM5	0.946745562	0.941176471	0.952380952
ST8SIA2 AND NOT-GRM6	0.956521739	1	0.916666667
MARCH11 AND NOT-GRM7	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-SLC25A4	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-CYP2S1	0.952380952	0.952380952	0.952380952
ST8SIA2 AND NOT-PCDHB1	0.936708861	1	0.880952381
MARCH11 AND NOT-GUCY2D	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-KCNIP3	0.958083832	0.963855422	0.952380952
ST8SIA2 AND NOT-SLC29A2	0.956521739	1	0.916666667
ST8SIA2 AND NOT-HTR1F	0.956521739	1	0.916666667
MARCH11 AND NOT-NAT8L	0.946745562	0.941176471	0.952380952
ST8SIA2 AND NOT-TREML1	0.943396226	1	0.892857143
ST8SIA2 AND NOT-AQP8	0.95	1	0.904761905
MARCH11 AND NOT-PAQR9	0.975609756	1	0.952380952
MARCH11 AND NOT-CATSPER3	0.963855422	0.975609756	0.952380952
ST8SIA2 AND NOT-CATSPER3	0.943396226	1	0.892857143
ST8SIA2 AND NOT-AQP4	0.95	1	0.904761905
ST8SIA2 AND NOT-AQP6	0.943396226	1	0.892857143
MARCH11 AND NOT-ITGAE	0.946745562	0.941176471	0.952380952
ST8SIA2 AND NOT-ITGB3	0.95	1	0.904761905
MARCH11 AND NOT-ITPR1	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-KCNA1	0.946745562	0.941176471	0.952380952
ST8SIA2 AND NOT-KCNA1	0.956521739	1	0.916666667
ST8SIA2 AND NOT-KCNA5	0.936708861	1	0.880952381
MARCH11 AND NOT-GALNT18	0.946745562	0.941176471	0.952380952
ST8SIA2 AND NOT-C1orf95	0.936708861	1	0.880952381
LHFPL4 AND NOT-SLC15A2	0.95	1	0.904761905
MARCH11 AND NOT-KCNF1	0.946745562	0.941176471	0.952380952
ST8SIA2 AND NOT-LHFPL3	0.956521739	1	0.916666667
ST8SIA2 AND NOT-KCNJ1	0.936708861	1	0.880952381
MARCH11 AND NOT-KCNJ4	0.958083832	0.963855422	0.952380952
MARCH11 AND NOT-KCNJ6	0.946745562	0.941176471	0.952380952
ST8SIA2 AND NOT-KCNJ6	0.936708861	1	0.880952381
MARCH11 AND NOT-KCNJ10	0.946745562	0.941176471	0.952380952
HMP19 AND NOT-KCNJ10	0.95	1	0.904761905
ST8SIA2 AND NOT-KCNJ10	0.95	1	0.904761905
MARCH11 AND NOT-KCNJ16	0.946745562	0.941176471	0.952380952
ST8SIA2 AND NOT-KCNJ16	0.956521739	1	0.916666667
ST8SIA2 AND NOT-KCNK1	0.956521739	1	0.916666667
MARCH11 AND NOT-KCNS1	0.952380952	0.952380952	0.952380952
MARCH11 AND NOT-KCNS2	0.946745562	0.941176471	0.952380952
ST8SIA2 AND NOT-KCNS2	0.956521739	1	0.916666667
ST8SIA2 AND NOT-KIR2DL4	0.95	1	0.904761905
MARCH11 AND NOT-ILDR2	0.951807229	0.963414634	0.94047619
ST8SIA2 AND NOT-CLEC12B	0.956521739	1	0.916666667
MARCH11 AND NOT-RPRML	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-FAM209B	0.963855422	0.975609756	0.952380952
ST8SIA2 AND NOT-FAM209B	0.95	1	0.904761905
MARCH11 AND NOT-SPATA31D1	0.957575758	0.975308642	0.94047619
ST8SIA2 AND NOT-OR51I1	0.936708861	1	0.880952381
MARCH11 AND NOT-TMEM200B	0.951807229	0.963414634	0.94047619
MARCH11 AND NOT-C11orf87	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-LMO7	0.96969697	0.987654321	0.952380952

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
CHST10 AND TNFRSF10A	0.9	0.88732394	0.913043478
JAM2 AND NOT-TNFRSF10A	0.916030534	0.96774194	0.869565217
PTPRZ1 AND NOT-TNFRSF10A	0.962962963	0.98484848	0.942028986
NDUFA4 AND NOT-TNFRSF10A	0.903703704	0.92424242	0.884057971
NLGN1 AND NLGN1	0.917293233	0.953125	0.884057971
ITGAV AND NOT-MUC1	0.907692308	0.96721311	0.855072464
TREM2 AND NOT-MUC1	0.90625	0.98305085	0.84057971
PRAF2 AND PRAF2	0.905109489	0.91176471	0.898550725
SYT11 AND MUC1	0.936170213	0.91666667	0.956521739
HSPA13 AND NOT-MUC1	0.904761905	1	0.826086957
TPST1 AND NOT-MUC1	0.938461538	1	0.884057971
AGPAT5 AND MUC1	0.905109489	0.91176471	0.898550725
CHST10 AND MUC1	0.9	0.88732394	0.913043478
PTPRZ1 AND NOT-MUC1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-NAALAD2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-NAALADL1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND KCNE3	0.985507246	0.98550725	0.985507246
CDH2 AND NOT-MRGPRX3	0.985507246	0.98550725	0.985507246
CDH2 AND NOT-RXFP2	0.985294118	1	0.971014493
CDH2 AND NOT-AADAC	0.970149254	1	0.942028986
CDH2 AND NOT-CYP3A4	0.985294118	1	0.971014493
CDH2 AND NOT-DSC3	0.985294118	1	0.971014493
CDH2 AND NOT-TMEM26	0.985294118	1	0.971014493
CDH2 AND NOT-FUT3	0.977777778	1	0.956521739
CDH2 AND ITGAV	0.970149254	1	0.942028986
CDH2 AND NDUFA4	0.977777778	1	0.956521739
CDH2 AND NOT-GABRQ	0.970149254	1	0.942028986
CDH2 AND NOT-OTOR	0.977777778	1	0.956521739
CDH2 AND NOT-SIGLEC6	0.970149254	1	0.942028986
GPM6B AND OST4	0.977777778	1	0.956521739
PTPRZ1 AND OST4	0.977777778	1	0.956521739
PTPRZ1 AND TMEM221	0.985507246	0.98550725	0.985507246
PTPRZ1 AND LRRC70	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-SMIM6	0.985294118	1	0.971014493
PTPRZ1 AND NOT-NEMP2	0.985294118	1	0.971014493
PTPRZ1 AND ARMCX4	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-BCL2L10	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-HCN4	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-TIMM23	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-ERVMER34-1	0.985294118	1	0.971014493
PTPRZ1 AND CDH4	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-PET117	0.985507246	0.98550725	0.985507246
PTPRZ1 AND CDH5	0.99270073	1	0.985507246
PTPRZ1 AND CDH6	0.985507246	0.98550725	0.985507246
PTPRZ1 AND 100505984?	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-APELA	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-MRLN	0.985507246	0.98550725	0.985507246
SYT11 AND NOT-TPBGL	0.977777778	1	0.956521739
PTPRZ1 AND NOT-TPBGL	0.977777778	1	0.956521739
PTPRZ1 AND NOT-SMLR1	0.99270073	1	0.985507246
PTPRZ1 AND TMEM178B	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-MICA	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-100507547?	0.985507246	0.98550725	0.985507246
PTPRZ1 AND GJC1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-UBA2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND ABCC9	0.985507246	0.98550725	0.985507246
PTPRZ1 AND SCAMP3	0.99270073	1	0.985507246
PTPRZ1 AND NOT-CDH8	0.985507246	0.98550725	0.985507246
CHST10 AND NOT-CDH8	0.970149254	1	0.942028986
PTPRZ1 AND MUC12	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-ATP9A	0.985507246	0.98550725	0.985507246
PTPRZ1 AND CDH10	0.985507246	0.98550725	0.985507246
PTPRZ1 AND TSPAN5	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-TSPAN1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND PREB	0.99270073	1	0.985507246
PTPRZ1 AND BCAP31	0.99270073	1	0.985507246
PTPRZ1 AND NOT-ADGRG2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-CDH17	0.985507246	0.98550725	0.985507246
PTPRZ1 AND LPAR6	0.977777778	1	0.956521739
PTPRZ1 AND NOT-CDH18	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-DHRS9	0.985507246	0.98550725	0.985507246
GPM6B AND CNIH1	0.977777778	1	0.956521739
PTPRZ1 AND NOT-TENM1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-101805491?	0.985507246	0.98550725	0.985507246
PTPRZ1 AND LHFPL2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND LHFP	0.985507246	0.98550725	0.985507246

Antigen Pair Logic	F1	Precision	Recall
ST8SIA2 AND NOT-XKRX	0.95	1	0.904761905
ST8SIA2 AND NOT-IFITM10	0.943396226	1	0.892857143
MARCH11 AND NOT-PPAPDC2	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-LRP2	0.958083832	0.963855422	0.952380952
RTN1 AND NOT-LRP2	0.936708861	1	0.880952381
ST8SIA2 AND NOT-LRP2	0.936708861	1	0.880952381
MARCH11 AND NOT-LRP4	0.952380952	0.952380952	0.952380952
HMP19 AND NOT-LRP4	0.951219512	0.975	0.928571429
ST8SIA2 AND NOT-LRP4	0.956521739	1	0.916666667
MARCH11 AND NOT-HAPLN4	0.946745562	0.941176471	0.952380952
ST8SIA2 AND NOT-LSAMP	0.943396226	1	0.892857143
ST8SIA2 AND NOT-LCN10	0.95	1	0.904761905
MARCH11 AND NOT-ADAM11	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-ART3	0.975609756	1	0.952380952
ST8SIA2 AND NOT-MME	0.95	1	0.904761905
ST8SIA2 AND NOT-ALDH6A1	0.936708861	1	0.880952381
MARCH11 AND NOT-MOG	0.952380952	0.952380952	0.952380952
ST8SIA3 AND NOT-MOG	0.963855422	0.975609756	0.952380952
HMP19 AND NOT-MOG	0.962962963	1	0.928571429
SCN3B AND NOT-MOG	0.952941176	0.941860465	0.964285714
ADGRB3 AND NOT-MOG	0.956521739	1	0.916666667
RTN1 AND NOT-MOG	0.963414634	0.9875	0.94047619
ST8SIA2 AND NOT-MOG	0.956521739	1	0.916666667
MARCH11 AND NOT-MINOS1	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-NDUFB8	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-NTRK2	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-ATP2B2	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-OPCML	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-CLDN11	0.951219512	0.975	0.928571429
MARCH11 AND NOT-GALNT9	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-PAM	0.951807229	0.963414634	0.94047619
MARCH11 AND NOT-NTM	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-C3orf18	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-KCNK9	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-CHST15	0.946745562	0.941176471	0.952380952
MARCH11 AND CUTA	0.951807229	0.963414634	0.94047619
MARCH11 AND NOT-GPRC5B	0.956521739	1	0.916666667
MARCH11 AND NOT-SLC26A4	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-TUBA8	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-SLCO1C1	0.946745562	0.941176471	0.952380952
MARCH11 AND SLC38A2	0.951219512	0.975	0.928571429
MARCH11 AND TOMM7	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-MANSC1	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-GRAMD1C	0.963855422	0.975609756	0.952380952
MARCH11 AND NOT-PIGG	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-RNF43	0.957055215	0.987341772	0.928571429
MARCH11 AND NOT-FBXW7	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-TMEM144	0.952380952	0.952380952	0.952380952
MARCH11 AND NOT-STYK1	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-GLT8D1	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-AJAP1	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-SLC7A10	0.952380952	0.952380952	0.952380952
MARCH11 AND NOT-PRRG1	0.951807229	0.963414634	0.94047619
MARCH11 AND NOT-PANX2	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-AGPAT3	0.958083832	0.963855422	0.952380952
MARCH11 AND NOT-SLC17A7	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-SLC45A4	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-TTYH1	0.951807229	0.963414634	0.94047619
MARCH11 AND NOT-ABHD6	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-LRTM1	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-HHATL	0.958083832	0.963855422	0.952380952
MARCH11 AND NOT-NCEH1	0.958083832	0.963855422	0.952380952
MARCH11 AND NOT-BBS4	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-ENPP5	0.957575758	0.975308642	0.94047619
MARCH11 AND NOT-EVA1C	0.963855422	0.975609756	0.952380952
MARCH11 AND NOT-CACNG8	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-LGR6	0.963855422	0.975609756	0.952380952
MARCH11 AND NOT-CTXN3	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-SLC22A23	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-SCN1A	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-SCN1B	0.958083832	0.963855422	0.952380952
MARCH11 AND NOT-SCN4B	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-SCN8A	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-SCNN1G	0.958083832	0.963855422	0.952380952
MARCH11 AND NOT-CX3CL1	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-XYLT2	0.951219512	0.975	0.928571429

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
PTPRZ1 AND ADAM8	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-TRIM13	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-ANGPTL7	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-GPA33	0.985507246	0.98550725	0.985507246
PTPRZ1 AND MFSD10	0.985507246	0.98550725	0.985507246
PTPRZ1 AND STX6	0.985507246	0.98550725	0.985507246
PTPRZ1 AND COQ7	0.985507246	0.98550725	0.985507246
GPM6B AND SLC35B1	0.970149254	1	0.942028986
PTPRZ1 AND SLC35B1	0.99270073	1	0.985507246
PTPRZ1 AND KCNMB2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND TIMM17B	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-SLC17A2	0.970149254	1	0.942028986
PTPRZ1 AND RAMP2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-RAMP3	0.985507246	0.98550725	0.985507246
GPM6B AND ZMPSTE24	0.970149254	1	0.942028986
PTPRZ1 AND ZMPSTE24	0.99270073	1	0.985507246
PTPRZ1 AND NOT-PRSS16	0.985507246	0.98550725	0.985507246
PTPRZ1 AND BET1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND LILRB2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-MARCH6	0.985507246	0.98550725	0.985507246
PTPRZ1 AND TCIRG1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-RTN3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-B3GALT5	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-CRISP3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND TMEM5	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-B3GNT3	0.970149254	1	0.942028986
PTPRZ1 AND NOT-CLEC4M	0.985507246	0.98550725	0.985507246
PTPRZ1 AND MRVI1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND TRDN	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-CACNG3	0.970149254	1	0.942028986
PTPRZ1 AND BTN2A2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-ATP8A1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-NDRG1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND PEMT	0.99270073	1	0.985507246
PTPRZ1 AND ST3GAL6	0.985507246	0.98550725	0.985507246
PTPRZ1 AND CDIPT	0.977777778	1	0.956521739
GPM6B AND PGRMC2	0.970149254	1	0.942028986
PTPRZ1 AND NOT-PGRMC2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND TMEM147	0.99270073	1	0.985507246
GPM6B AND LYPLA1	0.970149254	1	0.942028986
PTPRZ1 AND TIMM17A	0.99270073	1	0.985507246
PTPRZ1 AND NOT-N4BP2L2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND LRRN2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND GPNMB	0.985507246	0.98550725	0.985507246
PTPRZ1 AND MERTK	0.985507246	0.98550725	0.985507246
PTPRZ1 AND SLC30A9	0.985507246	0.98550725	0.985507246
PTPRZ1 AND CLGN	0.985507246	0.98550725	0.985507246
PTPRZ1 AND VTI1B	0.970149254	1	0.942028986
PTPRZ1 AND CRTAP	0.99270073	1	0.985507246
PTPRZ1 AND SEMA4B	0.985507246	0.98550725	0.985507246
PTPRZ1 AND DEAF1	0.985507246	0.98550725	0.985507246
GPM6B AND TM9SF1	0.970149254	1	0.942028986
PTPRZ1 AND TM9SF1	0.970149254	1	0.942028986
GPM6B AND PRDX4	0.985294118	1	0.971014493
PTPRZ1 AND PRDX4	0.985294118	1	0.971014493
PTPRZ1 AND ARL6IP5	0.970149254	1	0.942028986
PTPRZ1 AND NOT-HTATIP2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND AGPAT1	0.99270073	1	0.985507246
PTPRZ1 AND SLC35A1	0.977777778	1	0.956521739
PTPRZ1 AND SLC19A2	0.99270073	1	0.985507246
PTPRZ1 AND RABAC1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND SLC34A2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND IFITM2	0.977777778	1	0.956521739
PTPRZ1 AND NOT-SLCO1B1	0.985294118	1	0.971014493
PTPRZ1 AND NOT-ST6GALNAC2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND TGOLN2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND PDPN	0.985507246	0.98550725	0.985507246
LRRC37A3 AND NOT-SPINT2	0.970149254	1	0.942028986
PTPRZ1 AND NOT-SPINT2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-CXCR6	0.977777778	1	0.956521739
PTPRZ1 AND CD226	0.985507246	0.98550725	0.985507246
PTPRZ1 AND CGRRF1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND TNFSF13B	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-EBP	0.985507246	0.98550725	0.985507246
PTPRZ1 AND DLL3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-FUT9	0.985507246	0.98550725	0.985507246

Antigen Pair Logic	F1	Precision	Recall
MARCH11 AND NOT-WBSCR17	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-SFTPD	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-SGCD	0.951807229	0.963414634	0.94047619
MARCH11 AND NOT-SGCG	0.958083832	0.963855422	0.952380952
MARCH11 AND NOT-LMF1	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-FBXL17	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-SLC1A3	0.958083832	0.963855422	0.952380952
MARCH11 AND NOT-SLC3A1	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-SLC6A1	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-SLC6A13	0.963855422	0.975609756	0.952380952
MARCH11 AND NOT-LRTM2	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-SLC9A5	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-SLC15A2	0.952380952	0.952380952	0.952380952
MARCH11 AND NOT-SOX1	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-SPAM1	0.963855422	0.975609756	0.952380952
MARCH11 AND NOT-TAZ	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-TGFA	0.952380952	0.952380952	0.952380952
MARCH11 AND NOT-ICAM5	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-TRPC3	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-TYRP1	0.958083832	0.963855422	0.952380952
MARCH11 AND NOT-VIPR2	0.958083832	0.963855422	0.952380952
MARCH11 AND NOT-MYRF	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-RNF112	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-CACNA1C	0.958083832	0.963855422	0.952380952
MARCH11 AND CXCR4	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-CRELD1	0.946745562	0.941176471	0.952380952
NOT-MARCH11 AND GDAP1L1	0.943181818	0.902173913	0.988095238
MARCH11 AND NOT-FA2H	0.946745562	0.941176471	0.952380952
MARCH11 AND NKAIN1	0.952380952	0.952380952	0.952380952
MARCH11 AND NOT-C1orf115	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-PAQR6	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-TMPRSS5	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-TMEM47	0.951807229	0.963414634	0.94047619
MARCH11 AND NOT-CD99L2	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-SLC25A18	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-ESYT3	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-ATP13A4	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-ACSS1	0.96969697	0.987654321	0.952380952
MARCH11 AND NOT-AGPAT9	0.957055215	0.987341772	0.928571429
MARCH11 AND NOT-TMEM25	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-NDST2	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-CDC14B	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-SLC4A4	0.958083832	0.963855422	0.952380952
MARCH11 AND NOT-B3GALT2	0.958083832	0.963855422	0.952380952
MARCH11 AND NOT-TNFSF9	0.957575758	0.975308642	0.94047619
MARCH11 AND NOT-ADAM23	0.958083832	0.963855422	0.952380952
MARCH11 AND NOT-CCKBR	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-CACNA1I	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-LMLN	0.958083832	0.963855422	0.952380952
MARCH11 AND NOT-TMEM257	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-SLC16A7	0.951807229	0.963414634	0.94047619
MARCH11 AND NOT-BPIFB1	0.957055215	0.987341772	0.928571429
MARCH11 AND NOT-OPALIN	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-NRXN3	0.946745562	0.941176471	0.952380952
NOT-MARCH11 AND ACVR2B	0.931818182	0.891304348	0.976190476
MARCH11 AND NOT-SFXN5	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-FAM189A2	0.952380952	0.952380952	0.952380952
MARCH11 AND NOT-OPN4	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-ABCG2	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-SV2B	0.946745562	0.941176471	0.952380952
MARCH11 AND NOT-EXOG	0.952380952	0.952380952	0.952380952
MARCH11 AND NOT-SLC12A6	0.957055215	0.987341772	0.928571429
ASTN1 AND NOT-SLC15A2	0.952380952	0.952380952	0.952380952
HMP19 AND NOT-ATP1A2	0.936708861	1	0.880952381
ST8SIA2 AND NOT-ATP1A2	0.956521739	1	0.916666667
ST8SIA2 AND NOT-ATP1B2	0.95	1	0.904761905
NOT-ST8SIA2 AND NPY	0.937142857	0.901098901	0.976190476
ST8SIA2 AND NOT-NTRK2	0.956521739	1	0.916666667
ST8SIA2 AND NOT-ATP2B2	0.95	1	0.904761905
ST8SIA2 AND NOT-OMG	0.95	1	0.904761905
ST8SIA2 AND NOT-OPRK1	0.936708861	1	0.880952381
ST8SIA2 AND NOT-OR2C1	0.95	1	0.904761905
ST8SIA2 AND NOT-CD207	0.943396226	1	0.892857143
ST8SIA3 AND TMEM258	0.957575758	0.975308642	0.94047619
ST8SIA3 AND NOT-PRRT1	0.943396226	1	0.892857143
ST8SIA3 AND NOT-SLC4A4	0.936708861	1	0.880952381

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
PTPRZ1 AND NOT-RRH	0.985507246	0.98550725	0.985507246
PTPRZ1 AND CERS1	0.985507246	0.98550725	0.985507246
CERS1 AND PTTG1IP	0.970149254	1	0.942028986
PTPRZ1 AND NOT-NRG3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND YME1L1	0.970149254	1	0.942028986
PTPRZ1 AND PHTF1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND CHL1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-NUP50	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-HBS1L	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-SLC17A3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND VAMP5	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-OR5I1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-CYSLTR1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-CCR9	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-GJB6	0.985294118	1	0.971014493
PTPRZ1 AND NOT-CFTR	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-FAXDC2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-SLC26A1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND ADAM28	0.985507246	0.98550725	0.985507246
PTPRZ1 AND TSPAN9	0.985294118	1	0.971014493
PTPRZ1 AND NOT-USP19	0.985507246	0.98550725	0.985507246
PTPRZ1 AND HCST	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-CD300C	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-EDDM3A	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-NPFFR2	0.985294118	1	0.971014493
SYT11 AND NOT-GPR83	0.985294118	1	0.971014493
PTPRZ1 AND NOT-MMP24	0.985507246	0.98550725	0.985507246
PTPRZ1 AND LYVE1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-OCLM	0.99270073	1	0.985507246
PTPRZ1 AND JTB	0.970149254	1	0.942028986
PTPRZ1 AND CEACAM4	0.985507246	0.98550725	0.985507246
PTPRZ1 AND MAN1A2	0.977777778	1	0.956521739
PTPRZ1 AND PNPLA6	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-BTNL3	0.977777778	1	0.956521739
PTPRZ1 AND GPR75	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-AFG3L2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND KDELR1	0.985294118	1	0.971014493
PTPRZ1 AND STARD3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-SERINC3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND OS9	0.99270073	1	0.985507246
PTPRZ1 AND TMED2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND LMAN2	0.970149254	1	0.942028986
PTPRZ1 AND ERP29	0.977777778	1	0.956521739
PTPRZ1 AND UQCR11	0.985294118	1	0.971014493
PTPRZ1 AND SLC38A3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND ILVBL	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-SLC27A5	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-NAT2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND SLC27A3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND LILRB4	0.985507246	0.98550725	0.985507246
PTPRZ1 AND GLIPR1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND KDELR2	0.977777778	1	0.956521739
PTPRZ1 AND ATF7	0.985507246	0.98550725	0.985507246
PTPRZ1 AND TMED1	0.970149254	1	0.942028986
PTPRZ1 AND TDRKH	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-LECT1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND CYB561D2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-RER1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-ADAM30	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-ADAMTS13	0.985507246	0.98550725	0.985507246
PTPRZ1 AND CACFD1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND BTN2A1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-IL1RAPL1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-PLA2G16	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-HHLA2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-BVES	0.985507246	0.98550725	0.985507246
PTPRZ1 AND FICD	0.985507246	0.98550725	0.985507246
PTPRZ1 AND CHI3L2	0.985507246	0.98550725	0.985507246
SYT11 AND NOT-SLC2A6	0.977777778	1	0.956521739
PTPRZ1 AND NOT-CHIT1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-ADCY5	0.970149254	1	0.942028986
PTPRZ1 AND NOT-KLK8	0.985294118	1	0.971014493
PTPRZ1 AND GALNT5	0.985507246	0.98550725	0.985507246
PTPRZ1 AND PRAF2	0.99270073	1	0.985507246
PTPRZ1 AND SEC63	0.99270073	1	0.985507246
GPM6B AND RNF139	0.977777778	1	0.956521739

Antigen Pair Logic	F1	Precision	Recall
ST8SIA3 AND NOT-OPALIN	0.958083832	0.963855422	0.952380952
ST8SIA2 AND NOT-SLC45A2	0.943396226	1	0.892857143
ST8SIA2 AND NOT-CRIM1	0.95	1	0.904761905
ST8SIA2 AND ATP5F1	0.95	1	0.904761905
HMP19 AND NOT-TMEM144	0.943396226	1	0.892857143
HMP19 AND NOT-HHATL	0.962962963	1	0.928571429
HMP19 AND NOT-FA2H	0.951219512	0.975	0.928571429
HMP19 AND NOT-PAQR6	0.936708861	1	0.880952381
HMP19 AND NOT-OPALIN	0.962962963	1	0.928571429
ST8SIA2 AND SLC25A3	0.956521739	1	0.916666667
ST8SIA2 AND NOT-PI3	0.95	1	0.904761905
ST8SIA2 AND NOT-S1PR5	0.95	1	0.904761905
ST8SIA2 AND SLC38A2	0.943396226	1	0.892857143
ST8SIA2 AND NOT-SLC6A20	0.95	1	0.904761905
ST8SIA2 AND NOT-GRAMD1C	0.936708861	1	0.880952381
ST8SIA2 AND NOT-DCHS2	0.956521739	1	0.916666667
ST8SIA2 AND NOT-NXPE4	0.943396226	1	0.892857143
ST8SIA2 AND NOT-GDPD2	0.943396226	1	0.892857143
ST8SIA2 AND NOT-MS4A12	0.956521739	1	0.916666667
ST8SIA2 AND NOT-RHBDL2	0.95	1	0.904761905
ST8SIA2 AND NOT-TMEM144	0.956521739	1	0.916666667
ST8SIA2 AND NOT-STYK1	0.943396226	1	0.892857143
ST8SIA2 AND NOT-AVPR2	0.95	1	0.904761905
SCN3B AND NOT-HHATL	0.941860465	0.920454545	0.964285714
SCN3B AND NOT-SLC15A2	0.958579882	0.952941176	0.964285714
SCN3B AND NOT-FA2H	0.936416185	0.91011236	0.964285714
SCN3B AND DCHS1	0.952941176	0.941860465	0.964285714
SCN3B AND NOT-SLC4A4	0.943396226	1	0.892857143
SCN3B AND NOT-OPALIN	0.952941176	0.941860465	0.964285714
ST8SIA2 AND NOT-SLC22A11	0.95	1	0.904761905
ST8SIA2 AND NOT-PRRG1	0.95	1	0.904761905
NOT-ST8SIA2 AND DIABLO	0.947368421	0.931034483	0.964285714
DIABLO AND NOT-TMEM141	0.936708861	1	0.880952381
ST8SIA2 AND NOT-AGPAT3	0.956521739	1	0.916666667
ST8SIA2 AND NOT-C2orf83	0.936708861	1	0.880952381
ST8SIA2 AND NOT-SLC17A7	0.936708861	1	0.880952381
ST8SIA2 AND NOT-GJC2	0.956521739	1	0.916666667
RTN1 AND NOT-NIPAL3	0.943396226	1	0.892857143
ST8SIA2 AND NOT-TTYH1	0.95	1	0.904761905
ST8SIA2 AND NOT-KIAA1161	0.95	1	0.904761905
RTN1 AND NOT-HHATL	0.957575758	0.975308642	0.94047619
ST8SIA2 AND NOT-SORCS2	0.95	1	0.904761905
NOT-ST8SIA2 AND KIAA1549	0.947368421	0.931034483	0.964285714
ADGRB3 AND NOT-SLC4A4	0.936708861	1	0.880952381
ST8SIA2 AND NOT-PTPRD	0.95	1	0.904761905
ST8SIA2 AND NOT-RAD51B	0.943396226	1	0.892857143
ST8SIA2 AND NOT-CACNG8	0.95	1	0.904761905
ST8SIA2 AND NOT-CACNG6	0.943396226	1	0.892857143
ST8SIA2 AND NOT-CTXN3	0.956521739	1	0.916666667
RTN1 AND NOT-PAQR6	0.956521739	1	0.916666667
RTN1 AND DCHS1	0.957575758	0.975308642	0.94047619
RTN1 AND NOT-SLC4A4	0.936708861	1	0.880952381
RTN1 AND NOT-OPALIN	0.951807229	0.963414634	0.94047619
ST8SIA2 AND NOT-SCN1A	0.956521739	1	0.916666667
ST8SIA2 AND NOT-SCN1B	0.936708861	1	0.880952381
ST8SIA2 AND NOT-SCN4B	0.943396226	1	0.892857143
ST8SIA2 AND NOT-MS4A5	0.956521739	1	0.916666667
ST8SIA2 AND NOT-NDST4	0.956521739	1	0.916666667
ST8SIA2 AND NOT-CTAGE1	0.956521739	1	0.916666667
ST8SIA2 AND NOT-LMF1	0.956521739	1	0.916666667
ST8SIA2 AND NOT-P2RY12	0.95	1	0.904761905
ST8SIA2 AND NOT-ELOVL1	0.943396226	1	0.892857143
ST8SIA2 AND NOT-SLC13A3	0.95	1	0.904761905
ST8SIA2 AND NOT-CDCP1	0.95	1	0.904761905
ST8SIA2 AND NOT-LRRC19	0.956521739	1	0.916666667
ST8SIA2 AND NOT-SLC1A3	0.956521739	1	0.916666667
ST8SIA2 AND NOT-SLC5A1	0.95	1	0.904761905
ST8SIA2 AND NOT-SLC6A1	0.956521739	1	0.916666667
ST8SIA2 AND NOT-SLC6A4	0.95	1	0.904761905
ST8SIA2 AND NOT-SLC13A1	0.956521739	1	0.916666667
ST8SIA2 AND NOT-SSTR1	0.95	1	0.904761905
ST8SIA2 AND NOT-TGFA	0.956521739	1	0.916666667
ST8SIA2 AND NOT-TNF	0.943396226	1	0.892857143
ST8SIA2 AND NOT-TRHR	0.956521739	1	0.916666667
ST8SIA2 AND NOT-TRPC4	0.95	1	0.904761905
ST8SIA2 AND NOT-UPK1B	0.943396226	1	0.892857143

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
PTPRZ1 AND RNF139	0.977777778	1	0.956521739
PTPRZ1 AND GPR176	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-GPR45	0.970149254	1	0.942028986
PTPRZ1 AND NOT-PTGDR2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND MAN1B1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-SLC6A14	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NRM	0.970149254	1	0.942028986
PTPRZ1 AND NOT-KLHL2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND GLMP	0.985507246	0.98550725	0.985507246
PTPRZ1 AND HOGA1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND MGAT4B	0.985294118	1	0.971014493
PTPRZ1 AND NOT-CYP4F8	0.985507246	0.98550725	0.985507246
PTPRZ1 AND B4GALT7	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-CHRM2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND ADCY6	0.985507246	0.98550725	0.985507246
PTPRZ1 AND PTH2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND SLCO2B1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND CD300A	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-CHRM3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND CASC4	0.985507246	0.98550725	0.985507246
PTPRZ1 AND MGAT4A	0.99270073	1	0.985507246
PTPRZ1 AND NOT-TMC6	0.985507246	0.98550725	0.985507246
PTPRZ1 AND SLC46A1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND GLCCI1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND TMEM106A	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-SLC52A3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-CHRM4	0.985507246	0.98550725	0.985507246
PTPRZ1 AND TEX261	0.970149254	1	0.942028986
SCRG1 AND NOT-C1orf210	0.970149254	1	0.942028986
PTPRZ1 AND SCRG1	0.985294118	1	0.971014493
PTPRZ1 AND RNF13	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-SMIM12	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-CMTM1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-CHRNA2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-MFSD3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-CHRNA3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-PIK3IP1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-CHRNA4	0.985507246	0.98550725	0.985507246
PTPRZ1 AND ADCY7	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-SLC2A13	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-CHRNB3	0.99270073	1	0.985507246
PTPRZ1 AND NOT-CHRND	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-SLC22A9	0.977777778	1	0.956521739
PTPRZ1 AND NOT-CHRNE	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-ERMAP	0.985507246	0.98550725	0.985507246
PTPRZ1 AND LRRC37B	0.985507246	0.98550725	0.985507246
PTPRZ1 AND XKR4	0.985507246	0.98550725	0.985507246
PTPRZ1 AND SLC25A25	0.985294118	1	0.971014493
PTPRZ1 AND ELFN2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND TMEM132B	0.985507246	0.98550725	0.985507246
PTPRZ1 AND GALNT13	0.985507246	0.98550725	0.985507246
PTPRZ1 AND C1QTNF1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND C1QTNF6	0.985507246	0.98550725	0.985507246
PTPRZ1 AND TMEM123	0.99270073	1	0.985507246
GPM6B AND SMIM19	0.985294118	1	0.971014493
PTPRZ1 AND SMIM19	0.985294118	1	0.971014493
PTPRZ1 AND NOT-VASN	0.985507246	0.98550725	0.985507246
PTPRZ1 AND ADCY8	0.985507246	0.98550725	0.985507246
ITGAV AND NOT-SLC26A9	0.970149254	1	0.942028986
PTPRZ1 AND MARCH3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND OMA1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-SLC25A26	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-FCRL1	0.985294118	1	0.971014493
PTPRZ1 AND EVI5L	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-ADCY9	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-LYSMD3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-SLC22A12	0.985507246	0.98550725	0.985507246
PTPRZ1 AND LRRC3B	0.985507246	0.98550725	0.985507246
PTPRZ1 AND FAM210B	0.985507246	0.98550725	0.985507246
PTPRZ1 AND CMTM5	0.985507246	0.98550725	0.985507246
PTPRZ1 AND ABHD15	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-TLCD1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-MOGAT1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-PANX3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-SLC26A8	0.985507246	0.98550725	0.985507246
PTPRZ1 AND LYPD1	0.985507246	0.98550725	0.985507246

Antigen Pair Logic	F1	Precision	Recall
ST8SIA2 AND NOT-CLRN1	0.943396226	1	0.892857143
ST8SIA2 AND NOT-BEST1	0.95	1	0.904761905
ST8SIA2 AND NOT-MYRF	0.936708861	1	0.880952381
ST8SIA2 AND TMEM258	0.943396226	1	0.892857143
ST8SIA2 AND NOT-CA4	0.936708861	1	0.880952381
ST8SIA2 AND NOT-CACNA1A	0.956521739	1	0.916666667
ST8SIA2 AND NOT-FA2H	0.956521739	1	0.916666667
ST8SIA2 AND NOT-OR51B4	0.95	1	0.904761905
ST8SIA2 AND NOT-NOX5	0.95	1	0.904761905
ST8SIA2 AND NOT-TMC5	0.956521739	1	0.916666667
ST8SIA2 AND NOT-BTNL8	0.956521739	1	0.916666667
ST8SIA2 AND NOT-PAQR6	0.956521739	1	0.916666667
ST8SIA2 AND NOT-LRRC8E	0.956521739	1	0.916666667
ST8SIA2 AND NOT-ADAM33	0.956521739	1	0.916666667
ST8SIA2 AND NOT-SPX	0.95	1	0.904761905
ST8SIA2 AND NOT-ACSBG2	0.956521739	1	0.916666667
ST8SIA2 AND NOT-OR5V1	0.956521739	1	0.916666667
ST8SIA2 AND NOT-OR12D3	0.956521739	1	0.916666667
ST8SIA2 AND NOT-SPACA1	0.95	1	0.904761905
ST8SIA2 AND NOT-FZD9	0.943396226	1	0.892857143
ST8SIA2 AND NOT-SLC25A18	0.95	1	0.904761905
ST8SIA2 AND NOT-ESYT3	0.956521739	1	0.916666667
ST8SIA2 AND NOT-OR1G1	0.956521739	1	0.916666667
ST8SIA2 AND NOT-QRFPR	0.943396226	1	0.892857143
ST8SIA2 AND NOT-AGPAT9	0.943396226	1	0.892857143
ST8SIA2 AND NOT-CDC14B	0.936708861	1	0.880952381
ST8SIA2 AND NOT-SLC4A4	0.956521739	1	0.916666667
ST8SIA2 AND NOT-ADAM7	0.956521739	1	0.916666667
ST8SIA2 AND NOT-KCNH7	0.956521739	1	0.916666667
ST8SIA2 AND NOT-NUP210L	0.956521739	1	0.916666667
ST8SIA2 AND NOT-SYT12	0.943396226	1	0.892857143
ST8SIA2 AND NOT-NXPE3	0.936708861	1	0.880952381
ST8SIA2 AND NOT-VAPB	0.95	1	0.904761905
ST8SIA2 AND NOT-SLC38A5	0.936708861	1	0.880952381
ST8SIA2 AND NOT-OPALIN	0.956521739	1	0.916666667
ST8SIA2 AND NOT-TTYH2	0.936708861	1	0.880952381
ST8SIA2 AND NOT-SFXN5	0.956521739	1	0.916666667
ST8SIA2 AND NOT-ABCG2	0.943396226	1	0.892857143
ST8SIA2 AND NOT-SIGLEC6	0.95	1	0.904761905
ST8SIA2 AND NOT-CLCA2	0.956521739	1	0.916666667
ST8SIA2 AND NOT-TMEM63A	0.956521739	1	0.916666667
SV2A AND DCHS1	0.936416185	0.91011236	0.964285714
Lymphoma, Mantle-Cell (Mantle-Cell Lymphoma)			
CLECL1 AND NOT-EPHA3	0.904761905	0.826086957	1
CELSR1 AND NOT-ERBB2	0.923076923	0.9	0.947368421
QSOX2 AND NOT-ERBB2	0.902439024	0.840909091	0.973684211
CLECL1 AND NOT-CD160	0.926829268	0.863636364	1
CELSR1 AND NOT-SDC1	0.923076923	0.9	0.947368421
TNFRSF13C AND NOT-SDC1	0.904761905	0.826086957	1
CELSR1 AND NOT-ERBB3	0.923076923	0.9	0.947368421
TNFRSF13C AND NOT-FAP	0.915662651	0.844444444	1
SNN AND NOT-NCAM1	0.944444444	1	0.894736842
TNFRSF10A AND NOT-NCAM1	0.902439024	0.840909091	0.973684211
QSOX2 AND NOT-NCAM1	0.925	0.880952381	0.973684211
TNFRSF13C AND NOT-NCAM1	0.938271605	0.88372093	1
RHOT2 AND NOT-NCAM1	0.9	0.857142857	0.947368421
ORAI2 AND NOT-NCAM1	0.9	0.857142857	0.947368421
CLECL1 AND NOT-NCAM1	0.961038961	0.948717949	0.973684211
QSOX2 AND NOT-MSLN	0.925	0.880952381	0.973684211
QSOX2 AND NOT-ALK	0.948717949	0.925	0.973684211
MS4A1 AND NOT-SORL1	0.913580247	0.860465116	0.973684211
MS4A1 AND CELSR1	0.936708861	0.902439024	0.973684211
MS4A1 AND QSOX2	0.902439024	0.840909091	0.973684211
MS4A1 AND NOT-KIR3DL2	0.904761905	0.826086957	1
MS4A1 AND NOT-SCN3A	0.9	0.857142857	0.947368421
QSOX2 AND NOT-PSCA	0.913580247	0.860465116	0.973684211
TNFRSF13C AND NOT-PSCA	0.926829268	0.863636364	1
CLECL1 AND NOT-PSCA	0.904761905	0.826086957	1
CELSR1 AND NOT-MUC1	0.9	0.857142857	0.947368421
TNFRSF13C AND NOT-B4GALNT1	0.902439024	0.840909091	0.973684211
NOT-FAM210B AND TNFRSF10A	0.925	0.880952381	0.973684211
TNFRSF10A AND NOT-SPN	0.915662651	0.844444444	1
NOT-C4orf32 AND TNFRSF10A	0.935064935	0.923076923	0.947368421
TNFRSF10A AND NOT-FURIN	0.911392405	0.87804878	0.947368421
TNFRSF10A AND NOT-STAB1	0.904761905	0.826086957	1
TNFRSF10A AND NOT-SLC46A2	0.909090909	0.897435897	0.921052632

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
PTPRZ1 AND TM4SF18	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-GRIN3B	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-MRGPRF	0.985507246	0.98550725	0.985507246
PTPRZ1 AND SLC18B1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-CATSPER1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-MRGPRX2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-MRGPRX3	0.99270073	1	0.985507246
PTPRZ1 AND NOT-MRGPRX4	0.985507246	0.98550725	0.985507246
PTPRZ1 AND GALNT15	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-TMC1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-TMC2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-CLCN1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND CLCN2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND CLCN3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND ANTXR2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-GPR62	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-CLCN7	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-COMTD1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND SFXN2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-PDZD8	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-CALHM3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-OR5P2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-OR5P3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-GYLTL1B	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-SLC36A4	0.985507246	0.98550725	0.985507246
PTPRZ1 AND CLN3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-TMEM45B	0.985507246	0.98550725	0.985507246
PTPRZ1 AND CYP2R1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-AMICA1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-CLPS	0.985507246	0.98550725	0.985507246
PTPRZ1 AND TMEM52B	0.985507246	0.98550725	0.985507246
PTPRZ1 AND CLPTM1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-LRIG3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-TMEM132D	0.985507246	0.98550725	0.985507246
PTPRZ1 AND SLC15A4	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-TEX29	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-RXFP2	0.99270073	1	0.985507246
PTPRZ1 AND NOT-IFI27L1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND PLD4	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-ADSSL1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-GPHB5	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-SLC24A4	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-DEGS2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND CCR1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-CCR3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-CCR4	0.985507246	0.98550725	0.985507246
PTPRZ1 AND CCR5	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-C15orf27	0.985294118	1	0.971014493
PTPRZ1 AND NOT-CCR6	0.99270073	1	0.985507246
PTPRZ1 AND NOT-CCR8	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-ACKR2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND CMTM3	0.977777778	1	0.956521739
PTPRZ1 AND CMKLR1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-ZG16B	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-PAQR4	0.985507246	0.98550725	0.985507246
PTPRZ1 AND TMEM219	0.99270073	1	0.985507246
PTPRZ1 AND SLC38A10	0.985507246	0.98550725	0.985507246
PTPRZ1 AND CANT1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND TMEM132E	0.985507246	0.98550725	0.985507246
PTPRZ1 AND SEZ6	0.985507246	0.98550725	0.985507246
PTPRZ1 AND CYB5D2	0.985294118	1	0.971014493
PTPRZ1 AND NOT-SPNS2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-GJD3	0.985507246	0.98550725	0.985507246
GPM6B AND FAM210A	0.970149254	1	0.942028986
PTPRZ1 AND FAM210A	0.99270073	1	0.985507246
PTPRZ1 AND FAM69C	0.985507246	0.98550725	0.985507246
PTPRZ1 AND CLDND2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND ACER1	0.99270073	1	0.985507246
SYT11 AND NOT-IZUMO2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-IZUMO2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND CPT1C	0.985507246	0.98550725	0.985507246
PTPRZ1 AND CNGA3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND TMIGD2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-CNGA4	0.985507246	0.98550725	0.985507246
PTPRZ1 AND MFSD12	0.985507246	0.98550725	0.985507246
GPM6B AND NDUFA11	0.977777778	1	0.956521739

Antigen Pair Logic	F1	Precision	Recall
TNFRSF10A AND NOT-SLC19A1	0.925	0.880952381	0.973684211
QSOX2 AND NOT-GPC3	0.902439024	0.840909091	0.973684211
TNFRSF13C AND NOT-GPC3	0.915662651	0.844444444	1
QSOX2 AND NOT-MAGEA1	0.925	0.880952381	0.973684211
QSOX2 AND NOT-MUC17	0.906666667	0.918918919	0.894736842
TNFRSF13C AND NOT-MUC17	0.904761905	0.826086957	1
QSOX2 AND NOT-ITGB3	0.925	0.880952381	0.973684211
TNFRSF13C AND NOT-ITGB3	0.926829268	0.863636364	1
CLECL1 AND NOT-ITGB3	0.95	0.904761905	1
TNFRSF13C AND NOT-GPA33	0.915662651	0.844444444	1
QSOX2 AND NOT-TNFRSF8	0.947368421	0.947368421	0.947368421
TNFRSF13C AND NOT-TNFRSF8	0.938271605	0.88372093	1
CLECL1 AND NOT-TNFRSF8	0.948717949	0.925	0.973684211
CLECL1 AND NOT-CD33	0.926829268	0.863636364	1
CELSR1 AND NOT-EPHA2	0.9	0.857142857	0.947368421
TNFRSF13C AND NOT-CSPG4	0.904761905	0.826086957	1
CLECL1 AND NOT-CSPG4	0.915662651	0.844444444	1
CLECL1 AND NOT-NAALADL1	0.938271605	0.88372093	1
CLECL1 AND NOT-KCNE3	0.926829268	0.863636364	1
PTPN1 AND NOT-KCNE3	0.915662651	0.844444444	1
TNFRSF13C AND NOT-TMEM221	0.915662651	0.844444444	1
QSOX2 AND NOT-TMEM221	0.936708861	0.902439024	0.973684211
TNFRSF13C AND NOT-SMIM6	0.915662651	0.844444444	1
TNFRSF13C AND NOT-NEMP2	0.95	0.904761905	1
CLECL1 AND NOT-NEMP2	0.915662651	0.844444444	1
TNFRSF13C AND NOT-ERVMER34-1	0.974358974	0.95	1
CLECL1 AND NOT-ERVMER34-1	0.962025316	0.926829268	1
QSOX2 AND NOT-ERVMER34-1	0.948717949	0.925	0.973684211
TNFRSF13C AND NOT-CDH6	0.915662651	0.844444444	1
QSOX2 AND NOT-CDH7	0.935064935	0.923076923	0.947368421
TNFRSF13C AND NOT-MRLN	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SMLR1	0.936708861	0.902439024	0.973684211
TNFRSF13C AND NOT-100507547?	0.915662651	0.844444444	1
CLECL1 AND NOT-100507547?	0.918918919	0.944444444	0.894736842
TNFRSF13C AND NOT-ABCB6	0.987012987	0.974358974	1
CLECL1 AND NOT-KCNK7	0.938271605	0.88372093	1
QSOX2 AND NOT-KCNK7	0.948717949	0.925	0.973684211
TNFRSF13C AND NOT-CDH12	0.915662651	0.844444444	1
TNFRSF13C AND NOT-CDH15	0.926829268	0.863636364	1
QSOX2 AND NOT-CDH15	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-BCAP31	0.931506849	0.971428571	0.894736842
CLECL1 AND NOT-PLXNC1	0.936708861	0.902439024	0.973684211
CLECL1 AND NOT-PDZK1IP1	0.915662651	0.844444444	1
CLECL1 AND NOT-LPAR6	0.926829268	0.863636364	1
TNFRSF13C AND NOT-CHST4	0.915662651	0.844444444	1
TNFRSF13C AND NOT-DHRS9	0.926829268	0.863636364	1
CLECL1 AND NOT-DHRS9	0.926829268	0.863636364	1
SLC37A1 AND NOT-DHRS9	0.915662651	0.844444444	1
TNFRSF13C AND NOT-TENM1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-101805491?	0.915662651	0.844444444	1
TNFRSF13C AND NOT-DHRS2	0.926829268	0.863636364	1
QSOX2 AND NOT-DHRS2	0.947368421	0.947368421	0.947368421
CLECL1 AND NOT-MPZL2	0.926829268	0.863636364	1
CLECL1 AND NOT-KLRG1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-GPA33	0.915662651	0.844444444	1
CLECL1 AND NOT-IGSF6	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SIGMAR1	0.95	0.904761905	1
CLECL1 AND NOT-SIGMAR1	0.926829268	0.863636364	1
QSOX2 AND NOT-SIGMAR1	0.936708861	0.902439024	0.973684211
SLC37A1 AND NOT-SIGMAR1	0.935064935	0.923076923	0.947368421
CLECL1 AND NOT-LILRB2	0.915662651	0.844444444	1
TNFRSF13C AND NOT-NMUR1	0.915662651	0.844444444	1
QSOX2 AND NOT-B3GALT5	0.935064935	0.923076923	0.947368421
CLECL1 AND NOT-MRVI1	0.915662651	0.844444444	1
NOT-FAM210B AND PCGF3	0.96	0.972972973	0.947368421
TNFRSF13C AND NOT-PKDREJ	0.915662651	0.844444444	1
NOT-CDS1 AND CELSR1	0.935064935	0.923076923	0.947368421
CLECL1 AND NOT-ST3GAL6	0.926829268	0.863636364	1
CLECL1 AND NOT-WFDC2	0.915662651	0.844444444	1
CLECL1 AND NOT-CDIPT	0.962025316	0.926829268	1
CLECL1 AND NOT-CLEC10A	0.915662651	0.844444444	1
CLECL1 AND NOT-CRTAP	0.918918919	0.944444444	0.894736842
QSOX2 AND NOT-CRTAP	0.935064935	0.923076923	0.947368421
TNFRSF13C AND NOT-TM9SF1	0.918918919	0.944444444	0.894736842
NOT-ST6GALNAC2 AND SPPL3	0.936708861	0.902439024	0.973684211
CLECL1 AND NOT-ST6GALNAC2	0.938271605	0.88372093	1

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
PTPRZ1 AND NDUFA11	0.985294118	1	0.971014493
PTPRZ1 AND LRRC25	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-OR1I1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-ANKLE1	0.985507246	0.98550725	0.985507246
GPM6B AND B3GALT6	0.985294118	1	0.971014493
PTPRZ1 AND B3GALT6	0.977777778	1	0.956521739
PTPRZ1 AND NOT-SLC44A3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-CNTN1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-TMCO2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND RNF19B	0.985507246	0.98550725	0.985507246
PTPRZ1 AND TEDDM1	0.985507246	0.98550725	0.985507246
SYT11 AND NOT-SYT2	0.985294118	1	0.971014493
PTPRZ1 AND NOT-SYT2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-GOLT1A	0.970149254	1	0.942028986
PTPRZ1 AND NOT-TMEM125	0.970149254	1	0.942028986
PTPRZ1 AND DRAM2	0.99270073	1	0.985507246
PTPRZ1 AND C1orf162	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NKAIN4	0.985507246	0.98550725	0.985507246
SYT11 AND NOT-FITM2	0.977777778	1	0.956521739
PTPRZ1 AND COL4A2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND TBC1D20	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-PROKR2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND CST9L	0.985507246	0.98550725	0.985507246
PTPRZ1 AND EMID1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND TMEM150A	0.985507246	0.98550725	0.985507246
PTPRZ1 AND MBOAT2	0.977777778	1	0.956521739
PTPRZ1 AND NOT-CNTNAP5	0.985507246	0.98550725	0.985507246
SYT11 AND NOT-SGPP2	0.985294118	1	0.971014493
PTPRZ1 AND NOT-ACVR1C	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-LYPD6B	0.985294118	1	0.971014493
PTPRZ1 AND PQLC3	0.985294118	1	0.971014493
PTPRZ1 AND TMEM182	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-COL17A1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND DNAJC19	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-FAM3D	0.985294118	1	0.971014493
PTPRZ1 AND NOT-ZPLD1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND DCBLD2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND TPRA1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND SLC31A1	0.99270073	1	0.985507246
PTPRZ1 AND NOT-TMEM207	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-IL17RE	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-SYNPR	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-C4orf32	0.985507246	0.98550725	0.985507246
PTPRZ1 AND COX4I1	0.985294118	1	0.971014493
PTPRZ1 AND AASDH	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-TRAM1L1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-SLC9B2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-EMB	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-EGFLAM	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-UGT3A1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-TMEM174	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-COX7A1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND COX7A2	0.985294118	1	0.971014493
PTPRZ1 AND NOT-TAAR9	0.99270073	1	0.985507246
PTPRZ1 AND NOT-TAAR1	0.99270073	1	0.985507246
PTPRZ1 AND NOT-PACRG	0.985507246	0.98550725	0.985507246
PTPRZ1 AND B3GAT2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND COX8A	0.985294118	1	0.971014493
PTPRZ1 AND NOT-RAET1E	0.985294118	1	0.971014493
PTPRZ1 AND COX15	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-DPCR1	0.99270073	1	0.985507246
PTPRZ1 AND NOT-WBSCR28	0.985507246	0.98550725	0.985507246
SYT11 AND NOT-TMEM139	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-TMEM139	0.99270073	1	0.985507246
PTPRZ1 AND CPD	0.970149254	1	0.942028986
PTPRZ1 AND NOT-CLDN4	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-CLDN3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-CLDN23	0.985507246	0.98550725	0.985507246
PTPRZ1 AND CPT1A	0.985507246	0.98550725	0.985507246
PTPRZ1 AND TMEM68	0.985507246	0.98550725	0.985507246
PTPRZ1 AND CR1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND AGPAT6	0.985507246	0.98550725	0.985507246
PTPRZ1 AND LETM2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-HGSNAT	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-C9orf57	0.985507246	0.98550725	0.985507246
PTPRZ1 AND FAM69B	0.985507246	0.98550725	0.985507246

Antigen Pair Logic	F1	Precision	Recall
NOT-ST6GALNAC2 AND ETV6	0.961038961	0.948717949	0.973684211
NOT-ST6GALNAC2 AND ATP11B	0.929577465	1	0.868421053
NOT-ST6GALNAC2 AND FUT8	0.918918919	0.944444444	0.894736842
NOT-ST6GALNAC2 AND ADAM17	0.961038961	0.948717949	0.973684211
NOT-ST6GALNAC2 AND SNRNP40	0.986666667	1	0.973684211
CLECL1 AND NOT-TGOLN2	0.974358974	0.95	1
IL10RA AND NOT-TGOLN2	0.915662651	0.844444444	1
PTPN1 AND NOT-TGOLN2	0.962025316	0.926829268	1
TNFRSF13C AND NOT-DMRT2	0.915662651	0.844444444	1
CLECL1 AND NOT-CD226	0.926829268	0.863636364	1
CLECL1 AND NOT-TNFSF13B	0.926829268	0.863636364	1
TNFRSF13C AND NOT-DLL3	0.915662651	0.844444444	1
TNFRSF13C AND NOT-CLDN16	0.915662651	0.844444444	1
CLECL1 AND NOT-CLDN16	0.915662651	0.844444444	1
TNFRSF13C AND NOT-RRH	0.915662651	0.844444444	1
TNFRSF13C AND NOT-ADCY1	0.974358974	0.95	1
CLECL1 AND NOT-ADCY1	0.938271605	0.88372093	1
QSOX2 AND NOT-ADCY1	0.948717949	0.925	0.973684211
TNFRSF13C AND NOT-CERS1	0.95	0.904761905	1
TNFRSF13C AND NOT-NRG3	0.915662651	0.844444444	1
CLECL1 AND NOT-NUP50	0.961038961	0.948717949	0.973684211
CLECL1 AND NOT-ZMYND11	0.915662651	0.844444444	1
TNFRSF13C AND NOT-OR5I1	0.915662651	0.844444444	1
QSOX2 AND NOT-CCR9	0.929577465	1	0.868421053
TNFRSF13C AND NOT-GJB6	0.915662651	0.844444444	1
CLECL1 AND NOT-CEACAM3	0.95	0.904761905	1
TNFRSF13C AND NOT-SLC22A7	0.926829268	0.863636364	1
TNFRSF13C AND NOT-CD300C	0.936708861	0.902439024	0.973684211
CLECL1 AND NOT-CD300C	0.961038961	0.948717949	0.973684211
QSOX2 AND NOT-CD300C	0.935064935	0.923076923	0.947368421
TNFRSF13C AND NOT-NMU	0.915662651	0.844444444	1
CLECL1 AND NOT-CEACAM4	0.915662651	0.844444444	1
QSOX2 AND NOT-MMP24	0.936708861	0.902439024	0.973684211
TNFRSF13C AND NOT-OCLM	0.915662651	0.844444444	1
TNFRSF13C AND NOT-MAN1A2	0.962025316	0.926829268	1
CLECL1 AND NOT-MAN1A2	0.962025316	0.926829268	1
TNFRSF13C AND NOT-BTNL3	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SLC38A3	0.926829268	0.863636364	1
TNFRSF13C AND NOT-SLC27A4	0.948717949	0.925	0.973684211
QSOX2 AND NOT-SLC27A4	0.948717949	0.925	0.973684211
CLECL1 AND NOT-SLC27A3	0.935064935	0.923076923	0.947368421
TNFRSF13C AND NOT-KDELR3	0.915662651	0.844444444	1
TNFRSF13C AND NOT-TDRKH	0.95	0.904761905	1
CLECL1 AND NOT-TDRKH	0.948717949	0.925	0.973684211
QSOX2 AND NOT-TDRKH	0.935064935	0.923076923	0.947368421
CLECL1 AND NOT-LILRB3	0.961038961	0.948717949	0.973684211
PTPN1 AND NOT-LILRB3	0.926829268	0.863636364	1
TNFRSF13C AND NOT-ZPBP	0.935064935	0.923076923	0.947368421
QSOX2 AND NOT-ZPBP	0.918918919	0.944444444	0.894736842
CLECL1 AND NOT-TMEM115	0.947368421	0.947368421	0.947368421
TNFRSF13C AND NOT-ADAM30	0.926829268	0.863636364	1
TNFRSF13C AND NOT-PTPRT	0.926829268	0.863636364	1
QSOX2 AND NOT-PTPRT	0.961038961	0.948717949	0.973684211
TNFRSF13C AND NOT-IL1RAPL1	0.938271605	0.88372093	1
TNFRSF13C AND NOT-KLK8	0.926829268	0.863636364	1
TNFRSF13C AND NOT-GALNT6	0.926829268	0.863636364	1
CLECL1 AND NOT-GALNT6	0.938271605	0.88372093	1
TNFRSF13C AND NOT-GALNT5	0.915662651	0.844444444	1
TNFRSF13C AND NOT-RNF24	0.935064935	0.923076923	0.947368421
CLECL1 AND NOT-RNF24	0.947368421	0.947368421	0.947368421
TNFRSF13C AND NOT-PRRT2	0.915662651	0.844444444	1
TNFRSF13C AND NOT-PTGDR2	0.915662651	0.844444444	1
CLECL1 AND NOT-PTGDR2	0.938271605	0.88372093	1
QSOX2 AND NOT-PTGDR2	0.936708861	0.902439024	0.973684211
TNFRSF13C AND NOT-HRH3	0.926829268	0.863636364	1
CLECL1 AND NOT-HRH3	0.918918919	0.944444444	0.894736842
TNFRSF13C AND NOT-MRAP2	0.915662651	0.844444444	1
QSOX2 AND NOT-B4GALT7	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-CHRM2	0.936708861	0.902439024	0.973684211
CLECL1 AND NOT-ADCY7	0.938271605	0.88372093	1
TNFRSF13C AND NOT-CHRM3	0.915662651	0.844444444	1
CLECL1 AND NOT-CD300A	0.926829268	0.863636364	1
QSOX2 AND NOT-GPR182	0.918918919	0.944444444	0.894736842
CLECL1 AND NOT-MGAT4A	0.95	0.904761905	1
TNFRSF13C AND NOT-SLC46A1	0.915662651	0.844444444	1
CLECL1 AND NOT-SLC46A1	0.915662651	0.844444444	1

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
PTPRZ1 AND NOT-ADGRG4	0.985507246	0.98550725	0.985507246
PTPRZ1 AND PTCHD1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-CRHR1	0.985507246	0.98550725	0.985507246
GPM6A AND NOT-AADAC	0.970149254	1	0.942028986
GPM6B AND NOT-AADAC	0.99270073	1	0.985507246
APLP1 AND NOT-AADAC	0.977777778	1	0.956521739
NAT8L AND NOT-AADAC	0.970149254	1	0.942028986
LRRC37A3 AND NOT-AADAC	0.970149254	1	0.942028986
TMEM255A AND NOT-AADAC	0.977777778	1	0.956521739
PTPRZ1 AND NOT-AADAC	0.970149254	1	0.942028986
SCG2 AND NOT-AADAC	0.970149254	1	0.942028986
PTPRZ1 AND NOT-MUC17	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-ASB11	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-SELM	0.985507246	0.98550725	0.985507246
PTPRZ1 AND TMEM37	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-TRPM6	0.970149254	1	0.942028986
GPM6B AND ROMO1	0.977777778	1	0.956521739
PTPRZ1 AND ROMO1	0.977777778	1	0.956521739
PTPRZ1 AND NOT-WFDC10A	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-CST11	0.985507246	0.98550725	0.985507246
PTPRZ1 AND SIRPA	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-CRY2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND ADORA3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND SAMD8	0.985507246	0.98550725	0.985507246
PTPRZ1 AND CSF1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND FAM76B	0.985507246	0.98550725	0.985507246
PTPRZ1 AND CSF2RA	0.985507246	0.98550725	0.985507246
PTPRZ1 AND LAYN	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-TMEM86A	0.985507246	0.98550725	0.985507246
PTPRZ1 AND OR10A5	0.985507246	0.98550725	0.985507246
PTPRZ1 AND CSF3R	0.985507246	0.98550725	0.985507246
PTPRZ1 AND ALG10B	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-TMEM120B	0.985507246	0.98550725	0.985507246
PTPRZ1 AND B3GLCT	0.985507246	0.98550725	0.985507246
PTPRZ1 AND PRIMA1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND C14orf37	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-ISM2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-TMCO5A	0.99270073	1	0.985507246
PTPRZ1 AND NOT-CMTM4	0.985507246	0.98550725	0.985507246
PTPRZ1 AND CMTM2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-SLC22A31	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-TMED6	0.985507246	0.98550725	0.985507246
PTPRZ1 AND CSPG4	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-RBFOX3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND CD300LF	0.985507246	0.98550725	0.985507246
PTPRZ1 AND SLC47A2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-ODF4	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-SLC35G3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND TMEM199	0.985507246	0.98550725	0.985507246
PTPRZ1 AND DHRS13	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-TMC8	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-TMEM99	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-SMIM17	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-C19orf18	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-TMEM190	0.985507246	0.98550725	0.985507246
ITGAV AND NOT-TMC4	0.970149254	1	0.942028986
GPM6B AND DPY19L3	0.985294118	1	0.971014493
PTPRZ1 AND DPY19L3	0.970149254	1	0.942028986
PTPRZ1 AND NOT-FAM187B	0.99270073	1	0.985507246
PTPRZ1 AND NOT-ATP8B3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND SYT6	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-HFE2	0.985294118	1	0.971014493
PTPRZ1 AND NOT-FAM163A	0.977777778	1	0.956521739
PTPRZ1 AND NOT-MFSD4	0.977777778	1	0.956521739
PTPRZ1 AND NOT-PM20D1	0.985294118	1	0.971014493
PTPRZ1 AND SLC30A7	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-ADRA1A	0.985507246	0.98550725	0.985507246
PTPRZ1 AND CNIH3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND MANEAL	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-CTLA4	0.985294118	1	0.971014493
PTPRZ1 AND NOT-CLDN19	0.985507246	0.98550725	0.985507246
ITGAV AND NOT-C1orf210	0.970149254	1	0.942028986
LRRC37A3 AND NOT-C1orf210	0.970149254	1	0.942028986
PTPRZ1 AND NOT-C1orf210	0.99270073	1	0.985507246
NOT-C1orf210 AND FAM57A	0.970149254	1	0.942028986
CHST10 AND NOT-C1orf210	0.970149254	1	0.942028986

Antigen Pair Logic	F1	Precision	Recall
TNFRSF13C AND NOT-VSIG4	0.938271605	0.88372093	1
CLECL1 AND NOT-VSIG4	0.95	0.904761905	1
TNFRSF13C AND NOT-SLC52A3	0.974358974	0.95	1
QSOX2 AND NOT-SLC52A3	0.945945946	0.972222222	0.921052632
TNFRSF13C AND NOT-CHRM5	0.915662651	0.844444444	1
QSOX2 AND NOT-CHRM5	0.948717949	0.925	0.973684211
CLECL1 AND NOT-SCRG1	0.95	0.904761905	1
QSOX2 AND NOT-TEX261	0.948717949	0.925	0.973684211
TNFRSF13C AND NOT-CMTM1	0.938271605	0.88372093	1
TNFRSF13C AND NOT-CHRNA3	0.915662651	0.844444444	1
TNFRSF13C AND NOT-MFSD3	0.938271605	0.88372093	1
CLECL1 AND NOT-MFSD3	0.926829268	0.863636364	1
TNFRSF13C AND NOT-SDSL	0.915662651	0.844444444	1
CLECL1 AND NOT-SDSL	0.915662651	0.844444444	1
TNFRSF13C AND NOT-CHRNA4	0.915662651	0.844444444	1
CLECL1 AND NOT-CHRNA4	0.938271605	0.88372093	1
QSOX2 AND NOT-CHRNA4	0.935064935	0.923076923	0.947368421
TNFRSF13C AND NOT-ADCY8	0.915662651	0.844444444	1
CLECL1 AND NOT-CHRNB1	0.935064935	0.923076923	0.947368421
QSOX2 AND NOT-CHRNB1	0.947368421	0.947368421	0.947368421
TNFRSF13C AND NOT-CHRNE	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SLC22A9	0.915662651	0.844444444	1
TNFRSF13C AND NOT-CHRNG	0.962025316	0.926829268	1
CLECL1 AND NOT-CHRNG	0.918918919	0.944444444	0.894736842
QSOX2 AND NOT-CHRNG	0.933333333	0.945945946	0.921052632
TNFRSF13C AND NOT-PKD1L2	0.926829268	0.863636364	1
TNFRSF13C AND NOT-CSMD3	0.915662651	0.844444444	1
TNFRSF13C AND NOT-TMEM132B	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SLC5A11	0.945945946	0.972222222	0.921052632
QSOX2 AND NOT-SLC5A11	0.944444444	1	0.894736842
TNFRSF13C AND NOT-EVI5L	0.95	0.904761905	1
TNFRSF13C AND NOT-FAM210B	0.95	0.904761905	1
TNFRSF13C AND NOT-CMTM5	0.926829268	0.863636364	1
TNFRSF13C AND NOT-MOGAT1	0.962025316	0.926829268	1
TNFRSF13C AND NOT-LYPD1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-GRIN3A	0.915662651	0.844444444	1
TNFRSF13C AND NOT-CATSPER2	0.926829268	0.863636364	1
TNFRSF13C AND NOT-MRGPRX2	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SLC16A10	0.915662651	0.844444444	1
TNFRSF13C AND NOT-TMC1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-CLCN1	0.935064935	0.923076923	0.947368421
TNFRSF13C AND NOT-CLCN2	0.926829268	0.863636364	1
TNFRSF13C AND NOT-CLCN4	0.947368421	0.947368421	0.947368421
TNFRSF13C AND NOT-CLCNKB	0.915662651	0.844444444	1
TNFRSF13C AND NOT-COMTD1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-CALHM3	0.947368421	0.947368421	0.947368421
TNFRSF13C AND NOT-CLRN3	0.948717949	0.925	0.973684211
TNFRSF13C AND NOT-OR5P3	0.962025316	0.926829268	1
TNFRSF13C AND NOT-FAT3	0.915662651	0.844444444	1
TNFRSF13C AND NOT-TMEM52B	0.915662651	0.844444444	1
TNFRSF13C AND NOT-ANO4	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SLC24A4	0.915662651	0.844444444	1
TNFRSF13C AND NOT-CCR3	0.915662651	0.844444444	1
TNFRSF13C AND NOT-CCR8	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SLC38A10	0.95	0.904761905	1
TNFRSF13C AND NOT-WFIKKN2	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SEZ6	0.962025316	0.926829268	1
TNFRSF13C AND NOT-MIEF2	0.938271605	0.88372093	1
TNFRSF13C AND NOT-CNGA3	0.95	0.904761905	1
TNFRSF13C AND NOT-OR1I1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-GJB4	0.936708861	0.902439024	0.973684211
TNFRSF13C AND NOT-KLHDC7A	0.915662651	0.844444444	1
TNFRSF13C AND NOT-NKAIN4	0.915662651	0.844444444	1
TNFRSF13C AND NOT-CST9L	0.915662651	0.844444444	1
TNFRSF13C AND NOT-TMEM150A	0.916666667	0.970588235	0.868421053
TNFRSF13C AND NOT-CNTNAP5	0.915662651	0.844444444	1
TNFRSF13C AND NOT-AADAC	0.915662651	0.844444444	1
TNFRSF13C AND NOT-ACMSD	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SPATA3	0.926829268	0.863636364	1
TNFRSF13C AND NOT-TMEM198	0.962025316	0.926829268	1
TNFRSF13C AND NOT-ZPLD1	0.938271605	0.88372093	1
TNFRSF13C AND NOT-SLC31A1	0.938271605	0.88372093	1
TNFRSF13C AND NOT-SLCO6A1	0.926829268	0.863636364	1
TNFRSF13C AND NOT-ADORA1	0.926829268	0.863636364	1
TNFRSF13C AND COX7A2	0.915662651	0.844444444	1
TNFRSF13C AND NOT-TAAR9	0.915662651	0.844444444	1

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
PTPRZ1 AND NOT-SLC9B1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-PROM2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND ITPRIPL1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-KANSL1L	0.985507246	0.98550725	0.985507246
PTPRZ1 AND PLB1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-FAM132B	0.985507246	0.98550725	0.985507246
PTPRZ1 AND ARL6IP6	0.985507246	0.98550725	0.985507246
PTPRZ1 AND METTL21A	0.977777778	1	0.956521739
PTPRZ1 AND NOT-SLC23A3	0.970149254	1	0.942028986
PTPRZ1 AND NOT-GPBAR1	0.985294118	1	0.971014493
PTPRZ1 AND NOT-RMDN2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND GPR155	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-CCDC80	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-ADRA2B	0.985507246	0.98550725	0.985507246
PTPRZ1 AND XXYLT1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-CTSZ	0.985507246	0.98550725	0.985507246
PTPRZ1 AND IGSF11	0.985507246	0.98550725	0.985507246
PTPRZ1 AND CX3CR1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-PAQR3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND MGAT4D	0.985507246	0.98550725	0.985507246
PTPRZ1 AND THAP6	0.99270073	1	0.985507246
GPM6B AND SLC38A9	0.99270073	1	0.985507246
PTPRZ1 AND SLC38A9	0.99270073	1	0.985507246
PTPRZ1 AND SPINK13	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-SLC25A48	0.985507246	0.98550725	0.985507246
PTPRZ1 AND TMEM161B	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-MARVELD2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-BTNL9	0.985507246	0.98550725	0.985507246
PTPRZ1 AND CYBA	0.99270073	1	0.985507246
PTPRZ1 AND ARSK	0.985507246	0.98550725	0.985507246
PTPRZ1 AND CYBB	0.985507246	0.98550725	0.985507246
PTPRZ1 AND CCDC112	0.985507246	0.98550725	0.985507246
SYT11 AND NOT-NKAIN2	0.985294118	1	0.971014493
PTPRZ1 AND NOT-NKAIN2	0.985507246	0.98550725	0.985507246
GPM6B AND CCDC167	0.977777778	1	0.956521739
PTPRZ1 AND CCDC167	0.970149254	1	0.942028986
PTPRZ1 AND NOT-CYP1A2	0.99270073	1	0.985507246
PTPRZ1 AND NOT-CYP1B1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND ABCA13	0.985507246	0.98550725	0.985507246
SYT11 AND NOT-CLEC2L	0.977777778	1	0.956521739
PTPRZ1 AND VKORC1L1	0.977777778	1	0.956521739
PTPRZ1 AND NOT-CYP2A6	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-CYP2A7	0.985507246	0.98550725	0.985507246
PTPRZ1 AND ATP6V0E2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-CYP3A7	0.985294118	1	0.971014493
PTPRZ1 AND NOT-CYP2E1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-PEBP4	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-TMEM65	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-CYP2J2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-ANKRD46	0.985507246	0.98550725	0.985507246
SYT11 AND NOT-CYP3A4	0.985294118	1	0.971014493
GPM6A AND NOT-CYP3A4	0.985294118	1	0.971014493
PTPRZ1 AND NOT-CYP3A4	0.99270073	1	0.985507246
CHST10 AND NOT-CYP3A4	0.970149254	1	0.942028986
PTPRZ1 AND NOT-CYP3A5	0.985507246	0.98550725	0.985507246
PTPRZ1 AND KCNU1	0.985507246	0.98550725	0.985507246
SYT11 AND NOT-LINGO2	0.977777778	1	0.956521739
CHST10 AND NOT-LINGO2	0.970149254	1	0.942028986
PTPRZ1 AND NOT-CYP4B1	0.99270073	1	0.985507246
CHST10 AND NOT-CYP4B1	0.970149254	1	0.942028986
PTPRZ1 AND NOT-OR1Q1	0.977777778	1	0.956521739
PTPRZ1 AND NOT-CYP8B1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-FAAH2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-MOSPD2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND ARHGAP36	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-AWAT1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND CYP19A1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-SLC35G1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-CYP51A1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND DAD1	0.970149254	1	0.942028986
GPM6B AND TMTC3	0.970149254	1	0.942028986
PTPRZ1 AND NOT-TMTC3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-SLC5A8	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-GPR180	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-STOML3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND TMEM229B	0.985507246	0.98550725	0.985507246

Antigen Pair Logic	F1	Precision	Recall
TNFRSF13C AND NOT-TAAR1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-B3GAT2	0.935064935	0.923076923	0.947368421
TNFRSF13C AND NOT-DPCR1	0.962025316	0.926829268	1
TNFRSF13C AND NOT-TMEM139	0.915662651	0.844444444	1
TNFRSF13C AND NOT-OR6B1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-OR2F2	0.915662651	0.844444444	1
TNFRSF13C AND NOT-ADORA2B	0.915662651	0.844444444	1
TNFRSF13C AND NOT-CPD	0.915662651	0.844444444	1
TNFRSF13C AND NOT-COL26A1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SVOPL	0.926829268	0.863636364	1
TNFRSF13C AND NOT-SGCZ	0.915662651	0.844444444	1
TNFRSF13C AND NOT-RNF183	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SLITRK4	0.915662651	0.844444444	1
TNFRSF13C AND NOT-ADGRG4	0.974358974	0.95	1
TNFRSF13C AND NOT-CRHR1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-PTCHD1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-CRHR2	0.915662651	0.844444444	1
TNFRSF13C AND NOT-CHODL	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SLC32A1	0.962025316	0.926829268	1
TNFRSF13C AND NOT-SYT9	0.915662651	0.844444444	1
TNFRSF13C AND NOT-BEST3	0.915662651	0.844444444	1
TNFRSF13C AND NOT-ADRA1D	0.962025316	0.926829268	1
TNFRSF13C AND NOT-CMTM2	0.915662651	0.844444444	1
TNFRSF13C AND NOT-C16orf92	0.915662651	0.844444444	1
TNFRSF13C AND NOT-MGAT5B	0.962025316	0.926829268	1
TNFRSF13C AND NOT-RBFOX3	0.915662651	0.844444444	1
TNFRSF13C AND NOT-ODF4	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SLC35G3	0.915662651	0.844444444	1
TNFRSF13C AND NOT-ZNRF4	0.938271605	0.88372093	1
TNFRSF13C AND NOT-FAM187B	0.915662651	0.844444444	1
TNFRSF13C AND NOT-FAM163A	0.926829268	0.863636364	1
TNFRSF13C AND NOT-CNIH3	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SHISA4	0.95	0.904761905	1
TNFRSF13C AND NOT-ADIG	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SLC9B1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-NFAM1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-ITPRIPL1	0.95	0.904761905	1
TNFRSF13C AND NOT-ADRA2B	0.926829268	0.863636364	1
TNFRSF13C AND NOT-PLB1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SLC23A3	0.926829268	0.863636364	1
TNFRSF13C AND NOT-GPBAR1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-FAM19A4	0.915662651	0.844444444	1
TNFRSF13C AND NOT-PPM1L	0.915662651	0.844444444	1
TNFRSF13C AND NOT-CMTM8	0.915662651	0.844444444	1
TNFRSF13C AND NOT-IGSF11	0.938271605	0.88372093	1
TNFRSF13C AND NOT-KLB	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SLC25A48	0.915662651	0.844444444	1
TNFRSF13C AND NOT-ARSK	0.936708861	0.902439024	0.973684211
TNFRSF13C AND NOT-SLC2A12	0.95	0.904761905	1
TNFRSF13C AND NOT-CYP1A1	0.95	0.904761905	1
TNFRSF13C AND NOT-CYP1A2	0.915662651	0.844444444	1
TNFRSF13C AND NOT-CLEC2L	0.95	0.904761905	1
TNFRSF13C AND NOT-CYP3A7	0.915662651	0.844444444	1
TNFRSF13C AND NOT-CYP3A4	0.945945946	0.972222222	0.921052632
TNFRSF13C AND NOT-CYP3A5	0.915662651	0.844444444	1
TNFRSF13C AND NOT-TMEM74	0.915662651	0.844444444	1
TNFRSF13C AND NOT-LINGO2	0.915662651	0.844444444	1
TNFRSF13C AND NOT-FAAH2	0.915662651	0.844444444	1
TNFRSF13C AND NOT-CYP19A1	0.973684211	0.973684211	0.973684211
TNFRSF13C AND NOT-ZDHHC15	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SLC5A12	0.915662651	0.844444444	1
TNFRSF13C AND NOT-CCDC67	0.915662651	0.844444444	1
TNFRSF13C AND NOT-MFSD6L	0.947368421	0.947368421	0.947368421
TNFRSF13C AND NOT-RHBDL3	0.915662651	0.844444444	1
TNFRSF13C AND NOT-DCT	0.915662651	0.844444444	1
TNFRSF13C AND NOT-ADGRF3	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SPTSSB	0.915662651	0.844444444	1
TNFRSF13C AND NOT-CHST13	0.915662651	0.844444444	1
TNFRSF13C AND NOT-PKD1L1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-KCNG3	0.915662651	0.844444444	1
TNFRSF13C AND NOT-PTCRA	0.926829268	0.863636364	1
TNFRSF13C AND NOT-DHODH	0.95	0.904761905	1
TNFRSF13C AND NOT-DPP6	0.933333333	0.945945946	0.921052632
TNFRSF13C AND NOT-DRD1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-DRD3	0.915662651	0.844444444	1
TNFRSF13C AND NOT-DSC1	0.915662651	0.844444444	1

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
PTPRZ1 AND NOT-SYNE3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-FITM1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-TMEM30B	0.985507246	0.98550725	0.985507246
PTPRZ1 AND MDGA2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-TMEM92	0.99270073	1	0.985507246
PTPRZ1 AND RHBDL3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-TRPV3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND DCC	0.985507246	0.98550725	0.985507246
PTPRZ1 AND LPPR5	0.985507246	0.98550725	0.985507246
PTPRZ1 AND DENND1B	0.985507246	0.98550725	0.985507246
PTPRZ1 AND TOR1AIP2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND ACE	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-CNST	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-LRRN4	0.985507246	0.98550725	0.985507246
PTPRZ1 AND CABP7	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-ADGRF3	0.99270073	1	0.985507246
PTPRZ1 AND DDOST	0.977777778	1	0.956521739
FAM171B AND NOT-DSC3	0.970149254	1	0.942028986
PTPRZ1 AND NOT-FAM171B	0.970149254	1	0.942028986
FAM171B AND NOT-SIGLEC6	0.977777778	1	0.956521739
PTPRZ1 AND NOT-CLEC4F	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-SPTSSB	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-PKD1L1	0.99270073	1	0.985507246
PTPRZ1 AND NOT-SLC30A8	0.985507246	0.98550725	0.985507246
PTPRZ1 AND TMEM64	0.99270073	1	0.985507246
PTPRZ1 AND TMEM252	0.985507246	0.98550725	0.985507246
PTPRZ1 AND SSBP4	0.99270073	1	0.985507246
PTPRZ1 AND NOT-HTR3C	0.985507246	0.98550725	0.985507246
SYT11 AND NOT-KCNG3	0.977777778	1	0.956521739
PTPRZ1 AND NOT-KCNG3	0.977777778	1	0.956521739
PTPRZ1 AND FAM9C	0.985507246	0.98550725	0.985507246
GPM6B AND SPTSSA	0.985294118	1	0.971014493
PTPRZ1 AND SPTSSA	0.985294118	1	0.971014493
PTPRZ1 AND NOT-PTCRA	0.985507246	0.98550725	0.985507246
PTPRZ1 AND ABHD3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-DHCR7	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-DHCR24	0.985507246	0.98550725	0.985507246
PTPRZ1 AND DHODH	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-DIO1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-DIO2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-DNASE1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-AGER	0.985507246	0.98550725	0.985507246
PTPRZ1 AND DPAGT1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-DPP4	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-SLC26A3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-DRD4	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-DSC2	0.985507246	0.98550725	0.985507246
ITGAV AND NOT-DSC3	0.970149254	1	0.942028986
PTPRZ1 AND NOT-DSC3	0.985294118	1	0.971014493
PTPRZ1 AND DSCAM	0.985507246	0.98550725	0.985507246
PTPRZ1 AND JAG1	0.985294118	1	0.971014493
PTPRZ1 AND SLC26A2	0.977777778	1	0.956521739
PTPRZ1 AND DTNA	0.985507246	0.98550725	0.985507246
PTPRZ1 AND HBEGF	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-AGTR1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-AGTR2	0.985507246	0.98550725	0.985507246
GPM6B AND E2F5	0.985507246	0.98550725	0.985507246
PTPRZ1 AND E2F5	0.99270073	1	0.985507246
PTPRZ1 AND APLNR	0.985507246	0.98550725	0.985507246
PTPRZ1 AND GPR183	0.985507246	0.98550725	0.985507246
PTPRZ1 AND ECE1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-ECM1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-EDA	0.99270073	1	0.985507246
CHST10 AND NOT-EDA	0.970149254	1	0.942028986
PTPRZ1 AND S1PR1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND S1PR3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND EDNRA	0.985294118	1	0.971014493
PTPRZ1 AND EDNRB	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-PGAM5	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-B3GNT6	0.985507246	0.98550725	0.985507246
PTPRZ1 AND HIGD2A	0.985294118	1	0.971014493
PTPRZ1 AND NOT-EFNB2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND EFNB3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND EGF	0.985507246	0.98550725	0.985507246
PTPRZ1 AND CELSR3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-MEGF8	0.985507246	0.98550725	0.985507246

Antigen Pair Logic	F1	Precision	Recall
TNFRSF13C AND NOT-DSC3	0.926829268	0.863636364	1
TNFRSF13C AND NOT-DTNA	0.915662651	0.844444444	1
TNFRSF13C AND NOT-AGTR2	0.95	0.904761905	1
TNFRSF13C AND NOT-ECM1	0.926829268	0.863636364	1
TNFRSF13C AND NOT-PGAM5	0.938271605	0.88372093	1
TNFRSF13C AND NOT-B3GNT6	0.926829268	0.863636364	1
TNFRSF13C AND NOT-EFNB1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-EFNB3	0.915662651	0.844444444	1
TNFRSF13C AND NOT-EGF	0.915662651	0.844444444	1
TNFRSF13C AND NOT-PPAPDC1A	0.915662651	0.844444444	1
TNFRSF13C AND NOT-PLBD2	0.95	0.904761905	1
TNFRSF13C AND NOT-VSTM4	0.915662651	0.844444444	1
TNFRSF13C AND NOT-GRAMD2	0.926829268	0.863636364	1
TNFRSF13C AND NOT-CADM4	0.915662651	0.844444444	1
TNFRSF13C AND NOT-TMEM17	0.915662651	0.844444444	1
TNFRSF13C AND NOT-GABRR3	0.915662651	0.844444444	1
TNFRSF13C AND NOT-NRK	0.926829268	0.863636364	1
TNFRSF13C AND NOT-EPHA7	0.915662651	0.844444444	1
TNFRSF13C AND NOT-EPHB3	0.915662651	0.844444444	1
TNFRSF13C AND NOT-EPHB4	0.938271605	0.88372093	1
TNFRSF13C AND NOT-EPO	0.962025316	0.926829268	1
TNFRSF13C AND NOT-EXTL1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-F2RL2	0.938271605	0.88372093	1
TNFRSF13C AND NOT-FAP	0.915662651	0.844444444	1
TNFRSF13C AND NOT-MS4A15	0.915662651	0.844444444	1
TNFRSF13C AND NOT-HEPACAM	0.926829268	0.863636364	1
TNFRSF13C AND NOT-FCAR	0.938271605	0.88372093	1
TNFRSF13C AND NOT-RNF152	0.915662651	0.844444444	1
TNFRSF13C AND NOT-MS4A2	0.974358974	0.95	1
TNFRSF13C AND NOT-SLC39A12	0.915662651	0.844444444	1
TNFRSF13C AND NOT-OPN5	0.915662651	0.844444444	1
TNFRSF13C AND NOT-MMD2	0.915662651	0.844444444	1
TNFRSF13C AND NOT-FGF4	0.926829268	0.863636364	1
TNFRSF13C AND NOT-FGF6	0.915662651	0.844444444	1
TNFRSF13C AND NOT-FGF10	0.938271605	0.88372093	1
TNFRSF13C AND NOT-FGFR3	0.935064935	0.923076923	0.947368421
TNFRSF13C AND NOT-FGG	0.915662651	0.844444444	1
TNFRSF13C AND NOT-DOLK	0.938271605	0.88372093	1
TNFRSF13C AND NOT-ADGRL1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SV2C	0.915662651	0.844444444	1
TNFRSF13C AND NOT-MLC1	0.926829268	0.863636364	1
TNFRSF13C AND NOT-ASTN2	0.915662651	0.844444444	1
TNFRSF13C AND NOT-ACSL6	0.915662651	0.844444444	1
TNFRSF13C AND NOT-KCNH4	0.915662651	0.844444444	1
TNFRSF13C AND NOT-KCNH3	0.948717949	0.925	0.973684211
TNFRSF13C AND NOT-GPR161	0.938271605	0.88372093	1
TNFRSF13C AND NOT-SLC44A1	0.926829268	0.863636364	1
TNFRSF13C AND NOT-FOLR1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-OR52A1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SLC16A8	0.938271605	0.88372093	1
TNFRSF13C AND NOT-SEZ6L	0.915662651	0.844444444	1
TNFRSF13C AND NOT-HYAL4	0.915662651	0.844444444	1
TNFRSF13C AND NOT-LPAR3	0.915662651	0.844444444	1
TNFRSF13C AND NOT-FPR2	0.933333333	0.945945946	0.921052632
TNFRSF13C AND NOT-CLEC5A	0.938271605	0.88372093	1
TNFRSF13C AND NOT-FLRT1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-ADAM2	0.948717949	0.925	0.973684211
TNFRSF13C AND NOT-FUT2	0.95	0.904761905	1
TNFRSF13C AND NOT-FUT5	0.95	0.904761905	1
TNFRSF13C AND NOT-KDSR	0.95	0.904761905	1
TNFRSF13C AND NOT-CADM2	0.915662651	0.844444444	1
TNFRSF13C AND NOT-FAM26E	0.915662651	0.844444444	1
TNFRSF13C AND NOT-GABRA2	0.915662651	0.844444444	1
TNFRSF13C AND NOT-COL24A1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-GABRB1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-TMEM196	0.915662651	0.844444444	1
TNFRSF13C AND NOT-CDRT15L2	0.96	0.972972973	0.947368421
TNFRSF13C AND NOT-GABRD	0.915662651	0.844444444	1
TNFRSF13C AND NOT-GABRE	0.915662651	0.844444444	1
TNFRSF13C AND NOT-GABRG1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-GABRR1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-GABRR2	0.915662651	0.844444444	1
TNFRSF13C AND NOT-UTS2B	0.916666667	0.970588235	0.868421053
TNFRSF13C AND NOT-TMEM59L	0.962025316	0.926829268	1
TNFRSF13C AND NOT-KLK5	0.95	0.904761905	1
TNFRSF13C AND NOT-GALR1	0.926829268	0.863636364	1

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
SYT11 AND NOT-MEGF9	0.977777778	1	0.956521739
PTPRZ1 AND NOT-MEGF9	0.985507246	0.98550725	0.985507246
PTPRZ1 AND EGFR	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-MPZL3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-DNAH10	0.985507246	0.98550725	0.985507246
PTPRZ1 AND METTL7B	0.985507246	0.98550725	0.985507246
PTPRZ1 AND MYRFL	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-PLBD2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-VSTM4	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-FAM24B	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-MCEMP1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-CYP4Z1	0.985294118	1	0.971014493
PTPRZ1 AND ABCA1	0.985294118	1	0.971014493
SYT11 AND KRTCAP2	0.970149254	1	0.942028986
GPM6A AND KRTCAP2	0.970149254	1	0.942028986
PTPRZ1 AND NOT-APLF	0.985507246	0.98550725	0.985507246
GOLM1 AND NOT-KRTCAP3	0.970149254	1	0.942028986
PTPRZ1 AND NOT-KRTCAP3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND ELN	0.99270073	1	0.985507246
PTPRZ1 AND NOT-TMEM17	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-PLD6	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-SLC39A11	0.985507246	0.98550725	0.985507246
PTPRZ1 AND RDM1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND EMP1	0.985294118	1	0.971014493
PTPRZ1 AND NOT-SPNS3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND EMP3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-TIGIT	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-TMEM154	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-TMEM192	0.985507246	0.98550725	0.985507246
PTPRZ1 AND TAPT1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND DNAJC18	0.985507246	0.98550725	0.985507246
PTPRZ1 AND ENG	0.985507246	0.98550725	0.985507246
PTPRZ1 AND ENPEP	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-TSNARE1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND HTRA4	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-C9orf91	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-CCDC107	0.985507246	0.98550725	0.985507246
PTPRZ1 AND SUSD3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-CT83	0.985507246	0.98550725	0.985507246
PTPRZ1 AND SLC25A43	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-NRK	0.985507246	0.98550725	0.985507246
SYT11 AND NOT-ANO5	0.985294118	1	0.971014493
PTPRZ1 AND NOT-ANO5	0.985507246	0.98550725	0.985507246
PTPRZ1 AND STOM	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-EPHA1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-EPHA4	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-EPHA8	0.985507246	0.98550725	0.985507246
PTPRZ1 AND EPHB1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND EPHB2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND SLC44A5	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-EPHB4	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-EPHB6	0.985507246	0.98550725	0.985507246
PTPRZ1 AND STX2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND CLN8	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-EPO	0.977777778	1	0.956521739
PTPRZ1 AND EPOR	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-LVRN	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-SLC36A1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-ERBB3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-ERBB4	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-EREG	0.985507246	0.98550725	0.985507246
PTPRZ1 AND ERF	0.985507246	0.98550725	0.985507246
PTPRZ1 AND ERN1	0.977777778	1	0.956521739
PTPRZ1 AND NOT-ABCA2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-ETFDH	0.985507246	0.98550725	0.985507246
PTPRZ1 AND EVI2B	0.985507246	0.98550725	0.985507246
PTPRZ1 AND EXT1	0.970149254	1	0.942028986
PTPRZ1 AND EXTL2	0.99270073	1	0.985507246
GPM6B AND F2R	0.99270073	1	0.985507246
NOT-C1orf186 AND F2R	0.970149254	1	0.942028986
PTPRZ1 AND F2R	0.985294118	1	0.971014493
PTPRZ1 AND ALCAM	0.985507246	0.98550725	0.985507246
PTPRZ1 AND F3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND ACSL3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-FAP	0.985507246	0.98550725	0.985507246
PTPRZ1 AND LRRC55	0.985507246	0.98550725	0.985507246

Antigen Pair Logic	F1	Precision	Recall
TNFRSF13C AND NOT-LY6G6F	0.915662651	0.844444444	1
TNFRSF13C AND NOT-TMIE	0.915662651	0.844444444	1
TNFRSF13C AND NOT-WFDC11	0.915662651	0.844444444	1
TNFRSF13C AND NOT-TAS2R41	0.915662651	0.844444444	1
TNFRSF13C AND NOT-FAM205A	0.915662651	0.844444444	1
TNFRSF13C AND NOT-LRRTM2	0.973684211	0.973684211	0.973684211
TNFRSF13C AND NOT-CFAP61	0.938271605	0.88372093	1
TNFRSF13C AND NOT-LRIT1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-ABCA12	0.926829268	0.863636364	1
TNFRSF13C AND NOT-OR2M4	0.915662651	0.844444444	1
TNFRSF13C AND NOT-OR2L2	0.915662651	0.844444444	1
TNFRSF13C AND NOT-KCNG2	0.961038961	0.948717949	0.973684211
TNFRSF13C AND NOT-SLC13A4	0.926829268	0.863636364	1
TNFRSF13C AND NOT-GALNT8	0.926829268	0.863636364	1
TNFRSF13C AND NOT-OR5K1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-OR8B8	0.962025316	0.926829268	1
TNFRSF13C AND NOT-OR8B2	0.915662651	0.844444444	1
TNFRSF13C AND NOT-OR7C2	0.95	0.904761905	1
TNFRSF13C AND NOT-HS6ST3	0.938271605	0.88372093	1
TNFRSF13C AND NOT-GDNF	0.915662651	0.844444444	1
TNFRSF13C AND NOT-ADGRF1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-OR2H1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-OR1J2	0.915662651	0.844444444	1
TNFRSF13C AND NOT-GGCX	0.948717949	0.925	0.973684211
TNFRSF13C AND NOT-GHSR	0.915662651	0.844444444	1
TNFRSF13C AND NOT-GJA8	0.926829268	0.863636364	1
TNFRSF13C AND NOT-PKD2L2	0.926829268	0.863636364	1
TNFRSF13C AND NOT-DKKL1	0.974358974	0.95	1
TNFRSF13C AND NOT-KCNH5	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SIGLEC9	0.915662651	0.844444444	1
TNFRSF13C AND NOT-GPR78	0.915662651	0.844444444	1
TNFRSF13C AND NOT-PCDH17	0.915662651	0.844444444	1
TNFRSF13C AND NOT-GLP1R	0.915662651	0.844444444	1
TNFRSF13C AND NOT-GLRB	0.915662651	0.844444444	1
TNFRSF13C AND NOT-CECR6	0.926829268	0.863636364	1
TNFRSF13C AND NOT-PCLO	0.915662651	0.844444444	1
TNFRSF13C AND NOT-WFDC10B	0.915662651	0.844444444	1
TNFRSF13C AND NOT-GP2	0.915662651	0.844444444	1
TNFRSF13C AND NOT-GP9	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SLCO3A1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-GPM6B	0.961038961	0.948717949	0.973684211
TNFRSF13C AND NOT-IFNL1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-GPR3	0.915662651	0.844444444	1
TNFRSF13C AND NOT-GPR4	0.962025316	0.926829268	1
TNFRSF13C AND NOT-XCR1	0.926829268	0.863636364	1
TNFRSF13C AND NOT-OR8D2	0.938271605	0.88372093	1
TNFRSF13C AND NOT-SLC39A5	0.915662651	0.844444444	1
TNFRSF13C AND NOT-GPR12	0.926829268	0.863636364	1
TNFRSF13C AND NOT-ZDHHC22	0.915662651	0.844444444	1
TNFRSF13C AND NOT-OR4N4	0.915662651	0.844444444	1
TNFRSF13C AND NOT-GPR15	0.926829268	0.863636364	1
TNFRSF13C AND NOT-TMEM235	0.915662651	0.844444444	1
TNFRSF13C AND NOT-FAM171A2	0.987012987	0.974358974	1
TNFRSF13C AND NOT-SLC13A5	0.915662651	0.844444444	1
TNFRSF13C AND NOT-TMEM145	0.926829268	0.863636364	1
TNFRSF13C AND NOT-NKPD1	0.926829268	0.863636364	1
TNFRSF13C AND NOT-GPR21	0.95	0.904761905	1
TNFRSF13C AND NOT-VSTM1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-LPAR4	0.938271605	0.88372093	1
TNFRSF13C AND NOT-GPR25	0.915662651	0.844444444	1
TNFRSF13C AND NOT-DNAJC5G	0.915662651	0.844444444	1
TNFRSF13C AND NOT-CDH19	0.926829268	0.863636364	1
TNFRSF13C AND NOT-LSMEM1	0.918918919	0.944444444	0.894736842
TNFRSF13C AND NOT-SCARA5	0.915662651	0.844444444	1
TNFRSF13C AND NOT-CRB2	0.961038961	0.948717949	0.973684211
TNFRSF13C AND NOT-GPR39	0.95	0.904761905	1
TNFRSF13C AND NOT-CFAP47	0.926829268	0.863636364	1
TNFRSF13C AND NOT-TUSC5	0.974358974	0.95	1
TNFRSF13C AND NOT-GRIA1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-GRIA2	0.915662651	0.844444444	1
TNFRSF13C AND NOT-GRID2	0.95	0.904761905	1
TNFRSF13C AND NOT-GRIK2	0.915662651	0.844444444	1
TNFRSF13C AND NOT-TMPRSS11E	0.915662651	0.844444444	1
TNFRSF13C AND NOT-GRIN2A	0.915662651	0.844444444	1
TNFRSF13C AND NOT-GRIN2B	0.915662651	0.844444444	1
TNFRSF13C AND NOT-GRIN2D	0.915662651	0.844444444	1

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
PTPRZ1 AND NOT-TMEM26	0.99270073	1	0.985507246
PTPRZ1 AND NOT-FAT2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-C10orf35	0.985507246	0.98550725	0.985507246
CHST10 AND NOT-C10orf35	0.970149254	1	0.942028986
PTPRZ1 AND TMEM218	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-TPCN2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND MPEG1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-MS4A15	0.985507246	0.98550725	0.985507246
PTPRZ1 AND ABCA3	0.970149254	1	0.942028986
PTPRZ1 AND NOT-LRTOMT	0.985507246	0.98550725	0.985507246
PTPRZ1 AND FBN2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND HEPACAM	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-FCAR	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-FCER1A	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-MS4A2	0.99270073	1	0.985507246
PTPRZ1 AND NOT-FCER2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-MARCH8	0.985507246	0.98550725	0.985507246
PTPRZ1 AND REEP3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-FAM171A1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-SLC39A12	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-ADGRG5	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-CES5A	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-FAM26D	0.985507246	0.98550725	0.985507246
PTPRZ1 AND FAM162B	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-OPN5	0.985507246	0.98550725	0.985507246
PTPRZ1 AND ADGRF5	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-TMEM217	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-PI16	0.985507246	0.98550725	0.985507246
PTPRZ1 AND C6orf89	0.99270073	1	0.985507246
PTPRZ1 AND LEMD2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND FCGR3B	0.985507246	0.98550725	0.985507246
PTPRZ1 AND RNF182	0.985507246	0.98550725	0.985507246
PTPRZ1 AND FCGRT	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-SDK1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND DAGLB	0.985507246	0.98550725	0.985507246
PTPRZ1 AND THSD7A	0.985507246	0.98550725	0.985507246
GPM6B AND TMED4	0.977777778	1	0.956521739
PTPRZ1 AND TMED4	0.977777778	1	0.956521739
PTPRZ1 AND FCN2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-KIAA1324L	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-FDPS	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-HS3ST5	0.970149254	1	0.942028986
PTPRZ1 AND NOT-GPRC6A	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-ADGRF2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND LHFPL5	0.985507246	0.98550725	0.985507246
GPM6B AND ALDH3A2	0.977777778	1	0.956521739
PTPRZ1 AND ALDH3A2	0.977777778	1	0.956521739
PTPRZ1 AND NOT-FGF6	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-FGF10	0.985507246	0.98550725	0.985507246
PTPRZ1 AND FGFR1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-CLCA4	0.985507246	0.98550725	0.985507246
PTPRZ1 AND FKBP1A	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NLGN4Y	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-LMTK2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND ADGRL1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND SLITRK3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NLGN1	0.970149254	1	0.942028986
PTPRZ1 AND SACM1L	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-ZP1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-SCAP	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-P2RX2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND DIP2C	0.985507246	0.98550725	0.985507246
PTPRZ1 AND SORCS3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-SV2C	0.985294118	1	0.971014493
PTPRZ1 AND NOT-PCNX	0.985507246	0.98550725	0.985507246
PTPRZ1 AND ABCB7	0.977777778	1	0.956521739
PTPRZ1 AND KLHDC10	0.985507246	0.98550725	0.985507246
PTPRZ1 AND TMCC1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-ENDOD1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND TBC1D2B	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NFASC	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-ATP10B	0.985507246	0.98550725	0.985507246
PTPRZ1 AND PLXND1	0.985294118	1	0.971014493
PTPRZ1 AND ANKLE2	0.99270073	1	0.985507246
PTPRZ1 AND NOT-LRCH1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND GRAMD4	0.985507246	0.98550725	0.985507246

Antigen Pair Logic	F1	Precision	Recall
TNFRSF13C AND NOT-GRM1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-GRM2	0.915662651	0.844444444	1
TNFRSF13C AND NOT-GRM3	0.926829268	0.863636364	1
TNFRSF13C AND NOT-GRM4	0.95	0.904761905	1
TNFRSF13C AND NOT-GRM7	0.915662651	0.844444444	1
TNFRSF13C AND NOT-CYP2S1	0.926829268	0.863636364	1
TNFRSF13C AND NOT-GUCA2B	0.915662651	0.844444444	1
TNFRSF13C AND NOT-GUCY2F	0.95	0.904761905	1
TNFRSF13C AND NOT-NPC1L1	0.926829268	0.863636364	1
TNFRSF13C AND NOT-PCDHB1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-LRP12	0.926829268	0.863636364	1
TNFRSF13C AND NOT-HAS1	0.926829268	0.863636364	1
TNFRSF13C AND NOT-HCRTR2	0.938271605	0.88372093	1
TNFRSF13C AND NOT-HFE	0.95	0.904761905	1
TNFRSF13C AND NOT-ERVW-1	0.926829268	0.863636364	1
TNFRSF13C AND NOT-HGF	0.926829268	0.863636364	1
TNFRSF13C AND HLA-DRA	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SLC29A2	0.962025316	0.926829268	1
TNFRSF13C AND NOT-HRH1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-HRH2	0.915662651	0.844444444	1
TNFRSF13C AND NOT-HTR2C	0.915662651	0.844444444	1
TNFRSF13C AND NOT-HTR3A	0.936708861	0.902439024	0.973684211
TNFRSF13C AND NOT-HTR4	0.95	0.904761905	1
TNFRSF13C AND NOT-HTR7	0.915662651	0.844444444	1
TNFRSF13C AND NOT-CHSY3	0.948717949	0.925	0.973684211
TNFRSF13C AND NOT-FFAR4	0.95	0.904761905	1
TNFRSF13C AND NOT-ICAM4	0.95890411	1	0.921052632
TNFRSF13C AND NOT-TMEM225	0.915662651	0.844444444	1
TNFRSF13C AND NOT-B4GALNT4	0.926829268	0.863636364	1
TNFRSF13C AND NOT-TMEM52	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SLC35E4	0.915662651	0.844444444	1
TNFRSF13C AND NOT-LRRC66	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SLC35D3	0.915662651	0.844444444	1
TNFRSF13C AND NOT-COX8C	0.915662651	0.844444444	1
TNFRSF13C AND NOT-KCNT2	0.915662651	0.844444444	1
TNFRSF13C AND NOT-MOGAT3	0.926829268	0.863636364	1
TNFRSF13C AND NOT-LRRTM1	0.926829268	0.863636364	1
TNFRSF13C AND NOT-SERINC2	0.926829268	0.863636364	1
TNFRSF13C AND NOT-SLC6A18	0.915662651	0.844444444	1
TNFRSF13C AND NOT-RGSL1	0.95	0.904761905	1
TNFRSF13C AND NOT-FASLG	0.915662651	0.844444444	1
TNFRSF13C AND NOT-IL12B	0.915662651	0.844444444	1
TNFRSF13C AND NOT-IL12RB2	0.962025316	0.926829268	1
TNFRSF13C AND NOT-AQP6	0.915662651	0.844444444	1
TNFRSF13C AND NOT-INSRR	0.926829268	0.863636364	1
TNFRSF13C AND NOT-ITGA5	0.938271605	0.88372093	1
TNFRSF13C AND NOT-ABCC6	0.915662651	0.844444444	1
TNFRSF13C AND NOT-ITGB3	0.926829268	0.863636364	1
TNFRSF13C AND NOT-KCNA1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-KCNA4	0.961038961	0.948717949	0.973684211
TNFRSF13C AND NOT-KCNB1	0.938271605	0.88372093	1
TNFRSF13C AND NOT-KCNC2	0.948717949	0.925	0.973684211
TNFRSF13C AND NOT-IGFL1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-KCND3	0.926829268	0.863636364	1
TNFRSF13C AND NOT-KCNE1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-LHFPL4	0.926829268	0.863636364	1
TNFRSF13C AND NOT-TMEM110	0.935064935	0.923076923	0.947368421
TNFRSF13C AND NOT-KCNF1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-KCNG1	0.929577465	1	0.868421053
TNFRSF13C AND NOT-KCNH1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-KCNJ4	0.915662651	0.844444444	1
TNFRSF13C AND NOT-KCNJ12	0.938271605	0.88372093	1
TNFRSF13C AND NOT-KCNJ14	0.915662651	0.844444444	1
TNFRSF13C AND NOT-KCNK2	0.938271605	0.88372093	1
TNFRSF13C AND NOT-KCNK3	0.915662651	0.844444444	1
TNFRSF13C AND NOT-KCNN3	0.926829268	0.863636364	1
TNFRSF13C AND NOT-KIR2DL4	0.95	0.904761905	1
TNFRSF13C AND NOT-KIR2DS4	0.986666667	1	0.973684211
TNFRSF13C AND NOT-KIR3DL2	0.987012987	0.974358974	1
TNFRSF13C AND NOT-ILDR2	0.915662651	0.844444444	1
TNFRSF13C AND NOT-CLEC12B	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SMIM1	0.938271605	0.88372093	1
TNFRSF13C AND NOT-TMEM212	0.926829268	0.863636364	1
TNFRSF13C AND NOT-OR51I1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-OR52D1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-LHCGR	0.915662651	0.844444444	1

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
PTPRZ1 AND CLCC1	0.977777778	1	0.956521739
PTPRZ1 AND NOT-MLANA	0.985507246	0.98550725	0.985507246
PTPRZ1 AND STAB1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND SLC35D1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND GANAB	0.977777778	1	0.956521739
PTPRZ1 AND NOT-ATP11B	0.985507246	0.98550725	0.985507246
PTPRZ1 AND ARL6IP1	0.985507246	0.98550725	0.985507246
SYT11 AND NOT-ABCB9	0.970149254	1	0.942028986
SYT11 AND NOT-ABCA6	0.977777778	1	0.956521739
SYT11 AND NOT-LPCAT4	0.99270073	1	0.985507246
SYT11 AND NOT-CDRT15L2	0.985507246	0.98550725	0.985507246
SYT11 AND NOT-CYP4X1	0.99270073	1	0.985507246
SYT11 AND NOT-IL1RAPL2	0.985507246	0.98550725	0.985507246
SYT11 AND NOT-SMIM24	0.985507246	0.98550725	0.985507246
SYT11 AND SPCS1	0.970149254	1	0.942028986
SYT11 AND SLC25A5	0.99270073	1	0.985507246
SYT11 AND NOT-HCN1	0.985294118	1	0.971014493
SYT11 AND ITGAV	0.977777778	1	0.956521739
SYT11 AND NOT-KCNA1	0.985294118	1	0.971014493
SYT11 AND NOT-KCNC4	0.970149254	1	0.942028986
SYT11 AND NOT-KCNJ12	0.977777778	1	0.956521739
SYT11 AND NOT-KCNS1	0.99270073	1	0.985507246
SYT11 AND NOT-KEL	0.977777778	1	0.956521739
SYT11 AND NOT-KIR3DL2	0.985507246	0.98550725	0.985507246
SYT11 AND NOT-FAM209B	0.985507246	0.98550725	0.985507246
SYT11 AND NOT-LMO7	0.977777778	1	0.956521739
SYT11 AND NOT-ADAM11	0.99270073	1	0.985507246
SYT11 AND NDUFB5	0.977777778	1	0.956521739
SYT11 AND NOT-NTF3	0.985294118	1	0.971014493
SYT11 AND NOT-OCA2	0.985507246	0.98550725	0.985507246
SYT11 AND NOT-OPRK1	0.970149254	1	0.942028986
SYT11 AND NOT-OR1F1	0.99270073	1	0.985507246
SYT11 AND NOT-ANO7	0.985507246	0.98550725	0.985507246
SYT11 AND NOT-ATL1	0.985507246	0.98550725	0.985507246
SYT11 AND TMX2	0.970149254	1	0.942028986
SYT11 AND TMEM14C	0.970149254	1	0.942028986
SYT11 AND NOT-ATP8A2	0.985294118	1	0.971014493
SYT11 AND NOT-SMIM11	0.985507246	0.98550725	0.985507246
SYT11 AND SLC38A2	0.977777778	1	0.956521739
SYT11 AND NOT-MRAP	0.977777778	1	0.956521739
SYT11 AND NOT-OTOR	0.970149254	1	0.942028986
SYT11 AND NOT-PTGER3	0.99270073	1	0.985507246
SYT11 AND NOT-PTPRR	0.970149254	1	0.942028986
PTPRZ1 AND SYT11	0.985294118	1	0.971014493
SYT11 AND SELK	0.977777778	1	0.956521739
SYT11 AND NOT-ABCG4	0.985294118	1	0.971014493
SYT11 AND NOT-WBSCR17	0.970149254	1	0.942028986
SYT11 AND NOT-SYNDIG1L	0.970149254	1	0.942028986
SYT11 AND NOT-SLC10A1	0.977777778	1	0.956521739
SYT11 AND NOT-SLC15A1	0.977777778	1	0.956521739
SYT11 AND NOT-VAMP1	0.99270073	1	0.985507246
SYT11 AND NOT-TSPAN8	0.99270073	1	0.985507246
SYT11 AND NOT-TRPV1	0.99270073	1	0.985507246
SYT11 AND WRB	0.985507246	0.98550725	0.985507246
SYT11 AND PTTG1IP	0.985294118	1	0.971014493
SYT11 AND NOT-PQLC1	0.985294118	1	0.971014493
SYT11 AND NOT-AGPAT9	0.970149254	1	0.942028986
SYT11 AND NOT-DGKE	0.985294118	1	0.971014493
SYT11 AND ATP6V0E1	0.970149254	1	0.942028986
SYT11 AND NOT-SYNGR3	0.985507246	0.98550725	0.985507246
SYT11 AND NOT-VAPB	0.985294118	1	0.971014493
SYT11 AND NOT-CD8B	0.985507246	0.98550725	0.985507246
SYT11 AND NOT-CYP7B1	0.985507246	0.98550725	0.985507246
SYT11 AND NOT-SIGLEC6	0.99270073	1	0.985507246
SYT11 AND CD63	0.970149254	1	0.942028986
PTPRZ1 AND NUP210	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-FLT3	0.985294118	1	0.971014493
PTPRZ1 AND SEL1L3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-FLT3LG	0.985507246	0.98550725	0.985507246
PTPRZ1 AND ASTN2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-ATP11A	0.985507246	0.98550725	0.985507246
PTPRZ1 AND DENND5A	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-ADGRL2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-FMO2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND ADGRL3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND FMO3	0.985507246	0.98550725	0.985507246

Antigen Pair Logic	F1	Precision	Recall
TNFRSF13C AND NOT-C11orf87	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SERTM1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-C3orf80	0.915662651	0.844444444	1
TNFRSF13C AND NOT-IFITM10	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SPINK6	0.915662651	0.844444444	1
TNFRSF13C AND NOT-CTXN1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-ERVFRD-1	0.926829268	0.863636364	1
TNFRSF13C AND NOT-FAM19A1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-MAG	0.915662651	0.844444444	1
TNFRSF13C AND NOT-ARSB	0.948717949	0.925	0.973684211
TNFRSF13C AND NOT-MAS1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-ARSE	0.915662651	0.844444444	1
TNFRSF13C AND NOT-MC2R	0.915662651	0.844444444	1
TNFRSF13C AND NOT-ARSF	0.926829268	0.863636364	1
TNFRSF13C AND NOT-MC4R	0.938271605	0.88372093	1
TNFRSF13C AND NOT-ADAM11	0.915662651	0.844444444	1
TNFRSF13C AND NOT-MLN	0.95	0.904761905	1
TNFRSF13C AND NOT-ASGR1	0.926829268	0.863636364	1
TNFRSF13C AND NOT-MOG	0.915662651	0.844444444	1
TNFRSF13C AND NOT-MPL	0.938271605	0.88372093	1
TNFRSF13C AND NOT-MPZ	0.938271605	0.88372093	1
TNFRSF13C AND NOT-SMCO3	0.915662651	0.844444444	1
TNFRSF13C AND NOT-LRRC52	0.936708861	0.902439024	0.973684211
TNFRSF13C AND NOT-SMIM4	0.915662651	0.844444444	1
TNFRSF13C AND NOT-PTCHD4	0.915662651	0.844444444	1
TNFRSF13C AND NOT-MSMB	0.915662651	0.844444444	1
TNFRSF13C AND NOT-MST1R	0.938271605	0.88372093	1
TNFRSF13C AND NOT-NCAM1	0.938271605	0.88372093	1
TNFRSF13C AND NOT-NDUFB3	0.915662651	0.844444444	1
TNFRSF13C AND NOT-ATP1A2	0.915662651	0.844444444	1
TNFRSF13C AND NOT-ATP1A3	0.95	0.904761905	1
TNFRSF13C AND NOT-ATP1A4	0.95	0.904761905	1
TNFRSF13C AND NOT-NGFR	0.915662651	0.844444444	1
TNFRSF13C AND NOT-NPY	0.948717949	0.925	0.973684211
TNFRSF13C AND NOT-NPPA	0.987012987	0.974358974	1
TNFRSF13C AND NOT-NPR3	0.938271605	0.88372093	1
TNFRSF13C AND NOT-NT5E	0.931506849	0.971428571	0.894736842
TNFRSF13C AND NOT-NTF3	0.915662651	0.844444444	1
TNFRSF13C AND NOT-NTRK1	0.926829268	0.863636364	1
TNFRSF13C AND NOT-ATP2B3	0.915662651	0.844444444	1
TNFRSF13C AND NOT-OPRL1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-OR1F1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-OR3A1	0.938271605	0.88372093	1
TNFRSF13C AND NOT-OR3A2	0.926829268	0.863636364	1
TNFRSF13C AND NOT-GALNT9	0.926829268	0.863636364	1
TNFRSF13C AND NOT-IL22	0.915662651	0.844444444	1
TNFRSF13C AND NOT-CALY	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SLC45A1	0.938271605	0.88372093	1
TNFRSF13C AND NOT-KCNK4	0.915662651	0.844444444	1
TNFRSF13C AND NOT-TAS2R3	0.926829268	0.863636364	1
TNFRSF13C AND NOT-TAS2R16	0.936708861	0.902439024	0.973684211
TNFRSF13C AND NOT-TAS2R13	0.915662651	0.844444444	1
TNFRSF13C AND NOT-TAS2R14	0.915662651	0.844444444	1
TNFRSF13C AND NOT-CDON	0.926829268	0.863636364	1
TNFRSF13C AND NOT-PCDHGC3	0.938271605	0.88372093	1
TNFRSF13C AND NOT-SLC45A2	0.915662651	0.844444444	1
TNFRSF13C AND NOT-DUSP13	0.915662651	0.844444444	1
TNFRSF13C AND NOT-AIG1	0.948717949	0.925	0.973684211
TNFRSF13C AND NOT-NBAS	0.936708861	0.902439024	0.973684211
TNFRSF13C AND NOT-CD244	0.915662651	0.844444444	1
TNFRSF13C AND NOT-ATP8A2	0.915662651	0.844444444	1
TNFRSF13C AND NOT-TUBA8	0.915662651	0.844444444	1
TNFRSF13C AND SLC25A3	0.915662651	0.844444444	1
TNFRSF13C AND NOT-PLA2G2A	0.915662651	0.844444444	1
TNFRSF13C AND NOT-IL17D	0.915662651	0.844444444	1
TNFRSF13C AND NOT-ADAM22	0.915662651	0.844444444	1
TNFRSF13C AND NOT-FXYD4	0.948717949	0.925	0.973684211
TNFRSF13C AND NOT-GPR84	0.926829268	0.863636364	1
TNFRSF13C AND NOT-IL20RA	0.915662651	0.844444444	1
TNFRSF13C AND NOT-IL20RB	0.915662651	0.844444444	1
TNFRSF13C AND NOT-CDHR5	0.915662651	0.844444444	1
TNFRSF13C AND NOT-ATP7B	0.915662651	0.844444444	1
TNFRSF13C AND NOT-GPR88	0.926829268	0.863636364	1
TNFRSF13C AND NOT-TREM2	0.95	0.904761905	1
TNFRSF13C AND NOT-GPR85	0.915662651	0.844444444	1
TNFRSF13C AND NOT-WNT4	0.915662651	0.844444444	1

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
GPM6B AND NEMP1	0.970149254	1	0.942028986
PTPRZ1 AND NEMP1	0.970149254	1	0.942028986
PTPRZ1 AND NOT-FMO5	0.985507246	0.98550725	0.985507246
PTPRZ1 AND SLC9A8	0.985507246	0.98550725	0.985507246
PTPRZ1 AND DPY19L1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND SUN1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NCSTN	0.970149254	1	0.942028986
PTPRZ1 AND NOT-KCNH3	0.99270073	1	0.985507246
PTPRZ1 AND CRB1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-TMED3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-SLC7A8	0.985507246	0.98550725	0.985507246
PTPRZ1 AND GPR161	0.985507246	0.98550725	0.985507246
PTPRZ1 AND SLC35A3	0.99270073	1	0.985507246
PTPRZ1 AND ABCB10	0.99270073	1	0.985507246
PTPRZ1 AND NOT-ABCB9	0.970149254	1	0.942028986
PTPRZ1 AND NOT-ABCA5	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-ICMT	0.985507246	0.98550725	0.985507246
PTPRZ1 AND TRAM1	0.99270073	1	0.985507246
GPM6B AND SEC11A	0.977777778	1	0.956521739
PTPRZ1 AND SEC11A	0.977777778	1	0.956521739
GPM6B AND SEC61G	0.970149254	1	0.942028986
GPM6B AND LEPROTL1	0.977777778	1	0.956521739
PTPRZ1 AND LEPROTL1	0.977777778	1	0.956521739
PTPRZ1 AND NOT-FOLR1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND MACF1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND TMEM131	0.977777778	1	0.956521739
PTPRZ1 AND NOT-LRRC8B	0.985507246	0.98550725	0.985507246
PTPRZ1 AND SLC39A14	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NNT	0.985507246	0.98550725	0.985507246
PTPRZ1 AND MMD	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-SLC16A8	0.985507246	0.98550725	0.985507246
PTPRZ1 AND SEZ6L	0.985507246	0.98550725	0.985507246
PTPRZ1 AND ATP6V0A2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-HYAL4	0.985507246	0.98550725	0.985507246
PTPRZ1 AND TSPAN12	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-TSPAN15	0.985507246	0.98550725	0.985507246
PTPRZ1 AND PIGN	0.985507246	0.98550725	0.985507246
PTPRZ1 AND CLDN14	0.985507246	0.98550725	0.985507246
PTPRZ1 AND FPR1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-VSIG2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-TMEM50A	0.985507246	0.98550725	0.985507246
PTPRZ1 AND LEMD3	0.99270073	1	0.985507246
PTPRZ1 AND NOT-OPN3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND FPR3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND CLEC5A	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-NTSR2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND KCNE5	0.985507246	0.98550725	0.985507246
PTPRZ1 AND CA14	0.985507246	0.98550725	0.985507246
PTPRZ1 AND RABGAP1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND PLD3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-TMEM2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND KCNE4	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-FRRS1L	0.970149254	1	0.942028986
PTPRZ1 AND SDF2L1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND IL17RA	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-FLRT3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-FLRT2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND FLRT1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-FKBP8	0.985507246	0.98550725	0.985507246
PTPRZ1 AND BCL2L13	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-LAMP5	0.985507246	0.98550725	0.985507246
PTPRZ1 AND PANX1	0.99270073	1	0.985507246
PTPRZ1 AND CLDN15	0.985507246	0.98550725	0.985507246
PTPRZ1 AND FJX1	0.977777778	1	0.956521739
PTPRZ1 AND ALOX5AP	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-MS4A6E	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-ABCA4	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-FUT1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-FUT2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-FUT3	0.985294118	1	0.971014493
PTPRZ1 AND FUT4	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-FUT6	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-FNDC5	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-FUT7	0.99270073	1	0.985507246
PTPRZ1 AND HEPACAM2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND TECRL	0.985507246	0.98550725	0.985507246

Antigen Pair Logic	F1	Precision	Recall
TNFRSF13C AND NOT-NLGN3	0.926829268	0.863636364	1
TNFRSF13C AND NOT-PON3	0.915662651	0.844444444	1
TNFRSF13C AND NOT-FAM105A	0.915662651	0.844444444	1
TNFRSF13C AND NOT-EQTN	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SLC6A20	0.926829268	0.863636364	1
TNFRSF13C AND NOT-LY6K	0.938271605	0.88372093	1
TNFRSF13C AND NOT-FAM20A	0.926829268	0.863636364	1
TNFRSF13C AND NOT-DCHS2	0.915662651	0.844444444	1
TNFRSF13C AND NOT-PQLC2	0.962025316	0.926829268	1
TNFRSF13C AND NOT-HRASLS2	0.926829268	0.863636364	1
TNFRSF13C AND NOT-ACPP	0.938271605	0.88372093	1
TNFRSF13C AND NOT-TMEM51	0.938271605	0.88372093	1
TNFRSF13C AND NOT-SLC6A15	0.915662651	0.844444444	1
TNFRSF13C AND NOT-EVA1B	0.926829268	0.863636364	1
TNFRSF13C AND NOT-KIRREL	0.938271605	0.88372093	1
TNFRSF13C AND NOT-TMEM40	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SLC35C1	0.987012987	0.974358974	1
TNFRSF13C AND NOT-VNN3	0.915662651	0.844444444	1
TNFRSF13C AND NOT-STYK1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SVOP	0.915662651	0.844444444	1
TNFRSF13C AND NOT-POMGNT1	0.938271605	0.88372093	1
TNFRSF13C AND NOT-FAR2	0.915662651	0.844444444	1
TNFRSF13C AND NOT-TENM3	0.926829268	0.863636364	1
TNFRSF13C AND NOT-TEX2	0.938271605	0.88372093	1
TNFRSF13C AND NOT-GABRQ	0.915662651	0.844444444	1
TNFRSF13C AND NOT-ACSS2	0.926829268	0.863636364	1
TNFRSF13C AND NOT-AJAP1	0.961038961	0.948717949	0.973684211
TNFRSF13C AND NOT-PCDHGA11	0.915662651	0.844444444	1
TNFRSF13C AND NOT-PCDHB12	0.915662651	0.844444444	1
TNFRSF13C AND NOT-PCDHA10	0.938271605	0.88372093	1
TNFRSF13C AND NOT-SLC7A10	0.95	0.904761905	1
TNFRSF13C AND NOT-PRRG2	0.926829268	0.863636364	1
TNFRSF13C AND NOT-TMPRSS4	0.962025316	0.926829268	1
TNFRSF13C AND NOT-NDUFA4L2	0.95	0.904761905	1
TNFRSF13C AND NOT-OTOR	0.948717949	0.925	0.973684211
TNFRSF13C AND NOT-C2orf83	0.915662651	0.844444444	1
TNFRSF13C AND NOT-CEACAM19	0.915662651	0.844444444	1
TNFRSF13C AND NOT-CHRNA10	0.915662651	0.844444444	1
TNFRSF13C AND NOT-ENTPD7	0.95	0.904761905	1
TNFRSF13C AND NOT-ANO2	0.936708861	0.902439024	0.973684211
TNFRSF13C AND NOT-TMEM63C	0.915662651	0.844444444	1
TNFRSF13C AND NOT-JPH2	0.915662651	0.844444444	1
TNFRSF13C AND NOT-GJC2	0.915662651	0.844444444	1
TNFRSF13C AND NOT-PTAFR	0.962025316	0.926829268	1
TNFRSF13C AND NOT-TAS2R38	0.915662651	0.844444444	1
TNFRSF13C AND NOT-PTGDR	0.915662651	0.844444444	1
TNFRSF13C AND NOT-PTGER1	0.948717949	0.925	0.973684211
TNFRSF13C AND NOT-PTGER3	0.926829268	0.863636364	1
TNFRSF13C AND NOT-JPH3	0.915662651	0.844444444	1
TNFRSF13C AND NOT-PTGFR	0.926829268	0.863636364	1
TNFRSF13C AND NOT-PTGIS	0.915662651	0.844444444	1
TNFRSF13C AND NOT-KIAA1324	0.915662651	0.844444444	1
TNFRSF13C AND NOT-PCDH10	0.915662651	0.844444444	1
TNFRSF13C AND NOT-ISLR2	0.962025316	0.926829268	1
TNFRSF13C AND NOT-DPP10	0.915662651	0.844444444	1
TNFRSF13C AND NOT-KIAA1549	0.915662651	0.844444444	1
TNFRSF13C AND NOT-BRINP2	0.915662651	0.844444444	1
TNFRSF13C AND NOT-G6PC2	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SLC4A5	0.926829268	0.863636364	1
TNFRSF13C AND NOT-CADM3	0.915662651	0.844444444	1
TNFRSF13C AND NOT-PTPRH	0.926829268	0.863636364	1
TNFRSF13C AND NOT-PVR	0.926829268	0.863636364	1
TNFRSF13C AND NOT-DNASE2B	0.915662651	0.844444444	1
TNFRSF13C AND NOT-BCHE	0.915662651	0.844444444	1
TNFRSF13C AND NOT-ACE2	0.915662651	0.844444444	1
TNFRSF13C AND NOT-CACNG8	0.926829268	0.863636364	1
TNFRSF13C AND NOT-CACNG6	0.915662651	0.844444444	1
TNFRSF13C AND NOT-HRH4	0.915662651	0.844444444	1
TNFRSF13C AND NOT-LGR6	0.915662651	0.844444444	1
TNFRSF13C AND NOT-PRPH2	0.938271605	0.88372093	1
TNFRSF13C AND NOT-RHAG	0.95	0.904761905	1
TNFRSF13C AND NOT-RHCE	0.973684211	0.973684211	0.973684211
TNFRSF13C AND NOT-RHO	0.962025316	0.926829268	1
TNFRSF13C AND NOT-EDA2R	0.915662651	0.844444444	1
TNFRSF13C AND NOT-CDH26	0.938271605	0.88372093	1
TNFRSF13C AND NOT-SLC5A7	0.915662651	0.844444444	1

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
PTPRZ1 AND NOT-FUT8	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-EPHX4	0.985507246	0.98550725	0.985507246
PTPRZ1 AND KDSR	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-ACKR1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND LCLAT1	0.970149254	1	0.942028986
PTPRZ1 AND FZD2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND CERS6	0.99270073	1	0.985507246
PTPRZ1 AND NOT-IFI6	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-G6PC	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-FAM26E	0.985294118	1	0.971014493
PTPRZ1 AND ZDHHC24	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-NAALADL2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND TMEM256	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-ZDHHC23	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-TMEM86B	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-GABBR1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-CCDC108	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-TMCO4	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-MCOLN2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-GABRA2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND INAFM1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-GABRA5	0.985507246	0.98550725	0.985507246
PTPRZ1 AND CNEP1R1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND SYT14	0.985507246	0.98550725	0.985507246
PTPRZ1 AND GABRB1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-TMEM196	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-CDRT15L2	0.99270073	1	0.985507246
PTPRZ1 AND NOT-STEAP1B	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-GABRB3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-ST6GALNAC3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-MFSD8	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-GABRG1	0.970149254	1	0.942028986
PTPRZ1 AND NOT-GABRG2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND GABRG3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-NPB	0.985507246	0.98550725	0.985507246
PTPRZ1 AND SERINC5	0.985507246	0.98550725	0.985507246
PTPRZ1 AND FRMD3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-C1orf101	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-GABRR2	0.977777778	1	0.956521739
PTPRZ1 AND NOT-UTS2B	0.985507246	0.98550725	0.985507246
PTPRZ1 AND TMEM59L	0.985507246	0.98550725	0.985507246
GPM6B AND BRI3	0.977777778	1	0.956521739
PTPRZ1 AND BRI3	0.985294118	1	0.971014493
PTPRZ1 AND SLC39A6	0.970149254	1	0.942028986
PTPRZ1 AND BAMBI	0.985507246	0.98550725	0.985507246
PTPRZ1 AND RHBDD3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-KLK5	0.985294118	1	0.971014493
PTPRZ1 AND NOT-GALC	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-TPSG1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-TMEM184B	0.985507246	0.98550725	0.985507246
PTPRZ1 AND METTL7A	0.985507246	0.98550725	0.985507246
PTPRZ1 AND YIPF3	0.970149254	1	0.942028986
PTPRZ1 AND NOT-ABHD14A	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-LETMD1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND GALR1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND TMEM186	0.985507246	0.98550725	0.985507246
PTPRZ1 AND RNF19A	0.985507246	0.98550725	0.985507246
GPM6B AND GALNT1	0.99270073	1	0.985507246
PTPRZ1 AND GALNT1	0.99270073	1	0.985507246
PTPRZ1 AND OLFML2B	0.985507246	0.98550725	0.985507246
PTPRZ1 AND TMEM158	0.985507246	0.98550725	0.985507246
PTPRZ1 AND C1orf43	0.970149254	1	0.942028986
PTPRZ1 AND NOT-LY6G6F	0.985507246	0.98550725	0.985507246
GPM6B AND ZDHHC5	0.985294118	1	0.971014493
PTPRZ1 AND ZDHHC5	0.985294118	1	0.971014493
PTPRZ1 AND NOT-NALCN	0.985507246	0.98550725	0.985507246
PTPRZ1 AND TMIE	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-WFDC11	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-WFDC9	0.985294118	1	0.971014493
PTPRZ1 AND NOT-MRGPRX1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-TAS2R39	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-TAS2R41	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-TAS2R50	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-FAM205A	0.99270073	1	0.985507246
PTPRZ1 AND PVRL3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-ADGRA2	0.985507246	0.98550725	0.985507246

Antigen Pair Logic	F1	Precision	Recall
TNFRSF13C AND NOT-CTXN3	0.926829268	0.863636364	1
TNFRSF13C AND NOT-RTN2	0.938271605	0.88372093	1
TNFRSF13C AND NOT-BDNF	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SCN1A	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SCN1B	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SCN2B	0.926829268	0.863636364	1
TNFRSF13C AND NOT-SCN3A	0.935064935	0.923076923	0.947368421
TNFRSF13C AND NOT-SCN8A	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SCN9A	0.915662651	0.844444444	1
TNFRSF13C AND NOT-CDH23	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SLC39A8	0.936708861	0.902439024	0.973684211
TNFRSF13C AND NOT-MS4A5	0.915662651	0.844444444	1
TNFRSF13C AND NOT-ABCG8	0.926829268	0.863636364	1
TNFRSF13C AND NOT-CHST8	0.938271605	0.88372093	1
TNFRSF13C AND NOT-SFTPC	0.926829268	0.863636364	1
TNFRSF13C AND NOT-PIRT	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SGCD	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SMIM10	0.915662651	0.844444444	1
TNFRSF13C AND NOT-CSMD1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-NDST4	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SHH	0.936708861	0.902439024	0.973684211
TNFRSF13C AND NOT-HRCT1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-MARC1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-P2RY12	0.915662651	0.844444444	1
TNFRSF13C AND NOT-IL25	0.948717949	0.925	0.973684211
TNFRSF13C AND NOT-ST3GAL4	0.915662651	0.844444444	1
TNFRSF13C AND NOT-ST3GAL3	0.936708861	0.902439024	0.973684211
TNFRSF13C AND NOT-PCDH20	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SLC1A6	0.926829268	0.863636364	1
TNFRSF13C AND NOT-SLC2A4	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SLC5A2	0.95	0.904761905	1
TNFRSF13C AND NOT-SLC6A2	0.918918919	0.944444444	0.894736842
TNFRSF13C AND NOT-SLC6A4	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SLC6A8	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SLC6A9	0.926829268	0.863636364	1
TNFRSF13C AND NOT-SLC6A11	0.938271605	0.88372093	1
TNFRSF13C AND NOT-SLC8A2	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SLC7A4	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SLC8A1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SLC13A1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SLC16A2	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SLC18A2	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SLC22A1	0.926829268	0.863636364	1
TNFRSF13C AND NOT-SLC22A2	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SLC22A4	0.915662651	0.844444444	1
TNFRSF13C AND NOT-DLK2	0.915662651	0.844444444	1
TNFRSF13C AND NOT-CYP4F12	0.938271605	0.88372093	1
TNFRSF13C AND NOT-SMPD1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SSTR1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SSTR2	0.974358974	0.95	1
TNFRSF13C AND NOT-BST1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SYT4	0.936708861	0.902439024	0.973684211
TNFRSF13C AND NOT-SYT5	0.915662651	0.844444444	1
TNFRSF13C AND NOT-TACR3	0.926829268	0.863636364	1
TNFRSF13C AND NOT-TFR2	0.926829268	0.863636364	1
TNFRSF13C AND NOT-TSPAN8	0.915662651	0.844444444	1
TNFRSF13C AND NOT-TRPC4	0.915662651	0.844444444	1
TNFRSF13C AND NOT-TSHR	0.926829268	0.863636364	1
TNFRSF13C AND NOT-CLRN1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-VIPR1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-ZAN	0.915662651	0.844444444	1
TNFRSF13C AND NOT-WNT7B	0.926829268	0.863636364	1
TNFRSF13C AND NOT-CA12	0.926829268	0.863636364	1
TNFRSF13C AND NOT-CACNA1C	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SLC30A4	0.936708861	0.902439024	0.973684211
TNFRSF13C AND NOT-SEMA3B	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SMIM2	0.915662651	0.844444444	1
TNFRSF13C AND NOT-TRPM8	0.915662651	0.844444444	1
TNFRSF13C AND NOT-PRRG4	0.915662651	0.844444444	1
TNFRSF13C AND NOT-ELOVL6	0.962025316	0.926829268	1
TNFRSF13C AND NOT-ALG12	0.938271605	0.88372093	1
TNFRSF13C AND NOT-DHRS11	0.936708861	0.902439024	0.973684211
TNFRSF13C AND NOT-OR2H2	0.915662651	0.844444444	1
TNFRSF13C AND NOT-OR51B2	0.938271605	0.88372093	1
TNFRSF13C AND NOT-NOX5	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SRD5A3	0.926829268	0.863636364	1

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
PTPRZ1 AND NOT-C2CD2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND UNC50	0.985507246	0.98550725	0.985507246
PTPRZ1 AND DHRS7B	0.970149254	1	0.942028986
PTPRZ1 AND RNF167	0.985507246	0.98550725	0.985507246
PTPRZ1 AND TENM4	0.985507246	0.98550725	0.985507246
PTPRZ1 AND LRP10	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-CYP4X1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND SUSD5	0.985507246	0.98550725	0.985507246
PTPRZ1 AND ATRNL1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-CFAP61	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-HERC4	0.985507246	0.98550725	0.985507246
GPM6B AND TCTN3	0.970149254	1	0.942028986
PTPRZ1 AND TCTN3	0.977777778	1	0.956521739
PTPRZ1 AND LRRC32	0.985507246	0.98550725	0.985507246
PTPRZ1 AND PCDHB5	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-STEAP2	0.985507246	0.98550725	0.985507246
GPM6B AND TMEM251	0.977777778	1	0.956521739
PTPRZ1 AND TMEM251	0.977777778	1	0.956521739
PTPRZ1 AND OR1C1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND OR1A2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND OR2B6	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-OR1J4	0.985294118	1	0.971014493
PTPRZ1 AND B3GAT3	0.99270073	1	0.985507246
PTPRZ1 AND NOT-OR2M4	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-OR2L2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND CLEC4E	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-PLA2G2D	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-IL1RAPL2	0.977777778	1	0.956521739
CHST10 AND NOT-IL1RAPL2	0.970149254	1	0.942028986
PTPRZ1 AND GBGT1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-OR7A17	0.985507246	0.98550725	0.985507246
PTPRZ1 AND SEZ6L2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND OR10J1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-OR8B8	0.970149254	1	0.942028986
PTPRZ1 AND NOT-OR8G1	0.977777778	1	0.956521739
PTPRZ1 AND NOT-OR10A3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND SLC17A5	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-CNNM4	0.985507246	0.98550725	0.985507246
PTPRZ1 AND CNNM3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND MYOF	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-GCNT1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND CHIC2	0.977777778	1	0.956521739
PTPRZ1 AND NOT-GCNT2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-TSPAN16	0.985294118	1	0.971014493
PTPRZ1 AND NOT-OR11A1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-GREM1	0.970149254	1	0.942028986
PTPRZ1 AND NOT-OR8B2	0.977777778	1	0.956521739
PTPRZ1 AND TBL2	0.99270073	1	0.985507246
PTPRZ1 AND NOT-OR7C2	0.985294118	1	0.971014493
PTPRZ1 AND NOT-BEST4	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-OR4D1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-ADGRF1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-OR2J2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-OR2H1	0.99270073	1	0.985507246
PTPRZ1 AND GGCX	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-B4GALT1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-STEAP1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND GGT5	0.970149254	1	0.942028986
PTPRZ1 AND NOT-GIPR	0.985507246	0.98550725	0.985507246
PTPRZ1 AND GJA1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND GPR160	0.985507246	0.98550725	0.985507246
PTPRZ1 AND GJA3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-KCNV1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-GJA4	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-NPTN	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-GJA5	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-ATP2C1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND SIGLEC7	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-GJA8	0.977777778	1	0.956521739
PTPRZ1 AND NOT-GJB1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-NSG1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND GHITM	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-GJB2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-TSPAN13	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-GJB3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND B3GAT1	0.985507246	0.98550725	0.985507246

Antigen Pair Logic	F1	Precision	Recall
TNFRSF13C AND NOT-TMC5	0.915662651	0.844444444	1
TNFRSF13C AND NOT-ZDHHC11	0.962025316	0.926829268	1
TNFRSF13C AND NOT-TM4SF20	0.948717949	0.925	0.973684211
TNFRSF13C AND NOT-THSD4	0.915662651	0.844444444	1
TNFRSF13C AND NOT-CALCR	0.915662651	0.844444444	1
TNFRSF13C AND NOT-TMC7	0.938271605	0.88372093	1
TNFRSF13C AND NOT-DNAJC22	0.938271605	0.88372093	1
TNFRSF13C AND NOT-NRSN2	0.915662651	0.844444444	1
TNFRSF13C AND NOT-TRPM3	0.948717949	0.925	0.973684211
TNFRSF13C AND NOT-LRRC8E	0.915662651	0.844444444	1
TNFRSF13C AND NOT-FRAS1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-TMEM254	0.961038961	0.948717949	0.973684211
TNFRSF13C AND NOT-ULBP2	0.915662651	0.844444444	1
TNFRSF13C AND NOT-BPIFB2	0.938271605	0.88372093	1
TNFRSF13C AND NOT-PDCD1LG2	0.915662651	0.844444444	1
TNFRSF13C AND NOT-C6orf25	0.987012987	0.974358974	1
TNFRSF13C AND NOT-TMEM121	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SPX	0.915662651	0.844444444	1
TNFRSF13C AND NOT-APOL4	0.926829268	0.863636364	1
TNFRSF13C AND NOT-TAS1R1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-KCNH6	0.947368421	0.947368421	0.947368421
TNFRSF13C AND NOT-ST8SIA2	0.915662651	0.844444444	1
TNFRSF13C AND NOT-GPR63	0.915662651	0.844444444	1
TNFRSF13C AND NOT-AMN	0.926829268	0.863636364	1
TNFRSF13C AND NOT-OR5V1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-OR2B2	0.915662651	0.844444444	1
TNFRSF13C AND NOT-VANGL1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-ELOVL3	0.926829268	0.863636364	1
TNFRSF13C AND NOT-GSG1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SLC4A9	0.915662651	0.844444444	1
TNFRSF13C AND NOT-CALN1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-FAM57B	0.96	0.972972973	0.947368421
TNFRSF13C AND NOT-SLC25A18	0.961038961	0.948717949	0.973684211
TNFRSF13C AND NOT-GPR61	0.962025316	0.926829268	1
TNFRSF13C AND NOT-SLC25A2	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SPATA9	0.938271605	0.88372093	1
TNFRSF13C AND NOT-OR1G1	0.926829268	0.863636364	1
TNFRSF13C AND NOT-B3GNT5	0.95	0.904761905	1
TNFRSF13C AND NOT-ADGRV1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-QRFPR	0.915662651	0.844444444	1
TNFRSF13C AND NOT-EVA1A	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SLC37A3	0.936708861	0.902439024	0.973684211
TNFRSF13C AND NOT-TMEM246	0.915662651	0.844444444	1
TNFRSF13C AND NOT-TMEM101	0.931506849	0.971428571	0.894736842
TNFRSF13C AND NOT-MEGF11	0.962025316	0.926829268	1
TNFRSF13C AND NOT-CASR	0.926829268	0.863636364	1
TNFRSF13C AND NOT-ST6GAL2	0.915662651	0.844444444	1
TNFRSF13C AND NOT-DGAT2	0.96	0.972972973	0.947368421
TNFRSF13C AND NOT-ABHD1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-FUT10	0.936708861	0.902439024	0.973684211
TNFRSF13C AND NOT-AGPAT9	0.926829268	0.863636364	1
TNFRSF13C AND NOT-GALR3	0.915662651	0.844444444	1
TNFRSF13C AND NOT-TMEM25	0.915662651	0.844444444	1
TNFRSF13C AND NOT-PLXDC2	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SLC43A1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-ITGA10	0.915662651	0.844444444	1
TNFRSF13C AND NOT-CHST1	1	1	1
TNFRSF13C AND NOT-CNTNAP4	0.915662651	0.844444444	1
TNFRSF13C AND NOT-UNC5C	0.915662651	0.844444444	1
TNFRSF13C AND NOT-KCNK5	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SLC4A4	0.915662651	0.844444444	1
TNFRSF13C AND NOT-TNFSF14	0.915662651	0.844444444	1
TNFRSF13C AND NOT-ADAM23	0.926829268	0.863636364	1
TNFRSF13C AND NOT-ADAM21	0.915662651	0.844444444	1
TNFRSF13C AND NOT-TNFRSF10C	0.915662651	0.844444444	1
TNFRSF13C AND NOT-IL18RAP	0.915662651	0.844444444	1
TNFRSF13C AND NOT-GALR2	0.926829268	0.863636364	1
TNFRSF13C AND NOT-INPP4B	0.915662651	0.844444444	1
TNFRSF13C AND NOT-CCKAR	0.915662651	0.844444444	1
TNFRSF13C AND NOT-CACNA1I	0.938271605	0.88372093	1
TNFRSF13C AND NOT-P4HA2	0.915662651	0.844444444	1
TNFRSF13C AND NOT-PPP1R3F	0.915662651	0.844444444	1
TNFRSF13C AND NOT-KCNH7	0.948717949	0.925	0.973684211
TNFRSF13C AND NOT-UCN2	0.915662651	0.844444444	1
TNFRSF13C AND NOT-UNC5A	0.974358974	0.95	1
TNFRSF13C AND NOT-NAT8	0.938271605	0.88372093	1

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
PTPRZ1 AND ST6GALNAC4	0.985507246	0.98550725	0.985507246
PTPRZ1 AND CACNG4	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-GJB5	0.985507246	0.98550725	0.985507246
PTPRZ1 AND SLC39A1	0.985294118	1	0.971014493
PTPRZ1 AND NOT-SIGLEC9	0.985507246	0.98550725	0.985507246
PTPRZ1 AND SIGLEC8	0.985507246	0.98550725	0.985507246
PTPRZ1 AND GPR82	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-HCAR1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-C5AR2	0.985507246	0.98550725	0.985507246
GPM6B AND SERP1	0.985294118	1	0.971014493
PTPRZ1 AND SERP1	0.985294118	1	0.971014493
PTPRZ1 AND COQ2	0.985294118	1	0.971014493
PTPRZ1 AND NOT-GPR162	0.985507246	0.98550725	0.985507246
PTPRZ1 AND TNFRSF21	0.977777778	1	0.956521739
PTPRZ1 AND PCDH17	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-GCLC	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-PGAP2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-PCDH11X	0.99270073	1	0.985507246
PTPRZ1 AND GOLIM4	0.985507246	0.98550725	0.985507246
PTPRZ1 AND PCSK1N	0.985507246	0.98550725	0.985507246
PTPRZ1 AND KCNMB4	0.985507246	0.98550725	0.985507246
GPM6B AND TOR1B	0.970149254	1	0.942028986
PTPRZ1 AND TOR1B	0.985294118	1	0.971014493
PTPRZ1 AND GLG1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-GLP1R	0.99270073	1	0.985507246
PTPRZ1 AND NOT-GLRA1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND GLRA2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-TOR2A	0.985507246	0.98550725	0.985507246
PTPRZ1 AND CECR6	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-GLRB	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-PCLO	0.985507246	0.98550725	0.985507246
CHST10 AND NOT-PCLO	0.970149254	1	0.942028986
PTPRZ1 AND NOT-GNRHR	0.985507246	0.98550725	0.985507246
PTPRZ1 AND GNS	0.99270073	1	0.985507246
PTPRZ1 AND GOLGB1	0.970149254	1	0.942028986
ITGAV AND NOT-GP1BA	0.977777778	1	0.956521739
PTPRZ1 AND NOT-GP1BA	0.985294118	1	0.971014493
PTPRZ1 AND NOT-GP2	0.985294118	1	0.971014493
PTPRZ1 AND NOT-GPD2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND SLCO1B3	0.985507246	0.98550725	0.985507246
GPM6A AND SPCS1	0.970149254	1	0.942028986
GPM6A AND ATRAID	0.985294118	1	0.971014493
GPM6A AND TMEM14C	0.970149254	1	0.942028986
GPM6A AND ATP5F1	0.985294118	1	0.971014493
GPM6A AND SLC38A2	0.977777778	1	0.956521739
PTPRZ1 AND GPM6A	0.985507246	0.98550725	0.985507246
GPM6A AND NOT-UGT2B15	0.970149254	1	0.942028986
GPM6A AND PTTG1IP	0.985294118	1	0.971014493
GPM6A AND ATP6V0E1	0.970149254	1	0.942028986
GPM6A AND CD63	0.99270073	1	0.985507246
GPM6A AND LAPTM4A	0.99270073	1	0.985507246
GPM6B AND SPCS1	0.970149254	1	0.942028986
GPM6B AND TMEM208	0.985294118	1	0.971014493
GPM6B AND ALG5	0.970149254	1	0.942028986
GPM6B AND SEC61A1	0.970149254	1	0.942028986
GPM6B AND ITGAE	0.985294118	1	0.971014493
GPM6B AND SMIM20	0.970149254	1	0.942028986
GPM6B AND MFAP3	0.977777778	1	0.956521739
GPM6B AND OXA1L	0.99270073	1	0.985507246
GPM6B AND TMED5	0.985294118	1	0.971014493
GPM6B AND DERL2	0.977777778	1	0.956521739
GPM6B AND GOLT1B	0.970149254	1	0.942028986
GPM6B AND IER3IP1	0.985294118	1	0.971014493
GPM6B AND TMEM69	0.985294118	1	0.971014493
GPM6B AND WDR83OS	0.985294118	1	0.971014493
GPM6B AND UBE2J1	0.977777778	1	0.956521739
GPM6B AND TMEM138	0.985294118	1	0.971014493
GPM6B AND CUTA	0.970149254	1	0.942028986
GPM6B AND ATP5F1	0.985294118	1	0.971014493
GPM6B AND PIGC	0.977777778	1	0.956521739
GPM6B AND PIGF	0.977777778	1	0.956521739
GPM6B AND ACP1	0.977777778	1	0.956521739
GPM6B AND STX18	0.970149254	1	0.942028986
GPM6B AND TMCO1	0.970149254	1	0.942028986
GPM6B AND IMPAD1	0.970149254	1	0.942028986
GPM6B AND PRCP	0.970149254	1	0.942028986

Antigen Pair Logic	F1	Precision	Recall
TNFRSF13C AND NOT-RHBDL1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-CCRL2	0.931506849	0.971428571	0.894736842
TNFRSF13C AND NOT-FER1L5	0.938271605	0.88372093	1
TNFRSF13C AND NOT-GPRC5A	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SLC13A2	0.926829268	0.863636364	1
TNFRSF13C AND NOT-CLDN10	0.95	0.904761905	1
TNFRSF13C AND NOT-CLDN6	0.962025316	0.926829268	1
TNFRSF13C AND NOT-CLDN2	0.915662651	0.844444444	1
TNFRSF13C AND NOT-CLDN9	0.915662651	0.844444444	1
TNFRSF13C AND NOT-PIGQ	0.962025316	0.926829268	1
TNFRSF13C AND NOT-SLC6A5	0.944444444	1	0.894736842
TNFRSF13C AND NOT-SLC28A2	0.915662651	0.844444444	1
TNFRSF13C AND NOT-HTR3B	0.973684211	0.973684211	0.973684211
TNFRSF13C AND NOT-MARVELD3	0.915662651	0.844444444	1
TNFRSF13C AND NOT-DSEL	0.915662651	0.844444444	1
TNFRSF13C AND NOT-CDHR1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-TMEM169	0.915662651	0.844444444	1
TNFRSF13C AND NOT-REEP6	0.938271605	0.88372093	1
TNFRSF13C AND NOT-TAAR2	0.915662651	0.844444444	1
TNFRSF13C AND NOT-GPR55	0.936708861	0.902439024	0.973684211
TNFRSF13C AND NOT-OPALIN	0.915662651	0.844444444	1
TNFRSF13C AND NOT-MGAM2	0.95	0.904761905	1
TNFRSF13C AND NOT-NDST3	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SLC22A6	0.938271605	0.88372093	1
TNFRSF13C AND NOT-SLC22A13	0.948717949	0.925	0.973684211
TNFRSF13C AND NOT-CD101	0.938271605	0.88372093	1
TNFRSF13C AND NOT-LRRC4B	0.938271605	0.88372093	1
TNFRSF13C AND NOT-ABCC12	0.915662651	0.844444444	1
TNFRSF13C AND NOT-NTN1	0.938271605	0.88372093	1
TNFRSF13C AND NOT-KCNK6	0.938271605	0.88372093	1
TNFRSF13C AND NOT-ABCG2	0.95	0.904761905	1
TNFRSF13C AND NOT-TNFRSF8	0.938271605	0.88372093	1
TNFRSF13C AND NOT-NCR1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-SLC4A8	0.915662651	0.844444444	1
TNFRSF13C AND NOT-ENTPD2	0.938271605	0.88372093	1
TNFRSF13C AND NOT-IGDCC3	0.938271605	0.88372093	1
TNFRSF13C AND HERPUD1	0.915662651	0.844444444	1
TNFRSF13C AND NOT-ECE2	0.915662651	0.844444444	1
TNFRSF13C AND NOT-MFAP3L	0.915662651	0.844444444	1
TNFRSF13C AND NOT-KIAA0319	0.915662651	0.844444444	1
TNFRSF13C AND NOT-LPPR4	0.915662651	0.844444444	1
TNFRSF13C AND NOT-HS3ST2	0.915662651	0.844444444	1
TNFRSF13C AND NOT-FGF19	0.915662651	0.844444444	1
CLECL1 AND NOT-EVI5L	0.938271605	0.88372093	1
QSOX2 AND NOT-EVI5L	0.948717949	0.925	0.973684211
NOT-FAM210B AND SPPL3	0.936708861	0.902439024	0.973684211
CLECL1 AND NOT-FAM210B	0.962025316	0.926829268	1
QSOX2 AND NOT-FAM210B	0.936708861	0.902439024	0.973684211
NOT-FAM210B AND ETV6	0.948717949	0.925	0.973684211
NOT-FAM210B AND ST6GALNAC4	0.947368421	0.947368421	0.947368421
NOT-FAM210B AND IL10RA	0.936708861	0.902439024	0.973684211
NOT-FAM210B AND APH1A	0.948717949	0.925	0.973684211
NOT-FAM210B AND SLC37A1	0.948717949	0.925	0.973684211
NOT-FAM210B AND TRPM7	0.948717949	0.925	0.973684211
NOT-FAM210B AND ERGIC1	0.947368421	0.947368421	0.947368421
NOT-FAM210B AND SPG7	0.945945946	0.972222222	0.921052632
NOT-FAM210B AND ST14	0.918918919	0.944444444	0.894736842
NOT-FAM210B AND ADAM17	0.961038961	0.948717949	0.973684211
ORAI2 AND NOT-FAM210B	0.936708861	0.902439024	0.973684211
NOT-FAM210B AND SNN	0.947368421	0.947368421	0.947368421
NOT-FAM210B AND CELSR1	0.936708861	0.902439024	0.973684211
CLECL1 AND NOT-CMTM5	0.915662651	0.844444444	1
CLECL1 AND NOT-MOGAT1	0.948717949	0.925	0.973684211
QSOX2 AND NOT-MOGAT1	0.947368421	0.947368421	0.947368421
CLECL1 AND NOT-LYPD1	0.938271605	0.88372093	1
QSOX2 AND NOT-MAS1L	0.948717949	0.925	0.973684211
CLECL1 AND NOT-CATSPER1	0.926829268	0.863636364	1
QSOX2 AND NOT-CATSPER1	0.948717949	0.925	0.973684211
CLECL1 AND NOT-SLC16A10	0.948717949	0.925	0.973684211
QSOX2 AND NOT-UBE2J2	0.961038961	0.948717949	0.973684211
NOT-ELOVL6 AND CLCN6	0.948717949	0.925	0.973684211
CLECL1 AND NOT-FAM24A	0.926829268	0.863636364	1
QSOX2 AND NOT-FAM24A	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-GYLTL1B	0.935064935	0.923076923	0.947368421
CLECL1 AND NOT-AMICA1	0.926829268	0.863636364	1
NOT-ADSSL1 AND CELSR1	0.962025316	0.926829268	1

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
GPM6B AND PARL	0.977777778	1	0.956521739
GPM6B AND ETNK1	0.970149254	1	0.942028986
GPM6B AND SLC30A6	0.977777778	1	0.956521739
GPM6B AND VIMP	0.985294118	1	0.971014493
GPM6B AND EMC3	0.977777778	1	0.956521739
GPM6B AND PLGRKT	0.970149254	1	0.942028986
GPM6B AND DIABLO	0.99270073	1	0.985507246
GPM6B AND C5orf15	0.970149254	1	0.942028986
GPM6B AND MAVS	0.970149254	1	0.942028986
PTPRZ1 AND GPM6B	0.985507246	0.98550725	0.985507246
GPM6B AND PEX2	0.970149254	1	0.942028986
GPM6B AND SELK	0.977777778	1	0.956521739
GPM6B AND SRR	0.977777778	1	0.956521739
GPM6B AND C5orf28	0.970149254	1	0.942028986
GPM6B AND SLC30A5	0.985294118	1	0.971014493
GPM6B AND BNIP2	0.985507246	0.98550725	0.985507246
GPM6B AND SOAT1	0.970149254	1	0.942028986
GPM6B AND SSR2	0.970149254	1	0.942028986
GPM6B AND HSPA13	0.99270073	1	0.985507246
GPM6B AND TMBIM6	0.970149254	1	0.942028986
GPM6B AND PTTG1IP	0.985294118	1	0.971014493
GPM6B AND ZNF7	0.970149254	1	0.942028986
GPM6B AND ALG8	0.985294118	1	0.971014493
GPM6B AND DERL1	0.977777778	1	0.956521739
GPM6B AND TTC13	0.970149254	1	0.942028986
GPM6B AND KIAA1715	0.970149254	1	0.942028986
GPM6B AND CANX	0.970149254	1	0.942028986
GPM6B AND SPNS1	0.985294118	1	0.971014493
GPM6B AND TMEM126A	0.970149254	1	0.942028986
GPM6B AND SLC37A3	0.970149254	1	0.942028986
GPM6B AND JAGN1	0.977777778	1	0.956521739
GPM6B AND ZNF559	0.977777778	1	0.956521739
GPM6B AND ATP6V0E1	0.970149254	1	0.942028986
GPM6B AND NOT-CD1B	0.977777778	1	0.956521739
GPM6B AND XPR1	0.970149254	1	0.942028986
GPM6B AND ORMDL1	0.970149254	1	0.942028986
GPM6B AND TMEM203	0.985294118	1	0.971014493
GPM6B AND CDYL	0.970149254	1	0.942028986
GPM6B AND NOT-SIGLEC6	0.985507246	0.98550725	0.985507246
GPM6B AND STX8	0.970149254	1	0.942028986
PTPRZ1 AND IFNL2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-IFNL1	0.970149254	1	0.942028986
PTPRZ1 AND NOT-OR51B5	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-OR5J2	0.99270073	1	0.985507246
PTPRZ1 AND NOT-GPR3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-XCR1	0.985294118	1	0.971014493
PTPRZ1 AND NOT-SLC25A45	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-OR8D1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-OR8D2	0.99270073	1	0.985507246
PTPRZ1 AND NPBWR1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND TMEM80	0.985507246	0.98550725	0.985507246
PTPRZ1 AND SLC22A24	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-OR10A4	0.985507246	0.98550725	0.985507246
PTPRZ1 AND CD163L1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-B4GALNT3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-CXCR3	0.970149254	1	0.942028986
PTPRZ1 AND DPY19L2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND GXYLT1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-TMPRSS12	0.99270073	1	0.985507246
PTPRZ1 AND PRLHR	0.985507246	0.98550725	0.985507246
PTPRZ1 AND GPR137C	0.985507246	0.98550725	0.985507246
PTPRZ1 AND ZDHHC22	0.985507246	0.98550725	0.985507246
PTPRZ1 AND UTS2R	0.985507246	0.98550725	0.985507246
PTPRZ1 AND C16orf91	0.985294118	1	0.971014493
PTPRZ1 AND NOT-TMEM235	0.977777778	1	0.956521739
PTPRZ1 AND NOT-ANG	0.985507246	0.98550725	0.985507246
PTPRZ1 AND MILR1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-C17orf78	0.99270073	1	0.985507246
PTPRZ1 AND GPR17	0.985507246	0.98550725	0.985507246
PTPRZ1 AND SLC26A11	0.977777778	1	0.956521739
PTPRZ1 AND GDPD1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-TMEM105	0.985507246	0.98550725	0.985507246
ITGAV AND NOT-GPR18	0.977777778	1	0.956521739
PTPRZ1 AND NOT-GPR18	0.985507246	0.98550725	0.985507246
PTPRZ1 AND TMEM145	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-SIGLECL1	0.970149254	1	0.942028986

Antigen Pair Logic	F1	Precision	Recall
CLECL1 AND NOT-CCR1	0.918918919	0.944444444	0.894736842
CLECL1 AND NOT-CCR3	0.95	0.904761905	1
CLECL1 AND NOT-ZG16B	0.915662651	0.844444444	1
QSOX2 AND NOT-WFIKKN2	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-SEZ6	0.948717949	0.925	0.973684211
CLECL1 AND NOT-GJD3	0.926829268	0.863636364	1
QSOX2 AND NOT-GJD3	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-ACER1	0.948717949	0.925	0.973684211
CLECL1 AND NOT-LRRC25	0.926829268	0.863636364	1
CLECL1 AND NOT-CYP4F22	0.915662651	0.844444444	1
NOT-SLC44A3 AND CELSR1	0.961038961	0.948717949	0.973684211
QSOX2 AND NOT-KLHDC7A	0.935064935	0.923076923	0.947368421
QSOX2 AND NOT-CST9L	0.936708861	0.902439024	0.973684211
CLECL1 AND NOT-CST9	0.915662651	0.844444444	1
QSOX2 AND NOT-TMEM150A	0.947368421	0.947368421	0.947368421
QSOX2 AND NOT-ACMSD	0.948717949	0.925	0.973684211
QSOX2 AND NOT-SPATA3	0.948717949	0.925	0.973684211
CLECL1 AND NOT-TMEM198	0.938271605	0.88372093	1
QSOX2 AND NOT-TMEM198	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-LRRC15	0.936708861	0.902439024	0.973684211
CLECL1 AND NOT-TPRA1	0.948717949	0.925	0.973684211
QSOX2 AND NOT-TPRA1	0.947368421	0.947368421	0.947368421
CLECL1 AND NOT-SLC31A1	0.95	0.904761905	1
PTPN1 AND NOT-SLC31A1	0.915662651	0.844444444	1
CLECL1 AND NOT-SLC31A2	0.974358974	0.95	1
NOT-SLC31A2 AND ST6GALNAC4	0.935064935	0.923076923	0.947368421
NOT-SLC31A2 AND ADAM17	0.95	0.904761905	1
NOT-SLC31A2 AND SNRNP40	0.974358974	0.95	1
CLECL1 AND NOT-C4orf32	0.947368421	0.947368421	0.947368421
NOT-C4orf32 AND RNF149	0.931506849	0.971428571	0.894736842
IFNGR2 AND NOT-C4orf32	0.933333333	0.945945946	0.921052632
NOT-C4orf32 AND MARS	0.918918919	0.944444444	0.894736842
MCTP2 AND NOT-C4orf32	0.901408451	0.96969697	0.842105263
NOT-C4orf32 AND ADAM17	0.947368421	0.947368421	0.947368421
NOT-C4orf32 AND TNFRSF10A	0.935064935	0.923076923	0.947368421
QSOX2 AND NOT-UGT3A1	0.948717949	0.925	0.973684211
QSOX2 AND NOT-DPCR1	0.961038961	0.948717949	0.973684211
QSOX2 AND NOT-OR6B1	0.935064935	0.923076923	0.947368421
CLECL1 AND NOT-CPD	0.938271605	0.88372093	1
CLECL1 AND NOT-SVOPL	0.938271605	0.88372093	1
QSOX2 AND NOT-SVOPL	0.948717949	0.925	0.973684211
CELSR1 AND NOT-CLDN4	0.935064935	0.923076923	0.947368421
QSOX2 AND NOT-ADGRG4	0.948717949	0.925	0.973684211
QSOX2 AND NOT-CRHR1	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-PTCHD1	0.918918919	0.944444444	0.894736842
CLECL1 AND NOT-SLC32A1	0.926829268	0.863636364	1
QSOX2 AND NOT-SLC32A1	0.961038961	0.948717949	0.973684211
CLECL1 AND NOT-CSF2RA	0.95	0.904761905	1
CLECL1 AND NOT-ADRA1D	0.938271605	0.88372093	1
QSOX2 AND NOT-ADRA1D	0.948717949	0.925	0.973684211
CLECL1 AND NOT-CMTM2	0.926829268	0.863636364	1
CLECL1 AND NOT-CSPG4	0.915662651	0.844444444	1
CLECL1 AND NOT-MGAT5B	0.926829268	0.863636364	1
QSOX2 AND NOT-ATP8B3	0.948717949	0.925	0.973684211
CLECL1 AND NOT-SYT6	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-SYT6	0.936708861	0.902439024	0.973684211
CLECL1 AND NOT-NFAM1	0.926829268	0.863636364	1
QSOX2 AND NOT-ADRA2B	0.935064935	0.923076923	0.947368421
CLECL1 AND NOT-PLB1	0.926829268	0.863636364	1
CLECL1 AND NOT-GPR155	0.936708861	0.902439024	0.973684211
CLECL1 AND NOT-CMTM8	0.926829268	0.863636364	1
NOT-CMTM8 AND SNRNP40	0.926829268	0.863636364	1
CLECL1 AND NOT-IGSF11	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-IGSF11	0.947368421	0.947368421	0.947368421
CLECL1 AND NOT-CYBB	0.915662651	0.844444444	1
CLECL1 AND NOT-RNF217	0.915662651	0.844444444	1
NOT-RNF217 AND ADAM17	0.915662651	0.844444444	1
CLECL1 AND NOT-CYP1B1	0.926829268	0.863636364	1
CLECL1 AND NOT-LINGO2	0.974358974	0.95	1
QSOX2 AND NOT-LINGO2	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-FREM1	0.936708861	0.902439024	0.973684211
CLECL1 AND NOT-MOSPD2	0.918918919	0.944444444	0.894736842
QSOX2 AND NOT-CYP19A1	0.916666667	0.970588235	0.868421053
CLECL1 AND NOT-DAG1	0.915662651	0.844444444	1
CLECL1 AND NOT-TMEM30B	0.915662651	0.844444444	1
CLECL1 AND NOT-DBH	0.915662651	0.844444444	1

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
CHST10 AND NOT-SIGLECL1	0.970149254	1	0.942028986
PTPRZ1 AND NOT-GPR20	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-VSTM1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND TMEM150B	0.99270073	1	0.985507246
PTPRZ1 AND NOT-OR2L13	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-GPR22	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-EPHA10	0.985507246	0.98550725	0.985507246
PTPRZ1 AND LPAR4	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-SLC25A34	0.985507246	0.98550725	0.985507246
PTPRZ1 AND SIRPB2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-GPR25	0.985507246	0.98550725	0.985507246
PTPRZ1 AND CCDC141	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-GPR27	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-CDH19	0.985507246	0.98550725	0.985507246
PTPRZ1 AND DLL1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND SLC9A9	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-EPHA6	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-GPER1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-IGSF10	0.985507246	0.98550725	0.985507246
PTPRZ1 AND CYP4V2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-RNF175	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-ARL10	0.985507246	0.98550725	0.985507246
PTPRZ1 AND RNF180	0.985507246	0.98550725	0.985507246
PTPRZ1 AND DCBLD1	0.977777778	1	0.956521739
PTPRZ1 AND NOT-SCARA5	0.985507246	0.98550725	0.985507246
PTPRZ1 AND TRIQK	0.985507246	0.98550725	0.985507246
PTPRZ1 AND DPY19L4	0.970149254	1	0.942028986
PTPRZ1 AND CRB2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-GPR39	0.985507246	0.98550725	0.985507246
PTPRZ1 AND ATP11C	0.985294118	1	0.971014493
PTPRZ1 AND NOT-CFAP47	0.985507246	0.98550725	0.985507246
PTPRZ1 AND P2RY8	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-TUSC5	0.985294118	1	0.971014493
PTPRZ1 AND NOT-TRIM59	0.977777778	1	0.956521739
PTPRZ1 AND NOT-ANK1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND GRIA1	0.985507246	0.98550725	0.985507246
GRIA2 AND SLC25A3	0.970149254	1	0.942028986
PTPRZ1 AND NOT-GRID1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND COA3	0.977777778	1	0.956521739
PTPRZ1 AND TMEM176B	0.985507246	0.98550725	0.985507246
PTPRZ1 AND GRID2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND SPCS1	0.977777778	1	0.956521739
PTPRZ1 AND NOT-TMEM14A	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-GRIK1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND FLVCR1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-TMPRSS11E	0.985507246	0.98550725	0.985507246
PTPRZ1 AND MAGEH1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND GRIK2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND GRIK4	0.985507246	0.98550725	0.985507246
PTPRZ1 AND SLC43A3	0.985507246	0.98550725	0.985507246
PTPRZ1 AND GRIK5	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-GRIN2A	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-GRIN2B	0.99270073	1	0.985507246
PTPRZ1 AND GRIN2D	0.985507246	0.98550725	0.985507246
PTPRZ1 AND C1GALT1C1	0.970149254	1	0.942028986
PTPRZ1 AND GRINA	0.985507246	0.98550725	0.985507246
PTPRZ1 AND CNIH4	0.985294118	1	0.971014493
PTPRZ1 AND NOT-ANPEP	0.985507246	0.98550725	0.985507246
PTPRZ1 AND TMEM208	0.970149254	1	0.942028986
PTPRZ1 AND DNAJC15	0.985507246	0.98550725	0.985507246
PTPRZ1 AND CLEC2D	0.985507246	0.98550725	0.985507246
PTPRZ1 AND CD274	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-GRM2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-GRM3	0.99270073	1	0.985507246
PTPRZ1 AND NOT-GRM6	0.977777778	1	0.956521739
PTPRZ1 AND NOT-SLC25A4	0.985507246	0.98550725	0.985507246
PTPRZ1 AND SLC25A5	0.99270073	1	0.985507246
PTPRZ1 AND NOT-CYP2S1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-ZDHHC1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-ZDHHC8	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-GRHL1	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-GUCY2C	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-ICOS	0.985507246	0.98550725	0.985507246
PTPRZ1 AND NOT-GUCY2F	0.985507246	0.98550725	0.985507246
PTPRZ1 AND ALG5	0.977777778	1	0.956521739
PTPRZ1 AND NOT-NPC1L1	0.977777778	1	0.956521739

Antigen Pair Logic	F1	Precision	Recall
CLECL1 AND NOT-TRPV3	0.915662651	0.844444444	1
CLECL1 AND NOT-DCT	0.926829268	0.863636364	1
CLECL1 AND NOT-OXER1	0.961038961	0.948717949	0.973684211
CLECL1 AND NOT-SPTSSB	0.926829268	0.863636364	1
CLECL1 AND NOT-CHST13	0.938271605	0.88372093	1
CLECL1 AND NOT-SGMS2	0.915662651	0.844444444	1
CLECL1 AND NOT-PKD1L1	0.915662651	0.844444444	1
CLECL1 AND NOT-GIMAP1	0.926829268	0.863636364	1
CLECL1 AND NOT-SPTSSA	0.918918919	0.944444444	0.894736842
CLECL1 AND NOT-PTCRA	0.974358974	0.95	1
CLECL1 AND NOT-DHODH	0.933333333	0.945945946	0.921052632
CLECL1 AND NOT-DPP4	0.926829268	0.863636364	1
CLECL1 AND NOT-HBEGF	0.915662651	0.844444444	1
CLECL1 AND NOT-ECM1	0.938271605	0.88372093	1
CLECL1 AND NOT-PGAM5	0.938271605	0.88372093	1
CLECL1 AND NOT-DHRS7C	0.915662651	0.844444444	1
CLECL1 AND NOT-EMP1	0.915662651	0.844444444	1
CLECL1 AND NOT-ADGRE1	0.926829268	0.863636364	1
CLECL1 AND NOT-ERN1	0.926829268	0.863636364	1
CLECL1 AND NOT-ABCA3	0.926829268	0.863636364	1
CLECL1 AND NOT-EXTL3	0.938271605	0.88372093	1
CLECL1 AND NOT-F2RL1	0.926829268	0.863636364	1
CLECL1 AND NOT-MS4A15	0.915662651	0.844444444	1
CLECL1 AND NOT-HEPACAM	0.915662651	0.844444444	1
CLECL1 AND NOT-FCAR	0.915662651	0.844444444	1
CLECL1 AND NOT-C6orf89	0.948717949	0.925	0.973684211
CLECL1 AND NOT-DOLK	0.938271605	0.88372093	1
CLECL1 AND NOT-CD93	0.926829268	0.863636364	1
CLECL1 AND NOT-MLC1	0.918918919	0.944444444	0.894736842
CLECL1 AND NOT-FLT3	0.938271605	0.88372093	1
CLECL1 AND NOT-ACSL6	0.926829268	0.863636364	1
CLECL1 AND NOT-KCNH3	0.936708861	0.902439024	0.973684211
CLECL1 AND NOT-SLC44A1	0.95	0.904761905	1
CLECL1 AND NOT-NNT	0.915662651	0.844444444	1
CLECL1 AND NOT-SLC16A8	0.915662651	0.844444444	1
CLECL1 AND NOT-FPR2	0.926829268	0.863636364	1
CLECL1 AND NOT-CLEC5A	0.915662651	0.844444444	1
CLECL1 AND NOT-IL17RA	0.938271605	0.88372093	1
CLECL1 AND NOT-BCL2L13	0.918918919	0.944444444	0.894736842
CLECL1 AND NOT-FUT5	0.926829268	0.863636364	1
CLECL1 AND NOT-KDSR	0.918918919	0.944444444	0.894736842
CLECL1 AND NOT-FAM26E	0.926829268	0.863636364	1
CLECL1 AND NOT-CCDC108	0.938271605	0.88372093	1
CLECL1 AND NOT-TMEM186	0.915662651	0.844444444	1
CLECL1 AND NOT-TMEM158	0.915662651	0.844444444	1
CLECL1 AND NOT-LY6G6F	0.961038961	0.948717949	0.973684211
CLECL1 AND NOT-HIGD1A	0.961038961	0.948717949	0.973684211
CLECL1 AND NOT-LRIT1	0.938271605	0.88372093	1
CLECL1 AND NOT-GIMAP2	0.926829268	0.863636364	1
CLECL1 AND NOT-PCDHB5	0.915662651	0.844444444	1
CLECL1 AND NOT-KCNG2	0.948717949	0.925	0.973684211
CLECL1 AND NOT-OR10A3	0.915662651	0.844444444	1
CLECL1 AND NOT-MYOF	0.95	0.904761905	1
CLECL1 AND NOT-GGCX	0.948717949	0.925	0.973684211
CLECL1 AND NOT-SIGLEC7	0.926829268	0.863636364	1
CLECL1 AND NOT-KCNH5	0.915662651	0.844444444	1
CLECL1 AND NOT-SIGLEC9	0.938271605	0.88372093	1
CLECL1 AND NOT-GPR78	0.935064935	0.923076923	0.947368421
CLECL1 AND NOT-GLRA1	0.926829268	0.863636364	1
CLECL1 AND NOT-CECR6	0.962025316	0.926829268	1
CLECL1 AND NOT-GNS	0.962025316	0.926829268	1
CLECL1 AND NOT-GP9	0.926829268	0.863636364	1
CLECL1 AND NOT-SLCO3A1	0.926829268	0.863636364	1
CLECL1 AND NOT-GPR4	0.938271605	0.88372093	1
CLECL1 AND NOT-FAM171A2	0.938271605	0.88372093	1
CLECL1 AND NOT-SLC13A5	0.915662651	0.844444444	1
CLECL1 AND NOT-IZUMO1	0.915662651	0.844444444	1
CLECL1 AND NOT-SLC9A9	0.926829268	0.863636364	1
CLECL1 AND NOT-CRB2	0.95	0.904761905	1
CLECL1 AND NOT-GPR39	0.915662651	0.844444444	1
CLECL1 AND NOT-FFAR2	0.95	0.904761905	1
CLECL1 AND NOT-GRID2	0.926829268	0.863636364	1
CLECL1 AND NOT-DEXI	0.915662651	0.844444444	1
CLECL1 AND NOT-ANPEP	0.961038961	0.948717949	0.973684211
CLECL1 AND NOT-GRIN1	0.915662651	0.844444444	1
CLECL1 AND NOT-CNIH4	0.974358974	0.95	1

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
PTPRZ1 AND GNL2	0.985507246	0.98550725	0.985507246
PTPRZ1 AND GUSB	0.970149254	1	0.942028986
PTPRZ1 AND HILPDA	0.985294118	1	0.971014493
PTPRZ1 AND SEC61A1	0.970149254	1	0.942028986
ITGAV AND NOT-SLC39A2	0.970149254	1	0.942028986
PTPRZ1 AND NOT-SLC39A2	0.977777778	1	0.956521739
PTPRZ1 AND SLC40A1	0.977777778	1	0.956521739
PTPRZ1 AND NOT-HCRTR1	0.970149254	1	0.942028986
PTPRZ1 AND HEXB	0.977777778	1	0.956521739
PTPRZ1 AND HLA-B	0.99270073	1	0.985507246
PTPRZ1 AND HLA-DMA	0.985294118	1	0.971014493
PTPRZ1 AND HLA-DRA	0.977777778	1	0.956521739
PTPRZ1 AND HLA-DRB1	0.970149254	1	0.942028986
PTPRZ1 AND HLA-E	0.99270073	1	0.985507246
PTPRZ1 AND HLA-F	0.977777778	1	0.956521739
PTPRZ1 AND HLA-G	0.99270073	1	0.985507246
PTPRZ1 AND NOT-SERPINA9	0.99270073	1	0.985507246
APLP1 AND PTTG1IP	0.970149254	1	0.942028986
PTPRZ1 AND TNC	0.970149254	1	0.942028986
PTPRZ1 AND NOT-ICAM4	0.977777778	1	0.956521739
PTPRZ1 AND NOT-AQP8	0.99270073	1	0.985507246
CHST10 AND NOT-AQP8	0.970149254	1	0.942028986
PTPRZ1 AND IFNGR2	0.99270073	1	0.985507246
PTPRZ1 AND NOT-IHH	0.977777778	1	0.956521739
PTPRZ1 AND NOT-FASLG	0.99270073	1	0.985507246
PTPRZ1 AND CXCR1	0.977777778	1	0.956521739
PTPRZ1 AND CXCR2	0.99270073	1	0.985507246
PTPRZ1 AND NOT-PRAC2	0.985294118	1	0.971014493
PTPRZ1 AND INPP4A	0.99270073	1	0.985507246
PTPRZ1 AND INSL3	0.99270073	1	0.985507246
PTPRZ1 AND ITGAE	0.985294118	1	0.971014493
ITGAV AND NOT-C1orf186	0.977777778	1	0.956521739
ITGAV AND NOT-DUOX2	0.970149254	1	0.942028986
ITGAV AND GOLM1	0.970149254	1	0.942028986
ITGAV AND NOT-MRAP	0.970149254	1	0.942028986
ITGAV AND TMEM9B	0.970149254	1	0.942028986
PTPRZ1 AND ITGAV	0.985294118	1	0.971014493
ITGAV AND NOT-TSPAN8	0.977777778	1	0.956521739
ITGAV AND WRB	0.970149254	1	0.942028986
ITGAV AND TMX1	0.970149254	1	0.942028986
ITGAV AND PKD2L1	0.970149254	1	0.942028986
ITGAV AND NOT-CD28	0.970149254	1	0.942028986
ITGAV AND NOT-SIGLEC6	0.970149254	1	0.942028986
PTPRZ1 AND NOT-KCNA4	0.985294118	1	0.971014493
PTPRZ1 AND TMEM179B	0.99270073	1	0.985507246
LRRC37A3 AND NOT-EPCAM	0.970149254	1	0.942028986
LRRC37A3 AND NOT-SLC28A3	0.970149254	1	0.942028986
PTPRZ1 AND TMEM205	0.99270073	1	0.985507246
PTPRZ1 AND KCNH2	0.985294118	1	0.971014493
CHST10 AND NOT-KCNS1	0.970149254	1	0.942028986
PTPRZ1 AND NOT-KCNS2	0.970149254	1	0.942028986
PTPRZ1 AND KDR	0.977777778	1	0.956521739
PTPRZ1 AND NOT-KIR3DL2	0.99270073	1	0.985507246
CHST10 AND NOT-KIR3DL2	0.970149254	1	0.942028986
PTPRZ1 AND NOT-SPATA31D1	0.985294118	1	0.971014493
PTPRZ1 AND NOT-OR51I1	0.985294118	1	0.971014493
PTPRZ1 AND LAMP1	0.970149254	1	0.942028986
PTPRZ1 AND ARHGAP1	0.985294118	1	0.971014493
PTPRZ1 AND LBR	0.985294118	1	0.971014493
PTPRZ1 AND NOT-LCT	0.985294118	1	0.971014493
CHST10 AND NOT-C11orf87	0.970149254	1	0.942028986
CHST10 AND NOT-FAM174B	0.970149254	1	0.942028986
PTPRZ1 AND C4orf3	0.977777778	1	0.956521739
PTPRZ1 AND NOT-XKRX	0.977777778	1	0.956521739
PTPRZ1 AND NOT-LRP2	0.985294118	1	0.971014493
CHST10 AND NOT-HAPLN4	0.970149254	1	0.942028986
PTPRZ1 AND LTBP3	0.970149254	1	0.942028986
PTPRZ1 AND NOT-DUOXA2	0.985294118	1	0.971014493
PTPRZ1 AND NOT-LTK	0.985294118	1	0.971014493
PTPRZ1 AND NOT-MAL	0.970149254	1	0.942028986
PTPRZ1 AND MARS	0.985294118	1	0.971014493
PTPRZ1 AND MCL1	0.985294118	1	0.971014493
CHST10 AND NOT-ADAM11	0.970149254	1	0.942028986
PTPRZ1 AND NOT-ART4	0.985294118	1	0.971014493
PTPRZ1 AND MFAP3	0.977777778	1	0.956521739
PTPRZ1 AND MGAT2	0.985294118	1	0.971014493

Antigen Pair Logic	F1	Precision	Recall
CLECL1 AND NOT-GRM2	0.915662651	0.844444444	1
CLECL1 AND NOT-GRM3	0.938271605	0.88372093	1
CLECL1 AND NOT-ZDHHC1	0.926829268	0.863636364	1
CLECL1 AND NOT-GPR171	0.926829268	0.863636364	1
CLECL1 AND NOT-LRP12	0.938271605	0.88372093	1
CLECL1 AND NOT-HFE	0.948717949	0.925	0.973684211
CLECL1 AND NOT-MR1	0.945945946	0.972222222	0.921052632
CLECL1 AND NOT-HMOX2	0.974358974	0.95	1
CLECL1 AND NOT-CLEC4D	0.915662651	0.844444444	1
CLECL1 AND NOT-ICAM4	0.929577465	1	0.868421053
CLECL1 AND NOT-VSIG1	0.945945946	0.972222222	0.921052632
CLECL1 AND NOT-FASLG	0.926829268	0.863636364	1
CLECL1 AND NOT-IL6R	0.926829268	0.863636364	1
CLECL1 AND NOT-IL7R	0.926829268	0.863636364	1
CLECL1 AND NOT-CXCR1	0.961038961	0.948717949	0.973684211
CLECL1 AND NOT-AQP2	0.915662651	0.844444444	1
CLECL1 AND NOT-IL12RB1	0.935064935	0.923076923	0.947368421
CLECL1 AND NOT-IL12RB2	0.95	0.904761905	1
CLECL1 AND NOT-AQP3	0.926829268	0.863636364	1
CLECL1 AND NOT-ITGA5	0.974358974	0.95	1
CLECL1 AND NOT-ITGB3	0.95	0.904761905	1
CLECL1 AND NOT-ITGB5	0.926829268	0.863636364	1
CLECL1 AND NOT-ITGB6	0.936708861	0.902439024	0.973684211
CLECL1 AND NOT-KCNC2	0.936708861	0.902439024	0.973684211
CLECL1 AND NOT-IGFL1	0.95	0.904761905	1
CLECL1 AND NOT-KIR2DL4	0.926829268	0.863636364	1
CLECL1 AND NOT-KIR2DS4	0.961038961	0.948717949	0.973684211
CLECL1 AND NOT-KIR3DL2	0.974358974	0.95	1
CLECL1 AND NOT-SMIM1	0.935064935	0.923076923	0.947368421
CLECL1 AND NOT-ARHGAP1	0.926829268	0.863636364	1
CLECL1 AND NOT-ERVFRD-1	0.938271605	0.88372093	1
CLECL1 AND NOT-ARSB	0.926829268	0.863636364	1
CLECL1 AND NOT-MAL	0.936708861	0.902439024	0.973684211
CLECL1 AND NOT-CD46	0.938271605	0.88372093	1
CLECL1 AND NOT-MLN	0.926829268	0.863636364	1
CLECL1 AND NOT-ASGR1	0.95	0.904761905	1
CLECL1 AND NOT-SMIM4	0.926829268	0.863636364	1
CLECL1 AND NOT-NCAM1	0.961038961	0.948717949	0.973684211
CLECL1 AND NOT-NPY	0.935064935	0.923076923	0.947368421
CLECL1 AND NOT-NT5E	0.918918919	0.944444444	0.894736842
CLECL1 AND NOT-NTSR1	0.901408451	0.96969697	0.842105263
CLECL1 AND NOT-P2RX7	0.936708861	0.902439024	0.973684211
CLECL1 AND NOT-P2RY2	0.95	0.904761905	1
CLECL1 AND NOT-CD207	0.938271605	0.88372093	1
CLECL1 AND NOT-CALY	0.935064935	0.923076923	0.947368421
CLECL1 AND NOT-KCNK4	0.926829268	0.863636364	1
CLECL1 AND NOT-TAS2R14	0.936708861	0.902439024	0.973684211
CLECL1 AND NOT-TRAT1	0.926829268	0.863636364	1
CLECL1 AND NOT-CLEC4A	0.945945946	0.972222222	0.921052632
CLECL1 AND NOT-PCDH1	0.938271605	0.88372093	1
CLECL1 AND NOT-PCSK5	0.926829268	0.863636364	1
CLECL1 AND NOT-CLEC1B	0.926829268	0.863636364	1
CLECL1 AND NOT-ZDHHC3	0.938271605	0.88372093	1
CLECL1 AND NOT-FAM198B	0.938271605	0.88372093	1
CLECL1 AND NOT-MS4A4A	0.936708861	0.902439024	0.973684211
CLECL1 AND NOT-AIG1	0.961038961	0.948717949	0.973684211
CLECL1 AND NOT-TRPV2	0.987012987	0.974358974	1
CLECL1 AND NOT-PDE3B	0.95	0.904761905	1
CLECL1 AND NOT-GDE1	0.936708861	0.902439024	0.973684211
CLECL1 AND NOT-NBAS	0.948717949	0.925	0.973684211
CLECL1 AND NOT-CD244	0.926829268	0.863636364	1
CLECL1 AND NOT-ATP5G3	0.938271605	0.88372093	1
CLECL1 AND NOT-S1PR5	0.926829268	0.863636364	1
CLECL1 AND NOT-TREM2	0.926829268	0.863636364	1
CLECL1 AND NOT-NLGN3	0.926829268	0.863636364	1
CLECL1 AND NOT-FAM105A	0.926829268	0.863636364	1
CLECL1 AND NOT-LY6K	0.926829268	0.863636364	1
CLECL1 AND NOT-QPCTL	0.938271605	0.88372093	1
CLECL1 AND NOT-CDKAL1	0.938271605	0.88372093	1
CLECL1 AND NOT-LPCAT2	0.926829268	0.863636364	1
CLECL1 AND NOT-SLC35F6	0.962025316	0.926829268	1
CLECL1 AND NOT-ACPP	0.961038961	0.948717949	0.973684211
CLECL1 AND NOT-TMEM51	0.95	0.904761905	1
CLECL1 AND NOT-SLC35C1	0.96	0.972972973	0.947368421
CLECL1 AND NOT-CSGALNACT2	0.962025316	0.926829268	1
CLECL1 AND NOT-CHST12	0.947368421	0.947368421	0.947368421

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
PTPRZ1 AND MGAT5	0.977777778	1	0.956521739
PTPRZ1 AND MGP	0.985294118	1	0.971014493
PTPRZ1 AND MPG	0.977777778	1	0.956521739
PTPRZ1 AND NOT-SMCO3	0.99270073	1	0.985507246
NOT-C1orf186 AND CHST11	0.970149254	1	0.942028986
NOT-C1orf186 AND SLC35C2	0.970149254	1	0.942028986
NOT-C1orf186 AND APMAP	0.970149254	1	0.942028986
PTPRZ1 AND NOT-C1orf186	0.99270073	1	0.985507246
NOT-C1orf186 AND VAMP4	0.977777778	1	0.956521739
CHST10 AND NOT-C1orf186	0.970149254	1	0.942028986
PTPRZ1 AND NOT-PTCHD4	0.99270073	1	0.985507246
PTPRZ1 AND ASPH	0.985294118	1	0.971014493
PTPRZ1 AND MTHFD1	0.99270073	1	0.985507246
PTPRZ1 AND NOT-MTNR1B	0.977777778	1	0.956521739
PTPRZ1 AND NDUFA3	0.99270073	1	0.985507246
PTPRZ1 AND NDUFA4	0.99270073	1	0.985507246
PTPRZ1 AND NDUFB1	0.99270073	1	0.985507246
PTPRZ1 AND NDUFB5	0.977777778	1	0.956521739
PTPRZ1 AND NDUFB6	0.985294118	1	0.971014493
ATP1B2 AND TMEM14C	0.970149254	1	0.942028986
PTPRZ1 AND ATP1B3	0.99270073	1	0.985507246
PTPRZ1 AND NOTCH4	0.99270073	1	0.985507246
PTPRZ1 AND ATP2A1	0.970149254	1	0.942028986
PTPRZ1 AND ATP2A2	0.985294118	1	0.971014493
PTPRZ1 AND NOT-NTF3	0.985294118	1	0.971014493
CHST10 AND NOT-ATP2B3	0.970149254	1	0.942028986
PTPRZ1 AND NOT-OR1F1	0.99270073	1	0.985507246
PTPRZ1 AND OXA1L	0.99270073	1	0.985507246
PTPRZ1 AND P2RX4	0.977777778	1	0.956521739
PTPRZ1 AND RRM2B	0.985294118	1	0.971014493
PTPRZ1 AND CHST11	0.977777778	1	0.956521739
PTPRZ1 AND PAM	0.99270073	1	0.985507246
PTPRZ1 AND NOT-TAS2R9	0.985294118	1	0.971014493
PTPRZ1 AND NOT-TAS2R13	0.99270073	1	0.985507246
PTPRZ1 AND TMED5	0.99270073	1	0.985507246
PTPRZ1 AND SLC35C2	0.977777778	1	0.956521739
PTPRZ1 AND DERL2	0.977777778	1	0.956521739
PTPRZ1 AND TMED7	0.970149254	1	0.942028986
PTPRZ1 AND FIS1	0.985294118	1	0.971014493
PTPRZ1 AND GOLT1B	0.977777778	1	0.956521739
PTPRZ1 AND TXNDC12	0.99270073	1	0.985507246
PTPRZ1 AND SCCPDH	0.99270073	1	0.985507246
PTPRZ1 AND IER3IP1	0.99270073	1	0.985507246
PTPRZ1 AND CKLF	0.977777778	1	0.956521739
PTPRZ1 AND EMC4	0.99270073	1	0.985507246
PTPRZ1 AND SHISA5	0.99270073	1	0.985507246
PTPRZ1 AND TMEM69	0.99270073	1	0.985507246
PTPRZ1 AND NT5C3A	0.970149254	1	0.942028986
PTPRZ1 AND TBC1D7	0.99270073	1	0.985507246
GOLM1 AND NOT-SCNN1A	0.970149254	1	0.942028986
PTPRZ1 AND BFAR	0.99270073	1	0.985507246
PTPRZ1 AND ERGIC2	0.985294118	1	0.971014493
PTPRZ1 AND TIMMDC1	0.985294118	1	0.971014493
PTPRZ1 AND NOT-GCNT4	0.99270073	1	0.985507246
PTPRZ1 AND ARMCX1	0.985294118	1	0.971014493
PTPRZ1 AND CHST15	0.977777778	1	0.956521739
PTPRZ1 AND ATRAID	0.985294118	1	0.971014493
PTPRZ1 AND WDR83OS	0.985294118	1	0.971014493
PTPRZ1 AND TMEM14C	0.985294118	1	0.971014493
PTPRZ1 AND TMEM138	0.985294118	1	0.971014493
PTPRZ1 AND ARMCX3	0.99270073	1	0.985507246
PTPRZ1 AND CUTA	0.970149254	1	0.942028986
GPR158 AND ATP5F1	0.977777778	1	0.956521739
PTPRZ1 AND ATP5F1	0.985294118	1	0.971014493
PTPRZ1 AND PIGT	0.99270073	1	0.985507246
PTPRZ1 AND ERGIC3	0.985294118	1	0.971014493
PTPRZ1 AND SLC25A39	0.977777778	1	0.956521739
PTPRZ1 AND DHRS7	0.970149254	1	0.942028986
PTPRZ1 AND TMBIM4	0.99270073	1	0.985507246
PTPRZ1 AND PTRH2	0.99270073	1	0.985507246
PTPRZ1 AND FKBP7	0.99270073	1	0.985507246
PTPRZ1 AND DNAJB11	0.99270073	1	0.985507246
PTPRZ1 AND PECAM1	0.99270073	1	0.985507246
PTPRZ1 AND NOT-ASIC5	0.970149254	1	0.942028986
PTPRZ1 AND SLC25A3	0.99270073	1	0.985507246
PTPRZ1 AND ATP6V0C	0.99270073	1	0.985507246

Antigen Pair Logic	F1	Precision	Recall
CLECL1 AND NOT-POMGNT1	0.938271605	0.88372093	1
CLECL1 AND NOT-FLVCR2	0.974358974	0.95	1
CLECL1 AND NOT-FAR2	0.926829268	0.863636364	1
CLECL1 AND NOT-TMEM30A	0.974358974	0.95	1
CLECL1 AND NOT-PAG1	0.95	0.904761905	1
CLECL1 AND NOT-TEX2	0.95	0.904761905	1
CLECL1 AND NOT-ACSS2	0.95	0.904761905	1
CLECL1 AND NOT-PSEN2	0.918918919	0.944444444	0.894736842
CLECL1 AND NOT-PANX2	0.938271605	0.88372093	1
CLECL1 AND NOT-SUCNR1	0.926829268	0.863636364	1
CLECL1 AND NOT-UGGT1	0.95	0.904761905	1
CLECL1 AND NOT-FAM20C	0.962025316	0.926829268	1
CLECL1 AND NOT-ENTPD7	0.936708861	0.902439024	0.973684211
CLECL1 AND NOT-MAN1C1	0.926829268	0.863636364	1
CLECL1 AND NOT-ADGRG6	0.926829268	0.863636364	1
CLECL1 AND NOT-SNX14	0.933333333	0.945945946	0.921052632
CLECL1 AND NOT-PTAFR	0.926829268	0.863636364	1
CLECL1 AND NOT-PTGDR	0.926829268	0.863636364	1
CLECL1 AND NOT-PTGER2	0.936708861	0.902439024	0.973684211
CLECL1 AND NOT-ESYT2	0.926829268	0.863636364	1
CLECL1 AND NOT-PCDH10	0.926829268	0.863636364	1
CLECL1 AND NOT-ISLR2	0.938271605	0.88372093	1
CLECL1 AND NOT-HAMP	0.938271605	0.88372093	1
CLECL1 AND NOT-PTPRA	0.936708861	0.902439024	0.973684211
CLECL1 AND NOT-SLC46A2	0.974358974	0.95	1
CLECL1 AND NOT-PVR	0.95	0.904761905	1
CLECL1 AND NOT-LGR6	0.926829268	0.863636364	1
CLECL1 AND NOT-RHD	0.926829268	0.863636364	1
CLECL1 AND NOT-ROM1	0.926829268	0.863636364	1
CLECL1 AND NOT-RTN1	0.926829268	0.863636364	1
CLECL1 AND NOT-SCN3A	0.935064935	0.923076923	0.947368421
CLECL1 AND NOT-SDHC	0.974358974	0.95	1
CLECL1 AND NOT-SELPLG	0.926829268	0.863636364	1
CLECL1 AND NOT-CDH23	0.936708861	0.902439024	0.973684211
CLECL1 AND NOT-SLC39A8	0.948717949	0.925	0.973684211
CLECL1 AND NOT-MS4A6A	0.926829268	0.863636364	1
CLECL1 AND NOT-GPR135	0.926829268	0.863636364	1
CLECL1 AND NOT-P2RY12	0.926829268	0.863636364	1
CLECL1 AND NOT-SLC4A3	0.933333333	0.945945946	0.921052632
CLECL1 AND NOT-SLC1A5	0.95	0.904761905	1
CLECL1 AND NOT-SLC8A1	0.938271605	0.88372093	1
CLECL1 AND NOT-SLC22A4	0.926829268	0.863636364	1
CLECL1 AND NOT-SMPD1	0.926829268	0.863636364	1
CLECL1 AND NOT-SIGLEC1	0.926829268	0.863636364	1
CLECL1 AND NOT-SORL1	0.933333333	0.945945946	0.921052632
CLECL1 AND NOT-SPN	0.926829268	0.863636364	1
CLECL1 AND NOT-BPI	0.918918919	0.944444444	0.894736842
CLECL1 AND NOT-SSR3	0.933333333	0.945945946	0.921052632
CLECL1 AND NOT-SSTR5	0.938271605	0.88372093	1
CLECL1 AND NOT-BST1	0.926829268	0.863636364	1
CLECL1 AND NOT-TGFBR3	0.926829268	0.863636364	1
CLECL1 AND NOT-TLR5	0.926829268	0.863636364	1
CLECL1 AND NOT-TRPC4	0.938271605	0.88372093	1
CLECL1 AND NOT-FZD5	0.95	0.904761905	1
CLECL1 AND NOT-PRRG4	0.926829268	0.863636364	1
CLECL1 AND NOT-ELOVL6	0.95	0.904761905	1
CLECL1 AND NOT-ALG12	0.936708861	0.902439024	0.973684211
CLECL1 AND NOT-TMEM204	0.948717949	0.925	0.973684211
CLECL1 AND NOT-ATP8B4	0.926829268	0.863636364	1
CLECL1 AND NOT-DNAJC22	0.938271605	0.88372093	1
CLECL1 AND NOT-TMEM254	0.961038961	0.948717949	0.973684211
CLECL1 AND NOT-DNAJC5	0.974358974	0.95	1
CLECL1 AND NOT-TAS1R1	0.935064935	0.923076923	0.947368421
CLECL1 AND NOT-KIAA1715	0.901408451	0.96969697	0.842105263
CLECL1 AND NOT-HM13	0.95	0.904761905	1
CLECL1 AND NOT-LRRC3	0.936708861	0.902439024	0.973684211
CLECL1 AND NOT-VANGL1	0.926829268	0.863636364	1
CLECL1 AND NOT-SFXN3	0.938271605	0.88372093	1
CLECL1 AND NOT-DYSF	0.926829268	0.863636364	1
CLECL1 AND NOT-CRISPLD2	0.926829268	0.863636364	1
CLECL1 AND NOT-OR1G1	0.938271605	0.88372093	1
CLECL1 AND NOT-B3GNT5	0.95	0.904761905	1
CLECL1 AND NOT-LRRC8C	0.961038961	0.948717949	0.973684211
CLECL1 AND NOT-YIPF4	0.974358974	0.95	1
CLECL1 AND NOT-C2orf40	0.974358974	0.95	1
CLECL1 AND NOT-ATRN	0.938271605	0.88372093	1

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
PTPRZ1 AND PIGF	0.977777778	1	0.956521739
PTPRZ1 AND ACP1	0.985294118	1	0.971014493
PTPRZ1 AND NOT-TM6SF2	0.985294118	1	0.971014493
PTPRZ1 AND ATP6V0B	0.99270073	1	0.985507246
PTPRZ1 AND ATP6AP1	0.977777778	1	0.956521739
PTPRZ1 AND NOT-ATP7B	0.985294118	1	0.971014493
PTPRZ1 AND SLC38A2	0.977777778	1	0.956521739
PTPRZ1 AND DNAJC10	0.970149254	1	0.942028986
PTPRZ1 AND TMCO1	0.977777778	1	0.956521739
PTPRZ1 AND NDUFB11	0.985294118	1	0.971014493
PTPRZ1 AND NOT-RNF186	0.99270073	1	0.985507246
PTPRZ1 AND NOT-SLC6A20	0.970149254	1	0.942028986
PTPRZ1 AND SLC35F2	0.977777778	1	0.956521739
PTPRZ1 AND LEPROT	0.99270073	1	0.985507246
PTPRZ1 AND CTSA	0.970149254	1	0.942028986
PTPRZ1 AND PPIB	0.970149254	1	0.942028986
PTPRZ1 AND CMTM6	0.977777778	1	0.956521739
PTPRZ1 AND TMEM161A	0.985294118	1	0.971014493
PTPRZ1 AND NOT-RHBDL2	0.970149254	1	0.942028986
PTPRZ1 AND PIGX	0.985294118	1	0.971014493
PTPRZ1 AND TMEM70	0.99270073	1	0.985507246
PTPRZ1 AND NOT-HRASLS2	0.977777778	1	0.956521739
PTPRZ1 AND TMEM255A	0.977777778	1	0.956521739
TMEM255A AND WRB	0.970149254	1	0.942028986
TMEM255A AND NOT-SIGLEC6	0.977777778	1	0.956521739
PTPRZ1 AND SLC35A5	0.977777778	1	0.956521739
PTPRZ1 AND TMEM248	0.985294118	1	0.971014493
PTPRZ1 AND TMEM45A	0.985294118	1	0.971014493
ZDHHC4 AND NOT-TSPAN8	0.970149254	1	0.942028986
PTPRZ1 AND TMEM33	0.970149254	1	0.942028986
PTPRZ1 AND TMEM39A	0.985294118	1	0.971014493
PTPRZ1 AND NOT-TMEM144	0.977777778	1	0.956521739
PTPRZ1 AND AGPAT5	0.977777778	1	0.956521739
PTPRZ1 AND DRAM1	0.970149254	1	0.942028986
PTPRZ1 AND LAPTM4B	0.977777778	1	0.956521739
PTPRZ1 AND CSGALNACT2	0.99270073	1	0.985507246
PTPRZ1 AND PRCP	0.977777778	1	0.956521739
PTPRZ1 AND PARL	0.985294118	1	0.971014493
PTPRZ1 AND ETNK1	0.985294118	1	0.971014493
PTPRZ1 AND ZDHHC7	0.977777778	1	0.956521739
PTPRZ1 AND FLVCR2	0.985294118	1	0.971014493
PTPRZ1 AND TMEM127	0.977777778	1	0.956521739
PTPRZ1 AND SLC30A6	0.977777778	1	0.956521739
PTPRZ1 AND TMEM30A	0.970149254	1	0.942028986
PTPRZ1 AND NGLY1	0.99270073	1	0.985507246
PTPRZ1 AND RNF130	0.970149254	1	0.942028986
PTPRZ1 AND PAG1	0.970149254	1	0.942028986
PTPRZ1 AND VIMP	0.985294118	1	0.971014493
PTPRZ1 AND GLT8D1	0.970149254	1	0.942028986
PTPRZ1 AND EMC3	0.970149254	1	0.942028986
PTPRZ1 AND PLGRKT	0.970149254	1	0.942028986
PTPRZ1 AND ECHDC1	0.985294118	1	0.971014493
PTPRZ1 AND TMEM126B	0.985294118	1	0.971014493
PTPRZ1 AND NOT-GABRQ	0.985294118	1	0.971014493
PTPRZ1 AND BCAP29	0.970149254	1	0.942028986
PTPRZ1 AND NOT-PCDHGB6	0.99270073	1	0.985507246
PTPRZ1 AND NOT-GKN1	0.985294118	1	0.971014493
PTPRZ1 AND NOT-TRPV5	0.985294118	1	0.971014493
PTPRZ1 AND NOT-TMPRSS15	0.985294118	1	0.971014493
PTPRZ1 AND DIABLO	0.970149254	1	0.942028986
PTPRZ1 AND PSEN1	0.970149254	1	0.942028986
PTPRZ1 AND NOT-TMPRSS4	0.985294118	1	0.971014493
PTPRZ1 AND NPDC1	0.985294118	1	0.971014493
PTPRZ1 AND TMEM9B	0.99270073	1	0.985507246
PTPRZ1 AND EMC7	0.977777778	1	0.956521739
CHST10 AND NOT-CELF4	0.970149254	1	0.942028986
PTPRZ1 AND NOT-OTOR	0.99270073	1	0.985507246
PTPRZ1 AND NOT-C2orf83	0.970149254	1	0.942028986
PTPRZ1 AND C5orf15	0.977777778	1	0.956521739
PTPRZ1 AND ACKR3	0.985294118	1	0.971014493
PTPRZ1 AND NOT-SLC17A7	0.977777778	1	0.956521739
PTPRZ1 AND ATP13A1	0.970149254	1	0.942028986
PTPRZ1 AND SLC44A2	0.985294118	1	0.971014493
PTPRZ1 AND SNX14	0.99270073	1	0.985507246
PTPRZ1 AND NOT-PTGDR	0.99270073	1	0.985507246
PTPRZ1 AND NOT-PTGER3	0.985294118	1	0.971014493

Antigen Pair Logic	F1	Precision	Recall
CLECL1 AND NOT-CASR	0.926829268	0.863636364	1
CLECL1 AND NOT-DGAT2	0.931506849	0.971428571	0.894736842
CLECL1 AND NOT-ADGRE3	0.926829268	0.863636364	1
CLECL1 AND NOT-PIGO	0.95	0.904761905	1
CLECL1 AND NOT-AGPAT9	0.95	0.904761905	1
CLECL1 AND NOT-PLXDC2	0.926829268	0.863636364	1
CLECL1 AND NOT-SMIM3	0.935064935	0.923076923	0.947368421
CLECL1 AND NOT-VAMP4	0.947368421	0.947368421	0.947368421
CLECL1 AND NOT-TNFSF14	0.962025316	0.926829268	1
CLECL1 AND NOT-TNFSF10	0.926829268	0.863636364	1
CLECL1 AND NOT-ADAM9	0.938271605	0.88372093	1
CLECL1 AND NOT-TNFRSF10C	0.948717949	0.925	0.973684211
CLECL1 AND NOT-IL18RAP	0.926829268	0.863636364	1
CLECL1 AND NOT-IL1RL2	0.938271605	0.88372093	1
CLECL1 AND NOT-GALR2	0.938271605	0.88372093	1
CLECL1 AND NOT-INPP4B	0.926829268	0.863636364	1
CLECL1 AND NOT-CACNA1I	0.935064935	0.923076923	0.947368421
CLECL1 AND NOT-CCRL2	0.926829268	0.863636364	1
CLECL1 AND NOT-YIF1B	0.95	0.904761905	1
CLECL1 AND NOT-PIGQ	0.938271605	0.88372093	1
CLECL1 AND NOT-CD2	0.926829268	0.863636364	1
CLECL1 AND NOT-CD3D	0.926829268	0.863636364	1
CLECL1 AND NOT-PLXNA4	0.926829268	0.863636364	1
CLECL1 AND NOT-SYT12	0.926829268	0.863636364	1
CLECL1 AND NOT-REEP6	0.926829268	0.863636364	1
CLECL1 AND NOT-MS4A3	0.935064935	0.923076923	0.947368421
CLECL1 AND NOT-B4GALT6	0.95	0.904761905	1
CLECL1 AND NOT-CD163	0.95	0.904761905	1
CLECL1 AND NOT-NDST3	0.936708861	0.902439024	0.973684211
CLECL1 AND NOT-NRXN1	0.926829268	0.863636364	1
CLECL1 AND NOT-CD101	0.926829268	0.863636364	1
CLECL1 AND NOT-CD28	0.926829268	0.863636364	1
CLECL1 AND NOT-KCNK6	0.962025316	0.926829268	1
CLECL1 AND NOT-ABCG2	0.95	0.904761905	1
CLECL1 AND NOT-TNFRSF8	0.948717949	0.925	0.973684211
CLECL1 AND NOT-CD33	0.926829268	0.863636364	1
CLECL1 AND NOT-CD36	0.936708861	0.902439024	0.973684211
CLECL1 AND NOT-SCAMP1	0.938271605	0.88372093	1
CLECL1 AND NOT-CD68	0.926829268	0.863636364	1
CLECL1 AND NOT-TM9SF4	0.926829268	0.863636364	1
CLECL1 AND NOT-MFAP3L	0.926829268	0.863636364	1
CLECL1 AND NOT-TRIL	0.926829268	0.863636364	1
NOT-DAG1 AND CELSR1	0.95	0.904761905	1
QSOX2 AND NOT-TMEM92	0.935064935	0.923076923	0.947368421
QSOX2 AND NOT-TRPV3	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-SLC16A11	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-PTCRA	0.973684211	0.973684211	0.973684211
QSOX2 AND NOT-PGAM5	0.948717949	0.925	0.973684211
QSOX2 AND NOT-SLC51A	0.935064935	0.923076923	0.947368421
QSOX2 AND NOT-EPO	0.961038961	0.948717949	0.973684211
QSOX2 AND NOT-F10	0.935064935	0.923076923	0.947368421
QSOX2 AND NOT-FCAR	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-MS4A2	0.948717949	0.925	0.973684211
QSOX2 AND NOT-ADGRG3	0.948717949	0.925	0.973684211
QSOX2 AND NOT-FGF10	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-MLC1	0.935064935	0.923076923	0.947368421
QSOX2 AND NOT-FLT3LG	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-NCSTN	0.961038961	0.948717949	0.973684211
QSOX2 AND NOT-IL17RA	0.948717949	0.925	0.973684211
QSOX2 AND NOT-BCL2L13	0.935064935	0.923076923	0.947368421
QSOX2 AND NOT-FAM26E	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-CCDC108	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-GABRA4	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-GABRD	0.935064935	0.923076923	0.947368421
QSOX2 AND NOT-LY6G6F	0.961038961	0.948717949	0.973684211
QSOX2 AND NOT-TAS2R41	0.947368421	0.947368421	0.947368421
QSOX2 AND NOT-SLITRK5	0.935064935	0.923076923	0.947368421
QSOX2 AND NOT-OR2M4	0.935064935	0.923076923	0.947368421
QSOX2 AND NOT-KCNG2	0.961038961	0.948717949	0.973684211
QSOX2 AND NOT-GCGR	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-OR8B8	0.918918919	0.944444444	0.894736842
QSOX2 AND NOT-HS6ST3	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-B4GALT1	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-DKKL1	0.961038961	0.948717949	0.973684211
QSOX2 AND NOT-SIGLEC9	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-GPR78	0.96	0.972972973	0.947368421

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
PTPRZ1 AND CYP20A1	0.977777778	1	0.956521739
PTPRZ1 AND MAVS	0.970149254	1	0.942028986
GPR158 AND NOT-UGT2B15	0.977777778	1	0.956521739
GPR158 AND PTTG1IP	0.977777778	1	0.956521739
PTPRZ1 AND PTPN2	0.99270073	1	0.985507246
PTPRZ1 AND NOT-SLAMF7	0.985294118	1	0.971014493
PTPRZ1 AND PEX2	0.977777778	1	0.956521739
PTPRZ1 AND MS4A7	0.970149254	1	0.942028986
PTPRZ1 AND JAM2	0.985294118	1	0.971014493
PTPRZ1 AND OSTC	0.985294118	1	0.971014493
PTPRZ1 AND SELK	0.977777778	1	0.956521739
PTPRZ1 AND NOT-ACE2	0.99270073	1	0.985507246
PTPRZ1 AND NOT-CACNG6	0.977777778	1	0.956521739
PTPRZ1 AND RBM3	0.99270073	1	0.985507246
PTPRZ1 AND NOT-RHAG	0.99270073	1	0.985507246
PTPRZ1 AND RPN1	0.970149254	1	0.942028986
PTPRZ1 AND RPN2	0.99270073	1	0.985507246
PTPRZ1 AND RYK	0.99270073	1	0.985507246
PTPRZ1 AND CCL4	0.977777778	1	0.956521739
PTPRZ1 AND NOT-CCL22	0.985294118	1	0.971014493
PTPRZ1 AND SRR	0.977777778	1	0.956521739
PTPRZ1 AND SDCBP	0.985294118	1	0.971014493
PTPRZ1 AND MANBAL	0.99270073	1	0.985507246
PTPRZ1 AND MCUR1	0.977777778	1	0.956521739
PTPRZ1 AND ADGRL4	0.970149254	1	0.942028986
PTPRZ1 AND NOT-SEMA4A	0.977777778	1	0.956521739
PTPRZ1 AND MS4A6A	0.970149254	1	0.942028986
PTPRZ1 AND C5orf28	0.970149254	1	0.942028986
PTPRZ1 AND PCNXL4	0.99270073	1	0.985507246
PTPRZ1 AND SGCB	0.985294118	1	0.971014493
PTPRZ1 AND NOT-NDST4	0.977777778	1	0.956521739
PTPRZ1 AND NOT-SYNDIG1L	0.985294118	1	0.971014493
PTPRZ1 AND NOT-TMPRSS3	0.977777778	1	0.956521739
PTPRZ1 AND NOT-LRRC19	0.99270073	1	0.985507246
PTPRZ1 AND SLC30A5	0.985294118	1	0.971014493
PTPRZ1 AND NOT-SLAMF1	0.99270073	1	0.985507246
PTPRZ1 AND SLC3A2	0.985294118	1	0.971014493
PTPRZ1 AND NOT-SLC5A1	0.99270073	1	0.985507246
PTPRZ1 AND MPPE1	0.977777778	1	0.956521739
PTPRZ1 AND SLC5A3	0.985294118	1	0.971014493
PTPRZ1 AND NOT-SLC6A2	0.985294118	1	0.971014493
PTPRZ1 AND NOT-SLC6A4	0.985294118	1	0.971014493
PTPRZ1 AND NOT-SLC10A1	0.977777778	1	0.956521739
PTPRZ1 AND NOT-SLC18A2	0.985294118	1	0.971014493
PTPRZ1 AND SLC20A1	0.99270073	1	0.985507246
PTPRZ1 AND BMPR1A	0.970149254	1	0.942028986
PTPRZ1 AND NOT-SLC22A1	0.99270073	1	0.985507246
PTPRZ1 AND NOT-CYP4F12	0.977777778	1	0.956521739
PTPRZ1 AND BNIP2	0.99270073	1	0.985507246
PTPRZ1 AND SOAT1	0.977777778	1	0.956521739
PTPRZ1 AND NOT-SRD5A2	0.977777778	1	0.956521739
PTPRZ1 AND NOT-BPI	0.99270073	1	0.985507246
PTPRZ1 AND SSR2	0.970149254	1	0.942028986
PTPRZ1 AND SSR3	0.977777778	1	0.956521739
PTPRZ1 AND HSPA13	0.99270073	1	0.985507246
PTPRZ1 AND STX4	0.970149254	1	0.942028986
PTPRZ1 AND SURF4	0.99270073	1	0.985507246
PTPRZ1 AND VAMP7	0.970149254	1	0.942028986
PTPRZ1 AND NOT-SYP	0.970149254	1	0.942028986
PTPRZ1 AND TAPBP	0.985294118	1	0.971014493
PTPRZ1 AND TGFBI	0.977777778	1	0.956521739
PTPRZ1 AND NOT-ICAM5	0.99270073	1	0.985507246
PTPRZ1 AND NOT-TM4SF4	0.977777778	1	0.956521739
PTPRZ1 AND TMPO	0.99270073	1	0.985507246
PTPRZ1 AND NOT-TYR	0.985294118	1	0.971014493
PTPRZ1 AND NOT-UPK1B	0.977777778	1	0.956521739
PTPRZ1 AND NOT-UGT2B15	0.985294118	1	0.971014493
PTPRZ1 AND NOT-WNT7A	0.977777778	1	0.956521739
PTPRZ1 AND NOT-WNT9B	0.985294118	1	0.971014493
PTPRZ1 AND WRB	0.99270073	1	0.985507246
PTPRZ1 AND PTTG1IP	0.99270073	1	0.985507246
PTPRZ1 AND ZNF7	0.977777778	1	0.956521739
PTPRZ1 AND TMEM50B	0.99270073	1	0.985507246
PTPRZ1 AND ZNF138	0.977777778	1	0.956521739
PTPRZ1 AND NOT-SLC30A3	0.970149254	1	0.942028986
PTPRZ1 AND TVP23A	0.970149254	1	0.942028986

Antigen Pair Logic	F1	Precision	Recall
QSOX2 AND NOT-GLRA1	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-GP9	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-GPM6B	0.948717949	0.925	0.973684211
QSOX2 AND NOT-GPR4	0.948717949	0.925	0.973684211
QSOX2 AND NOT-FAM171A2	0.961038961	0.948717949	0.973684211
QSOX2 AND NOT-GPR20	0.918918919	0.944444444	0.894736842
QSOX2 AND NOT-CRB2	0.947368421	0.947368421	0.947368421
QSOX2 AND NOT-GPR39	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-TUSC5	0.96	0.972972973	0.947368421
QSOX2 AND NOT-GRID2	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-ANPEP	0.948717949	0.925	0.973684211
QSOX2 AND NOT-GRIK4	0.948717949	0.925	0.973684211
QSOX2 AND NOT-GRIN1	0.948717949	0.925	0.973684211
QSOX2 AND NOT-CNIH4	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-GRM3	0.948717949	0.925	0.973684211
QSOX2 AND NOT-GRM4	0.948717949	0.925	0.973684211
QSOX2 AND NOT-CYP2S1	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-GUCY2F	0.935064935	0.923076923	0.947368421
QSOX2 AND NOT-HFE	0.935064935	0.923076923	0.947368421
QSOX2 AND NOT-SLC29A2	0.933333333	0.945945946	0.921052632
QSOX2 AND NOT-B4GALNT4	0.935064935	0.923076923	0.947368421
QSOX2 AND NOT-MOGAT3	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-IL12RB2	0.948717949	0.925	0.973684211
QSOX2 AND NOT-INSRR	0.935064935	0.923076923	0.947368421
QSOX2 AND NOT-KCNC2	0.947368421	0.947368421	0.947368421
QSOX2 AND NOT-IGFL1	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-KCND3	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-KIR2DL4	0.947368421	0.947368421	0.947368421
QSOX2 AND NOT-KIR2DS4	0.947368421	0.947368421	0.947368421
QSOX2 AND NOT-KIR3DL2	0.973684211	0.973684211	0.973684211
QSOX2 AND NOT-LCAT	0.935064935	0.923076923	0.947368421
QSOX2 AND NOT-SPINK6	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-ERVFRD-1	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-MC5R	0.948717949	0.925	0.973684211
QSOX2 AND NOT-MOG	0.948717949	0.925	0.973684211
QSOX2 AND NOT-CD200	0.918918919	0.944444444	0.894736842
QSOX2 AND NOT-PTCHD4	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-MTHFD1	0.948717949	0.925	0.973684211
QSOX2 AND NOT-ATP1A4	0.948717949	0.925	0.973684211
QSOX2 AND NOT-NPY	0.935064935	0.923076923	0.947368421
QSOX2 AND NOT-NT5E	0.929577465	1	0.868421053
QSOX2 AND NOT-NUCB1	0.948717949	0.925	0.973684211
QSOX2 AND NOT-OPCML	0.935064935	0.923076923	0.947368421
QSOX2 AND NOT-OR3A1	0.935064935	0.923076923	0.947368421
QSOX2 AND NOT-CALY	0.933333333	0.945945946	0.921052632
QSOX2 AND NOT-KCNK4	0.961038961	0.948717949	0.973684211
QSOX2 AND NOT-TAS2R3	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-TAS2R16	0.935064935	0.923076923	0.947368421
QSOX2 AND NOT-TAS2R14	0.947368421	0.947368421	0.947368421
QSOX2 AND NOT-CLEC4A	0.931506849	0.971428571	0.894736842
QSOX2 AND NOT-CDON	0.935064935	0.923076923	0.947368421
QSOX2 AND NOT-PRLH	0.933333333	0.945945946	0.921052632
QSOX2 AND NOT-TMEM69	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-TEX264	0.935064935	0.923076923	0.947368421
QSOX2 AND NOT-AIG1	0.947368421	0.947368421	0.947368421
QSOX2 AND NOT-TRPV2	0.948717949	0.925	0.973684211
QSOX2 AND NOT-GDE1	0.948717949	0.925	0.973684211
QSOX2 AND NOT-NBAS	0.948717949	0.925	0.973684211
QSOX2 AND NOT-ATP5G3	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-PI3	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-PKHD1	0.948717949	0.925	0.973684211
QSOX2 AND NOT-IL20RA	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-NLGN3	0.948717949	0.925	0.973684211
QSOX2 AND NOT-TAS2R5	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-POR	0.948717949	0.925	0.973684211
QSOX2 AND NOT-QPCTL	0.948717949	0.925	0.973684211
QSOX2 AND NOT-BEST2	0.935064935	0.923076923	0.947368421
QSOX2 AND NOT-RETSAT	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-PIGX	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-SLC35F6	0.948717949	0.925	0.973684211
QSOX2 AND NOT-ACPP	0.935064935	0.923076923	0.947368421
QSOX2 AND NOT-SLC35C1	0.973684211	0.973684211	0.973684211
QSOX2 AND NOT-AVPR2	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-CHST12	0.95890411	1	0.921052632
QSOX2 AND NOT-POMGNT1	0.948717949	0.925	0.973684211
QSOX2 AND NOT-FLVCR2	0.948717949	0.925	0.973684211

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
PTPRZ1 AND NOT-BSND	0.985294118	1	0.971014493
PTPRZ1 AND DDR1	0.977777778	1	0.956521739
PTPRZ1 AND PXDN	0.970149254	1	0.942028986
PTPRZ1 AND ALG8	0.985294118	1	0.971014493
PTPRZ1 AND REEP5	0.99270073	1	0.985507246
PTPRZ1 AND ATG9A	0.970149254	1	0.942028986
PTPRZ1 AND APOO	0.99270073	1	0.985507246
PTPRZ1 AND DERL1	0.977777778	1	0.956521739
PTPRZ1 AND MFSD11	0.99270073	1	0.985507246
PTPRZ1 AND TMEM243	0.970149254	1	0.942028986
PTPRZ1 AND TMEM43	0.99270073	1	0.985507246
PTPRZ1 AND NOT-OR2H2	0.99270073	1	0.985507246
PTPRZ1 AND NOT-FCRL2	0.977777778	1	0.956521739
PTPRZ1 AND NOT-NOX5	0.99270073	1	0.985507246
PTPRZ1 AND C17orf62	0.985294118	1	0.971014493
PTPRZ1 AND ATP13A3	0.985294118	1	0.971014493
PTPRZ1 AND TTC13	0.99270073	1	0.985507246
PTPRZ1 AND ADIPOR2	0.985294118	1	0.971014493
PTPRZ1 AND RHBDF2	0.970149254	1	0.942028986
PTPRZ1 AND TMEM204	0.985294118	1	0.971014493
PTPRZ1 AND MANEA	0.970149254	1	0.942028986
PTPRZ1 AND NOT-C1orf115	0.985294118	1	0.971014493
PTPRZ1 AND TXNDC15	0.985294118	1	0.971014493
PTPRZ1 AND ST7	0.99270073	1	0.985507246
PTPRZ1 AND NOT-ZDHHC11	0.985294118	1	0.971014493
PTPRZ1 AND NOT-CALCR	0.970149254	1	0.942028986
PTPRZ1 AND UXS1	0.985294118	1	0.971014493
PTPRZ1 AND ORAI2	0.977777778	1	0.956521739
PTPRZ1 AND NOT-BPIFB2	0.99270073	1	0.985507246
PTPRZ1 AND CD276	0.985294118	1	0.971014493
PTPRZ1 AND NOT-SPX	0.970149254	1	0.942028986
PTPRZ1 AND MLF2	0.977777778	1	0.956521739
PTPRZ1 AND CLPTM1L	0.99270073	1	0.985507246
PTPRZ1 AND SGPP1	0.977777778	1	0.956521739
PTPRZ1 AND TMX1	0.985294118	1	0.971014493
PTPRZ1 AND YIPF5	0.99270073	1	0.985507246
PTPRZ1 AND APOLD1	0.977777778	1	0.956521739
PTPRZ1 AND NOT-PVRL4	0.99270073	1	0.985507246
PTPRZ1 AND NIPA2	0.977777778	1	0.956521739
PTPRZ1 AND NOT-ACSBG2	0.985294118	1	0.971014493
PTPRZ1 AND NOT-MADCAM1	0.970149254	1	0.942028986
PTPRZ1 AND NOT-OR12D3	0.970149254	1	0.942028986
PTPRZ1 AND NOT-SPACA1	0.977777778	1	0.956521739
PTPRZ1 AND CANX	0.970149254	1	0.942028986
PTPRZ1 AND CAPZA2	0.970149254	1	0.942028986
PTPRZ1 AND DDX59	0.99270073	1	0.985507246
PTPRZ1 AND ARMC10	0.977777778	1	0.956521739
PTPRZ1 AND TM2D2	0.985294118	1	0.971014493
PTPRZ1 AND STARD3NL	0.99270073	1	0.985507246
PTPRZ1 AND SPNS1	0.977777778	1	0.956521739
PTPRZ1 AND ITFG3	0.985294118	1	0.971014493
PTPRZ1 AND ANTXR1	0.977777778	1	0.956521739
PTPRZ1 AND FAR1	0.985294118	1	0.971014493
PTPRZ1 AND TMEM126A	0.977777778	1	0.956521739
PTPRZ1 AND SLC37A3	0.977777778	1	0.956521739
PTPRZ1 AND JAGN1	0.977777778	1	0.956521739
PTPRZ1 AND ZNF559	0.977777778	1	0.956521739
PTPRZ1 AND HIATL1	0.985294118	1	0.971014493
PTPRZ1 AND SLC35B4	0.977777778	1	0.956521739
PTPRZ1 AND CIRH1A	0.985294118	1	0.971014493
PTPRZ1 AND COX14	0.985294118	1	0.971014493
PTPRZ1 AND TMEM128	0.977777778	1	0.956521739
PTPRZ1 AND TMEM60	0.970149254	1	0.942028986
PTPRZ1 AND DEGS1	0.99270073	1	0.985507246
PTPRZ1 AND CAV1	0.977777778	1	0.956521739
PTPRZ1 AND NOT-CAV3	0.977777778	1	0.956521739
PTPRZ1 AND PPAP2A	0.977777778	1	0.956521739
PTPRZ1 AND VAMP8	0.977777778	1	0.956521739
PTPRZ1 AND B4GALT4	0.970149254	1	0.942028986
PTPRZ1 AND GPAA1	0.977777778	1	0.956521739
PTPRZ1 AND NOT-TNFSF9	0.985294118	1	0.971014493
PTPRZ1 AND NOT-ADAM21	0.970149254	1	0.942028986
PTPRZ1 AND NOT-ADAM18	0.977777778	1	0.956521739
PTPRZ1 AND CDS2	0.970149254	1	0.942028986
PTPRZ1 AND CD164	0.985294118	1	0.971014493
PTPRZ1 AND PEX11B	0.985294118	1	0.971014493

Antigen Pair Logic	F1	Precision	Recall
QSOX2 AND NOT-AJAP1	0.947368421	0.947368421	0.947368421
QSOX2 AND NOT-PCDHA10	0.947368421	0.947368421	0.947368421
QSOX2 AND NOT-PRRG2	0.918918919	0.944444444	0.894736842
QSOX2 AND NOT-SUCNR1	0.948717949	0.925	0.973684211
QSOX2 AND NOT-NDUFA4L2	0.948717949	0.925	0.973684211
QSOX2 AND NOT-ANO2	0.935064935	0.923076923	0.947368421
QSOX2 AND NOT-TRPC7	0.935064935	0.923076923	0.947368421
QSOX2 AND NOT-PTGER1	0.933333333	0.945945946	0.921052632
QSOX2 AND NOT-PTGER4	0.948717949	0.925	0.973684211
QSOX2 AND NOT-LRTM1	0.933333333	0.945945946	0.921052632
QSOX2 AND NOT-MAVS	0.935064935	0.923076923	0.947368421
QSOX2 AND NOT-ISLR2	0.948717949	0.925	0.973684211
QSOX2 AND NOT-PTPRH	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-PVRL1	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-PRPH2	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-RHCE	0.973684211	0.973684211	0.973684211
QSOX2 AND NOT-ROM1	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-SLC39A8	0.948717949	0.925	0.973684211
QSOX2 AND NOT-POPDC3	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-NDST4	0.948717949	0.925	0.973684211
QSOX2 AND NOT-SHH	0.935064935	0.923076923	0.947368421
QSOX2 AND NOT-IL25	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-CYP3A43	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-ST3GAL2	0.935064935	0.923076923	0.947368421
QSOX2 AND NOT-PORCN	0.961038961	0.948717949	0.973684211
QSOX2 AND NOT-SLC1A5	0.948717949	0.925	0.973684211
QSOX2 AND NOT-SLC5A2	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-SLC6A2	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-LRTM2	0.935064935	0.923076923	0.947368421
QSOX2 AND NOT-SLC9A2	0.935064935	0.923076923	0.947368421
QSOX2 AND NOT-SLC9A3	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-SMPD2	0.916666667	0.970588235	0.868421053
QSOX2 AND NOT-SORL1	0.935064935	0.923076923	0.947368421
QSOX2 AND NOT-SOX1	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-SSTR5	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-TRPC5	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-UPK1B	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-CACNA1E	0.935064935	0.923076923	0.947368421
QSOX2 AND NOT-FZD5	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-VKORC1	0.948717949	0.925	0.973684211
QSOX2 AND NOT-ALG12	0.948717949	0.925	0.973684211
QSOX2 AND NOT-SLC52A2	0.935064935	0.923076923	0.947368421
QSOX2 AND NOT-DNAJC22	0.948717949	0.925	0.973684211
QSOX2 AND NOT-NRSN2	0.935064935	0.923076923	0.947368421
QSOX2 AND NOT-TRPM3	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-MLLT10	0.931506849	0.971428571	0.894736842
QSOX2 AND NOT-PDCD1LG2	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-C6orf25	0.973684211	0.973684211	0.973684211
QSOX2 AND NOT-LY6G6C	0.918918919	0.944444444	0.894736842
QSOX2 AND NOT-TMEM121	0.948717949	0.925	0.973684211
QSOX2 AND NOT-MLF2	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-KCNH6	0.933333333	0.945945946	0.921052632
QSOX2 AND NOT-ST8SIA2	0.935064935	0.923076923	0.947368421
QSOX2 AND NOT-OR51E2	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-HM13	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-LRRC3	0.947368421	0.947368421	0.947368421
QSOX2 AND NOT-AKAP1	0.948717949	0.925	0.973684211
QSOX2 AND NOT-SLC10A3	0.918918919	0.944444444	0.894736842
QSOX2 AND NOT-GSG1	0.948717949	0.925	0.973684211
QSOX2 AND NOT-FAM57B	0.947368421	0.947368421	0.947368421
QSOX2 AND NOT-SLC25A18	0.935064935	0.923076923	0.947368421
QSOX2 AND NOT-GPR61	0.948717949	0.925	0.973684211
QSOX2 AND NOT-OR1G1	0.948717949	0.925	0.973684211
QSOX2 AND NOT-ADGRV1	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-LRRC8C	0.935064935	0.923076923	0.947368421
QSOX2 AND NOT-YIPF4	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-MEGF11	0.961038961	0.948717949	0.973684211
QSOX2 AND NOT-CASR	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-TPST1	0.933333333	0.945945946	0.921052632
QSOX2 AND NOT-DGAT2	0.945945946	0.972222222	0.921052632
QSOX2 AND NOT-ADGRE3	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-PIGO	0.948717949	0.925	0.973684211
QSOX2 AND NOT-DEGS1	0.933333333	0.945945946	0.921052632
QSOX2 AND NOT-PPAP2C	0.935064935	0.923076923	0.947368421
QSOX2 AND NOT-TNFSF14	0.948717949	0.925	0.973684211
QSOX2 AND NOT-IL1RL2	0.948717949	0.925	0.973684211

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
PTPRZ1 AND NOT-TRPA1	0.977777778	1	0.956521739
PTPRZ1 AND ATP6V0E1	0.977777778	1	0.956521739
PTPRZ1 AND PKD2L1	0.99270073	1	0.985507246
PTPRZ1 AND TMEM263	0.970149254	1	0.942028986
PTPRZ1 AND NOT-DUOXA1	0.970149254	1	0.942028986
PTPRZ1 AND CEP95	0.985294118	1	0.971014493
PTPRZ1 AND NOT-CD1A	0.99270073	1	0.985507246
PTPRZ1 AND NOT-CD1B	0.977777778	1	0.956521739
PTPRZ1 AND SLC39A13	0.985294118	1	0.971014493
PTPRZ1 AND AIFM1	0.985294118	1	0.971014493
PTPRZ1 AND NOT-HTR3B	0.985294118	1	0.971014493
PTPRZ1 AND NOT-CD3G	0.977777778	1	0.956521739
PTPRZ1 AND NOT-TIMD4	0.99270073	1	0.985507246
PTPRZ1 AND NOT-GGTLC1	0.99270073	1	0.985507246
PTPRZ1 AND XPR1	0.970149254	1	0.942028986
PTPRZ1 AND MTDH	0.977777778	1	0.956521739
PTPRZ1 AND NOT-CD5	0.985294118	1	0.971014493
PTPRZ1 AND NOT-GCNT3	0.99270073	1	0.985507246
PTPRZ1 AND G6PC3	0.985294118	1	0.971014493
PTPRZ1 AND DTD1	0.970149254	1	0.942028986
PTPRZ1 AND NOT-CD8B	0.99270073	1	0.985507246
PTPRZ1 AND NOT-TAAR2	0.99270073	1	0.985507246
PTPRZ1 AND ORAI3	0.970149254	1	0.942028986
PTPRZ1 AND PIGM	0.99270073	1	0.985507246
PTPRZ1 AND NOT-MS4A1	0.977777778	1	0.956521739
PTPRZ1 AND GTF3C3	0.977777778	1	0.956521739
PTPRZ1 AND B4GALT5	0.985294118	1	0.971014493
PTPRZ1 AND NOT-OPALIN	0.985294118	1	0.971014493
PTPRZ1 AND NOT-CD22	0.985294118	1	0.971014493
PTPRZ1 AND NOT-GLP2R	0.970149254	1	0.942028986
PTPRZ1 AND SFXN1	0.977777778	1	0.956521739
PTPRZ1 AND TMEM203	0.985294118	1	0.971014493
PTPRZ1 AND CDYL	0.977777778	1	0.956521739
PTPRZ1 AND NOT-SIGLEC6	0.99270073	1	0.985507246
PTPRZ1 AND STX8	0.970149254	1	0.942028986
PTPRZ1 AND CHST10	0.970149254	1	0.942028986
PTPRZ1 AND EEF1E1	0.985294118	1	0.971014493
PTPRZ1 AND EI24	0.985294118	1	0.971014493
PTPRZ1 AND ATP5J2	0.99270073	1	0.985507246
PTPRZ1 AND PDIA4	0.985294118	1	0.971014493
PTPRZ1 AND CD63	0.99270073	1	0.985507246
PTPRZ1 AND HERPUD1	0.985294118	1	0.971014493
PTPRZ1 AND LAPTM4A	0.99270073	1	0.985507246
PTPRZ1 AND CD81	0.985294118	1	0.971014493
PTPRZ1 AND SUSD6	0.977777778	1	0.956521739
PTPRZ1 AND ADGRE5	0.99270073	1	0.985507246
PTPRZ1 AND PTDSS1	0.99270073	1	0.985507246
PTPRZ1 AND TOMM70A	0.977777778	1	0.956521739
PTPRZ1 AND NOT-ATP2C2	0.970149254	1	0.942028986
PTPRZ1 AND HELZ	0.977777778	1	0.956521739
PTPRZ1 AND NOT-SLC23A1	0.985294118	1	0.971014493
CHST10 AND NOT-SCN2B	0.970149254	1	0.942028986
CHST10 AND NOT-VAMP1	0.970149254	1	0.942028986
CHST10 AND NOT-UGT2B15	0.970149254	1	0.942028986
CNTNAP1 AND WRB	0.970149254	1	0.942028986
CHST10 AND NOT-SIGLEC6	0.970149254	1	0.942028986
CHST10 AND CD63	0.970149254	1	0.942028986
CHST10 AND LAPTM4A	0.970149254	1	0.942028986
Carcinoma, Ductal, Breast (Breast C)			
CACFD1 AND NOT-TMED6	0.901960784	0.95833333	0.851851852
CACFD1 AND NOT-TMEM61	0.9	0.97826087	0.833333333
CACFD1 AND NOT-CDHR2	0.901960784	0.95833333	0.851851852
NOT-TMEM246 AND CACFD1	0.907407407	0.90740741	0.907407407
CACFD1 AND NOT-SLC4A4	0.909090909	1	0.833333333
NOT-SLC4A4 AND C10orf35	0.914285714	0.94117647	0.888888889
RNF121 AND NOT-PTCH1	0.903846154	0.94	0.87037037
Carcinoma, Renal Cell (Renal)			
NOT-SCARA5 AND CAV2	0.909090909	1	0.833333333
Carcinoma, Pancreatic Ductal (Pancreas)			
CTRB2 AND NOT-CDH18	0.9	1	0.818181818
TM4SF4 AND NOT-ABCC2	0.909090909	0.90909091	0.909090909
TM4SF4 AND NOT-TMEM56	0.909090909	0.90909091	0.909090909
TM4SF4 AND NOT-CYP3A7	0.909090909	0.90909091	0.909090909
TM4SF4 AND NOT-CYP2J2	0.909090909	0.90909091	0.909090909
TM4SF4 AND NOT-CLYBL	0.909090909	0.90909091	0.909090909
TM4SF4 AND NOT-SLC35D1	0.909090909	0.90909091	0.909090909

Antigen Pair Logic	F1	Precision	Recall
QSOX2 AND NOT-GALR2	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-CACNA1I	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-KCNH7	0.947368421	0.947368421	0.947368421
QSOX2 AND NOT-NAT8	0.947368421	0.947368421	0.947368421
QSOX2 AND NOT-FER1L5	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-YIF1B	0.948717949	0.925	0.973684211
QSOX2 AND NOT-CLDN2	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-SLC6A5	0.929577465	1	0.868421053
QSOX2 AND NOT-SLC28A1	0.948717949	0.925	0.973684211
QSOX2 AND NOT-SYT12	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-LMBRD2	0.948717949	0.925	0.973684211
QSOX2 AND NOT-LEMD1	0.948717949	0.925	0.973684211
QSOX2 AND NOT-OPALIN	0.947368421	0.947368421	0.947368421
QSOX2 AND NOT-NDST3	0.935064935	0.923076923	0.947368421
QSOX2 AND NOT-ACPT	0.948717949	0.925	0.973684211
QSOX2 AND NOT-NRXN1	0.935064935	0.923076923	0.947368421
QSOX2 AND NOT-SLC22A13	0.935064935	0.923076923	0.947368421
QSOX2 AND NOT-ABCG2	0.936708861	0.902439024	0.973684211
QSOX2 AND NOT-TNFRSF8	0.947368421	0.947368421	0.947368421
QSOX2 AND NOT-TRIL	0.936708861	0.902439024	0.973684211
PTPN1 AND NOT-PTCRA	0.974358974	0.95	1
CD83 AND NOT-PTCRA	0.947368421	0.947368421	0.947368421
CELSR1 AND NOT-EDNRA	0.935064935	0.923076923	0.947368421
CELSR1 AND NOT-EFNA5	0.918918919	0.944444444	0.894736842
SNN AND NOT-EGF	0.918918919	0.944444444	0.894736842
CELSR1 AND NOT-EMP1	0.947368421	0.947368421	0.947368421
NOT-MGST1 AND ETV6	0.947368421	0.947368421	0.947368421
PTPN1 AND NOT-MS4A2	0.974358974	0.95	1
ORAI2 AND NOT-MS4A2	0.986666667	1	0.973684211
NOT-FGFR2 AND CELSR1	0.936708861	0.902439024	0.973684211
PTPN1 AND NOT-DOLK	0.987012987	0.974358974	1
SPG7 AND NOT-DOLK	0.933333333	0.945945946	0.921052632
TMEM55B AND NOT-DOLK	0.936708861	0.902439024	0.973684211
CELSR1 AND NOT-DOLK	0.935064935	0.923076923	0.947368421
NOT-TEX2 AND ANKLE2	0.96	0.972972973	0.947368421
NOT-ELOVL6 AND ANKLE2	0.948717949	0.925	0.973684211
NOT-AGPAT9 AND ATP11B	0.918918919	0.944444444	0.894736842
NUP210 AND NOT-KIR3DL2	0.918918919	0.944444444	0.894736842
NUP210 AND NOT-TRPV2	0.944444444	1	0.894736842
PTPN1 AND NOT-FMO5	0.918918919	0.944444444	0.894736842
PTPN1 AND NOT-FPR2	0.948717949	0.925	0.973684211
PTPN1 AND NOT-CLEC5A	0.926829268	0.863636364	1
PTPN1 AND NOT-SDF2L1	0.933333333	0.945945946	0.921052632
PTPN1 AND NOT-FUT5	0.926829268	0.863636364	1
NOT-MYOF AND ADAM17	0.962025316	0.926829268	1
NOT-MGST1 AND ST6GALNAC4	0.933333333	0.945945946	0.921052632
NOT-ACSS2 AND ST6GALNAC4	0.916666667	0.970588235	0.868421053
NOT-SLC22A4 AND ST6GALNAC4	0.973684211	0.973684211	0.973684211
SNN AND NOT-CECR6	0.95890411	1	0.921052632
CELSR1 AND NOT-PCLO	0.947368421	0.947368421	0.947368421
NOT-GPM6B AND CELSR1	0.961038961	0.948717949	0.973684211
SPG7 AND NOT-CCR10	0.916666667	0.970588235	0.868421053
NOT-ELOVL6 AND SLC6A16	0.938271605	0.88372093	1
NOT-ANPEP AND SLC37A1	0.936708861	0.902439024	0.973684211
NOT-CNIH4 AND SLC37A1	0.987012987	0.974358974	1
SPG7 AND NOT-CNIH4	0.933333333	0.945945946	0.921052632
CELSR1 AND NOT-SLC39A2	0.933333333	0.945945946	0.921052632
CELSR1 AND NOT-IGSF3	0.935064935	0.923076923	0.947368421
ICAM3 AND NOT-SORL1	0.947368421	0.947368421	0.947368421
ORAI2 AND NOT-ICAM4	0.931506849	0.971428571	0.894736842
SPG7 AND NOT-SERINC2	0.916666667	0.970588235	0.868421053
SLC37A1 AND NOT-IL6R	0.938271605	0.88372093	1
IL10RA AND NOT-TRPV2	0.974358974	0.95	1
IL10RA AND NOT-CHST12	0.933333333	0.945945946	0.921052632
IL10RA AND NOT-SORL1	0.945945946	0.972222222	0.921052632
ADAM17 AND NOT-ITGA5	0.948717949	0.925	0.973684211
CELSR1 AND NOT-IGFL1	0.933333333	0.945945946	0.921052632
NRROS AND NOT-MARCO	0.929577465	1	0.868421053
KCNN4 AND NOT-VKORC1	0.936708861	0.902439024	0.973684211
SIDT1 AND NOT-KIR3DL2	0.95	0.904761905	1
CXCR4 AND NOT-KIR3DL2	0.926829268	0.863636364	1
ORAI2 AND NOT-KIR3DL2	0.961038961	0.948717949	0.973684211
CELSR1 AND NOT-SAMD5	0.933333333	0.945945946	0.921052632
SNN AND NOT-LRP3	0.933333333	0.945945946	0.921052632
NOT-LTBR AND SLC37A1	0.926829268	0.863636364	1
NOT-KITLG AND CELSR1	0.936708861	0.902439024	0.973684211

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
TM4SF4 AND NOT-G6PC	0.909090909	0.90909091	0.909090909
TM4SF4 AND GJA1	0.9	1	0.818181818
CTRB2 AND NPTN	0.9	1	0.818181818
TM4SF4 AND NOT-ABCC6	0.909090909	0.90909091	0.909090909
CTRB2 AND NOT-KCNN2	0.9	1	0.818181818
CTRB2 AND RTN4	0.9	1	0.818181818
CTRB2 AND NOT-TMEM63C	0.9	1	0.818181818
CTRB2 AND SDCBP	0.9	1	0.818181818
TM4SF4 AND NOT-DNAJC22	0.909090909	0.90909091	0.909090909
TM4SF4 AND NOT-NAT8	0.909090909	0.90909091	0.909090909
TM4SF4 AND NOT-SLC28A1	0.909090909	0.90909091	0.909090909
TM4SF4 AND NOT-ENTPD5	0.952380952	1	0.909090909
TM4SF4 AND CD63	0.9	1	0.818181818
TM4SF4 AND NOT-SLC23A1	0.909090909	0.90909091	0.909090909
Melanoma (Melanoma)			
MLANA	1	1	1
PMEL	0.952380952	1	0.909090909
MLANA	1	1	1
TMEM147	0.9	1	0.818181818
CIRH1A	0.952380952	1	0.909090909
ROMO1	0.952380952	1	0.909090909
MANBAL	0.952380952	1	0.909090909
FDPS	1	1	1
SFXN3	0.952380952	1	0.909090909
OS9	0.952380952	1	0.909090909
ERGIC3	0.9	1	0.818181818
TMEM138	0.952380952	1	0.909090909
NOT-FCN2 AND DBI	0.909090909	0.90909091	0.909090909
OS9 AND OS9	0.9	1	0.818181818
KIAA1549L AND DBI	0.9	1	0.818181818
MLANA AND MLANA	1	1	1
TMED3 AND DBI	0.9	1	0.818181818
SFXN3 AND NOT-DBI	0.952380952	1	0.909090909
GPR19 AND DBI	0.9	1	0.818181818
NOT-ITGA2B AND DBI	0.952380952	1	0.909090909
NOT-VSTM1 AND DBI	0.9	1	0.818181818
BNIP1 AND DBI	0.909090909	0.90909091	0.909090909
NOT-CD300LB AND DBI	0.952380952	1	0.909090909
MANBAL AND DBI	0.952380952	1	0.909090909
TMEM138 AND DBI	0.9	1	0.818181818
NPC1 AND DBI	0.952380952	1	0.909090909
NOT-CIRH1A AND CIRH1A	0.952380952	1	0.909090909
NOT-ERGIC3 AND ERGIC3	0.9	1	0.818181818
CTSA AND DBI	0.9	1	0.818181818
ADIPOR2 AND DBI	0.9	1	0.818181818
SLC5A6 AND DBI	0.952380952	1	0.909090909
MC1R AND DBI	0.9	1	0.818181818
TMEM147 AND TMEM147	0.9	1	0.818181818
EXT1 AND DBI	0.909090909	0.90909091	0.909090909
TNFSF9 AND DBI	0.952380952	1	0.909090909
C11orf24 AND DBI	0.9	1	0.818181818
NOT-PMEL AND PMEL	0.952380952	1	0.909090909
BCAP31 AND DBI	0.952380952	1	0.909090909
FDPS AND DBI	0.952380952	1	0.909090909
PNKD AND DBI	0.952380952	1	0.909090909
ROMO1 AND NOT-ROMO1	0.952380952	1	0.909090909
NAT14 AND DBI	0.9	1	0.818181818
MLANA AND BNIP3L	1	1	1
TMEM138 AND TMEM138	0.952380952	1	0.909090909
CIRH1A AND NOT-BNIP3L	0.9	1	0.818181818
NOT-PMEL AND PMEL	0.952380952	1	0.909090909
FDPS AND NOT-BNIP3L	0.952380952	1	0.909090909
ROMO1 AND NOT-BNIP3L	0.9	1	0.818181818
DENND5A AND ENPP2	0.9	1	0.818181818
NOT-TAAR8 AND ENPP2	0.956521739	0.91666667	1
MLANA AND MLANA	1	1	1
NOT-NNAT AND ENPP2	0.9	1	0.818181818
NOT-TM4SF5 AND ENPP2	1	1	1
NOT-WSCD2 AND ENPP2	0.952380952	1	0.909090909
NOT-SERPINA9 AND ENPP2	0.952380952	1	0.909090909
GPR137B AND ENPP2	0.956521739	0.91666667	1
NOT-TMEM95 AND ENPP2	1	1	1
NOT-ABCC8 AND ENPP2	0.9	1	0.818181818
NOT-C2orf82 AND ENPP2	0.952380952	1	0.909090909
CIRH1A AND ENPP2	0.952380952	1	0.909090909
NOT-SLC5A5 AND ENPP2	1	1	1

Antigen Pair Logic	F1	Precision	Recall
NOT-MGST1 AND ADAM17	0.947368421	0.947368421	0.947368421
CELSR1 AND NOT-MST1R	0.944444444	1	0.894736842
SNN AND NOT-NCAM1	0.944444444	1	0.894736842
P2RX5 AND NOT-SORL1	0.918918919	0.944444444	0.894736842
NOT-PCSK5 AND SNN	0.947368421	0.947368421	0.947368421
NOT-FAM198B AND ADAM17	0.936708861	0.902439024	0.973684211
NOT-FAM198B AND SNRNP40	0.961038961	0.948717949	0.973684211
PTPN1 AND NOT-TRPV2	0.974358974	0.95	1
CELSR1 AND NOT-DUOX1	0.935064935	0.923076923	0.947368421
SLC37A1 AND NOT-ST3GAL4	0.918918919	0.944444444	0.894736842
SLC37A1 AND NOT-SORL1	0.914285714	1	0.842105263
SLC37A1 AND NOT-HM13	0.926829268	0.863636364	1
NOT-AGPAT9 AND SLC37A1	0.926829268	0.863636364	1
CELSR1 AND NOT-LY6K	0.933333333	0.945945946	0.921052632
SPG7 AND NOT-TRPM4	0.944444444	1	0.894736842
NOT-TRPM4 AND CELSR1	0.95890411	1	0.921052632
SIDT1 AND NOT-SORL1	0.936708861	0.902439024	0.973684211
NOT-LPCAT2 AND ADAM17	0.926829268	0.863636364	1
NOT-ELOVL6 AND BANP	0.935064935	0.923076923	0.947368421
CELSR1 AND NOT-TMEM51	0.96	0.972972973	0.947368421
CELSR1 AND NOT-ANO1	0.918918919	0.944444444	0.894736842
NOT-ANO10 AND CELSR1	0.974358974	0.95	1
CELSR1 AND NOT-SPTLC3	0.935064935	0.923076923	0.947368421
PTPN1 AND NOT-SLC35C1	0.947368421	0.947368421	0.947368421
RHOT2 AND NOT-SLC35C1	0.933333333	0.945945946	0.921052632
PTPN1 AND NOT-CHST12	0.933333333	0.945945946	0.921052632
NOT-ELOVL6 AND NGLY1	0.961038961	0.948717949	0.973684211
NOT-TEX2 AND SPG7	0.973684211	0.973684211	0.973684211
NOT-TEX2 AND ST14	0.931506849	0.971428571	0.894736842
NOT-TEX2 AND CELSR1	0.962025316	0.926829268	1
NOT-ACSS2 AND ERGIC1	0.918918919	0.944444444	0.894736842
NOT-ACSS2 AND ADAM17	0.933333333	0.945945946	0.921052632
NOT-FAM20C AND ERGIC1	0.933333333	0.945945946	0.921052632
NOT-FAM20C AND ADAM17	0.961038961	0.948717949	0.973684211
SNN AND NOT-FAM20C	0.945945946	0.972222222	0.921052632
NOT-FAM20C AND SNRNP40	0.973684211	0.973684211	0.973684211
NOT-CRISPLD2 AND ERGIC1	0.935064935	0.923076923	0.947368421
NOT-MTUS1 AND CELSR1	0.936708861	0.902439024	0.973684211
CELSR1 AND NOT-PTN	0.947368421	0.947368421	0.947368421
PTPN1 AND NOT-C5AR1	0.926829268	0.863636364	1
PTPN1 AND NOT-VKORC1	0.962025316	0.926829268	1
NOT-SLC46A2 AND ADAM17	0.948717949	0.925	0.973684211
CELSR1 AND NOT-PVR	0.933333333	0.945945946	0.921052632
CELSR1 AND NOT-SCNN1B	0.935064935	0.923076923	0.947368421
NOT-SLC6A8 AND ST14	0.935064935	0.923076923	0.947368421
NOT-SLC6A8 AND CELSR1	0.986666667	1	0.973684211
NOT-SLC18A2 AND ADAM17	0.938271605	0.88372093	1
NOT-SLC22A4 AND ADAM17	0.962025316	0.926829268	1
NOT-SLC22A4 AND SNRNP40	0.938271605	0.88372093	1
CELSR1 AND NOT-SLC22A5	0.914285714	1	0.842105263
CD1D AND NOT-SORL1	0.931506849	0.971428571	0.894736842
SPG7 AND NOT-C6orf25	0.933333333	0.945945946	0.921052632
NOT-ELOVL6 AND ST14	0.96	0.972972973	0.947368421
CELSR1 AND NOT-ELOVL4	0.916666667	0.970588235	0.868421053
NOT-CRISPLD2 AND ADAM17	0.938271605	0.88372093	1
NOT-B3GNT5 AND ADAM17	0.901408451	0.96969697	0.842105263
NOT-PLXDC2 AND ADAM17	0.947368421	0.947368421	0.947368421
ADAM17 AND NOT-MARCO	0.916666667	0.970588235	0.868421053
CELSR1 AND NOT-TMPRSS2	0.901408451	0.96969697	0.842105263
NOT-XK AND CELSR1	0.931506849	0.971428571	0.894736842
CELSR1 AND NOT-CA12	0.947368421	0.947368421	0.947368421
CXCR4 AND CELSR1	0.936708861	0.902439024	0.973684211
NOT-FZD5 AND CELSR1	0.973684211	0.973684211	0.973684211
ORAI2 AND NOT-ELOVL6	0.961038961	0.948717949	0.973684211
NOT-ELOVL6 AND CELSR1	0.974358974	0.95	1
NOT-C2orf40 AND FKRP	0.938271605	0.88372093	1
NOT-RNF128 AND CELSR1	0.936708861	0.902439024	0.973684211
CELSR1 AND NOT-ERMP1	0.918918919	0.944444444	0.894736842
NOT-CRISPLD2 AND SNRNP40	0.962025316	0.926829268	1
CELSR1 AND NOT-KREMEN1	0.935064935	0.923076923	0.947368421
CELSR1 AND NOT-FAM213A	0.931506849	0.971428571	0.894736842
NOT-TMEM246 AND CELSR1	0.947368421	0.947368421	0.947368421
NOT-TMEM101 AND SNRNP40	0.933333333	0.945945946	0.921052632
NOT-PLXDC2 AND SNRNP40	0.935064935	0.923076923	0.947368421
NOT-LRP11 AND CELSR1	0.938271605	0.88372093	1
NOT-SMIM3 AND SNRNP40	0.914285714	1	0.842105263

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
NOT-SLC34A2 AND ENPP2	1	1	1
NOT-PTH2 AND ENPP2	1	1	1
AUP1 AND ENPP2	1	1	1
NOT-VTCN1 AND ENPP2	0.9	1	0.818181818
NOT-CLDN19 AND ENPP2	1	1	1
C14orf2 AND ENPP2	0.9	1	0.818181818
NOT-TMEM235 AND ENPP2	0.952380952	1	0.909090909
MC1R AND NOT-MC1R	0.9	1	0.818181818
TMEM147 AND ENPP2	0.9	1	0.818181818
NOT-ATP13A5 AND ENPP2	0.9	1	0.818181818
NOT-NTRK3 AND ENPP2	0.9	1	0.818181818
NOT-PTCH2 AND ENPP2	0.956521739	0.91666667	1
NOT-PMEL AND PMEL	0.952380952	1	0.909090909
NOT-C8B AND ENPP2	0.952380952	1	0.909090909
NOT-AVPR1B AND ENPP2	1	1	1
NOT-KCNB1 AND ENPP2	0.956521739	0.91666667	1
NOT-FCN2 AND POLG2	0.9	1	0.818181818
MLANA AND POLG2	0.9	1	0.818181818
CIRH1A AND POLG2	0.952380952	1	0.909090909
SLC45A2 AND NOT-POLG2	0.952380952	1	0.909090909
NOT-FCN2 AND DDB2	1	1	1
OS9 AND OS9	0.9	1	0.818181818
NOT-MLANA AND DDB2	0.916666667	0.84615385	1
TMED3 AND DDB2	0.9	1	0.818181818
SFXN3 AND NOT-DDB2	0.952380952	1	0.909090909
NOT-CD300LB AND DDB2	1	1	1
MANBAL AND DDB2	0.952380952	1	0.909090909
APMAP AND DDB2	0.909090909	0.90909091	0.909090909
TMEM138 AND DDB2	0.952380952	1	0.909090909
NOT-CIRH1A AND CIRH1A	0.952380952	1	0.909090909
ERGIC3 AND NOT-DDB2	0.9	1	0.818181818
SLC45A2 AND NOT-DDB2	0.909090909	0.90909091	0.909090909
NOT-GP6 AND DDB2	0.9	1	0.818181818
MC1R AND NOT-MC1R	0.9	1	0.818181818
EDNRB AND DDB2	0.9	1	0.818181818
TMEM147 AND TMEM147	0.9	1	0.818181818
NOT-PMEL AND PMEL	0.952380952	1	0.909090909
BCAP31 AND DDB2	0.952380952	1	0.909090909
FDPS AND DDB2	1	1	1
ROMO1 AND DDB2	0.952380952	1	0.909090909
STARD3NL AND DDB2	0.9	1	0.818181818
MLANA AND BSG	1	1	1
NOT-CIRH1A AND CIRH1A	0.952380952	1	0.909090909
SLC45A2 AND NOT-BSG	0.9	1	0.818181818
MC1R AND NOT-MC1R	0.9	1	0.818181818
TMEM147 AND TMEM147	0.9	1	0.818181818
NOT-PMEL AND PMEL	0.952380952	1	0.909090909
NOT-SCN4A AND CDKN3	0.952380952	1	0.909090909
MLANA AND CDKN3	1	1	1
NOT-ISM2 AND CDKN3	0.952380952	1	0.909090909
SLC45A2 AND NOT-CDKN3	0.952380952	1	0.909090909
NOT-PMEL AND PMEL	0.952380952	1	0.909090909
NDUFA1 AND CDKN3	0.9	1	0.818181818
STARD3NL AND CDKN3	0.9	1	0.818181818
OS9 AND OS9	0.9	1	0.818181818
KIAA1549L AND NOT-CSH1	0.952380952	1	0.909090909
MLANA AND NOT-CSH1	1	1	1
SFXN3 AND CSH1	0.9	1	0.818181818
GPR19 AND NOT-CSH1	0.909090909	0.90909091	0.909090909
PPT2 AND NOT-CSH1	0.9	1	0.818181818
BNIP1 AND NOT-CSH1	0.952380952	1	0.909090909
CLCN5 AND NOT-CSH1	0.916666667	0.84615385	1
NSDHL AND NOT-CSH1	0.9	1	0.818181818
TMEM138 AND TMEM138	0.952380952	1	0.909090909
LSAMP AND NOT-CSH1	0.9	1	0.818181818
NPC1 AND NOT-CSH1	1	1	1
CIRH1A AND NOT-CSH1	0.952380952	1	0.909090909
NOT-ERGIC3 AND ERGIC3	0.9	1	0.818181818
NOT-CSH1 AND MMP15	0.9	1	0.818181818
GPR107 AND NOT-CSH1	0.9	1	0.818181818
SDC3 AND NOT-CSH1	0.9	1	0.818181818
SLC45A2 AND NOT-CSH1	0.952380952	1	0.909090909
SLC5A6 AND NOT-CSH1	0.952380952	1	0.909090909
MC1R AND NOT-CSH1	0.9	1	0.818181818
TMEM147 AND NOT-CSH1	0.9	1	0.818181818
EXT1 AND NOT-CSH1	0.952380952	1	0.909090909

Antigen Pair Logic	F1	Precision	Recall
CELSR1 AND NOT-PPAP2C	0.935064935	0.923076923	0.947368421
CELSR1 AND NOT-PEX11A	0.96	0.972972973	0.947368421
CELSR1 AND NOT-CLDN8	0.945945946	0.972222222	0.921052632
MS4A1 AND CELSR1	0.936708861	0.902439024	0.973684211
NOT-B4GALT6 AND SNRNP40	0.962025316	0.926829268	1
NOT-B4GALT6 AND CELSR1	0.936708861	0.902439024	0.973684211
CELSR1 AND NOT-FAM189A2	0.935064935	0.923076923	0.947368421
CELSR1 AND NOT-TRIL	0.931506849	0.971428571	0.894736842
Sarcoma (Sarcoma)			
ADAM12 AND NOT-ERBB4	0.941176471	0.941176471	0.941176471
NOT-ST6GALNAC2 AND AXL	0.903225806	1	0.823529412
ADAM12 AND NOT-ERBB3	0.909090909	0.9375	0.882352941
ADAM12 AND NOT-NAALAD2	0.96969697	1	0.941176471
TRAM2 AND NOT-TMEM265	0.903225806	1	0.823529412
ADAM12 AND NOT-PDZK1IP1	0.903225806	1	0.823529412
TRAM2 AND NOT-LPCAT3	0.903225806	1	0.823529412
CD163L1 AND NOT-PRSS16	0.903225806	1	0.823529412
ADAM12 AND NOT-PRSS16	0.944444444	0.894736842	1
TRAM2 AND NOT-PRSS16	0.903225806	1	0.823529412
HEXB AND NOT-ATP8A1	0.903225806	1	0.823529412
TRAM2 AND NOT-ATP8A1	0.903225806	1	0.823529412
ADAM12 AND NOT-ST3GAL6	0.918918919	0.85	1
ADAM12 AND NOT-WFDC2	0.971428571	0.944444444	1
TRAM2 AND NOT-CDS1	0.903225806	1	0.823529412
TRAM2 AND NOT-LRRN2	0.903225806	1	0.823529412
NOT-ST6GALNAC2 AND PROCR	0.9375	1	0.882352941
ADAM12 AND NOT-SLC34A2	0.914285714	0.888888889	0.941176471
NOT-ST6GALNAC2 AND CSF1	0.903225806	1	0.823529412
NOT-ST6GALNAC2 AND SLC38A6	0.903225806	1	0.823529412
NOT-ST6GALNAC2 AND ACE	0.909090909	0.9375	0.882352941
NOT-ST6GALNAC2 AND CD93	0.909090909	0.9375	0.882352941
PLXND1 AND NOT-ST6GALNAC2	0.903225806	1	0.823529412
NOT-ST6GALNAC2 AND LRRC32	0.903225806	1	0.823529412
NOT-ST6GALNAC2 AND MYOF	0.9375	1	0.882352941
NOT-ST6GALNAC2 AND DSE	0.903225806	1	0.823529412
HEXB AND NOT-ST6GALNAC2	0.903225806	1	0.823529412
NOT-ST6GALNAC2 AND ITGB5	0.9375	1	0.882352941
NOT-ST6GALNAC2 AND LTBR	0.903225806	1	0.823529412
NOT-ST6GALNAC2 AND LTC4S	0.903225806	1	0.823529412
NOT-ST6GALNAC2 AND PMP22	0.909090909	0.9375	0.882352941
NOT-ST6GALNAC2 AND DRAM1	0.9375	1	0.882352941
NOT-ST6GALNAC2 AND AXL	0.903225806	1	0.823529412
NOT-ST6GALNAC2 AND ACKR3	0.9375	1	0.882352941
NOT-ST6GALNAC2 AND TGFBI	0.903225806	1	0.823529412
NOT-ST6GALNAC2 AND THBD	0.903225806	1	0.823529412
NOT-ST6GALNAC2 AND REEP4	0.909090909	0.9375	0.882352941
ADAM12 AND NOT-ST6GALNAC2	0.971428571	0.944444444	1
NOT-ST6GALNAC2 AND SLC2A10	0.903225806	1	0.823529412
NOT-ST6GALNAC2 AND PHLDB2	0.909090909	0.9375	0.882352941
NOT-ST6GALNAC2 AND MPZL1	0.903225806	1	0.823529412
NOT-ST6GALNAC2 AND VAMP3	0.903225806	1	0.823529412
NOT-ST6GALNAC2 AND ACVRL1	0.903225806	1	0.823529412
TRAM2 AND NOT-ST6GALNAC2	0.903225806	1	0.823529412
ADAM12 AND NOT-SPINT2	0.903225806	1	0.823529412
TRAM2 AND NOT-CFTR	0.903225806	1	0.823529412
TRAM2 AND NOT-MMP24	0.903225806	1	0.823529412
ADAM12 AND JTB	0.941176471	0.941176471	0.941176471
OS9 AND NOT-GALNT3	0.903225806	1	0.823529412
TRAM2 AND NOT-ERP29	0.903225806	1	0.823529412
TRAM2 AND NOT-SLC27A2	0.903225806	1	0.823529412
ADAM12 AND NOT-ABCB8	0.914285714	0.888888889	0.941176471
TRAM2 AND NOT-KLK8	0.903225806	1	0.823529412
ADAM12 AND NOT-MRAP2	0.909090909	0.9375	0.882352941
ADAM12 AND NOT-MAL2	0.941176471	0.941176471	0.941176471
ADAM12 AND NOT-SLC26A7	0.909090909	0.9375	0.882352941
ADAM12 AND NOT-SLC5A11	0.914285714	0.888888889	0.941176471
TRAM2 AND NOT-GRIN3B	0.903225806	1	0.823529412
ADAM12 AND NOT-COMTD1	0.918918919	0.85	1
TRAM2 AND NOT-COMTD1	0.903225806	1	0.823529412
ADAM12 AND NOT-SFXN2	0.909090909	0.9375	0.882352941
TRAM2 AND NOT-PDZD8	0.903225806	1	0.823529412
TRAM2 AND NOT-GYLTL1B	0.903225806	1	0.823529412
TRAM2 AND NOT-GPHB5	0.903225806	1	0.823529412
ADAM12 AND NOT-C15orf27	0.914285714	0.888888889	0.941176471
TRAM2 AND NOT-TMEM170A	0.903225806	1	0.823529412
TRAM2 AND NOT-ACER1	0.903225806	1	0.823529412

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
TNFSF9 AND NOT-CSH1	0.952380952	1	0.909090909
TYRP1 AND NOT-CSH1	0.9	1	0.818181818
C11orf24 AND NOT-CSH1	0.9	1	0.818181818
NOT-PMEL AND PMEL	0.952380952	1	0.909090909
BCAP31 AND NOT-CSH1	0.9	1	0.818181818
FDPS AND NOT-CSH1	0.9	1	0.818181818
IL12RB2 AND NOT-CSH1	0.9	1	0.818181818
PNKD AND NOT-CSH1	0.952380952	1	0.909090909
LPPR4 AND NOT-CSH1	0.9	1	0.818181818
NAT14 AND NOT-CSH1	0.9	1	0.818181818
MLANA AND NOT-FOLR1	1	1	1
NOT-FOLR1 AND SLC4A11	0.9	1	0.818181818
CTDNEP1 AND NOT-FOLR1	0.9	1	0.818181818
NOT-FOLR1 AND IL20RB	0.9	1	0.818181818
STX18 AND NOT-FOLR1	0.9	1	0.818181818
KIAA1024 AND NOT-FOLR1	0.9	1	0.818181818
CLCN5 AND NOT-FOLR1	0.9	1	0.818181818
MANBAL AND NOT-FOLR1	0.9	1	0.818181818
RER1 AND NOT-FOLR1	0.9	1	0.818181818
LMAN2L AND NOT-FOLR1	0.9	1	0.818181818
NPC1 AND NOT-FOLR1	0.9	1	0.818181818
NOT-FOLR1 AND DAPL1	0.9	1	0.818181818
TMEM189 AND NOT-FOLR1	0.9	1	0.818181818
YIPF1 AND NOT-FOLR1	0.9	1	0.818181818
GALNS AND NOT-FOLR1	0.9	1	0.818181818
SLC45A2 AND NOT-FOLR1	0.952380952	1	0.909090909
MC1R AND NOT-FOLR1	0.9	1	0.818181818
PTPRS AND NOT-FOLR1	0.9	1	0.818181818
TYRP1 AND NOT-FOLR1	0.9	1	0.818181818
NOT-PMEL AND PMEL	0.952380952	1	0.909090909
FDPS AND NOT-FOLR1	0.9	1	0.818181818
NOT-FOLR1 AND WNT5A	0.9	1	0.818181818
MARVELD1 AND NOT-FOLR1	0.952380952	1	0.909090909
OS9 AND OS9	0.9	1	0.818181818
MLANA AND NOT-GNB2	1	1	1
PPT2 AND NOT-GNB2	0.9	1	0.818181818
MANBAL AND NOT-GNB2	0.952380952	1	0.909090909
TMEM138 AND GNB2	0.9	1	0.818181818
CIRH1A AND GNB2	0.952380952	1	0.909090909
SLC45A2 AND NOT-GNB2	0.952380952	1	0.909090909
SLC5A6 AND NOT-GNB2	0.952380952	1	0.909090909
MC1R AND NOT-GNB2	0.9	1	0.818181818
SLC35A4 AND NOT-GNB2	0.9	1	0.818181818
TMEM147 AND TMEM147	0.9	1	0.818181818
EXT1 AND NOT-GNB2	0.909090909	0.90909091	0.909090909
TNFSF9 AND NOT-GNB2	0.909090909	0.90909091	0.909090909
NOT-PMEL AND PMEL	0.952380952	1	0.909090909
BCAP31 AND NOT-GNB2	0.952380952	1	0.909090909
IL12RB2 AND NOT-GNB2	0.9	1	0.818181818
ROMO1 AND NOT-GNB2	0.952380952	1	0.909090909
MLANA AND CSF3R	1	1	1
OXA1L AND NOT-CSF3R	0.952380952	1	0.909090909
PPT2 AND NOT-CSF3R	0.9	1	0.818181818
PARL AND NOT-CSF3R	0.9	1	0.818181818
LMAN2L AND NOT-CSF3R	0.909090909	0.90909091	0.909090909
CIRH1A AND CSF3R	0.952380952	1	0.909090909
AUP1 AND NOT-CSF3R	0.9	1	0.818181818
SLC45A2 AND NOT-CSF3R	0.952380952	1	0.909090909
SLC5A6 AND NOT-CSF3R	0.952380952	1	0.909090909
MC1R AND NOT-CSF3R	0.9	1	0.818181818
TMEM147 AND CSF3R	0.9	1	0.818181818
NOT-PMEL AND PMEL	0.952380952	1	0.909090909
IL12RB2 AND NOT-CSF3R	0.9	1	0.818181818
NOT-SCN10A AND BMPR1A	0.9	1	0.818181818
PQLC1 AND NOT-BMPR1A	0.9	1	0.818181818
OS9 AND NOT-BMPR1A	0.9	1	0.818181818
NOT-FUT6 AND BMPR1A	0.952380952	1	0.909090909
DENND5A AND BMPR1A	0.9	1	0.818181818
NOT-TAAR8 AND BMPR1A	0.909090909	0.90909091	0.909090909
NOT-SCN4A AND BMPR1A	0.952380952	1	0.909090909
MLANA AND NOT-BMPR1A	1	1	1
NOT-TM4SF5 AND BMPR1A	0.952380952	1	0.909090909
NOT-SERPINA9 AND BMPR1A	0.9	1	0.818181818
NOT-CHRNA2 AND BMPR1A	0.952380952	1	0.909090909
NOT-TMEM150B AND BMPR1A	0.952380952	1	0.909090909
SFXN3 AND BMPR1A	0.952380952	1	0.909090909

Antigen Pair Logic	F1	Precision	Recall
ADAM12 AND NOT-CNR1	0.914285714	0.888888889	0.941176471
TRAM2 AND NOT-CNR1	0.903225806	1	0.823529412
TRAM2 AND NOT-SLC44A3	0.903225806	1	0.823529412
ADAM12 AND NOT-TMEM125	0.941176471	0.941176471	0.941176471
PLD3 AND NOT-SGPP2	0.903225806	1	0.823529412
CD163L1 AND NOT-SGPP2	0.903225806	1	0.823529412
ADAM12 AND NOT-SGPP2	0.944444444	0.894736842	1
TRAM2 AND NOT-COL17A1	0.903225806	1	0.823529412
TRAM2 AND NOT-RAET1E	0.903225806	1	0.823529412
TRAM2 AND NOT-COX15	0.903225806	1	0.823529412
CD163L1 AND NOT-TMEM139	0.903225806	1	0.823529412
ADAM12 AND NOT-TMEM139	0.971428571	0.944444444	1
TRAM2 AND NOT-TMEM139	0.903225806	1	0.823529412
ADAM12 AND NOT-CLDN4	0.914285714	0.888888889	0.941176471
TRAM2 AND NOT-CLDN4	0.903225806	1	0.823529412
ADAM12 AND NOT-CLDN3	0.971428571	0.944444444	1
ADAM12 AND NOT-AADAC	0.96969697	1	0.941176471
TRAM2 AND NOT-TRPM6	0.903225806	1	0.823529412
ADAM12 AND NOT-MUC15	0.96969697	1	0.941176471
ADAM12 AND NOT-TMC4	0.909090909	0.9375	0.882352941
ADAM12 AND NOT-MFSD4	0.914285714	0.888888889	0.941176471
ADAM12 AND NOT-PROM2	0.914285714	0.888888889	0.941176471
TRAM2 AND NOT-CXADR	0.903225806	1	0.823529412
ADAM12 AND NOT-KLB	0.914285714	0.888888889	0.941176471
ADAM12 AND NOT-MARVELD2	0.971428571	0.944444444	1
TRAM2 AND NOT-MARVELD2	0.903225806	1	0.823529412
ADAM12 AND NOT-CYP2J2	0.941176471	0.941176471	0.941176471
TRAM2 AND NOT-CYP2J2	0.903225806	1	0.823529412
ADAM12 AND NOT-FAAH2	0.909090909	0.9375	0.882352941
ADAM12 AND NOT-SYNE4	0.944444444	0.894736842	1
TRAM2 AND NOT-SYNE4	0.903225806	1	0.823529412
HEXB AND NOT-IFNLR1	0.903225806	1	0.823529412
ADAM12 AND NOT-IFNLR1	0.909090909	0.9375	0.882352941
ADAM12 AND NOT-PAQR7	0.914285714	0.888888889	0.941176471
CD163L1 AND NOT-CLYBL	0.903225806	1	0.823529412
ADAM12 AND NOT-CLYBL	0.914285714	0.888888889	0.941176471
TRAM2 AND NOT-AGER	0.903225806	1	0.823529412
ADAM12 AND NOT-DSC3	0.909090909	0.9375	0.882352941
ADAM12 AND NOT-E2F5	0.918918919	0.85	1
TRAM2 AND NOT-E2F5	0.903225806	1	0.823529412
ADAM12 AND NOT-EDNRB	0.918918919	0.85	1
TRAM2 AND NOT-EFNB3	0.903225806	1	0.823529412
EMP3 AND NOT-MPZL3	0.9375	1	0.882352941
HEXB AND NOT-MPZL3	0.909090909	0.9375	0.882352941
ADAM12 AND NOT-GRAMD2	1	1	1
TRAM2 AND NOT-GRAMD2	0.903225806	1	0.823529412
ADAM12 AND NOT-KRTCAP3	0.971428571	0.944444444	1
EMP3 AND NOT-GALNT3	0.9375	1	0.882352941
EMP3 AND NOT-GCNT2	0.909090909	0.9375	0.882352941
EMP3 AND NOT-C16orf54	0.9375	1	0.882352941
EMP3 AND NOT-HBD	0.9375	1	0.882352941
EMP3 AND NOT-HACD4	0.903225806	1	0.823529412
EMP3 AND NOT-NIPAL3	0.903225806	1	0.823529412
TGFBI AND NOT-TMEM154	0.903225806	1	0.823529412
TRAM2 AND NOT-EPHA1	0.903225806	1	0.823529412
TRAM2 AND NOT-CERS3	0.903225806	1	0.823529412
ADAM12 AND NOT-ERBB3	0.909090909	0.9375	0.882352941
ADAM12 AND NOT-ERBB4	0.941176471	0.941176471	0.941176471
ADAM12 AND NOT-EXTL1	0.918918919	0.85	1
TRAM2 AND NOT-F2RL1	0.903225806	1	0.823529412
TRAM2 AND NOT-GJD4	0.903225806	1	0.823529412
ADAM12 AND NOT-LRTOMT	0.941176471	0.941176471	0.941176471
TRAM2 AND NOT-ADGRF4	0.903225806	1	0.823529412
ADAM12 AND NOT-FGFR3	0.944444444	0.894736842	1
TRAM2 AND NOT-FGFR3	0.903225806	1	0.823529412
TRAM2 AND NOT-LMTK2	0.903225806	1	0.823529412
HEXB AND NOT-ENPP4	0.9375	1	0.882352941
TRAM2 AND NOT-ENPP4	0.903225806	1	0.823529412
ADAM12 AND NOT-PLA2R1	0.903225806	1	0.823529412
TRAM2 AND NOT-CNOT1	0.903225806	1	0.823529412
ADAM12 AND NOT-MLC1	0.914285714	0.888888889	0.941176471
CD163L1 AND NOT-FMO5	0.903225806	1	0.823529412
ADAM12 AND NOT-ABCB9	0.909090909	0.9375	0.882352941
ADAM12 AND NOT-FOLR1	0.941176471	0.941176471	0.941176471
ADAM12 AND NOT-HYAL4	0.918918919	0.85	1
TRAM2 AND NOT-HYAL4	0.903225806	1	0.823529412

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
GPR137B AND BMPR1A	0.909090909	0.90909091	0.909090909
NOT-GRPR AND BMPR1A	0.952380952	1	0.909090909
OXA1L AND BMPR1A	0.952380952	1	0.909090909
PARL AND BMPR1A	0.952380952	1	0.909090909
BNIP1 AND NOT-BMPR1A	0.9	1	0.818181818
NOT-CTRB2 AND BMPR1A	0.9	1	0.818181818
NOT-NPFFR1 AND BMPR1A	0.9	1	0.818181818
NOT-HGFAC AND BMPR1A	0.952380952	1	0.909090909
NOT-CD300LB AND BMPR1A	0.9	1	0.818181818
MANBAL AND NOT-BMPR1A	0.9	1	0.818181818
NOT-CLDN7 AND BMPR1A	0.9	1	0.818181818
NOT-RTP3 AND BMPR1A	0.9	1	0.818181818
VANGL2 AND NOT-BMPR1A	0.9	1	0.818181818
NOT-SLC22A8 AND BMPR1A	0.952380952	1	0.909090909
NOT-ADAM29 AND BMPR1A	0.909090909	0.90909091	0.909090909
NOT-NMUR2 AND BMPR1A	0.952380952	1	0.909090909
NOT-TMEM95 AND BMPR1A	0.952380952	1	0.909090909
TMEM138 AND BMPR1A	0.9	1	0.818181818
NOT-ISM2 AND BMPR1A	0.952380952	1	0.909090909
NOT-TEX38 AND BMPR1A	0.952380952	1	0.909090909
CIRH1A AND NOT-BMPR1A	0.952380952	1	0.909090909
NOT-SLC5A5 AND BMPR1A	0.952380952	1	0.909090909
DERL2 AND BMPR1A	0.9	1	0.818181818
NOT-SLC6A13 AND BMPR1A	0.9	1	0.818181818
NOT-SLC34A2 AND BMPR1A	0.952380952	1	0.909090909
NOT-GIPR AND BMPR1A	0.952380952	1	0.909090909
NOT-PTH2 AND BMPR1A	0.952380952	1	0.909090909
AUP1 AND BMPR1A	0.952380952	1	0.909090909
NOT-GJD2 AND BMPR1A	0.952380952	1	0.909090909
YIF1B AND NOT-BMPR1A	0.9	1	0.818181818
NOT-CACNG6 AND BMPR1A	0.952380952	1	0.909090909
NOT-SLC34A3 AND BMPR1A	0.952380952	1	0.909090909
TMEM184B AND NOT-BMPR1A	0.9	1	0.818181818
NOT-ATCAY AND BMPR1A	0.952380952	1	0.909090909
NOT-CLDN19 AND BMPR1A	0.952380952	1	0.909090909
NOT-DNAH10 AND BMPR1A	0.9	1	0.818181818
NOT-CAV3 AND BMPR1A	0.9	1	0.818181818
YIPF1 AND NOT-BMPR1A	0.9	1	0.818181818
NOT-CLPS AND BMPR1A	0.9	1	0.818181818
NOT-SLC4A1 AND BMPR1A	0.952380952	1	0.909090909
DAD1 AND BMPR1A	0.9	1	0.818181818
MC1R AND NOT-MC1R	0.9	1	0.818181818
NOT-HTR4 AND BMPR1A	0.952380952	1	0.909090909
NOT-CALHM1 AND BMPR1A	0.952380952	1	0.909090909
NOT-TMPRSS6 AND BMPR1A	0.9	1	0.818181818
TMEM147 AND NOT-BMPR1A	0.9	1	0.818181818
EXT1 AND NOT-BMPR1A	0.9	1	0.818181818
NOT-PTCH2 AND BMPR1A	0.952380952	1	0.909090909
NOT-FTHL17 AND BMPR1A	0.952380952	1	0.909090909
NOT-TMEM105 AND BMPR1A	0.952380952	1	0.909090909
NOT-PCDHGB5 AND BMPR1A	0.9	1	0.818181818
NOT-PMEL AND PMEL	0.952380952	1	0.909090909
BCAP31 AND BCAP31	0.952380952	1	0.909090909
ABHD12 AND NOT-BMPR1A	0.9	1	0.818181818
NOT-OR51M1 AND BMPR1A	0.9	1	0.818181818
FDPS AND NOT-BMPR1A	0.952380952	1	0.909090909
NOT-AMHR2 AND BMPR1A	0.9	1	0.818181818
ROMO1 AND NOT-BMPR1A	0.9	1	0.818181818
NOT-GRIK5 AND BMPR1A	0.952380952	1	0.909090909
NOT-MADCAM1 AND BMPR1A	0.9	1	0.818181818
NOT-C8B AND BMPR1A	0.9	1	0.818181818
CELSR2 AND NOT-BMPR1A	0.9	1	0.818181818
NOT-AVPR1B AND BMPR1A	0.952380952	1	0.909090909
STARD3NL AND BMPR1A	0.9	1	0.818181818
PQLC1 AND NOT-MOS	0.952380952	1	0.909090909
BCS1L AND NOT-MOS	0.9	1	0.818181818
MLANA AND NOT-MOS	1	1	1
SFXN3 AND NOT-MOS	0.9	1	0.818181818
PPT2 AND NOT-MOS	0.9	1	0.818181818
PIGW AND NOT-MOS	0.9	1	0.818181818
HSD3B7 AND NOT-MOS	0.9	1	0.818181818
MANBAL AND NOT-MOS	0.952380952	1	0.909090909
TMEM138 AND NOT-MOS	0.9	1	0.818181818
CIRH1A AND NOT-MOS	0.9	1	0.818181818
TIMM22 AND NOT-MOS	0.9	1	0.818181818
TMEM189 AND NOT-MOS	0.952380952	1	0.909090909

Antigen Pair Logic	F1	Precision	Recall
CD163L1 AND NOT-VSIG2	0.903225806	1	0.823529412
ADAM12 AND PLD3	0.918918919	0.85	1
ADAM12 AND NOT-FUT1	0.971428571	0.944444444	1
TRAM2 AND NOT-FUT1	0.903225806	1	0.823529412
ADAM12 AND NOT-FUT2	0.9375	1	0.882352941
ADAM12 AND NOT-FUT3	0.96969697	1	0.941176471
ADAM12 AND NOT-FUT5	0.941176471	0.941176471	0.941176471
ADAM12 AND NOT-ZDHHC23	0.914285714	0.888888889	0.941176471
TRAM2 AND NOT-ZDHHC23	0.903225806	1	0.823529412
ADAM12 AND NOT-GABRP	0.941176471	0.941176471	0.941176471
TRAM2 AND NOT-SERINC5	0.903225806	1	0.823529412
TRAM2 AND NOT-FRMD3	0.903225806	1	0.823529412
ADAM12 AND NOT-KLK5	1	1	1
ADAM12 AND NOT-PARM1	0.9375	1	0.882352941
NOT-PARM1 AND ACVRL1	0.903225806	1	0.823529412
HEXB AND NOT-GALNT3	0.9375	1	0.882352941
TGFBI AND NOT-GALNT3	0.903225806	1	0.823529412
ADAM12 AND NOT-GALNT3	0.941176471	0.941176471	0.941176471
TRAM2 AND NOT-GALNT3	0.903225806	1	0.823529412
TRAM2 AND NOT-NALCN	0.903225806	1	0.823529412
HEXB AND NOT-HIGD1A	0.903225806	1	0.823529412
TRAM2 AND NOT-TKFC	0.903225806	1	0.823529412
ADAM12 AND NOT-LRRTM2	0.9375	1	0.882352941
NOT-F11R AND MYOF	0.918918919	0.85	1
HEXB AND NOT-GCNT2	0.9375	1	0.882352941
ADAM12 AND NOT-GCNT2	0.941176471	0.941176471	0.941176471
ADAM12 AND NOT-NSG1	0.9375	1	0.882352941
TRAM2 AND NOT-NSG1	0.903225806	1	0.823529412
ADAM12 AND NOT-GP1BA	0.9375	1	0.882352941
TRAM2 AND NOT-SLCO4A1	0.903225806	1	0.823529412
GPR1 AND NOT-DAPL1	0.903225806	1	0.823529412
CD163L1 AND NOT-EPCAM	0.903225806	1	0.823529412
CD163L1 AND NOT-SMIM22	0.903225806	1	0.823529412
CD163L1 AND NOT-P2RY2	0.903225806	1	0.823529412
CD163L1 AND NOT-F11R	0.903225806	1	0.823529412
CD163L1 AND NOT-PLLP	0.903225806	1	0.823529412
CD163L1 AND NOT-ABCB1	0.903225806	1	0.823529412
CD163L1 AND NOT-GRAMD1C	0.903225806	1	0.823529412
CD163L1 AND NOT-SMPD3	0.903225806	1	0.823529412
CD163L1 AND NOT-ACSS2	0.903225806	1	0.823529412
CD163L1 AND NOT-ENPP5	0.903225806	1	0.823529412
CD163L1 AND NOT-SMPD2	0.903225806	1	0.823529412
CD163L1 AND NOT-ST14	0.903225806	1	0.823529412
CD163L1 AND NOT-MMEL1	0.903225806	1	0.823529412
CD163L1 AND NOT-CRB3	0.903225806	1	0.823529412
CD163L1 AND NOT-ATP2C2	0.903225806	1	0.823529412
TRAM2 AND NOT-MCHR1	0.903225806	1	0.823529412
ADAM12 AND NOT-PRRT3	0.918918919	0.85	1
ADAM12 AND NOT-RNF180	0.96969697	1	0.941176471
TRAM2 AND NOT-GPR35	0.903225806	1	0.823529412
ADAM12 AND NOT-GPR37	0.918918919	0.85	1
ADAM12 AND NOT-ILDR1	0.944444444	0.894736842	1
TRAM2 AND NOT-ILDR1	0.903225806	1	0.823529412
ADAM12 AND NOT-TUSC5	0.944444444	0.894736842	1
ADAM12 AND NOT-TMPRSS11E	0.9375	1	0.882352941
ADAM12 AND NOT-HAS3	0.941176471	0.941176471	0.941176471
TRAM2 AND NOT-HAS3	0.903225806	1	0.823529412
HEXB AND NOT-HBD	0.9375	1	0.882352941
ADAM12 AND NOT-HBD	0.941176471	0.941176471	0.941176471
TRAM2 AND NOT-HDAC2	0.903225806	1	0.823529412
HEXB AND NOT-LY75	0.903225806	1	0.823529412
HEXB AND NOT-F11R	0.96969697	1	0.941176471
HEXB AND NOT-ABCB1	0.941176471	0.941176471	0.941176471
HEXB AND NOT-MFSD6	0.903225806	1	0.823529412
HEXB AND NOT-NIPAL3	0.903225806	1	0.823529412
HEXB AND NOT-MARC1	0.903225806	1	0.823529412
HEXB AND NOT-SPINT1	0.941176471	0.941176471	0.941176471
HEXB AND NOT-XK	0.903225806	1	0.823529412
HEXB AND NOT-CLMN	0.903225806	1	0.823529412
ADAM12 AND HLA-B	0.941176471	0.941176471	0.941176471
ADAM12 AND HLA-C	0.914285714	0.888888889	0.941176471
ADAM12 AND NOT-HSD17B2	0.941176471	0.941176471	0.941176471
ADAM12 AND NOT-ICA1	0.914285714	0.888888889	0.941176471
ADAM12 AND NOT-NAT8L	0.918918919	0.85	1
TRAM2 AND NOT-ZDHHC21	0.903225806	1	0.823529412
ADAM12 AND NOT-FREM2	0.941176471	0.941176471	0.941176471

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
YIF1B AND NOT-MOS	0.9	1	0.818181818
SLC45A2 AND NOT-MOS	0.952380952	1	0.909090909
MC1R AND NOT-MOS	0.9	1	0.818181818
SLC35A4 AND NOT-MOS	0.9	1	0.818181818
TMEM147 AND TMEM147	0.9	1	0.818181818
EXT1 AND NOT-MOS	0.9	1	0.818181818
TNFSF9 AND NOT-MOS	0.909090909	0.90909091	0.909090909
NDST1 AND NOT-MOS	0.9	1	0.818181818
NOT-PMEL AND PMEL	0.952380952	1	0.909090909
BCAP31 AND BCAP31	0.952380952	1	0.909090909
FDPS AND NOT-MOS	0.952380952	1	0.909090909
IL12RB2 AND NOT-MOS	0.9	1	0.818181818
CELSR2 AND NOT-MOS	0.952380952	1	0.909090909
MLANA AND NOT-BMX	1	1	1
SLC45A2 AND NOT-BMX	0.952380952	1	0.909090909
NOT-PMEL AND PMEL	0.952380952	1	0.909090909
IL12RB2 AND NOT-BMX	0.9	1	0.818181818
PQLC1 AND NOT-DDC	0.9	1	0.818181818
OS9 AND NOT-DDC	0.9	1	0.818181818
TMEM187 AND NOT-DDC	0.952380952	1	0.909090909
MLANA AND NOT-DDC	1	1	1
PIGL AND NOT-DDC	0.9	1	0.818181818
SFXN3 AND NOT-DDC	0.9	1	0.818181818
CTDNEP1 AND NOT-DDC	0.9	1	0.818181818
PPT2 AND NOT-DDC	0.9	1	0.818181818
BNIP1 AND NOT-DDC	0.9	1	0.818181818
CLCN5 AND NOT-DDC	0.952380952	1	0.909090909
RPN1 AND NOT-DDC	0.9	1	0.818181818
MANBAL AND NOT-DDC	0.9	1	0.818181818
YIF1A AND NOT-DDC	0.9	1	0.818181818
VANGL2 AND NOT-DDC	0.9	1	0.818181818
APMAP AND NOT-DDC	0.9	1	0.818181818
RER1 AND NOT-DDC	0.9	1	0.818181818
LMAN2L AND NOT-DDC	0.952380952	1	0.909090909
NPC1 AND NOT-DDC	0.909090909	0.90909091	0.909090909
CIRH1A AND NOT-DDC	0.952380952	1	0.909090909
TMEM189 AND NOT-DDC	0.9	1	0.818181818
TMEM184B AND NOT-DDC	0.9	1	0.818181818
FAM174B AND NOT-DDC	0.9	1	0.818181818
YIPF1 AND NOT-DDC	0.9	1	0.818181818
SLC45A2 AND NOT-DDC	0.952380952	1	0.909090909
SLC5A6 AND NOT-DDC	0.9	1	0.818181818
HSD17B12 AND NOT-DDC	0.9	1	0.818181818
MC1R AND NOT-DDC	0.9	1	0.818181818
PTPRS AND NOT-DDC	0.9	1	0.818181818
TMEM147 AND NOT-DDC	0.9	1	0.818181818
EXT1 AND NOT-DDC	0.9	1	0.818181818
TNFSF9 AND NOT-DDC	0.9	1	0.818181818
NDST1 AND NOT-DDC	0.9	1	0.818181818
C11orf24 AND NOT-DDC	0.9	1	0.818181818
NOT-PMEL AND PMEL	0.952380952	1	0.909090909
BCAP31 AND NOT-DDC	0.9	1	0.818181818
ABHD12 AND NOT-DDC	0.9	1	0.818181818
FDPS AND NOT-DDC	0.952380952	1	0.909090909
IL12RB2 AND NOT-DDC	0.9	1	0.818181818
PNKD AND NOT-DDC	0.9	1	0.818181818
PRRT3 AND NOT-DDC	0.9	1	0.818181818
ROMO1 AND NOT-DDC	0.9	1	0.818181818
MARVELD1 AND NOT-DDC	0.9	1	0.818181818
GCNT2 AND NOT-DDC	0.9	1	0.818181818
CELSR2 AND NOT-DDC	0.9	1	0.818181818
NOT-CHST4 AND DST	0.9	1	0.818181818
DENND5A AND DST	0.9	1	0.818181818
NOT-SCN4A AND DST	0.909090909	0.90909091	0.909090909
MLANA AND NOT-DST	0.952380952	1	0.909090909
NOT-TM4SF5 AND DST	1	1	1
NOT-KCNJ6 AND DST	0.909090909	0.90909091	0.909090909
NOT-CHRNA2 AND DST	0.909090909	0.90909091	0.909090909
NOT-TMEM150B AND DST	0.952380952	1	0.909090909
NOT-ITGA2B AND DST	0.952380952	1	0.909090909
NOT-SYT2 AND DST	0.952380952	1	0.909090909
GPR137B AND DST	0.909090909	0.90909091	0.909090909
OXA1L AND DST	0.952380952	1	0.909090909
PARL AND DST	1	1	1
NOT-NPFFR1 AND DST	0.952380952	1	0.909090909
NOT-TMPRSS2 AND DST	0.909090909	0.90909091	0.909090909

Antigen Pair Logic	F1	Precision	Recall
TRAM2 AND NOT-ZACN	0.903225806	1	0.823529412
TRAM2 AND NOT-PRAC2	0.903225806	1	0.823529412
TRAM2 AND NOT-ITGA2	0.903225806	1	0.823529412
NOT-MAOB AND ITGB5	0.9375	1	0.882352941
ADAM12 AND NOT-ITGB6	0.903225806	1	0.823529412
ADAM12 AND NOT-ITGB8	0.971428571	0.944444444	1
ADAM12 AND NOT-KCNC4	0.914285714	0.888888889	0.941176471
TRAM2 AND NOT-KCNC4	0.903225806	1	0.823529412
ADAM12 AND NOT-KRT5	0.941176471	0.941176471	0.941176471
TRAM2 AND NOT-RPRML	0.903225806	1	0.823529412
ADAM12 AND NOT-LBR	0.971428571	0.944444444	1
TRAM2 AND NOT-LETM1	0.903225806	1	0.823529412
ADAM12 AND NOT-C14orf180	0.9375	1	0.882352941
TRAM2 AND NOT-C14orf180	0.903225806	1	0.823529412
ADAM12 AND NOT-KLHL31	0.96969697	1	0.941176471
TRAM2 AND NOT-LTA	0.903225806	1	0.823529412
NOT-F11R AND LTBR	0.914285714	0.888888889	0.941176471
ADAM12 AND NOT-ERVFRD-1	0.944444444	0.894736842	1
ADAM12 AND NOT-LTF	0.918918919	0.85	1
ADAM12 AND NOT-TACSTD2	0.909090909	0.9375	0.882352941
ADAM12 AND NOT-EPCAM	0.971428571	0.944444444	1
TRAM2 AND NOT-EPCAM	0.903225806	1	0.823529412
TGFBI AND NOT-MAL	0.903225806	1	0.823529412
TRAM2 AND NOT-MAL	0.903225806	1	0.823529412
PMP22 AND NOT-MAOB	0.9375	1	0.882352941
ADAM12 AND NOT-MAOB	0.9375	1	0.882352941
ADAM12 AND NOT-MGAT3	0.944444444	0.894736842	1
TRAM2 AND NOT-MGAT3	0.903225806	1	0.823529412
ADAM12 AND NOT-MPZ	0.903225806	1	0.823529412
ADAM12 AND NOT-SMIM22	0.971428571	0.944444444	1
TRAM2 AND NOT-SMIM22	0.903225806	1	0.823529412
TRAM2 AND NOT-MTHFD1	0.903225806	1	0.823529412
ADAM12 AND NOT-MTNR1B	1	1	1
ADAM12 AND NOT-NNAT	0.918918919	0.85	1
ADAM12 AND NOT-ATP1B2	0.941176471	0.941176471	0.941176471
ADAM12 AND NOT-NPY5R	0.903225806	1	0.823529412
TRAM2 AND NOT-NTRK3	0.903225806	1	0.823529412
ADAM12 AND NOT-GPR143	0.914285714	0.888888889	0.941176471
TRAM2 AND NOT-ATP4A	0.903225806	1	0.823529412
ADAM12 AND NOT-P2RY2	0.971428571	0.944444444	1
TRAM2 AND NOT-P2RY2	0.903225806	1	0.823529412
TRAM2 AND NOT-ATP6V0A4	0.903225806	1	0.823529412
TGFBI AND NOT-F11R	0.909090909	0.9375	0.882352941
ADAM12 AND NOT-F11R	0.971428571	0.944444444	1
TRAM2 AND NOT-F11R	0.903225806	1	0.823529412
ADAM12 AND NOT-PLLP	0.971428571	0.944444444	1
TRAM2 AND NOT-PLLP	0.903225806	1	0.823529412
ADAM12 AND NOT-ATP8A2	0.918918919	0.85	1
TRAM2 AND NOT-ATP8A2	0.903225806	1	0.823529412
ADAM12 AND NOT-PF4	0.918918919	0.85	1
ADAM12 AND NOT-ABCB1	0.918918919	0.85	1
TRAM2 AND NOT-ABCB1	0.903225806	1	0.823529412
ADAM12 AND NOT-PIGR	0.941176471	0.941176471	0.941176471
PMP22 AND NOT-SEMA6A	0.9375	1	0.882352941
NOT-MARC1 AND PMP22	0.903225806	1	0.823529412
ADAM12 AND NOT-IL20RA	0.903225806	1	0.823529412
ADAM12 AND NOT-GPR87	0.918918919	0.85	1
ADAM12 AND NOT-ATP7B	0.9375	1	0.882352941
TRAM2 AND NOT-FAM105A	0.903225806	1	0.823529412
ADAM12 AND NOT-SDK2	0.909090909	0.9375	0.882352941
ADAM12 AND NOT-GRAMD1C	0.918918919	0.85	1
TRAM2 AND NOT-GRAMD1C	0.903225806	1	0.823529412
ADAM12 AND NOT-GDPD2	0.971428571	0.944444444	1
TRAM2 AND NOT-GDPD2	0.903225806	1	0.823529412
ADAM12 AND NOT-RNF43	0.914285714	0.888888889	0.941176471
TRAM2 AND NOT-RNF43	0.903225806	1	0.823529412
ADAM12 AND NOT-TMEM40	0.914285714	0.888888889	0.941176471
ADAM12 AND NOT-TMEM144	0.918918919	0.85	1
TRAM2 AND NOT-TMEM144	0.903225806	1	0.823529412
ADAM12 AND NOT-STYK1	0.96969697	1	0.941176471
ADAM12 AND NOT-VEZT	0.918918919	0.85	1
TRAM2 AND NOT-SLC39A4	0.903225806	1	0.823529412
ADAM12 AND NOT-ST6GALNAC1	0.944444444	0.894736842	1
TRAM2 AND NOT-ST6GALNAC1	0.903225806	1	0.823529412
TRAM2 AND NOT-PLGRKT	0.903225806	1	0.823529412
TRAM2 AND NOT-ACSS2	0.903225806	1	0.823529412

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
NOT-CLDN7 AND DST	0.952380952	1	0.909090909
NOT-ADAM29 AND DST	0.909090909	0.90909091	0.909090909
NOT-NMUR2 AND DST	0.952380952	1	0.909090909
NOT-FNDC5 AND DST	0.9	1	0.818181818
NOT-TMEM95 AND DST	0.952380952	1	0.909090909
TMEM138 AND DST	0.952380952	1	0.909090909
NOT-ISM2 AND DST	0.909090909	0.90909091	0.909090909
DERL2 AND DST	0.952380952	1	0.909090909
NOT-SLC34A2 AND DST	0.909090909	0.90909091	0.909090909
NOT-GIPR AND DST	0.952380952	1	0.909090909
AUP1 AND DST	1	1	1
NOT-GP2 AND DST	0.952380952	1	0.909090909
GUSB AND DST	0.909090909	0.90909091	0.909090909
NOT-CACNG6 AND DST	0.909090909	0.90909091	0.909090909
NOT-ATCAY AND DST	0.956521739	0.91666667	1
NOT-CLDN19 AND DST	0.909090909	0.90909091	0.909090909
NOT-CAV3 AND DST	0.909090909	0.90909091	0.909090909
NOT-CLPS AND DST	0.9	1	0.818181818
SLC45A2 AND NOT-DST	0.9	1	0.818181818
NOT-SLC4A1 AND DST	0.909090909	0.90909091	0.909090909
DAD1 AND DST	0.9	1	0.818181818
NOT-ATP13A5 AND DST	0.9	1	0.818181818
NOT-FFAR2 AND DST	0.909090909	0.90909091	0.909090909
COX8A AND DST	0.909090909	0.90909091	0.909090909
NOT-ODF4 AND DST	0.909090909	0.90909091	0.909090909
NOT-PTCH2 AND DST	0.909090909	0.90909091	0.909090909
NOT-TMEM105 AND DST	0.952380952	1	0.909090909
NOT-PMEL AND PMEL	0.952380952	1	0.909090909
BCAP31 AND DST	0.952380952	1	0.909090909
NOT-OR51M1 AND DST	0.9	1	0.818181818
NDUFA1 AND DST	0.952380952	1	0.909090909
NOT-MADCAM1 AND DST	0.909090909	0.90909091	0.909090909
NOT-TMEM252 AND DST	0.909090909	0.90909091	0.909090909
STARD3NL AND DST	0.9	1	0.818181818
PQLC1 AND NOT-CRYGC	0.952380952	1	0.909090909
OS9 AND OS9	0.9	1	0.818181818
BCS1L AND NOT-CRYGC	0.9	1	0.818181818
SLAMF8 AND NOT-CRYGC	0.9	1	0.818181818
TMEM187 AND NOT-CRYGC	0.916666667	0.84615385	1
FAM3A AND NOT-CRYGC	0.9	1	0.818181818
MLANA AND NOT-CRYGC	1	1	1
PIGL AND NOT-CRYGC	0.952380952	1	0.909090909
PQLC2 AND NOT-CRYGC	0.952380952	1	0.909090909
LARGE AND NOT-CRYGC	0.9	1	0.818181818
TMCO4 AND NOT-CRYGC	0.9	1	0.818181818
SLC1A4 AND NOT-CRYGC	0.9	1	0.818181818
SCAP AND NOT-CRYGC	0.9	1	0.818181818
SFXN3 AND NOT-CRYGC	0.952380952	1	0.909090909
PPT2 AND NOT-CRYGC	0.9	1	0.818181818
PIGW AND NOT-CRYGC	0.9	1	0.818181818
HGSNAT AND NOT-CRYGC	0.9	1	0.818181818
HSD3B7 AND NOT-CRYGC	0.9	1	0.818181818
ST7L AND NOT-CRYGC	0.9	1	0.818181818
BNIP1 AND NOT-CRYGC	0.952380952	1	0.909090909
CLCN5 AND NOT-CRYGC	0.916666667	0.84615385	1
MANBAL AND NOT-CRYGC	0.952380952	1	0.909090909
VANGL2 AND NOT-CRYGC	0.952380952	1	0.909090909
TMEM138 AND TMEM138	0.952380952	1	0.909090909
NPC1 AND NOT-CRYGC	1	1	1
NOT-CIRH1A AND CIRH1A	0.952380952	1	0.909090909
NOT-ERGIC3 AND ERGIC3	0.9	1	0.818181818
TIMM22 AND NOT-CRYGC	0.9	1	0.818181818
TMEM189 AND NOT-CRYGC	0.952380952	1	0.909090909
YIF1B AND NOT-CRYGC	0.9	1	0.818181818
TMEM184B AND NOT-CRYGC	0.952380952	1	0.909090909
FAM174B AND NOT-CRYGC	0.952380952	1	0.909090909
GPR107 AND NOT-CRYGC	0.9	1	0.818181818
SDC3 AND NOT-CRYGC	0.9	1	0.818181818
SLC45A2 AND NOT-CRYGC	0.952380952	1	0.909090909
SLC5A6 AND NOT-CRYGC	0.952380952	1	0.909090909
MC1R AND NOT-CRYGC	0.9	1	0.818181818
SLC35A4 AND NOT-CRYGC	0.9	1	0.818181818
ARMCX1 AND NOT-CRYGC	0.9	1	0.818181818
TMEM147 AND TMEM147	0.9	1	0.818181818
EXT1 AND NOT-CRYGC	0.952380952	1	0.909090909
TNFSF9 AND NOT-CRYGC	0.909090909	0.90909091	0.909090909

Antigen Pair Logic	F1	Precision	Recall
ADAM12 AND NOT-ACPP	0.918918919	0.85	1
TRAM2 AND NOT-ACPP	0.903225806	1	0.823529412
ADAM12 AND NOT-MRAP	0.918918919	0.85	1
TRAM2 AND NOT-MRAP	0.903225806	1	0.823529412
ADAM12 AND NOT-SLC7A10	0.944444444	0.894736842	1
ADAM12 AND NOT-TMPRSS4	0.941176471	0.941176471	0.941176471
TRAM2 AND NOT-PANX2	0.903225806	1	0.823529412
ADAM12 AND NOT-PYY	0.909090909	0.9375	0.882352941
TRAM2 AND NOT-PYY	0.903225806	1	0.823529412
TRAM2 AND NOT-SLC12A9	0.903225806	1	0.823529412
NOT-SMIM5 AND AGTRAP	0.903225806	1	0.823529412
ADAM12 AND NOT-ANO2	0.944444444	0.894736842	1
ADAM12 AND NOT-TMEM63C	0.918918919	0.85	1
TRAM2 AND NOT-NIPAL3	0.903225806	1	0.823529412
TRAM2 AND NOT-ADGRG6	0.903225806	1	0.823529412
TRAM2 AND NOT-SMAGP	0.903225806	1	0.823529412
TRAM2 AND NOT-ABHD6	0.903225806	1	0.823529412
ADAM12 AND NOT-KIAA1324	0.909090909	0.9375	0.882352941
ADAM12 AND NOT-SEMA6A	0.909090909	0.9375	0.882352941
TRAM2 AND NOT-SEMA6A	0.903225806	1	0.823529412
ADAM12 AND NOT-SEMA4G	0.909090909	0.9375	0.882352941
TRAM2 AND NOT-PTPRF	0.903225806	1	0.823529412
ADAM12 AND OSTC	0.914285714	0.888888889	0.941176471
ADAM12 AND NOT-ACE2	0.909090909	0.9375	0.882352941
ADAM12 AND NOT-SCN7A	0.941176471	0.941176471	0.941176471
OSMR AND NOT-SCN7A	0.903225806	1	0.823529412
TRAM2 AND NOT-SCN7A	0.903225806	1	0.823529412
ADAM12 AND NOT-SCNN1A	0.941176471	0.941176471	0.941176471
ADAM12 AND NOT-SCNN1B	0.914285714	0.888888889	0.941176471
TRAM2 AND NOT-SCNN1B	0.903225806	1	0.823529412
ADAM12 AND NOT-SCNN1G	0.941176471	0.941176471	0.941176471
ADAM12 AND NOT-CCL22	0.918918919	0.85	1
TRAM2 AND NOT-CCL22	0.903225806	1	0.823529412
ADAM12 AND NOT-CLSTN2	0.903225806	1	0.823529412
ADAM12 AND NOT-TNMD	0.944444444	0.894736842	1
NOT-MARC1 AND ITSN1	0.903225806	1	0.823529412
ADAM12 AND NOT-TMPRSS3	0.941176471	0.941176471	0.941176471
ADAM12 AND NOT-MARC1	0.9375	1	0.882352941
NOT-MARC1 AND ACVRL1	0.903225806	1	0.823529412
ADAM12 AND NOT-SLC4A3	0.971428571	0.944444444	1
ADAM12 AND NOT-SLC6A2	0.903225806	1	0.823529412
TRAM2 AND NOT-SLC6A2	0.903225806	1	0.823529412
TRAM2 AND NOT-SLC8A2	0.903225806	1	0.823529412
TRAM2 AND NOT-SLC12A2	0.903225806	1	0.823529412
ADAM12 AND NOT-SLC15A1	0.9375	1	0.882352941
ADAM12 AND NOT-SMPD2	0.918918919	0.85	1
TGFBI AND NOT-SPINT1	0.909090909	0.9375	0.882352941
ADAM12 AND NOT-SPINT1	0.971428571	0.944444444	1
TRAM2 AND NOT-SPINT1	0.903225806	1	0.823529412
ADAM12 AND NOT-ST14	0.971428571	0.944444444	1
TRAM2 AND NOT-ST14	0.903225806	1	0.823529412
TRAM2 AND NOT-STXBP2	0.903225806	1	0.823529412
TRAM2 AND NOT-BTC	0.903225806	1	0.823529412
ADAM12 AND NOT-SYT4	0.909090909	0.9375	0.882352941
ADAM12 AND NOT-TACR1	0.944444444	0.894736842	1
ADAM12 AND NOT-TSPAN8	0.909090909	0.9375	0.882352941
ADAM12 AND NOT-TRPC6	0.909090909	0.9375	0.882352941
ADAM12 AND NOT-UMOD	0.914285714	0.888888889	0.941176471
TRAM2 AND NOT-UPK3A	0.903225806	1	0.823529412
TRAM2 AND NOT-VIPR1	0.903225806	1	0.823529412
TRAM2 AND NOT-VIPR2	0.903225806	1	0.823529412
ADAM12 AND NOT-WNT6	0.914285714	0.888888889	0.941176471
ADAM12 AND NOT-WNT7A	0.903225806	1	0.823529412
ADAM12 AND NOT-XK	0.9375	1	0.882352941
ADAM12 AND NOT-DDR1	0.9375	1	0.882352941
TRAM2 AND NOT-PRRG4	0.903225806	1	0.823529412
ADAM12 AND NOT-SLC25A23	0.941176471	0.941176471	0.941176471
ADAM12 AND NOT-MMEL1	0.971428571	0.944444444	1
ADAM12 AND NOT-NOX5	0.914285714	0.888888889	0.941176471
TRAM2 AND NOT-KREMEN2	0.903225806	1	0.823529412
TRAM2 AND NOT-RNF128	0.903225806	1	0.823529412
ADAM12 AND NOT-CLMN	0.941176471	0.941176471	0.941176471
ADAM12 AND NOT-TMC5	0.941176471	0.941176471	0.941176471
TRAM2 AND NOT-TMC5	0.903225806	1	0.823529412
ADAM12 AND NOT-ERMP1	0.909090909	0.9375	0.882352941
ADAM12 AND NOT-CWH43	0.941176471	0.941176471	0.941176471

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
NDST1 AND NOT-CRYGC	0.952380952	1	0.909090909
SMIM12 AND NOT-CRYGC	0.9	1	0.818181818
C11orf24 AND NOT-CRYGC	0.9	1	0.818181818
NOT-PMEL AND PMEL	0.952380952	1	0.909090909
BCAP31 AND CRYGC	0.9	1	0.818181818
ABHD12 AND NOT-CRYGC	0.952380952	1	0.909090909
FDPS AND NOT-CRYGC	1	1	1
IL12RB2 AND NOT-CRYGC	0.9	1	0.818181818
ICMT AND NOT-CRYGC	0.9	1	0.818181818
PNKD AND NOT-CRYGC	0.952380952	1	0.909090909
PRRT3 AND NOT-CRYGC	0.952380952	1	0.909090909
ROMO1 AND CRYGC	0.952380952	1	0.909090909
YIPF3 AND NOT-CRYGC	0.9	1	0.818181818
CELSR2 AND NOT-CRYGC	0.952380952	1	0.909090909
MLANA AND DHODH	0.952380952	1	0.909090909
NOT-CHRNA2 AND DHODH	0.952380952	1	0.909090909
PPT2 AND NOT-DHODH	0.9	1	0.818181818
TMEM138 AND DHODH	0.9	1	0.818181818
SLC45A2 AND NOT-DHODH	0.952380952	1	0.909090909
MC1R AND NOT-DHODH	0.9	1	0.818181818
TMEM147 AND DHODH	0.9	1	0.818181818
NOT-PMEL AND PMEL	0.952380952	1	0.909090909
OS9 AND OS9	0.9	1	0.818181818
MLANA AND AES	0.9	1	0.818181818
SFXN3 AND AES	0.952380952	1	0.909090909
MANBAL AND AES	0.952380952	1	0.909090909
MC1R AND NOT-MC1R	0.9	1	0.818181818
TNFSF9 AND NOT-TNFSF9	0.909090909	0.90909091	0.909090909
NOT-PMEL AND PMEL	0.952380952	1	0.909090909
BCAP31 AND AES	0.9	1	0.818181818
FDPS AND AES	0.952380952	1	0.909090909
ROMO1 AND AES	0.952380952	1	0.909090909
OS9 AND NOT-HADH	0.9	1	0.818181818
MLANA AND HADH	0.952380952	1	0.909090909
SFXN3 AND NOT-HADH	0.9	1	0.818181818
MANBAL AND NOT-HADH	0.9	1	0.818181818
CIRH1A AND NOT-HADH	0.9	1	0.818181818
MC1R AND NOT-MC1R	0.9	1	0.818181818
NOT-PMEL AND PMEL	0.952380952	1	0.909090909
BCAP31 AND NOT-HADH	0.9	1	0.818181818
FDPS AND NOT-HADH	0.952380952	1	0.909090909
ROMO1 AND NOT-HADH	0.9	1	0.818181818
MLANA AND GPX7	1	1	1
CIRH1A AND GPX7	0.952380952	1	0.909090909
TMEM147 AND GPX7	0.9	1	0.818181818
OS9 AND OS9	0.9	1	0.818181818
NOT-SCN4A AND ART3	0.9	1	0.818181818
MLANA AND MLANA	1	1	1
NOT-CHRNA2 AND ART3	0.909090909	0.90909091	0.909090909
SFXN3 AND ART3	0.9	1	0.818181818
GPR137B AND ART3	0.909090909	0.90909091	0.909090909
OXA1L AND ART3	0.9	1	0.818181818
PARL AND ART3	0.952380952	1	0.909090909
NOT-ADAM29 AND ART3	0.909090909	0.90909091	0.909090909
NOT-NMUR2 AND ART3	0.909090909	0.90909091	0.909090909
TMEM138 AND ART3	0.9	1	0.818181818
NOT-CIRH1A AND CIRH1A	0.952380952	1	0.909090909
ERGIC3 AND ART3	0.9	1	0.818181818
DERL2 AND ART3	0.9	1	0.818181818
AUP1 AND ART3	0.952380952	1	0.909090909
MC1R AND NOT-MC1R	0.9	1	0.818181818
TMEM147 AND TMEM147	0.9	1	0.818181818
NOT-PMEL AND PMEL	0.952380952	1	0.909090909
FDPS AND ART3	0.952380952	1	0.909090909
ROMO1 AND ART3	0.9	1	0.818181818
OS9 AND OS9	0.9	1	0.818181818
NOT-FUT6 AND DDB1	0.9	1	0.818181818
NOT-ASIC5 AND DDB1	0.9	1	0.818181818
NOT-TAAR8 AND DDB1	0.9	1	0.818181818
MLANA AND DDB1	1	1	1
NOT-GPR137 AND DDB1	0.9	1	0.818181818
NOT-SYT13 AND DDB1	0.9	1	0.818181818
NOT-CYP2A6 AND DDB1	0.9	1	0.818181818
NOT-TM4SF5 AND DDB1	0.9	1	0.818181818
NOT-CHRNA2 AND DDB1	0.9	1	0.818181818
NOT-TMEM150B AND DDB1	0.9	1	0.818181818

Antigen Pair Logic	F1	Precision	Recall
TRAM2 AND NOT-DNAJC5	0.903225806	1	0.823529412
ADAM12 AND NOT-SPX	0.903225806	1	0.823529412
ADAM12 AND NOT-SLC25A18	0.903225806	1	0.823529412
ADAM12 AND NOT-FAM213A	0.903225806	1	0.823529412
ADAM12 AND NOT-PROM1	0.971428571	0.944444444	1
ADAM12 AND NOT-GLB1L2	0.9375	1	0.882352941
ADAM12 AND NOT-SLC13A2	0.914285714	0.888888889	0.941176471
ADAM12 AND NOT-CLDN8	0.918918919	0.85	1
ADAM12 AND NOT-SLC28A2	0.914285714	0.888888889	0.941176471
ADAM12 AND NOT-KIAA1919	0.918918919	0.85	1
ADAM12 AND NOT-MARVELD3	0.909090909	0.9375	0.882352941
ADAM12 AND NOT-GGTLC1	0.941176471	0.941176471	0.941176471
ADAM12 AND NOT-DAPL1	0.971428571	0.944444444	1
ADAM12 AND NOT-TMEM132C	0.96969697	1	0.941176471
ADAM12 AND NOT-CRB3	0.971428571	0.944444444	1
ADAM12 AND NOT-FAM189A2	0.903225806	1	0.823529412
ADAM12 AND NOT-SIGLEC6	0.971428571	0.944444444	1
ADAM12 AND NOT-NRG2	0.96969697	1	0.941176471
ADAM12 AND NOT-ENTPD3	0.941176471	0.941176471	0.941176471
ADAM12 AND NOT-KIAA0040	0.914285714	0.888888889	0.941176471
ADAM12 AND NOT-TMEM63A	0.918918919	0.85	1
ADAM12 AND NOT-CD79B	0.914285714	0.888888889	0.941176471
ADAM12 AND NOT-MFAP3L	0.971428571	0.944444444	1
ADAM12 AND NOT-CDH1	0.909090909	0.9375	0.882352941
TRAM2 AND NOT-ADGRV1	0.903225806	1	0.823529412
TRAM2 AND NOT-SLC10A7	0.903225806	1	0.823529412
TRAM2 AND NOT-MFSD7	0.903225806	1	0.823529412
TRAM2 AND NOT-YIPF4	0.903225806	1	0.823529412
TRAM2 AND NOT-AGPAT9	0.903225806	1	0.823529412
TRAM2 AND NOT-GALR3	0.903225806	1	0.823529412
TRAM2 AND NOT-ITGA8	0.903225806	1	0.823529412
TRAM2 AND NOT-KCNK5	0.903225806	1	0.823529412
TRAM2 AND NOT-PROM1	0.903225806	1	0.823529412
TRAM2 AND NOT-CLDN8	0.903225806	1	0.823529412
TRAM2 AND NOT-KCNQ4	0.903225806	1	0.823529412
TRAM2 AND NOT-SLC28A2	0.903225806	1	0.823529412
TRAM2 AND NOT-MARVELD3	0.903225806	1	0.823529412
TRAM2 AND NOT-CRB3	0.903225806	1	0.823529412
TRAM2 AND NOT-TMEM183A	0.903225806	1	0.823529412
TRAM2 AND NOT-GPR37L1	0.903225806	1	0.823529412
TRAM2 AND NOT-SIGLEC6	0.903225806	1	0.823529412
TRAM2 AND NOT-ACVRL1	0.903225806	1	0.823529412
TRAM2 AND NOT-NRG2	0.903225806	1	0.823529412
TRAM2 AND NOT-TMEM63A	0.903225806	1	0.823529412
TRAM2 AND NOT-DGCR2	0.903225806	1	0.823529412
Carcinoma, Non-Small-Cell Lung (Lung NSCC)			
FAP AND NOT-LRRN4CL	0.9	1	0.818181818
FAP AND NOT-GALNT16	0.9	1	0.818181818
FAP AND NOT-PTGFR	0.9	1	0.818181818
FAP AND NOT-LCAT	0.9	1	0.818181818
FAP AND NOT-SFRP1	0.9	1	0.818181818
FAP AND NOT-ITGA7	0.9	1	0.818181818
FAP AND NOT-ABCB1	0.9	1	0.818181818
FAP AND NOT-RECK	0.9	1	0.818181818
FAP AND NOT-ENPP1	0.9	1	0.818181818
FAP AND NOT-TGFBR3	0.9	1	0.818181818
NOT-ACACB AND CD276	0.952380952	1	0.909090909
FZD2 AND NOT-GPA33	0.9	1	0.818181818
NOT-ACACB AND SDC1	0.9	1	0.818181818
SCAMP3 AND NOT-TMEM170B	0.9	1	0.818181818
OST4 AND SEMA4F	0.952380952	1	0.909090909
SCAMP3 AND NOT-SCAMP2	0.9	1	0.818181818
NOT-ACACB AND SCAMP3	0.9	1	0.818181818
SCAMP3 AND NOT-MAN2A2	0.9	1	0.818181818
SCAMP3 AND NOT-STX7	0.9	1	0.818181818
NOT-ACACB AND CALCRL	0.9	1	0.818181818
FZD2 AND NOT-GPA33	0.9	1	0.818181818
OSMR AND NOT-ABCA9	0.9	1	0.818181818
GGCX AND NOT-N4BP2L2	0.909090909	0.909090909	0.909090909
NOT-ACACB AND MERTK	0.952380952	1	0.909090909
DNAJB11 AND NOT-SLC30A9	0.9	1	0.818181818
TMEM39A AND NOT-SLC30A9	0.9	1	0.818181818
SEMA4F AND NOT-MAN2A2	0.9	1	0.818181818
NOT-ACACB AND YIF1A	0.9	1	0.818181818
NOT-ACACB AND KDELR1	0.9	1	0.818181818
NOT-ACACB AND TMED2	0.9	1	0.818181818

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
SFXN3 AND DDB1	0.9	1	0.818181818
NOT-GRPR AND DDB1	0.9	1	0.818181818
OXA1L AND DDB1	0.9	1	0.818181818
PARL AND DDB1	0.9	1	0.818181818
COX7A2L AND DDB1	0.9	1	0.818181818
NOT-HGFAC AND DDB1	0.9	1	0.818181818
MANBAL AND DDB1	0.9	1	0.818181818
NOT-B4GALNT3 AND DDB1	0.9	1	0.818181818
NOT-SLC22A8 AND DDB1	0.9	1	0.818181818
NOT-NMUR2 AND DDB1	0.9	1	0.818181818
NOT-TMEM95 AND DDB1	0.9	1	0.818181818
NOT-CIRH1A AND CIRH1A	0.952380952	1	0.909090909
ERGIC3 AND DDB1	0.9	1	0.818181818
NOT-SLC5A5 AND DDB1	0.9	1	0.818181818
DERL2 AND DDB1	0.9	1	0.818181818
NOT-SLC34A2 AND DDB1	0.9	1	0.818181818
NOT-GIPR AND DDB1	0.9	1	0.818181818
NOT-SLC22A14 AND DDB1	0.9	1	0.818181818
NOT-PTH2 AND DDB1	0.9	1	0.818181818
AUP1 AND DDB1	0.9	1	0.818181818
NOT-GJD2 AND DDB1	0.9	1	0.818181818
NOT-CACNG6 AND DDB1	0.9	1	0.818181818
NOT-SLC34A3 AND DDB1	0.9	1	0.818181818
NOT-ATCAY AND DDB1	0.9	1	0.818181818
NOT-CLDN19 AND DDB1	0.9	1	0.818181818
NOT-DNAH10 AND DDB1	0.9	1	0.818181818
SLC45A2 AND NOT-DDB1	0.952380952	1	0.909090909
NOT-CACNG5 AND DDB1	0.9	1	0.818181818
TMEM147 AND TMEM147	0.9	1	0.818181818
NOT-PTCH2 AND DDB1	0.9	1	0.818181818
NOT-TMEM105 AND DDB1	0.9	1	0.818181818
NOT-PMEL AND PMEL	0.952380952	1	0.909090909
NOT-MRGPRX4 AND DDB1	0.9	1	0.818181818
BCAP31 AND DDB1	0.9	1	0.818181818
FDPS AND DDB1	0.9	1	0.818181818
NOT-SLC10A2 AND DDB1	0.9	1	0.818181818
ROMO1 AND DDB1	0.952380952	1	0.909090909
NOT-GRIK5 AND DDB1	0.9	1	0.818181818
NOT-AVPR1B AND DDB1	0.9	1	0.818181818
PQLC1 AND NOT-GABRD	0.9	1	0.818181818
OS9 AND OS9	0.9	1	0.818181818
KIAA1549L AND NOT-GABRD	0.9	1	0.818181818
MLANA AND NOT-GABRD	1	1	1
PPT2 AND NOT-GABRD	0.9	1	0.818181818
NOT-CIRH1A AND CIRH1A	0.952380952	1	0.909090909
NOT-ERGIC3 AND ERGIC3	0.9	1	0.818181818
TMEM189 AND NOT-GABRD	0.9	1	0.818181818
SLC45A2 AND NOT-GABRD	0.952380952	1	0.909090909
TMEM147 AND TMEM147	0.9	1	0.818181818
EXT1 AND NOT-GABRD	0.9	1	0.818181818
NOT-PMEL AND PMEL	0.952380952	1	0.909090909
FDPS AND NOT-GABRD	0.9	1	0.818181818
IL12RB2 AND NOT-GABRD	0.9	1	0.818181818
DENND5A AND BMPR1B	0.9	1	0.818181818
NOT-SCN4A AND BMPR1B	0.9	1	0.818181818
MLANA AND MLANA	1	1	1
NOT-TM4SF5 AND BMPR1B	0.9	1	0.818181818
GPR137B AND BMPR1B	0.909090909	0.90909091	0.909090909
PPT2 AND NOT-BMPR1B	0.9	1	0.818181818
PARL AND BMPR1B	0.952380952	1	0.909090909
CIRH1A AND BMPR1B	0.952380952	1	0.909090909
DERL2 AND BMPR1B	0.952380952	1	0.909090909
AUP1 AND BMPR1B	0.952380952	1	0.909090909
SLC45A2 AND NOT-BMPR1B	0.952380952	1	0.909090909
MC1R AND NOT-BMPR1B	0.9	1	0.818181818
TMEM147 AND BMPR1B	0.9	1	0.818181818
NOT-PMEL AND PMEL	0.952380952	1	0.909090909
ROMO1 AND BMPR1B	0.952380952	1	0.909090909
NOT-GRIK5 AND BMPR1B	0.9	1	0.818181818
NOT-AVPR1B AND BMPR1B	0.9	1	0.818181818
MLANA AND GMFB	1	1	1
TMEM138 AND NOT-GMFB	0.952380952	1	0.909090909
CIRH1A AND GMFB	0.952380952	1	0.909090909
SLC5A6 AND NOT-GMFB	0.909090909	0.90909091	0.909090909
MC1R AND NOT-MC1R	0.9	1	0.818181818
TMEM147 AND GMFB	0.9	1	0.818181818

Antigen Pair Logic	F1	Precision	Recall
NOT-ACACB AND ILVBL	0.952380952	1	0.909090909
CLPTM1L AND NOT-SEC63	0.9	1	0.818181818
NOT-ACACB AND TLCD1	0.9	1	0.818181818
NOT-ACACB AND CLCN2	0.9	1	0.818181818
NOT-ACACB AND COL4A2	0.952380952	1	0.909090909
C5orf28 AND NOT-TMEM18	0.9	1	0.818181818
NOT-ACACB AND TMEM182	0.952380952	1	0.909090909
NOT-ACACB AND DCBLD2	0.9	1	0.818181818
LSR AND NOT-TMEM139	0.9	1	0.818181818
NOT-ACACB AND CSF1	0.9	1	0.818181818
DHRS13 AND NOT-MAN2A2	0.9	1	0.818181818
NOT-ACACB AND ACE	0.9	1	0.818181818
TMEM41A AND NOT-PAQR7	0.9	1	0.818181818
GGCX AND NOT-EDNRB	0.916666667	0.846153846	1
OSMR AND NOT-ADCY4	0.9	1	0.818181818
GGCX AND NOT-TAPT1	0.909090909	0.909090909	0.909090909
ATP6V0C AND NOT-TAPT1	0.9	1	0.818181818
NOT-ACACB AND EXT1	0.9	1	0.818181818
FAP AND NOT-LRRN4CL	0.9	1	0.818181818
FAP AND NOT-ITGA7	0.9	1	0.818181818
FAP AND NOT-LCAT	0.9	1	0.818181818
FAP AND NOT-ENPP1	0.9	1	0.818181818
FAP AND NOT-ABCB1	0.9	1	0.818181818
FAP AND NOT-PTGFR	0.9	1	0.818181818
FAP AND NOT-GALNT16	0.9	1	0.818181818
FAP AND NOT-SFRP1	0.9	1	0.818181818
FAP AND NOT-TGFBR3	0.9	1	0.818181818
FAP AND NOT-RECK	0.9	1	0.818181818
NOT-ACACB AND LEMD2	0.952380952	1	0.909090909
NOT-ACACB AND P2RX2	0.909090909	0.909090909	0.909090909
NOT-ACACB AND TMEM9	0.952380952	1	0.909090909
GALNT2 AND NOT-WFDC9	0.9	1	0.818181818
GALNT2 AND NOT-KCNS2	0.9	1	0.818181818
GALNT2 AND NOT-OR1F1	0.9	1	0.818181818
NOT-ACACB AND TKFC	0.909090909	0.909090909	0.909090909
NOT-ACACB AND B3GAT3	0.9	1	0.818181818
GGCX AND NOT-ACACB	1	1	1
GGCX AND NOT-ABCB1	1	1	1
GGCX AND NOT-LRRN3	0.9	1	0.818181818
GGCX AND NOT-SLC22A5	0.9	1	0.818181818
GGCX AND NOT-ANGPTL1	0.909090909	0.909090909	0.909090909
NOT-ACACB AND GJA5	0.9	1	0.818181818
NOT-ACACB AND TMEM97	0.9	1	0.818181818
TMEM41A AND NOT-AQP11	0.9	1	0.818181818
TMEM41A AND NOT-GPR22	0.9	1	0.818181818
NOT-ACACB AND TMEM176B	0.909090909	0.909090909	0.909090909
NOT-ACACB AND SLC35B2	0.952380952	1	0.909090909
NOT-ACACB AND JAG2	0.9	1	0.818181818
NOT-ACACB AND KCNJ8	0.9	1	0.818181818
NOT-ACACB AND OR51M1	0.909090909	0.909090909	0.909090909
NOT-ACACB AND LTBP3	0.9	1	0.818181818
NOT-ACACB AND EPCAM	0.9	1	0.818181818
NOT-ACACB AND NTRK1	0.952380952	1	0.909090909
NOT-ACACB AND TBC1D7	0.952380952	1	0.909090909
NOT-ACACB AND ATRAID	0.909090909	0.909090909	0.909090909
NOT-ACACB AND ACP2	0.952380952	1	0.909090909
NOT-ACACB AND IMPAD1	0.952380952	1	0.909090909
NOT-ACACB AND ZDHHC4	0.952380952	1	0.909090909
NOT-ACACB AND CDCA7L	0.9	1	0.818181818
NOT-ACACB AND PTK7	0.909090909	0.909090909	0.909090909
NOT-ACACB AND PVRL2	0.9	1	0.818181818
NOT-ACACB AND SRPRB	0.909090909	0.909090909	0.909090909
NOT-ACACB AND SDC1	0.9	1	0.818181818
NOT-ACACB AND LMBR1	0.9	1	0.818181818
NOT-ACACB AND C5orf28	0.909090909	0.909090909	0.909090909
NOT-ACACB AND GXYLT2	0.9	1	0.818181818
NOT-ACACB AND SLC35A2	0.909090909	0.909090909	0.909090909
NOT-ACACB AND MOGS	0.909090909	0.909090909	0.909090909
NOT-ACACB AND LRFN4	0.952380952	1	0.909090909
NOT-ACACB AND RNF26	0.9	1	0.818181818
NOT-ACACB AND TMEM185B	0.952380952	1	0.909090909
NOT-ACACB AND SLC39A7	0.909090909	0.909090909	0.909090909
NOT-ACACB AND CD276	0.952380952	1	0.909090909
NOT-ACACB AND AKAP1	0.9	1	0.818181818
NOT-ACACB AND PLVAP	0.952380952	1	0.909090909
NOT-ACACB AND ANTXR1	0.9	1	0.818181818

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
NOT-PMEL AND PMEL	0.952380952	1	0.909090909
MLANA AND NOT-CLDN4	0.952380952	1	0.909090909
CTDNEP1 AND NOT-CLDN4	0.9	1	0.818181818
PPT2 AND NOT-CLDN4	0.9	1	0.818181818
PIGW AND NOT-CLDN4	0.9	1	0.818181818
STX18 AND NOT-CLDN4	0.9	1	0.818181818
MANBAL AND NOT-CLDN4	0.9	1	0.818181818
RER1 AND NOT-CLDN4	0.952380952	1	0.909090909
LMAN2L AND NOT-CLDN4	0.952380952	1	0.909090909
YIPF1 AND NOT-CLDN4	0.9	1	0.818181818
GPR107 AND NOT-CLDN4	0.9	1	0.818181818
SLC45A2 AND NOT-CLDN4	0.952380952	1	0.909090909
SLC5A6 AND NOT-CLDN4	0.952380952	1	0.909090909
MC1R AND NOT-CLDN4	0.9	1	0.818181818
TMEM147 AND NOT-CLDN4	0.9	1	0.818181818
TNFSF9 AND NOT-CLDN4	0.952380952	1	0.909090909
TYRP1 AND NOT-CLDN4	0.9	1	0.818181818
C11orf24 AND NOT-CLDN4	0.9	1	0.818181818
NOT-PMEL AND PMEL	0.952380952	1	0.909090909
BCAP31 AND NOT-CLDN4	0.952380952	1	0.909090909
ABHD12 AND NOT-CLDN4	0.952380952	1	0.909090909
FDPS AND NOT-CLDN4	0.9	1	0.818181818
IL12RB2 AND NOT-CLDN4	0.9	1	0.818181818
PNKD AND NOT-CLDN4	0.909090909	0.90909091	0.909090909
MARVELD1 AND NOT-CLDN4	0.9	1	0.818181818
PQLC1 AND NOT-DRD3	0.909090909	0.90909091	0.909090909
OS9 AND NOT-DRD3	0.952380952	1	0.909090909
MLANA AND NOT-DRD3	1	1	1
PIGL AND NOT-DRD3	0.909090909	0.90909091	0.909090909
SFXN3 AND NOT-DRD3	0.952380952	1	0.909090909
PPT2 AND NOT-DRD3	0.9	1	0.818181818
HSD3B7 AND NOT-DRD3	0.9	1	0.818181818
MANBAL AND NOT-DRD3	0.952380952	1	0.909090909
TMEM138 AND TMEM138	0.952380952	1	0.909090909
LMAN2L AND NOT-DRD3	0.952380952	1	0.909090909
CIRH1A AND NOT-DRD3	0.952380952	1	0.909090909
ERGIC3 AND NOT-DRD3	0.9	1	0.818181818
TIMM22 AND NOT-DRD3	0.9	1	0.818181818
TMEM189 AND NOT-DRD3	0.952380952	1	0.909090909
TMEM184B AND NOT-DRD3	0.952380952	1	0.909090909
GPR107 AND NOT-DRD3	0.9	1	0.818181818
SLC45A2 AND NOT-DRD3	0.952380952	1	0.909090909
SLC5A6 AND NOT-DRD3	0.952380952	1	0.909090909
MC1R AND NOT-DRD3	0.9	1	0.818181818
TMEM147 AND NOT-DRD3	0.9	1	0.818181818
EXT1 AND NOT-DRD3	0.952380952	1	0.909090909
TNFSF9 AND NOT-DRD3	0.909090909	0.90909091	0.909090909
NDST1 AND NOT-DRD3	0.909090909	0.90909091	0.909090909
C11orf24 AND NOT-DRD3	0.9	1	0.818181818
NOT-PMEL AND PMEL	0.952380952	1	0.909090909
ABHD12 AND NOT-DRD3	0.909090909	0.90909091	0.909090909
FDPS AND NOT-DRD3	1	1	1
IL12RB2 AND NOT-DRD3	0.9	1	0.818181818
PNKD AND NOT-DRD3	0.952380952	1	0.909090909
ROMO1 AND NOT-DRD3	0.952380952	1	0.909090909
NOT-SCN4A AND PICK1	0.9	1	0.818181818
MLANA AND PICK1	0.952380952	1	0.909090909
NOT-CHRNA2 AND PICK1	0.9	1	0.818181818
SFXN3 AND PICK1	0.952380952	1	0.909090909
OXA1L AND PICK1	0.9	1	0.818181818
PARL AND PICK1	0.9	1	0.818181818
MANBAL AND NOT-PICK1	0.952380952	1	0.909090909
NOT-NMUR2 AND PICK1	0.9	1	0.818181818
NOT-TMEM95 AND PICK1	0.9	1	0.818181818
TMEM138 AND PICK1	0.9	1	0.818181818
NOT-ISM2 AND PICK1	0.9	1	0.818181818
AUP1 AND PICK1	0.9	1	0.818181818
NOT-ATCAY AND PICK1	0.9	1	0.818181818
SLC45A2 AND NOT-PICK1	0.952380952	1	0.909090909
MC1R AND NOT-PICK1	0.9	1	0.818181818
TNFSF9 AND NOT-PICK1	0.909090909	0.90909091	0.909090909
NOT-CFTR AND PICK1	0.9	1	0.818181818
NOT-PTCH2 AND PICK1	0.9	1	0.818181818
NOT-PMEL AND PMEL	0.952380952	1	0.909090909
BCAP31 AND PICK1	0.9	1	0.818181818
NDUFA1 AND PICK1	0.9	1	0.818181818

Antigen Pair Logic	F1	Precision	Recall
NOT-ACACB AND SOAT2	0.9	1	0.818181818
NOT-ACACB AND PPAP2C	0.952380952	1	0.909090909
NOT-ACACB AND TNFRSF18	0.909090909	0.909090909	0.909090909
NOT-ACACB AND MARVELD3	0.9	1	0.818181818
NETO2 AND NOT-KCNS2	0.9	1	0.818181818
ATP6V0C AND NOT-MAN2A2	0.952380952	1	0.909090909
TMEM206 AND NOT-MAN2A2	0.909090909	0.909090909	0.909090909
OSMR AND NOT-CRIM1	0.9	1	0.818181818
LSR AND NOT-SEMA4G	0.9	1	0.818181818
LSR AND NOT-CYP4F12	0.9	1	0.818181818
TMEM41A AND NOT-ENPP1	0.9	1	0.818181818
DNAJB11 AND NOT-KIAA1109	0.9	1	0.818181818
ATP6V0C AND PFN2	0.9	1	0.818181818
MOGS AND NOT-ABCB1	0.9	1	0.818181818
CLPTM1L AND NOT-ABCB1	0.9	1	0.818181818
TMEM41A AND NOT-ABCB1	0.9	1	0.818181818
ATP6V0C AND NOT-RNF130	0.9	1	0.818181818
ATP6V0C AND NOT-TNFRSF14	0.9	1	0.818181818
ATP6V0C AND NOT-CD101	0.9	1	0.818181818
ATP6V0C AND NOT-FIG4	0.9	1	0.818181818
OSMR AND NOT-FXYD1	0.9	1	0.818181818
TMEM41A AND NOT-VSIG10	0.9	1	0.818181818
PPAP2C AND NOT-PCDHA6	0.952380952	1	0.909090909
OSMR AND NOT-SCN7A	0.9	1	0.818181818
C5orf28 AND NOT-SMIM15	0.9	1	0.818181818
NOT-KIAA1109 AND C5orf28	0.9	1	0.818181818
OSMR AND NOT-TGFBR3	0.9	1	0.818181818
OSMR AND NOT-ACSS3	0.9	1	0.818181818
OSMR AND NOT-UPK3B	0.9	1	0.818181818
TMEM41A AND NOT-PRRT1	0.9	1	0.818181818
TMEM41A AND NOT-KIAA1109	0.9	1	0.818181818
OSMR AND NOT-KIAA1109	0.9	1	0.818181818
OSMR AND NOT-ANGPTL1	0.9	1	0.818181818
OSMR AND NOT-ADAMTS1	0.9	1	0.818181818
Leukemia, Myeloid, Acute (AML)			
FLT3	0.935222672	0.950617284	0.920318725
NOT-SPINK2 AND SPINK2	0.912863071	0.952380952	0.876494024
SLC22A16 AND NOT-SDC1	0.90513834	0.898039216	0.912350598
FUT4 AND NOT-SDC1	0.964285714	0.960474308	0.96812749
ELANE AND NOT-SDC1	0.909819639	0.915322581	0.90438247
NOT-SPINK2 AND SPINK2	0.912863071	0.952380952	0.876494024
MS4A3 AND NOT-SLAMF7	0.932270916	0.932270916	0.932270916
TMC8 AND NOT-SLAMF7	0.90513834	0.898039216	0.912350598
NOT-PFN2 AND NOT-SLAMF7	0.938461538	0.907063197	0.972111554
CD33 AND NOT-SLAMF7	0.916996047	0.909803922	0.924302789
NOT-SPINK2 AND SPINK2	0.912863071	0.952380952	0.876494024
FUT4 AND NOT-SLAMF7	0.971544715	0.991701245	0.952191235
NRROS AND NOT-SLAMF7	0.940936864	0.9625	0.920318725
P2RY8 AND NOT-SLAMF7	0.909774436	0.861209964	0.964143426
HCST AND NOT-SLAMF7	0.920454545	0.877256318	0.96812749
CXCL8 AND NOT-SLAMF7	0.905027933	0.84965035	0.96812749
TSPAN32 AND NOT-SLAMF7	0.947162427	0.930769231	0.964143426
P2RX1 AND NOT-SLAMF7	0.907063197	0.850174216	0.972111554
HBD AND NOT-SLAMF7	0.939096267	0.926356589	0.952191235
ITGA4 AND NOT-SLAMF7	0.910569106	0.929460581	0.892430279
NOT-TMTC3 AND CD44	0.901303538	0.846153846	0.964143426
NOT-HEG1 AND CD44	0.909090909	0.944206009	0.876494024
NOT-PFN2 AND CD44	0.911111111	0.851211073	0.980079681
NOT-EXT1 AND CD44	0.900934579	0.848591549	0.960159363
NOT-FAM69A AND CD44	0.944971537	0.902173913	0.992031873
NOT-GPR155 AND CD44	0.923976608	0.904580153	0.944223108
NOT-WLS AND CD44	0.908745247	0.869090909	0.952191235
NOT-SPINK2 AND SPINK2	0.912863071	0.952380952	0.876494024
NOT-BMPR1A AND CD44	0.912959381	0.887218045	0.940239044
SLC22A16 AND CD44	0.914285714	0.937238494	0.892430279
NOT-ARMCX2 AND CD44	0.922787194	0.875	0.976095618
NOT-SLC37A3 AND CD44	0.923076923	0.892193309	0.956175299
FUT4 AND CD44	0.903954802	0.857142857	0.956175299
FLT3 AND CD44	0.930894309	0.950207469	0.912350598
CD44 AND NOT-SMIM14	0.9140625	0.896551724	0.932270916
NOT-GAL3ST4 AND CD44	0.901960784	0.888030888	0.916334661
HBD AND CD44	0.903591682	0.85971223	0.952191235
NOT-PHLDB2 AND CD44	0.947775629	0.921052632	0.976095618
NOT-LRIG1 AND CD44	0.920222635	0.861111111	0.988047809
NOT-SPINK2 AND SPINK2	0.912863071	0.952380952	0.876494024
FLT3 AND NOT-IL11RA	0.940936864	0.9625	0.920318725

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
ROMO1 AND NOT-PICK1	0.952380952	1	0.909090909
NOT-STX1B AND PICK1	0.9	1	0.818181818
PQLC1 AND PSMB7	0.9	1	0.818181818
OS9 AND OS9	0.9	1	0.818181818
FAM3A AND PSMB7	0.9	1	0.818181818
MLANA AND PSMB7	1	1	1
PIGL AND PSMB7	0.9	1	0.818181818
LARGE AND PSMB7	0.9	1	0.818181818
GPR137B AND PSMB7	0.9	1	0.818181818
HGSNAT AND PSMB7	0.9	1	0.818181818
CLCN5 AND PSMB7	0.952380952	1	0.909090909
VANGL2 AND PSMB7	0.952380952	1	0.909090909
APMAP AND PSMB7	0.952380952	1	0.909090909
FAM134A AND PSMB7	0.952380952	1	0.909090909
TMEM138 AND PSMB7	0.9	1	0.818181818
NOT-CIRH1A AND CIRH1A	0.952380952	1	0.909090909
NOT-ERGIC3 AND ERGIC3	0.9	1	0.818181818
YIF1B AND PSMB7	0.9	1	0.818181818
TMEM184B AND PSMB7	0.952380952	1	0.909090909
SLC5A6 AND PSMB7	0.952380952	1	0.909090909
PTPRS AND PSMB7	0.9	1	0.818181818
TMEM147 AND TMEM147	0.9	1	0.818181818
EXT1 AND PSMB7	0.952380952	1	0.909090909
TNFSF9 AND PSMB7	0.909090909	0.90909091	0.909090909
NDST1 AND PSMB7	0.952380952	1	0.909090909
NOT-TMEM105 AND PSMB7	0.952380952	1	0.909090909
NOT-PMEL AND PMEL	0.952380952	1	0.909090909
CLCC1 AND PSMB7	0.9	1	0.818181818
ABHD12 AND PSMB7	0.952380952	1	0.909090909
FDPS AND PSMB7	0.952380952	1	0.909090909
ICMT AND PSMB7	0.9	1	0.818181818
PNKD AND PSMB7	0.909090909	0.90909091	0.909090909
PFKP AND PSMB7	0.9	1	0.818181818
ROMO1 AND PSMB7	0.952380952	1	0.909090909
GCNT2 AND PSMB7	0.9	1	0.818181818
CELSR2 AND PSMB7	0.952380952	1	0.909090909
PQLC1 AND NOT-CSF3	0.9	1	0.818181818
SLC35E3 AND NOT-CSF3	0.952380952	1	0.909090909
OS9 AND OS9	0.9	1	0.818181818
SLAMF8 AND NOT-CSF3	0.9	1	0.818181818
WFS1 AND NOT-CSF3	0.9	1	0.818181818
TMEM187 AND NOT-CSF3	0.952380952	1	0.909090909
MLANA AND NOT-CSF3	0.952380952	1	0.909090909
CXCL9 AND NOT-CSF3	0.9	1	0.818181818
PIGL AND NOT-CSF3	0.952380952	1	0.909090909
NOT-CSF3 AND NRM	0.9	1	0.818181818
PQLC2 AND NOT-CSF3	0.9	1	0.818181818
NOT-CSF3 AND CNNM4	0.9	1	0.818181818
SFXN3 AND CSF3	0.952380952	1	0.909090909
PPT2 AND NOT-CSF3	0.9	1	0.818181818
KIAA1024 AND NOT-CSF3	0.9	1	0.818181818
ST7L AND NOT-CSF3	0.9	1	0.818181818
BNIP1 AND NOT-CSF3	0.9	1	0.818181818
CLCN5 AND NOT-CSF3	0.952380952	1	0.909090909
SMIM4 AND NOT-CSF3	0.9	1	0.818181818
VANGL2 AND NOT-CSF3	0.9	1	0.818181818
TMEM138 AND TMEM138	0.952380952	1	0.909090909
NPC1 AND NOT-CSF3	0.952380952	1	0.909090909
NOT-CIRH1A AND CIRH1A	0.952380952	1	0.909090909
ERGIC3 AND CSF3	0.9	1	0.818181818
NOT-CSF3 AND CDH24	0.9	1	0.818181818
TMEM184B AND NOT-CSF3	0.9	1	0.818181818
FAM174B AND NOT-CSF3	0.9	1	0.818181818
NOT-CSF3 AND PIGQ	0.9	1	0.818181818
YIPF1 AND NOT-CSF3	0.909090909	0.90909091	0.909090909
SLC45A2 AND NOT-CSF3	0.952380952	1	0.909090909
SLC5A6 AND NOT-CSF3	0.9	1	0.818181818
NOT-CSF3 AND LRRC8E	0.9	1	0.818181818
ARMCX1 AND NOT-CSF3	0.9	1	0.818181818
TMEM147 AND TMEM147	0.9	1	0.818181818
EXT1 AND NOT-CSF3	0.9	1	0.818181818
TNFSF9 AND NOT-CSF3	0.9	1	0.818181818
CDC14B AND NOT-CSF3	0.9	1	0.818181818
TYRP1 AND NOT-CSF3	0.952380952	1	0.909090909
NDST1 AND NOT-CSF3	0.9	1	0.818181818
NOT-PMEL AND PMEL	0.952380952	1	0.909090909

Antigen Pair Logic	F1	Precision	Recall
P2RX1 AND NOT-IL11RA	0.908382066	0.889312977	0.928286853
NOT-SPINK2 AND SPINK2	0.912863071	0.952380952	0.876494024
NOT-FLT3 AND FLT3	0.935222672	0.950617284	0.920318725
TMC8 AND NOT-CD19	0.906504065	0.925311203	0.888446215
NOT-PFN2 AND NOT-CD19	0.933333333	0.946721311	0.920318725
NOT-SPINK2 AND SPINK2	0.912863071	0.952380952	0.876494024
NOT-FLT3 AND FLT3	0.935222672	0.950617284	0.920318725
NRROS AND NOT-CD19	0.933333333	0.946721311	0.920318725
P2RY8 AND NOT-CD19	0.911538462	0.881040892	0.944223108
TSPAN32 AND NOT-CD19	0.918604651	0.894339623	0.944223108
NOT-ATP9A AND NOT-CD19	0.90234375	0.885057471	0.920318725
HBD AND NOT-CD19	0.918811881	0.913385827	0.924302789
NOT-SPINK2 AND SPINK2	0.912863071	0.952380952	0.876494024
NOT-FLT3 AND FLT3	0.935222672	0.950617284	0.920318725
NOT-SPINK2 AND SPINK2	0.912863071	0.952380952	0.876494024
FUT4 AND NOT-MUC1	0.941883768	0.947580645	0.93625498
NOT-FLT3 AND FLT3	0.935222672	0.950617284	0.920318725
NOT-SPINK2 AND SPINK2	0.912863071	0.952380952	0.876494024
SLC22A16 AND NOT-MUC13	0.901574803	0.891050584	0.912350598
FUT4 AND NOT-MUC13	0.960474308	0.952941176	0.96812749
TMC8 AND NOT-CD22	0.904483431	0.885496183	0.924302789
NOT-PFN2 AND NOT-CD22	0.959223301	0.935606061	0.984063745
NOT-DIP2C AND NOT-CD22	0.903460838	0.832214765	0.988047809
NOT-SPINK2 AND SPINK2	0.912863071	0.952380952	0.876494024
NOT-FLT3 AND FLT3	0.935222672	0.950617284	0.920318725
NOT-CADM1 AND NOT-CD22	0.906309751	0.871323529	0.944223108
NRROS AND NOT-CD22	0.935742972	0.943319838	0.928286853
P2RY8 AND NOT-CD22	0.921052632	0.871886121	0.976095618
TSPAN32 AND NOT-CD22	0.931818182	0.888086643	0.980079681
NOT-ATP9A AND NOT-CD22	0.902857143	0.864963504	0.944223108
HBD AND NOT-CD22	0.92543021	0.889705882	0.964143426
NOT-SPINK2 AND SPINK2	0.912863071	0.952380952	0.876494024
ELANE AND NOT-EPHA3	0.90438247	0.90438247	0.90438247
NOT-SPINK2 AND CSPG4	0.912863071	0.952380952	0.876494024
NOT-SPINK2 AND MUC16	0.912863071	0.952380952	0.876494024
SLC22A16 AND NOT-MUC16	0.901574803	0.891050584	0.912350598
NOT-SPINK2 AND SPINK2	0.912863071	0.952380952	0.876494024
ELANE AND NOT-ROR1	0.902584493	0.900793651	0.90438247
NOT-SPINK2 AND SPINK2	0.912863071	0.952380952	0.876494024
NOT-FLT3 AND FLT3	0.935222672	0.950617284	0.920318725
NOT-SPINK2 AND SPINK2	0.912863071	0.952380952	0.876494024
NOT-SPINK2 AND SPINK2	0.912863071	0.952380952	0.876494024
NOT-FLT3 AND FLT3	0.935222672	0.950617284	0.920318725
NOT-SPINK2 AND TNFSF11	0.912863071	0.952380952	0.876494024
NOT-SPINK2 AND SPINK2	0.912863071	0.952380952	0.876494024
FLT3 AND NOT-FOLR2	0.935281837	0.98245614	0.892430279
ELANE AND NOT-FOLR2	0.904564315	0.943722944	0.868525896
SMIM3 AND NOT-FOLR2	0.928425358	0.953781513	0.90438247
NOT-SPINK2 AND SPINK2	0.912863071	0.952380952	0.876494024
MS4A3 AND NOT-TNFRSF17	0.920318725	0.920318725	0.920318725
TMC8 AND NOT-TNFRSF17	0.902970297	0.897637795	0.908366534
CD33 AND NOT-TNFRSF17	0.900195695	0.884615385	0.916334661
NOT-SPINK2 AND SPINK2	0.912863071	0.952380952	0.876494024
FUT4 AND NOT-TNFRSF17	0.965376782	0.9875	0.944223108
NRROS AND NOT-TNFRSF17	0.928716904	0.95	0.908366534
P2RY8 AND NOT-TNFRSF17	0.909433962	0.863799283	0.960159363
HCST AND NOT-TNFRSF17	0.901303538	0.846153846	0.964143426
TSPAN32 AND NOT-TNFRSF17	0.936170213	0.909774436	0.964143426
HBD AND NOT-TNFRSF17	0.922178988	0.901140684	0.944223108
ITGA4 AND NOT-TNFRSF17	0.91206544	0.93697479	0.888446215
NOT-SPINK2 AND TNFRSF8	0.912863071	0.952380952	0.876494024
NOT-SPINK2 AND SPINK2	0.912863071	0.952380952	0.876494024
NOT-FLT3 AND FLT3	0.935222672	0.950617284	0.920318725
NOT-SPINK2 AND SPINK2	0.912863071	0.952380952	0.876494024
NOT-SPINK2 AND SPINK2	0.912863071	0.952380952	0.876494024
FLT3 AND NOT-FOLH1	0.944785276	0.970588235	0.920318725
NOT-SPINK2 AND SPINK2	0.912863071	0.952380952	0.876494024
SLC22A16 AND NOT-ERBB4	0.903353057	0.89453125	0.912350598
NOT-SPINK2 AND SPINK2	0.912863071	0.952380952	0.876494024
SLC22A16 AND NOT-ERBB2	0.901574803	0.891050584	0.912350598
FUT4 AND NOT-ERBB2	0.960474308	0.952941176	0.96812749
ELANE AND NOT-ERBB2	0.902584493	0.900793651	0.90438247
SMIM24 AND NOT-KDR	0.912	0.915662651	0.908366534
LRRC70 AND NOT-KDR	0.938613861	0.933070866	0.944223108
NOT-SPINK2 AND SPINK2	0.912863071	0.952380952	0.876494024
FLT3 AND NOT-KDR	0.948665298	0.978813559	0.920318725

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
NOT-CSF3 AND TLCD1	0.952380952	1	0.909090909
BCAP31 AND NOT-CSF3	0.9	1	0.818181818
NOT-CSF3 AND TMEM110	0.9	1	0.818181818
CLCC1 AND NOT-CSF3	0.9	1	0.818181818
ABHD12 AND NOT-CSF3	0.9	1	0.818181818
PNKD AND NOT-CSF3	0.9	1	0.818181818
PRRT3 AND NOT-CSF3	0.9	1	0.818181818
ROMO1 AND NOT-CSF3	0.952380952	1	0.909090909
CELSR2 AND NOT-CSF3	0.9	1	0.818181818
PRADC1 AND NOT-CSF3	0.9	1	0.818181818
MLANA AND NOT-GCG	1	1	1
SFXN3 AND NOT-GCG	0.952380952	1	0.909090909
PPT2 AND NOT-GCG	0.9	1	0.818181818
SLC25A33 AND NOT-GCG	0.9	1	0.818181818
TMEM138 AND TMEM138	0.952380952	1	0.909090909
LMAN2L AND NOT-GCG	0.909090909	0.90909091	0.909090909
NPC1 AND NOT-GCG	0.952380952	1	0.909090909
CIRH1A AND NOT-GCG	0.952380952	1	0.909090909
TMEM189 AND NOT-GCG	0.952380952	1	0.909090909
SLC45A2 AND NOT-GCG	0.952380952	1	0.909090909
SLC5A6 AND NOT-GCG	0.952380952	1	0.909090909
MC1R AND NOT-GCG	0.9	1	0.818181818
SLC35A4 AND NOT-GCG	0.9	1	0.818181818
TNFSF9 AND NOT-GCG	0.952380952	1	0.909090909
NOT-PMEL AND PMEL	0.952380952	1	0.909090909
FDPS AND NOT-GCG	0.952380952	1	0.909090909
IL12RB2 AND NOT-GCG	0.9	1	0.818181818
PNKD AND NOT-GCG	0.952380952	1	0.909090909
FDPS AND NOT-NAALAD2	1	1	1
MLANA AND NOT-NAALAD2	1	1	1
FDPS AND NAALADL1	1	1	1
MLANA AND NOT-NAALADL1	1	1	1
MLANA AND KCNE3	1	1	1
AUP1 AND NOT-KCNE3	1	1	1
PARL AND NOT-KCNE3	1	1	1
FDPS AND CDH2	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND TMEM170B	1	1	1
AUP1 AND NOT-TMEM170B	1	1	1
PARL AND NOT-TMEM170B	1	1	1
FDPS AND NOT-TMEM221	1	1	1
MLANA AND NOT-TMEM221	1	1	1
NPC1 AND NOT-TMEM221	1	1	1
MLANA AND NOT-LRRC70	1	1	1
MLANA AND NEMP2	1	1	1
CLCN5 AND NOT-100132596?	1	1	1
MLANA AND NOT-100132596?	1	1	1
MLANA AND NOT-BCL2L10	1	1	1
NOT-MLANA AND CDH3	1	1	1
NOT-PMEL AND CDH3	1	1	1
MLANA AND NOT-HCN4	1	1	1
NPC1 AND NOT-HCN4	1	1	1
SLC35E3 AND NOT-HCN4	1	1	1
MLANA AND PIGK	1	1	1
MLANA AND TIMM23	1	1	1
FDPS AND NOT-ERVMER34-1	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND NOT-TMEM239	1	1	1
MLANA AND NOT-CDH4	1	1	1
MLANA AND MLANA	1	1	1
FDPS AND APELA	1	1	1
MLANA AND NOT-APELA	1	1	1
MLANA AND MLANA	1	1	1
FDPS AND NOT-MRLN	1	1	1
MLANA AND NOT-MRLN	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND NOT-SMLR1	1	1	1
NPC1 AND NOT-SMLR1	1	1	1
MLANA AND TMEM178B	1	1	1
MLANA AND MLANA	1	1	1
NOT-CHRNA2 AND 100507547?	1	1	1
MLANA AND 100507547?	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND UBA2	1	1	1
FDPS AND ABCC5	1	1	1
MLANA AND ABCC5	1	1	1

Antigen Pair Logic	F1	Precision	Recall
CXCL8 AND NOT-KDR	0.923364486	0.86971831	0.984063745
ELANE AND NOT-KDR	0.915322581	0.926530612	0.90438247
SMIM3 AND NOT-KDR	0.96812749	0.96812749	0.96812749
CD33 AND NOT-ENDOD1	0.909465021	0.940425532	0.880478088
CD33 AND NOT-NELL2	0.909803922	0.895752896	0.924302789
NOT-TMTC3 AND CD33	0.908396947	0.871794872	0.948207171
CD33 AND NOT-LY9	0.903100775	0.879245283	0.928286853
CD33 AND NOT-S1PR1	0.92371134	0.957264957	0.892430279
CD33 AND NOT-FAM200A	0.90625	0.888888889	0.924302789
CD33 AND NOT-CD1C	0.907258065	0.918367347	0.896414343
CD33 AND NOT-CDC14B	0.906976744	0.883018868	0.932270916
CD33 AND NOT-LPAR1	0.912280702	0.893129771	0.932270916
CD33 AND NOT-STX17	0.91322314	0.948497854	0.880478088
NOT-HEG1 AND CD33	0.918238994	0.969026549	0.87250996
CD33 AND NOT-TGFBI	0.903361345	0.955555556	0.856573705
CD33 AND NOT-IL7R	0.919449902	0.906976744	0.932270916
CD33 AND NOT-SLC44A2	0.934156379	0.965957447	0.90438247
CD33 AND NOT-PXYLP1	0.93081761	0.982300885	0.884462151
NOT-PFN2 AND CD33	0.917293233	0.868327402	0.972111554
CD33 AND NOT-ADAM19	0.911591356	0.899224806	0.924302789
CD33 AND NOT-TMEM140	0.938016529	0.974248927	0.90438247
CD33 AND NOT-SLC18B1	0.9	0.990430622	0.824701195
CD33 AND NOT-IFNLR1	0.924901186	0.917647059	0.932270916
CD33 AND NOT-INSIG2	0.913043478	0.905882353	0.920318725
NOT-EXT1 AND CD33	0.914396887	0.893536122	0.93625498
NOT-FAM69A AND CD33	0.96031746	0.956521739	0.964143426
CD33 AND NOT-RTN1	0.934693878	0.958158996	0.912350598
CD33 AND NOT-CPED1	0.901467505	0.951327434	0.856573705
CD33 AND NOT-ENPP5	0.906614786	0.885931559	0.928286853
CD33 AND NOT-SYNJ2BP	0.916334661	0.916334661	0.916334661
CD33 AND NOT-CCR7	0.900584795	0.881679389	0.920318725
CD33 AND NOT-PVRIG	0.904854369	0.882575758	0.928286853
CD33 AND NOT-C19orf12	0.904483431	0.885496183	0.924302789
CD33 AND NOT-GPR155	0.929859719	0.935483871	0.924302789
CD33 AND NOT-HLA-DOB	0.914396887	0.893536122	0.93625498
CD33 AND NOT-CD2	0.909448819	0.898832685	0.920318725
NOT-PTPRM AND CD33	0.900195695	0.884615385	0.916334661
CD33 AND NOT-ADTRP	0.927021696	0.91796875	0.93625498
CD33 AND NOT-LDLRAD4	0.905263158	0.959821429	0.856573705
CD33 AND NOT-SEC63	0.908	0.911646586	0.90438247
CD33 AND NOT-NPC1	0.901185771	0.894117647	0.908366534
NOT-WLS AND CD33	0.922772277	0.917322835	0.928286853
CD33 AND NOT-CD3G	0.900195695	0.884615385	0.916334661
CD33 AND NOT-SLC46A3	0.92555332	0.93495935	0.916334661
CD33 AND NOT-IL18R1	0.901803607	0.907258065	0.896414343
CD33 AND NOT-CMTM8	0.925311203	0.965367965	0.888446215
CD33 AND NOT-GIMAP1	0.901353965	0.87593985	0.928286853
CD33 AND NOT-ST3GAL5	0.926530612	0.949790795	0.90438247
CD33 AND NOT-CLCN3	0.906560636	0.904761905	0.908366534
CD33 AND NOT-PTPRO	0.923076923	0.9140625	0.932270916
CD33 AND NOT-IL21R	0.908737864	0.886363636	0.932270916
CD33 AND NOT-FAM171A1	0.917695473	0.94893617	0.888446215
CD33 AND NOT-NT5E	0.921529175	0.930894309	0.912350598
CD33 AND NOT-MAL	0.903885481	0.928571429	0.880478088
CD33 AND NOT-STEAP4	0.918163673	0.92	0.916334661
NOT-SPINK2 AND CD33	0.912863071	0.952380952	0.876494024
CD33 AND NOT-YIPF5	0.906976744	0.883018868	0.932270916
CD33 AND NOT-STX6	0.908382066	0.889312977	0.928286853
CD33 AND NOT-FAM134B	0.93442623	0.962025316	0.908366534
NOT-BMPR1A AND CD33	0.905222437	0.879699248	0.932270916
CD33 AND NOT-BMPR2	0.929460581	0.96969697	0.892430279
CD33 AND NOT-ARMCX2	0.915520629	0.903100775	0.928286853
CD33 AND NOT-ANKH	0.910891089	0.905511811	0.916334661
NOT-SLC37A3 AND CD33	0.939759036	0.947368421	0.932270916
CD33 AND NOT-CD28	0.911591356	0.899224806	0.924302789
CD33 AND NOT-FCRL3	0.903474903	0.876404494	0.932270916
CD33 AND NOT-SEMA4B	0.909090909	0.901960784	0.916334661
NOT-FLT3 AND FLT3	0.935222672	0.950617284	0.920318725
CD33 AND NOT-MGAT5	0.929166667	0.973799127	0.888446215
CD33 AND NOT-CASD1	0.905511811	0.894941634	0.916334661
CD33 AND NOT-MGAT4A	0.931726908	0.939271255	0.924302789
CD33 AND NOT-DPP4	0.935222672	0.950617284	0.920318725
CD33 AND NOT-FCMR	0.909090909	0.883458647	0.93625498
CD33 AND NOT-PAG1	0.906560636	0.904761905	0.908366534
CD33 AND NOT-KLRD1	0.900383142	0.867158672	0.93625498
CD33 AND NOT-RARRES3	0.932790224	0.954166667	0.912350598

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
NPC1 AND ABCC5	1	1	1
FDPS AND ABCB6	1	1	1
CLCN5 AND NOT-CDH7	1	1	1
FDPS AND NOT-CDH7	1	1	1
MLANA AND NOT-CDH7	1	1	1
FDPS AND NOT-ABCC9	1	1	1
MLANA AND NOT-ABCC9	1	1	1
MLANA AND SCAMP2	1	1	1
MLANA AND SCAMP3	1	1	1
FDPS AND NOT-MUC12	1	1	1
MLANA AND NOT-MUC12	1	1	1
MLANA AND NOT-PTPRU	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND MLANA	1	1	1
NPC1 AND ATP9A	1	1	1
MLANA AND MLANA	1	1	1
NPC1 AND NOT-CDH9	1	1	1
MLANA AND MLANA	1	1	1
NOT-CDH10 AND TLCD1	1	1	1
CLCN5 AND NOT-CDH10	1	1	1
FDPS AND NOT-CDH10	1	1	1
MLANA AND NOT-CDH10	1	1	1
NOT-CDH10 AND SUSD5	1	1	1
NOT-CDH10 AND KRT5	1	1	1
NPC1 AND NOT-CDH10	1	1	1
NOT-CDH10 AND IL20RB	1	1	1
SLC35E3 AND NOT-CDH10	1	1	1
NOT-CDH10 AND TYRP1	1	1	1
NOT-CDH10 AND DHRS11	1	1	1
TMEM187 AND NOT-CDH10	1	1	1
NOT-CDH10 AND LY6D	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND NOT-TSPAN5	1	1	1
MLANA AND TSPAN3	1	1	1
MLANA AND NOT-CDH11	1	1	1
MLANA AND NOT-TSPAN2	1	1	1
MLANA AND PREB	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND NOT-CLEC3A	1	1	1
MLANA AND NOT-ADGRG2	1	1	1
CLCN5 AND NOT-CDH16	1	1	1
FDPS AND NOT-CDH16	1	1	1
MLANA AND NOT-CDH16	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND MLANA	1	1	1
FDPS AND NOT-CDH17	1	1	1
MLANA AND NOT-CDH17	1	1	1
NPC1 AND NOT-CDH17	1	1	1
MLANA AND NOT-LPAR6	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND NOT-CHST4	1	1	1
CLCN5 AND NOT-CDH18	1	1	1
FDPS AND NOT-CDH18	1	1	1
MLANA AND NOT-CDH18	1	1	1
NPC1 AND NOT-CDH18	1	1	1
MLANA AND NOT-DHRS9	1	1	1
MLANA AND CNIH1	1	1	1
MLANA AND NOT-TENM1	1	1	1
MLANA AND NOT-101805491?	1	1	1
NPC1 AND NOT-101805491?	1	1	1
NOT-SLC34A2 AND LHFP	1	1	1
NOT-PTH2 AND LHFP	1	1	1
NOT-CHRNA2 AND LHFP	1	1	1
NOT-CD300LB AND LHFP	1	1	1
NOT-SYT2 AND LHFP	1	1	1
NOT-CLDN7 AND LHFP	1	1	1
NOT-SLC34A3 AND LHFP	1	1	1
NOT-ISM2 AND LHFP	1	1	1
NOT-ZNRF4 AND LHFP	1	1	1
NOT-CLDN19 AND LHFP	1	1	1
NOT-DNAH10 AND LHFP	1	1	1
NOT-FCN2 AND LHFP	1	1	1
FDPS AND LHFP	1	1	1
MLANA AND LHFP	1	1	1
NOT-CALHM1 AND LHFP	1	1	1

Antigen Pair Logic	F1	Precision	Recall
CD33 AND NOT-BTN3A1	0.919765166	0.903846154	0.93625498
CD33 AND NOT-FCRLA	0.901353965	0.87593985	0.928286853
CD33 AND NOT-PTCH1	0.918811881	0.913385827	0.924302789
CD33 AND NOT-GLCCI1	0.908730159	0.90513834	0.912350598
CD33 AND NOT-GPR171	0.917835671	0.923387097	0.912350598
CD33 AND NOT-C12orf49	0.909803922	0.895752896	0.924302789
CD33 AND NOT-KMO	0.903474903	0.876404494	0.932270916
CD33 AND NOT-MFAP3L	0.91015625	0.892720307	0.928286853
CD33 AND NOT-LPAR5	0.901734104	0.873134328	0.932270916
CD33 AND NOT-BTLA	0.906976744	0.883018868	0.932270916
CD33 AND NOT-PTGDR	0.912280702	0.893129771	0.932270916
CD33 AND NOT-BTN2A2	0.928286853	0.928286853	0.928286853
CD33 AND NOT-SLC17A5	0.901574803	0.891050584	0.912350598
CD33 AND NOT-EPHA4	0.927021696	0.91796875	0.93625498
CD33 AND NOT-MINOS1	0.934693878	0.958158996	0.912350598
NOT-TMCC1 AND CD33	0.922131148	0.949367089	0.896414343
CD33 AND NOT-SLC9A9	0.90729783	0.8984375	0.916334661
CD33 AND NOT-SLC4A7	0.905737705	0.932489451	0.880478088
CD33 AND NOT-SLC16A7	0.905811623	0.911290323	0.900398406
CD33 AND NOT-CHI3L2	0.934131737	0.936	0.932270916
CD33 AND NOT-CCR6	0.9	0.869888476	0.932270916
CD33 AND NOT-KIAA0040	0.919765166	0.903846154	0.93625498
CD33 AND NOT-SGPL1	0.909819639	0.915322581	0.90438247
NOT-TGFBR3 AND CD33	0.909853249	0.960176991	0.864541833
CD33 AND NOT-SIRPG	0.901734104	0.873134328	0.932270916
CD33 AND NOT-SMIM14	0.919587629	0.952991453	0.888446215
CD33 AND NOT-FAM73A	0.902111324	0.87037037	0.93625498
CD33 AND NOT-CYP2U1	0.917322835	0.906614786	0.928286853
NOT-VAPB AND CD33	0.93592233	0.912878788	0.960159363
CD33 AND NOT-SP4	0.913043478	0.905882353	0.920318725
CD33 AND NOT-GAL3ST4	0.90797546	0.932773109	0.884462151
CD33 AND NOT-HMOX2	0.916666667	0.913043478	0.920318725
CD33 AND NOT-BTN3A3	0.93495935	0.954356846	0.916334661
CD33 AND NOT-TMEM204	0.910852713	0.886792453	0.93625498
CD33 AND NOT-CD27	0.906614786	0.885931559	0.928286853
CD33 AND NOT-ITPR3	0.914396887	0.893536122	0.93625498
CD33 AND NOT-AMIGO2	0.931451613	0.942857143	0.920318725
CD33 AND NOT-TUBD1	0.919449902	0.906976744	0.932270916
CD33 AND NOT-TRAT1	0.903846154	0.873605948	0.93625498
CD33 AND NOT-RHBDD1	0.906976744	0.883018868	0.932270916
CD33 AND NOT-TMEM2	0.910505837	0.88973384	0.932270916
NOT-PHLDB2 AND CD33	0.935420744	0.919230769	0.952191235
CD33 AND NOT-SEC62	0.905349794	0.936170213	0.876494024
CD33 AND NOT-STX7	0.926441352	0.924603175	0.928286853
CD33 AND NOT-AQP3	0.904862579	0.963963964	0.852589641
CD33 AND NOT-LRRN3	0.926441352	0.924603175	0.928286853
CD33 AND NOT-RNF13	0.903732809	0.891472868	0.916334661
CD33 AND NOT-LRIG1	0.939516129	0.951020408	0.928286853
CD33 AND NOT-ANKRD46	0.905811623	0.911290323	0.900398406
CD33 AND NOT-MAN1C1	0.927966102	0.990950226	0.87250996
CD33 AND SMIM3	0.900398406	0.900398406	0.900398406
CD33 AND NOT-INPP4A	0.909090909	0.883458647	0.93625498
NOT-SPINK2 AND SPINK2	0.912863071	0.952380952	0.876494024
SLC22A16 AND NOT-PSCA	0.901574803	0.891050584	0.912350598
SMIM24 AND NOT-ERBB3	0.906930693	0.901574803	0.912350598
NOT-SPINK2 AND SPINK2	0.912863071	0.952380952	0.876494024
SLC22A16 AND NOT-ERBB3	0.901574803	0.891050584	0.912350598
FUT4 AND NOT-ERBB3	0.960474308	0.952941176	0.96812749
ELANE AND NOT-ERBB3	0.902584493	0.900793651	0.90438247
NOT-PHLDB2 AND NOT-ERBB3	0.903460838	0.832214765	0.988047809
NOT-SPINK2 AND SPINK2	0.912863071	0.952380952	0.876494024
SLC22A16 AND NOT-MET	0.902584493	0.900793651	0.90438247
FUT4 AND NOT-MET	0.952191235	0.952191235	0.952191235
ELANE AND NOT-MET	0.906504065	0.925311203	0.888446215
LRRC70 AND NOT-AXL	0.923076923	0.9140625	0.932270916
NOT-SPINK2 AND SPINK2	0.912863071	0.952380952	0.876494024
SLC22A16 AND NOT-AXL	0.90872211	0.925619835	0.892430279
FLT3 AND NOT-AXL	0.944329897	0.978632479	0.912350598
CXCL8 AND NOT-AXL	0.915254237	0.867857143	0.96812749
ELANE AND NOT-AXL	0.918367347	0.941422594	0.896414343
SMIM3 AND NOT-AXL	0.964143426	0.964143426	0.964143426
NOT-SPINK2 AND SPINK2	0.912863071	0.952380952	0.876494024
FLT3 AND IL13RA2	0.935222672	0.950617284	0.920318725
NOT-SPINK2 AND SPINK2	0.912863071	0.952380952	0.876494024
SLC22A16 AND NOT-EPHA2	0.901574803	0.891050584	0.912350598
ELANE AND NOT-EPHA2	0.908	0.911646586	0.90438247

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
NOT-GIPR AND LHFP	1	1	1
NOT-CACNG5 AND LHFP	1	1	1
NOT-GLRA2 AND LHFP	1	1	1
NOT-TMEM150B AND LHFP	1	1	1
NOT-GRIK5 AND LHFP	1	1	1
NOT-GRPR AND LHFP	1	1	1
NOT-HCRTR1 AND LHFP	1	1	1
NOT-TMEM95 AND LHFP	1	1	1
NOT-KCNB1 AND LHFP	1	1	1
NOT-TEX38 AND LHFP	1	1	1
NOT-KCNS2 AND LHFP	1	1	1
NOT-ASIC5 AND LHFP	1	1	1
AUP1 AND LHFP	1	1	1
NOT-AVPR1B AND LHFP	1	1	1
PARL AND LHFP	1	1	1
NOT-GPR137 AND LHFP	1	1	1
NOT-NMUR2 AND LHFP	1	1	1
NOT-GJD2 AND LHFP	1	1	1
NOT-CACNG6 AND LHFP	1	1	1
NOT-GAL3ST2 AND LHFP	1	1	1
NOT-SLC4A1 AND LHFP	1	1	1
NOT-SLC5A5 AND LHFP	1	1	1
NOT-ATCAY AND LHFP	1	1	1
NOT-CAV3 AND LHFP	1	1	1
NOT-PTCH2 AND LHFP	1	1	1
NOT-TM4SF5 AND LHFP	1	1	1
NOT-SLC22A8 AND LHFP	1	1	1
MLANA AND ALG3	1	1	1
MLANA AND MLANA	1	1	1
FDPS AND CALCRL	1	1	1
MLANA AND CALCRL	1	1	1
MLANA AND MLANA	1	1	1
FDPS AND TRIM13	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND NOT-ANGPTL7	1	1	1
FDPS AND KLRG1	1	1	1
MLANA AND MLANA	1	1	1
AUP1 AND NOT-KLRG1	1	1	1
PARL AND NOT-KLRG1	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND STX6	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND NOT-MSLN	1	1	1
AUP1 AND MSLN	1	1	1
MLANA AND NOT-KCNMB2	1	1	1
NPC1 AND NOT-KCNMB2	1	1	1
MLANA AND MLANA	1	1	1
NOT-SLC17A2 AND TLCD1	1	1	1
MLANA AND NOT-SLC17A2	1	1	1
NOT-SLC17A2 AND TNFRSF19	1	1	1
NOT-SLC17A2 AND TYRP1	1	1	1
MLANA AND ABCC4	1	1	1
MLANA AND IGSF6	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND NOT-RAMP1	1	1	1
NOT-CD300LB AND ZMPSTE24	1	1	1
NOT-FCN2 AND ZMPSTE24	1	1	1
FDPS AND ZMPSTE24	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND SIGMAR1	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND LILRB2	1	1	1
MLANA AND NOT-MARCH6	1	1	1
MLANA AND ADAM10	1	1	1
MLANA AND MLANA	1	1	1
FDPS AND NOT-NMUR1	1	1	1
MLANA AND NOT-NMUR1	1	1	1
CLCN5 AND NOT-B3GALT5	1	1	1
MLANA AND NOT-B3GALT5	1	1	1
NPC1 AND NOT-B3GALT5	1	1	1
MLANA AND NOT-CRISP3	1	1	1
FDPS AND NOT-FDPS	1	1	1
MLANA AND MLANA	1	1	1
AUP1 AND NOT-SIRPB1	1	1	1
PARL AND NOT-SIRPB1	1	1	1

Antigen Pair Logic	F1	Precision	Recall
NOT-SPINK2 AND SPINK2	0.912863071	0.952380952	0.876494024
FUT4 AND NOT-GPA33	0.956349206	0.95256917	0.960159363
NOT-SPINK2 AND SPINK2	0.912863071	0.952380952	0.876494024
NOT-FLT3 AND FLT3	0.935222672	0.950617284	0.920318725
NOT-SPINK2 AND SPINK2	0.912863071	0.952380952	0.876494024
SLC22A16 AND NOT-ST8SIA1	0.905811623	0.911290323	0.900398406
CXCL8 AND NOT-ST8SIA1	0.912149533	0.85915493	0.972111554
P2RX1 AND NOT-ST8SIA1	0.912878788	0.870036101	0.960159363
HBD AND NOT-ST8SIA1	0.91221374	0.875457875	0.952191235
ELANE AND NOT-ST8SIA1	0.9	0.903614458	0.896414343
NOT-SPINK2 AND ALK	0.912863071	0.952380952	0.876494024
NOT-SPINK2 AND CA9	0.912863071	0.952380952	0.876494024
NOT-HEG1 AND CD70	0.900608519	0.917355372	0.884462151
NOT-PFN2 AND CD70	0.911439114	0.848797251	0.984063745
NOT-SPINK2 AND SPINK2	0.912863071	0.952380952	0.876494024
NOT-FLT3 AND FLT3	0.935222672	0.950617284	0.920318725
NOT-MTUS1 AND CD70	0.920222635	0.861111111	0.988047809
MS4A3 AND NOT-CD160	0.906976744	0.883018868	0.932270916
NOT-PFN2 AND NOT-CD160	0.931818182	0.888086643	0.980079681
CD33 AND NOT-CD160	0.906976744	0.883018868	0.932270916
NOT-SPINK2 AND SPINK2	0.912863071	0.952380952	0.876494024
FUT4 AND NOT-CD160	0.902621723	0.851590106	0.960159363
NRROS AND NOT-CD160	0.928	0.931726908	0.924302789
TSPAN32 AND NOT-CD160	0.933333333	0.894160584	0.976095618
HBD AND NOT-CD160	0.935672515	0.916030534	0.956175299
NOT-MTUS1 AND NOT-CD160	0.928571429	0.879003559	0.984063745
TMC8 AND NOT-MS4A1	0.910891089	0.905511811	0.916334661
NOT-PFN2 AND NOT-MS4A1	0.95703125	0.938697318	0.976095618
NOT-DIP2C AND NOT-MS4A1	0.902752294	0.836734694	0.980079681
NOT-SPINK2 AND SPINK2	0.912863071	0.952380952	0.876494024
NOT-CADM1 AND NOT-MS4A1	0.905950096	0.874074074	0.940239044
NRROS AND NOT-MS4A1	0.93902439	0.958506224	0.920318725
P2RY8 AND NOT-MS4A1	0.918714556	0.874100719	0.96812749
TSPAN32 AND NOT-MS4A1	0.926553672	0.878571429	0.980079681
NOT-ATP9A AND NOT-MS4A1	0.903474903	0.876404494	0.932270916
HBD AND NOT-MS4A1	0.925143954	0.892592593	0.960159363
NOT-MTUS1 AND NOT-MS4A1	0.953488372	0.928301887	0.980079681
NOT-SPINK2 AND SPINK2	0.912863071	0.952380952	0.876494024
NOT-SPINK2 AND MSLN	0.912863071	0.952380952	0.876494024
NOT-SPINK2 AND ITGB3	0.912863071	0.952380952	0.876494024
NOT-FAM171A1 AND LRRC70	0.935817805	0.974137931	0.900398406
NOT-LRIG1 AND LRRC70	0.950298211	0.948412698	0.952191235
NOT-FAM69A AND LRRC70	0.938697318	0.904059041	0.976095618
NOT-PFN2 AND LRRC70	0.929791271	0.887681159	0.976095618
NOT-HEG1 AND LRRC70	0.926315789	0.982142857	0.876494024
NOT-PTPRK AND LRRC70	0.951076321	0.934615385	0.96812749
NOT-SLC37A3 AND LRRC70	0.939334638	0.923076923	0.956175299
LRRC70 AND NOT-AOC3	0.944	0.947791165	0.940239044
NOT-PHLDB2 AND LRRC70	0.954274354	0.952380952	0.956175299
NOT-VAPB AND LRRC70	0.937984496	0.913207547	0.964143426
SMIM3 AND NOT-CDH5	0.968	0.97188755	0.964143426
FLT3 AND NOT-MICA	0.920430108	1	0.852589641
FLT3 AND NOT-ABCC5	0.946502058	0.978723404	0.916334661
TSPAN32 AND NOT-BTN3A3	0.936902486	0.900735294	0.976095618
TSPAN32 AND NOT-BTN2A2	0.931818182	0.888086643	0.980079681
TSPAN32 AND NOT-BTN3A1	0.930581614	0.879432624	0.988047809
TSPAN32 AND NOT-CHI3L2	0.933837429	0.888489209	0.984063745
TSPAN32 AND NOT-MGAT4A	0.938931298	0.901098901	0.980079681
TSPAN32 AND NOT-CYP2U1	0.935361217	0.894545455	0.980079681
TSPAN32 AND NOT-SLAMF6	0.927102804	0.873239437	0.988047809
TSPAN32 AND NOT-FCRL3	0.942084942	0.913857678	0.972111554
TSPAN32 AND NOT-CCR6	0.926829268	0.875886525	0.984063745
TSPAN32 AND NOT-CCR7	0.936416185	0.906716418	0.96812749
TSPAN32 AND NOT-CR2	0.939163498	0.898181818	0.984063745
TSPAN32 AND NOT-GPR155	0.955165692	0.935114504	0.976095618
TSPAN32 AND NOT-BTLA	0.931818182	0.888086643	0.980079681
TSPAN32 AND NOT-IFNLR1	0.935606061	0.891696751	0.984063745
TSPAN32 AND NOT-GIMAP1	0.928030303	0.884476534	0.976095618
TSPAN32 AND NOT-DPP4	0.94049904	0.907407407	0.976095618
TSPAN32 AND NOT-TIGIT	0.928571429	0.879003559	0.984063745
TSPAN32 AND NOT-EPHA4	0.953846154	0.921933086	0.988047809
TSPAN32 AND NOT-TMEM2	0.927102804	0.873239437	0.988047809
TSPAN32 AND NOT-NCR3	0.921933086	0.864111498	0.988047809
TSPAN32 AND NOT-LRIG1	0.957364341	0.932075472	0.984063745
TSPAN32 AND NOT-SIT1	0.942748092	0.904761905	0.984063745
TSPAN32 AND NOT-SLC46A3	0.941860465	0.916981132	0.96812749

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
MLANA AND TMEM5	1	1	1
CLCN5 AND NOT-CLEC4M	1	1	1
FDPS AND NOT-CLEC4M	1	1	1
MLANA AND NOT-CLEC4M	1	1	1
FDPS AND TLR6	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND NOT-PKDREJ	1	1	1
MLANA AND NOT-TRDN	1	1	1
MLANA AND NOT-ABCA9	1	1	1
PARL AND ABCA9	1	1	1
AUP1 AND ABCA8	1	1	1
PARL AND ABCA8	1	1	1
FDPS AND NOT-CACNG3	1	1	1
MLANA AND NOT-CACNG3	1	1	1
NPC1 AND NOT-CACNG3	1	1	1
CLCN5 AND NOT-CACNG2	1	1	1
FDPS AND NOT-CACNG2	1	1	1
MLANA AND NOT-CACNG2	1	1	1
MLANA AND BTN3A3	1	1	1
MLANA AND BTN2A2	1	1	1
MLANA AND CEPT1	1	1	1
MLANA AND MLANA	1	1	1
FDPS AND NDRG1	1	1	1
MLANA AND MLANA	1	1	1
FDPS AND PEMT	1	1	1
MLANA AND PEMT	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND NOT-CDS1	1	1	1
AUP1 AND CDS1	1	1	1
MLANA AND NOT-CDSN	1	1	1
MLANA AND CDIPT	1	1	1
MLANA AND MLANA	1	1	1
NOT-PMEL AND TMEM147	1	1	1
MLANA AND LYPLA1	1	1	1
FDPS AND CD52	1	1	1
MLANA AND MLANA	1	1	1
NPC1 AND CD52	1	1	1
MLANA AND TIMM17A	1	1	1
MLANA AND NOT-LRRN2	1	1	1
NOT-SLC34A2 AND GPNMB	1	1	1
NOT-CFTR AND GPNMB	1	1	1
NOT-PTH2 AND GPNMB	1	1	1
NOT-CHRNA2 AND GPNMB	1	1	1
NOT-CD300LB AND GPNMB	1	1	1
NOT-SYT2 AND GPNMB	1	1	1
NOT-PROKR2 AND GPNMB	1	1	1
NOT-CLDN7 AND GPNMB	1	1	1
NOT-SLC34A3 AND GPNMB	1	1	1
NOT-ISM2 AND GPNMB	1	1	1
NOT-ZNRF4 AND GPNMB	1	1	1
NOT-CLDN19 AND GPNMB	1	1	1
NOT-CYP2A6 AND GPNMB	1	1	1
NOT-MFSD6L AND GPNMB	1	1	1
NOT-DNAH10 AND GPNMB	1	1	1
FDPS AND GPNMB	1	1	1
MLANA AND MLANA	1	1	1
NOT-FSHR AND GPNMB	1	1	1
NOT-CALHM1 AND GPNMB	1	1	1
NOT-GIPR AND GPNMB	1	1	1
NOT-AMHR2 AND GPNMB	1	1	1
NOT-CACNG5 AND GPNMB	1	1	1
NOT-CACNG4 AND GPNMB	1	1	1
NOT-GLRA2 AND GPNMB	1	1	1
NOT-TMEM105 AND GPNMB	1	1	1
NOT-TMEM150B AND GPNMB	1	1	1
NOT-DNAJC5G AND GPNMB	1	1	1
NOT-FFAR2 AND GPNMB	1	1	1
NOT-GRIK5 AND GPNMB	1	1	1
NOT-GRPR AND GPNMB	1	1	1
NOT-GUCA2B AND GPNMB	1	1	1
NOT-HCRTR1 AND GPNMB	1	1	1
NOT-HTR4 AND GPNMB	1	1	1
NOT-TMEM95 AND GPNMB	1	1	1
NOT-KCNB1 AND GPNMB	1	1	1
NOT-TEX38 AND GPNMB	1	1	1
NOT-KCNS2 AND GPNMB	1	1	1

Antigen Pair Logic	F1	Precision	Recall
TSPAN32 AND NOT-CLEC2D	0.923364486	0.86971831	0.984063745
TSPAN32 AND NOT-GPR171	0.931558935	0.890909091	0.976095618
TSPAN32 AND NOT-HLA-DOB	0.942965779	0.901818182	0.988047809
TSPAN32 AND NOT-VSIG1	0.930581614	0.879432624	0.988047809
TSPAN32 AND NOT-IL7R	0.937142857	0.897810219	0.980079681
TSPAN32 AND CXCL8	0.931558935	0.890909091	0.976095618
TSPAN32 AND NOT-KLRD1	0.934086629	0.885714286	0.988047809
NOT-FAM69A AND TSPAN32	0.970873786	0.946969697	0.996015936
TSPAN32 AND NOT-MGAT5	0.959064327	0.938931298	0.980079681
TSPAN32 AND NOT-MINOS1	0.95481336	0.941860465	0.96812749
TSPAN32 AND NOT-NT5E	0.934362934	0.906367041	0.964143426
TSPAN32 AND NOT-IL21R	0.942528736	0.907749077	0.980079681
TSPAN32 AND NOT-TRAT1	0.929791271	0.887681159	0.976095618
TSPAN32 AND NOT-TUBD1	0.955165692	0.935114504	0.976095618
TSPAN32 AND NOT-FAM134B	0.957746479	0.967479675	0.948207171
TSPAN32 AND NOT-LRRN3	0.942307692	0.910780669	0.976095618
TSPAN32 AND NOT-TMEM140	0.942574257	0.937007874	0.948207171
TSPAN32 AND NOT-SYNJ2BP	0.92952381	0.890510949	0.972111554
TSPAN32 AND NOT-SIRPG	0.935361217	0.894545455	0.980079681
TSPAN32 AND NOT-LPAR5	0.930320151	0.882142857	0.984063745
TSPAN32 AND NOT-SLC44A2	0.950884086	0.937984496	0.964143426
TSPAN32 AND NOT-PTGDR	0.939163498	0.898181818	0.984063745
TSPAN32 AND NOT-SLAMF7	0.947162427	0.930769231	0.964143426
TSPAN32 AND NOT-TNFRSF17	0.936170213	0.909774436	0.964143426
TSPAN32 AND NOT-SCN3A	0.929791271	0.887681159	0.976095618
TSPAN32 AND NOT-BMPR2	0.944664032	0.937254902	0.952191235
TSPAN32 AND NOT-SP4	0.938697318	0.904059041	0.976095618
TSPAN32 AND SPINK2	0.930526316	0.986607143	0.880478088
TSPAN32 AND NOT-TTC13	0.931558935	0.890909091	0.976095618
TSPAN32 AND NOT-STEAP4	0.942574257	0.937007874	0.948207171
TSPAN32 AND NOT-FCRL5	0.930056711	0.884892086	0.980079681
TSPAN32 AND NOT-SLC37A3	0.956175299	0.956175299	0.956175299
TSPAN32 AND NOT-FCRLA	0.94026975	0.910447761	0.972111554
TSPAN32 AND NOT-ADTRP	0.932330827	0.882562278	0.988047809
TSPAN32 AND NOT-LRCH3	0.935606061	0.891696751	0.984063745
TSPAN32 AND SMIM3	0.954455446	0.948818898	0.960159363
TSPAN32 AND NOT-CDC14B	0.926829268	0.875886525	0.984063745
TSPAN32 AND NOT-KMO	0.930320151	0.882142857	0.984063745
TSPAN32 AND NOT-ADAM19	0.936902486	0.900735294	0.976095618
TSPAN32 AND NOT-ST3GAL5	0.9348659	0.900369004	0.972111554
NOT-PHLDB2 AND TSPAN32	0.957364341	0.932075472	0.984063745
TSPAN32 AND NOT-FCMR	0.946564885	0.908424908	0.988047809
TSPAN32 AND NOT-VAPB	0.939393939	0.895306859	0.988047809
TSPAN32 AND NOT-CD8B	0.926553672	0.878571429	0.980079681
TSPAN32 AND NOT-MS4A1	0.926553672	0.878571429	0.980079681
TSPAN32 AND NOT-CD22	0.931818182	0.888086643	0.980079681
TSPAN32 AND NOT-KIAA0040	0.921933086	0.864111498	0.988047809
TSPAN32 AND NOT-CD72	0.936660269	0.903703704	0.972111554
TSPAN32 AND NOT-ARMCX2	0.938931298	0.901098901	0.980079681
NOT-ATP9A AND FUT4	0.951417004	0.967078189	0.93625498
FUT4 AND NOT-TSPAN3	0.960474308	0.952941176	0.96812749
CXCL8 AND NOT-CDH11	0.925373134	0.870175439	0.988047809
SMIM3 AND NOT-CDH11	0.942084942	0.913857678	0.972111554
FUT4 AND NOT-TSPAN1	0.960474308	0.952941176	0.96812749
FLT3 AND NOT-PREB	0.916129032	0.995327103	0.848605578
FUT4 AND NOT-CDH17	0.960474308	0.952941176	0.96812749
FLT3 AND NOT-LHFP	0.948665298	0.978813559	0.920318725
SMIM3 AND NOT-LHFP	0.949019608	0.934362934	0.964143426
FUT4 AND NOT-MPZL2	0.961770624	0.971544715	0.952191235
FUT4 AND NOT-TRIM13	0.95	0.995633188	0.908366534
FUT4 AND NOT-GPA33	0.956349206	0.95256917	0.960159363
NOT-LRIG1 AND MFSD10	0.934615385	0.903345725	0.96812749
NOT-VAPB AND MFSD10	0.92481203	0.87544484	0.980079681
NOT-ARMCX2 AND MFSD10	0.936660269	0.903703704	0.972111554
FLT3 AND NOT-COQ7	0.948665298	0.978813559	0.920318725
FUT4 AND NOT-COQ7	0.954091816	0.956	0.952191235
FUT4 AND NOT-PRSS16	0.960474308	0.952941176	0.96812749
FLT3 AND NOT-BET1	0.943157895	1	0.892430279
FUT4 AND NOT-BET1	0.95257732	0.987179487	0.920318725
FLT3 AND NOT-TMEM5	0.943632568	0.99122807	0.900398406
FUT4 AND NOT-B3GNT3	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-ABCA8	0.961923848	0.967741935	0.956175299
NOT-FAM69A AND NOT-ABCA8	0.955684008	0.925373134	0.988047809
HCST AND NOT-BTN3A3	0.941860465	0.916981132	0.96812749
HBD AND NOT-BTN3A3	0.945525292	0.923954373	0.96812749
NRROS AND NOT-BTN3A3	0.944785276	0.970588235	0.920318725

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
NOT-MUC3A AND GPNMB	1	1	1
NPC1 AND GPNMB	1	1	1
NOT-ASIC5 AND GPNMB	1	1	1
AUP1 AND GPNMB	1	1	1
NOT-AVPR1B AND GPNMB	1	1	1
PARL AND GPNMB	1	1	1
NOT-PCDHA10 AND GPNMB	1	1	1
NOT-GPR137 AND GPNMB	1	1	1
NOT-NMUR2 AND GPNMB	1	1	1
NOT-GJD2 AND GPNMB	1	1	1
NOT-LRFN2 AND GPNMB	1	1	1
NOT-CACNG6 AND GPNMB	1	1	1
NOT-GAL3ST2 AND GPNMB	1	1	1
NOT-SLC4A1 AND GPNMB	1	1	1
NOT-SLC5A5 AND GPNMB	1	1	1
NOT-RTP3 AND GPNMB	1	1	1
NOT-ATCAY AND GPNMB	1	1	1
NOT-CAV3 AND GPNMB	1	1	1
NOT-PTCH2 AND GPNMB	1	1	1
GPNMB AND NOT-MARCO	1	1	1
NOT-TM4SF5 AND GPNMB	1	1	1
NOT-SLC22A8 AND GPNMB	1	1	1
GPNMB AND NOT-OTOF	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND NOT-CLEC10A	1	1	1
FDPS AND SLC30A9	1	1	1
MLANA AND MLANA	1	1	1
NOT-FCN2 AND SLC9A6	1	1	1
FDPS AND SLC9A6	1	1	1
MLANA AND SLC9A6	1	1	1
MLANA AND NOT-CLGN	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND CRTAP	1	1	1
MLANA AND NOT-SEMA6B	1	1	1
FDPS AND SEMA4B	1	1	1
MLANA AND SEMA4B	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND HYOU1	1	1	1
NOT-CHRNA2 AND PROCR	1	1	1
NOT-CLDN19 AND PROCR	1	1	1
FDPS AND PROCR	1	1	1
MLANA AND MLANA	1	1	1
NOT-TMEM150B AND PROCR	1	1	1
NOT-TMEM95 AND PROCR	1	1	1
AUP1 AND PROCR	1	1	1
NOT-AVPR1B AND PROCR	1	1	1
PARL AND PROCR	1	1	1
NOT-NMUR2 AND PROCR	1	1	1
NOT-TM4SF5 AND PROCR	1	1	1
MLANA AND PRDX4	1	1	1
MLANA AND ARL6IP5	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND AGPAT1	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND SLC19A2	1	1	1
NOT-CAV3 AND SLC19A2	1	1	1
FDPS AND RABAC1	1	1	1
MLANA AND MLANA	1	1	1
NOT-SLC34A2 AND ADCY6	1	1	1
NOT-SLC34A2 AND TLCD1	1	1	1
CLCN5 AND NOT-SLC34A2	1	1	1
NOT-SLC34A2 AND SLC44A3	1	1	1
NOT-SLC34A2 AND LRRC15	1	1	1
NOT-SLC34A2 AND APCDD1	1	1	1
NOT-SLC34A2 AND VSIG10L	1	1	1
NOT-SLC34A2 AND PROM2	1	1	1
NOT-SLC34A2 AND ADRA2A	1	1	1
NOT-SLC34A2 AND CCDC80	1	1	1
NOT-SLC34A2 AND DSC3	1	1	1
NOT-SLC34A2 AND EFNB2	1	1	1
NOT-SLC34A2 AND ELN	1	1	1
NOT-SLC34A2 AND ENG	1	1	1
FDPS AND NOT-SLC34A2	1	1	1
MLANA AND NOT-SLC34A2	1	1	1
NOT-SLC34A2 AND ABI3BP	1	1	1
NOT-SLC34A2 AND PVRL3	1	1	1

Antigen Pair Logic	F1	Precision	Recall
NOT-PFN2 AND NOT-BTN3A3	0.92393321	0.864583333	0.992031873
HCST AND NOT-BTN2A2	0.928301887	0.88172043	0.980079681
FLT3 AND NOT-BTN2A2	0.956340956	1	0.916334661
P2RY8 AND NOT-BTN2A2	0.944123314	0.914179104	0.976095618
HBD AND NOT-BTN2A2	0.937743191	0.91634981	0.960159363
P2RX1 AND NOT-BTN2A2	0.941860465	0.916981132	0.96812749
NOT-PFN2 AND NOT-BTN2A2	0.95	0.918215613	0.984063745
FXYD5 AND NOT-BTN2A2	0.922222222	0.861591696	0.992031873
NOT-MTUS1 AND NOT-BTN2A2	0.96124031	0.935849057	0.988047809
FLT3 AND NOT-CEPT1	0.950413223	0.987124464	0.916334661
FUT4 AND NOT-CDS1	0.960474308	0.952941176	0.96812749
FLT3 AND NOT-CDSN	0.940936864	0.9625	0.920318725
FLT3 AND NOT-PGRMC2	0.938689218	1	0.884462151
FUT4 AND NOT-PGRMC2	0.933884298	0.969957082	0.900398406
FLT3 AND NOT-TIMM17A	0.944785276	0.970588235	0.920318725
FLT3 AND NOT-N4BP2L2	0.939958592	0.978448276	0.90438247
FLT3 AND NOT-GPNMB	0.928870293	0.977973568	0.884462151
FUT4 AND NOT-GPNMB	0.934156379	0.965957447	0.90438247
FLT3 AND NOT-SLC25A17	0.944785276	0.970588235	0.920318725
FUT4 AND NOT-SEMA4B	0.963709677	0.975510204	0.952191235
P2RX1 AND NOT-SEMA4B	0.954274354	0.952380952	0.956175299
SMIM3 AND NOT-PROCR	0.942386831	0.974468085	0.912350598
FLT3 AND NOT-HTATIP2	0.920770878	0.99537037	0.856573705
FLT3 AND NOT-POMT1	0.940936864	0.9625	0.920318725
NOT-ANKRD46 AND ERLIN1	0.940206186	0.974358974	0.908366534
NOT-TMTC3 AND ERLIN1	0.946280992	0.982832618	0.912350598
NOT-TMCC1 AND ERLIN1	0.931106472	0.978070175	0.888446215
NOT-HEG1 AND ERLIN1	0.933054393	0.982378855	0.888446215
NOT-PHLDB2 AND ERLIN1	0.958742633	0.945736434	0.972111554
NOT-HEG1 AND TNFSF13B	0.919491525	0.981900452	0.864541833
NOT-PHLDB2 AND TNFSF13B	0.946954813	0.934108527	0.960159363
FLT3 AND NOT-CSPG5	0.940936864	0.9625	0.920318725
FLT3 AND NOT-VAMP5	0.922746781	1	0.856573705
FUT4 AND NOT-CFTR	0.960474308	0.952941176	0.96812749
HCST AND NOT-CHI3L2	0.937381404	0.894927536	0.984063745
HCST AND NOT-MGAT4A	0.95256917	0.945098039	0.960159363
HCST AND NOT-CYP2U1	0.930056711	0.884892086	0.980079681
HCST AND NOT-CMTM8	0.938271605	0.970212766	0.908366534
HCST AND NOT-IFNLR1	0.937381404	0.894927536	0.984063745
HCST AND NOT-DPP4	0.953125	0.9348659	0.972111554
HCST AND NOT-LPAR1	0.933837429	0.888489209	0.984063745
HCST AND NOT-SMIM14	0.944	0.947791165	0.940239044
HCST AND NOT-EPHA4	0.923364486	0.86971831	0.984063745
NOT-EXT1 AND HCST	0.940038685	0.913533835	0.96812749
HCST AND NOT-TMEM2	0.953307393	0.931558935	0.976095618
HCST AND NOT-LRIG1	0.960784314	0.945945946	0.976095618
HCST AND NOT-SLC46A3	0.934615385	0.903345725	0.96812749
HCST AND NOT-HMOX2	0.92952381	0.890510949	0.972111554
HCST AND NOT-AMIGO2	0.943907157	0.917293233	0.972111554
HCST AND NOT-IL7R	0.935114504	0.897435897	0.976095618
HCST AND NOT-ITPR3	0.934086629	0.885714286	0.988047809
NOT-FAM69A AND HCST	0.9765625	0.957854406	0.996015936
HCST AND NOT-MINOS1	0.949019608	0.934362934	0.964143426
HCST AND NOT-NT5E	0.936660269	0.903703704	0.972111554
HCST AND NOT-TUBD1	0.92481203	0.87544484	0.980079681
NOT-PFN2 AND HCST	0.925650558	0.867595819	0.992031873
NOT-FAM134B AND HCST	0.944664032	0.937254902	0.952191235
HCST AND NOT-TMEM140	0.956	0.959839357	0.952191235
HCST AND NOT-MAN1C1	0.952772074	0.983050847	0.924302789
HCST AND NOT-SLC44A2	0.96	0.963855422	0.956175299
HCST AND NOT-PTCH1	0.936902486	0.900735294	0.976095618
NOT-HEG1 AND HCST	0.931106472	0.978070175	0.888446215
HCST AND NOT-ENPP5	0.935361217	0.894545455	0.980079681
HCST AND NOT-RARRES3	0.947162427	0.930769231	0.964143426
HCST AND NOT-RTN1	0.939805825	0.916666667	0.964143426
HCST AND NOT-BMPR2	0.948412698	0.944664032	0.952191235
NOT-TGFBR3 AND HCST	0.924686192	0.973568282	0.880478088
NOT-WLS AND HCST	0.936170213	0.909774436	0.964143426
NOT-SLC37A3 AND HCST	0.95481336	0.941860465	0.96812749
HCST AND NOT-ADTRP	0.940952381	0.901459854	0.984063745
NOT-PHLDB2 AND HCST	0.96124031	0.935849057	0.988047809
NOT-VAPB AND HCST	0.92936803	0.871080139	0.996015936
HCST AND NOT-KIAA0040	0.927102804	0.873239437	0.988047809
HCST AND NOT-ARMCX2	0.938931298	0.901098901	0.980079681
FUT4 AND NOT-CEACAM7	0.958579882	0.94921875	0.96812749
FLT3 AND NOT-BLCAP	0.922746781	1	0.856573705

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
NOT-SLC34A2 AND GJA1	1	1	1
NOT-SLC34A2 AND DLL1	1	1	1
NOT-SLC34A2 AND IFI27	1	1	1
NOT-SLC34A2 AND ITGA2	1	1	1
NOT-SLC34A2 AND JAG2	1	1	1
NOT-SLC34A2 AND KDR	1	1	1
NOT-SLC34A2 AND SAMD5	1	1	1
NOT-SLC34A2 AND ATP1A1	1	1	1
NPC1 AND NOT-SLC34A2	1	1	1
NOT-SLC34A2 AND CDON	1	1	1
NOT-SLC34A2 AND PDGFRA	1	1	1
NOT-SLC34A2 AND ENPP2	1	1	1
NOT-SLC34A2 AND PMP22	1	1	1
NOT-SLC34A2 AND IL20RB	1	1	1
NOT-SLC34A2 AND SLC47A1	1	1	1
NOT-SLC34A2 AND LGR4	1	1	1
NOT-SLC34A2 AND SLC35E3	1	1	1
NOT-SLC34A2 AND PCDHB14	1	1	1
NOT-SLC34A2 AND GKN1	1	1	1
NOT-SLC34A2 AND NTN4	1	1	1
NOT-SLC34A2 AND SORT1	1	1	1
NOT-SLC34A2 AND PERP	1	1	1
NOT-SLC34A2 AND SGCB	1	1	1
NOT-SLC34A2 AND SLC7A2	1	1	1
NOT-SLC34A2 AND TPBG	1	1	1
NOT-SLC34A2 AND TYRP1	1	1	1
NOT-SLC34A2 AND COL14A1	1	1	1
NOT-SLC34A2 AND VCAM1	1	1	1
NOT-SLC34A2 AND DHRS11	1	1	1
NOT-SLC34A2 AND TMEM231	1	1	1
NOT-SLC34A2 AND WLS	1	1	1
NOT-SLC34A2 AND PIGZ	1	1	1
TMEM187 AND NOT-SLC34A2	1	1	1
NOT-SLC34A2 AND ZNRF3	1	1	1
NOT-SLC34A2 AND LY6D	1	1	1
NOT-SLC34A2 AND OSMR	1	1	1
NOT-SLC34A2 AND DSEL	1	1	1
NOT-SLC34A2 AND REEP6	1	1	1
NOT-SLC34A2 AND CLCA2	1	1	1
NOT-SLC34A2 AND LPPR4	1	1	1
MLANA AND IFITM2	1	1	1
FDPS AND NOT-POMT1	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND NOT-SLCO1B1	1	1	1
NOT-CHRNA2 AND ST6GALNAC2	1	1	1
MLANA AND ST6GALNAC2	1	1	1
NPC1 AND ST6GALNAC2	1	1	1
AUP1 AND ST6GALNAC2	1	1	1
PARL AND ST6GALNAC2	1	1	1
MLANA AND ERLIN1	1	1	1
FDPS AND TGOLN2	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND MLANA	1	1	1
FDPS AND DMRT2	1	1	1
MLANA AND MLANA	1	1	1
FDPS AND CXCR6	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND MLANA	1	1	1
NOT-MLANA AND CGRRF1	1	1	1
MLANA AND MLANA	1	1	1
NOT-CHRNA2 AND CSPG5	1	1	1
MLANA AND MLANA	1	1	1
NPC1 AND CSPG5	1	1	1
AUP1 AND CSPG5	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND EBP	1	1	1
MLANA AND MLANA	1	1	1
CLCN5 AND NOT-CLDN16	1	1	1
FDPS AND NOT-CLDN16	1	1	1
MLANA AND NOT-CLDN16	1	1	1
FDPS AND NOT-FUT9	1	1	1
MLANA AND NOT-FUT9	1	1	1
FDPS AND NOT-RRH	1	1	1
MLANA AND NOT-RRH	1	1	1
MLANA AND NOT-CORIN	1	1	1

Antigen Pair Logic	F1	Precision	Recall
FUT4 AND NOT-BLCAP	0.922105263	0.977678571	0.87250996
FUT4 AND NOT-BTNL3	0.960474308	0.952941176	0.96812749
FLT3 AND NOT-AFG3L2	0.926315789	0.982142857	0.876494024
FUT4 AND NOT-AFG3L2	0.940206186	0.974358974	0.908366534
FLT3 AND NOT-LMAN2	0.906318083	1	0.828685259
FLT3 AND NOT-UQCR11	0.95257732	0.987179487	0.920318725
FUT4 AND NOT-UQCR11	0.950819672	0.978902954	0.924302789
FUT4 AND NOT-ILVBL	0.948616601	0.941176471	0.956175299
FUT4 AND NOT-NAT2	0.960474308	0.952941176	0.96812749
NOT-FAM69A AND GLIPR1	0.935114504	0.897435897	0.976095618
FLT3 AND NOT-KDELR2	0.948240166	0.987068966	0.912350598
FUT4 AND NOT-KDELR2	0.960985626	0.991525424	0.932270916
FUT4 AND NOT-KDELR3	0.964143426	0.964143426	0.964143426
SMIM3 AND NOT-KDELR3	0.938223938	0.91011236	0.96812749
FLT3 AND NOT-TDRKH	0.950617284	0.982978723	0.920318725
FLT3 AND NOT-B4GAT1	0.935550936	0.97826087	0.896414343
FLT3 AND NOT-SLC35D2	0.937759336	0.978354978	0.900398406
FUT4 AND NOT-CACFD1	0.956349206	0.95256917	0.960159363
NRROS AND NOT-BTN3A2	0.934156379	0.965957447	0.90438247
P2RY8 AND NOT-BTN3A1	0.935361217	0.894545455	0.980079681
HBD AND NOT-BTN3A1	0.932821497	0.9	0.96812749
NRROS AND NOT-BTN3A1	0.947368421	0.962962963	0.932270916
NOT-PFN2 AND NOT-BTN3A1	0.939622642	0.892473118	0.992031873
P2RX1 AND NOT-BTN2A1	0.932821497	0.9	0.96812749
FUT4 AND NOT-HHLA2	0.960474308	0.952941176	0.96812749
NOT-DIP2C AND NOT-CHI3L2	0.918819188	0.855670103	0.992031873
P2RY8 AND NOT-CHI3L2	0.935361217	0.894545455	0.980079681
HBD AND NOT-CHI3L2	0.947368421	0.927480916	0.96812749
P2RX1 AND NOT-CHI3L2	0.938461538	0.907063197	0.972111554
NOT-PFN2 AND NOT-CHI3L2	0.953846154	0.921933086	0.988047809
FXYD5 AND NOT-CHI3L2	0.931098696	0.874125874	0.996015936
NOT-NDRG2 AND NOT-CHI3L2	0.929791271	0.887681159	0.976095618
NOT-MTUS1 AND NOT-CHI3L2	0.957692308	0.925650558	0.992031873
NOT-PTPRK AND NOT-CHI3L2	0.930841121	0.876760563	0.992031873
NOT-FAM69A AND RNF139	0.943609023	0.893238434	1
NOT-PFN2 AND RNF24	0.939622642	0.892473118	0.992031873
NOT-DDR1 AND RNF24	0.930056711	0.884892086	0.980079681
FLT3 AND NOT-GPR176	0.940936864	0.9625	0.920318725
FUT4 AND NOT-MRAP2	0.960474308	0.952941176	0.96812749
NOT-FAM69A AND CMTM7	0.935361217	0.894545455	0.980079681
NOT-FAM69A AND NRM	0.962962963	0.942748092	0.984063745
NOT-PTPRK AND NRM	0.928301887	0.88172043	0.980079681
NOT-PHLDB2 AND NRM	0.9348659	0.900369004	0.972111554
FLT3 AND NOT-CHRM3	0.940936864	0.9625	0.920318725
FUT4 AND NOT-MGAT4A	0.973947896	0.97983871	0.96812749
NOT-PFN2 AND NOT-MGAT4A	0.930320151	0.882142857	0.984063745
FLT3 AND NOT-SCRG1	0.940936864	0.9625	0.920318725
FUT4 AND NOT-TMEM54	0.960474308	0.952941176	0.96812749
FLT3 AND NOT-CYP2U1	0.952380952	0.99137931	0.916334661
FUT4 AND NOT-CYP2U1	0.958250497	0.956349206	0.960159363
CXCL8 AND NOT-CYP2U1	0.933837429	0.888489209	0.984063745
P2RX1 AND NOT-CYP2U1	0.945525292	0.923954373	0.96812749
FUT4 AND NOT-SLC2A13	0.957746479	0.967479675	0.948207171
FUT4 AND NOT-MAL2	0.960474308	0.952941176	0.96812749
FLT3 AND NOT-XKR4	0.940936864	0.9625	0.920318725
FLT3 AND NOT-TMEM200A	0.946058091	0.987012987	0.908366534
FUT4 AND NOT-TMEM200A	0.96	0.963855422	0.956175299
FUT4 AND NOT-MARCH3	0.944785276	0.970588235	0.920318725
NOT-PFN2 AND NOT-FCRL1	0.956349206	0.95256917	0.960159363
NOT-PFN2 AND NOT-FCRL3	0.960629921	0.949416342	0.972111554
NOT-MTUS1 AND NOT-FCRL3	0.968503937	0.957198444	0.980079681
FLT3 AND NOT-LYSMD3	0.950617284	0.982978723	0.920318725
FUT4 AND NOT-LYSMD3	0.943469786	0.923664122	0.964143426
FLT3 AND NOT-CYYR1	0.931106472	0.978070175	0.888446215
SMIM3 AND NOT-TM4SF18	0.962376238	0.956692913	0.96812749
SMIM3 AND NOT-MRGPRF	0.954990215	0.938461538	0.972111554
FUT4 AND NOT-SLC18B1	0.913793103	0.995305164	0.844621514
P2RX1 AND NOT-SLC18B1	0.914893617	0.98173516	0.856573705
FLT3 AND NOT-SLC16A10	0.946502058	0.978723404	0.916334661
FUT4 AND NOT-CLCN3	0.973843058	0.983739837	0.964143426
FUT4 AND NOT-COMTD1	0.956521739	0.949019608	0.964143426
FUT4 AND NOT-CLRN3	0.960474308	0.952941176	0.96812749
FLT3 AND NOT-CPXM2	0.948665298	0.978813559	0.920318725
FUT4 AND NOT-TMEM45B	0.960474308	0.952941176	0.96812749
FLT3 AND NOT-CLN5	0.950207469	0.991341991	0.912350598
FUT4 AND NOT-NXPE1	0.958579882	0.94921875	0.96812749

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
MLANA AND NRG3	1	1	1
MLANA AND MLANA	1	1	1
NOT-MLANA AND NUDC	1	1	1
NOT-PMEL AND NUDC	1	1	1
FDPS AND NOT-YME1L1	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND PHTF1	1	1	1
MLANA AND CHL1	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND MLANA	1	1	1
CLCN5 AND NOT-SLC17A3	1	1	1
FDPS AND NOT-SLC17A3	1	1	1
MLANA AND NOT-SLC17A3	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND NOT-OR5I1	1	1	1
MLANA AND ADCY1	1	1	1
MLANA AND NOT-CYSLTR1	1	1	1
MLANA AND MLANA	1	1	1
NOT-CFTR AND TLCD1	1	1	1
CLCN5 AND NOT-CFTR	1	1	1
FDPS AND NOT-CFTR	1	1	1
MLANA AND NOT-CFTR	1	1	1
GALNS AND NOT-CFTR	1	1	1
ATP1A1 AND NOT-CFTR	1	1	1
NPC1 AND NOT-CFTR	1	1	1
NOT-CFTR AND CDON	1	1	1
NOT-CFTR AND TNFRSF19	1	1	1
SLC35E3 AND NOT-CFTR	1	1	1
NOT-CFTR AND WLS	1	1	1
TMEM187 AND NOT-CFTR	1	1	1
FDPS AND NOT-CGA	1	1	1
MLANA AND NOT-CGA	1	1	1
MLANA AND FAXDC2	1	1	1
FDPS AND NOT-CEACAM3	1	1	1
MLANA AND NOT-CEACAM3	1	1	1
TMEM187 AND NOT-CEACAM3	1	1	1
MLANA AND NOT-PGRMC1	1	1	1
FDPS AND NOT-SLC26A1	1	1	1
MLANA AND NOT-SLC26A1	1	1	1
MLANA AND MLANA	1	1	1
CLCN5 AND NOT-SLC22A7	1	1	1
FDPS AND NOT-SLC22A7	1	1	1
MLANA AND NOT-SLC22A7	1	1	1
MLANA AND NOT-TSPAN9	1	1	1
MLANA AND USP19	1	1	1
MLANA AND MLANA	1	1	1
NOT-CD300C AND TLCD1	1	1	1
CLCN5 AND NOT-CD300C	1	1	1
FDPS AND NOT-CD300C	1	1	1
MLANA AND NOT-CD300C	1	1	1
GUSB AND NOT-CD300C	1	1	1
NPC1 AND NOT-CD300C	1	1	1
AUP1 AND NOT-CD300C	1	1	1
GPR137B AND NOT-CD300C	1	1	1
TMEM187 AND NOT-CD300C	1	1	1
MLANA AND NOT-NMU	1	1	1
FDPS AND NOT-CEACAM7	1	1	1
MLANA AND NOT-CEACAM7	1	1	1
MLANA AND NOT-NPFFR2	1	1	1
MLANA AND MLANA	1	1	1
FDPS AND MMP24	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND NOT-LYVE1	1	1	1
NPC1 AND NOT-LYVE1	1	1	1
MLANA AND NOT-OCLM	1	1	1
MLANA AND YIF1A	1	1	1
MLANA AND MLANA	1	1	1
NOT-CHRNA2 AND ADCY2	1	1	1
MLANA AND NOT-ADCY2	1	1	1
AUP1 AND ADCY2	1	1	1
PARL AND ADCY2	1	1	1
MLANA AND MLANA	1	1	1
FDPS AND MAN1A2	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND PNPLA6	1	1	1

Antigen Pair Logic	F1	Precision	Recall
FUT4 AND NOT-LRIG3	0.960474308	0.952941176	0.96812749
FLT3 AND NOT-SLC15A4	0.947589099	1	0.900398406
FUT4 AND NOT-SLC51B	0.960474308	0.952941176	0.96812749
P2RY8 AND NOT-CCR6	0.937142857	0.897810219	0.980079681
NOT-PFN2 AND NOT-CCR6	0.955684008	0.925373134	0.988047809
NOT-MTUS1 AND NOT-CCR6	0.957692308	0.925650558	0.992031873
NOT-PFN2 AND NOT-CCR7	0.948176583	0.914814815	0.984063745
NOT-FAM69A AND CMTM3	0.930581614	0.879432624	0.988047809
NOT-HEG1 AND CMTM3	0.922746781	1	0.856573705
NOT-TGFBR3 AND CMTM3	0.920770878	0.99537037	0.856573705
NOT-PHLDB2 AND CMTM3	0.946745562	0.9375	0.956175299
FUT4 AND NOT-CANT1	0.952191235	0.952191235	0.952191235
FUT4 AND NOT-ZNF816	0.93852459	0.966244726	0.912350598
FUT4 AND NOT-SLC44A3	0.960474308	0.952941176	0.96812749
FLT3 AND RNF19B	0.93877551	0.962343096	0.916334661
FUT4 AND NOT-TMEM125	0.960474308	0.952941176	0.96812749
FLT3 AND NOT-DRAM2	0.945378151	1	0.896414343
NOT-FAM69A AND C1orf162	0.923364486	0.86971831	0.984063745
FUT4 AND NOT-COL4A2	0.96812749	0.96812749	0.96812749
NOT-FAM69A AND NOT-COL4A2	0.927643785	0.868055556	0.996015936
FLT3 AND NOT-TBC1D20	0.940695297	0.966386555	0.916334661
FLT3 AND NOT-PIGU	0.93877551	0.962343096	0.916334661
NOT-PFN2 AND EMID1	0.941860465	0.916981132	0.96812749
NOT-MTUS1 AND EMID1	0.953125	0.9348659	0.972111554
NOT-PTPRK AND EMID1	0.94026975	0.910447761	0.972111554
FUT4 AND NOT-SGPP2	0.9500998	0.952	0.948207171
FUT4 AND NOT-ACVR1C	0.97005988	0.972	0.96812749
SMIM3 AND NOT-ACVR1C	0.936416185	0.906716418	0.96812749
FUT4 AND NOT-FAM3D	0.960474308	0.952941176	0.96812749
FLT3 AND NOT-TMEM42	0.930526316	0.986607143	0.880478088
FUT4 AND NOT-SLC31A1	0.939203354	0.991150442	0.892430279
FLT3 AND NOT-SYNPR	0.940936864	0.9625	0.920318725
FLT3 AND NOT-C4orf32	0.931106472	0.978070175	0.888446215
FLT3 AND NOT-AASDH	0.9375	0.982532751	0.896414343
FLT3 AND NOT-EGFLAM	0.948665298	0.978813559	0.920318725
FUT4 AND NOT-TMEM171	0.960474308	0.952941176	0.96812749
NOT-FAM69A AND NOT-COX7A1	0.958015267	0.919413919	1
FLT3 AND NOT-COX10	0.93877551	0.962343096	0.916334661
FUT4 AND NOT-COX10	0.943548387	0.955102041	0.932270916
FLT3 AND NOT-COX11	0.909090909	0.995260664	0.836653386
FUT4 AND NOT-TMEM139	0.958579882	0.94921875	0.96812749
FUT4 AND NOT-CLDN4	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-CLDN3	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-CLDN7	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-CLDN23	0.958415842	0.952755906	0.964143426
FLT3 AND NOT-TMEM68	0.952380952	0.99137931	0.916334661
FUT4 AND NOT-TMEM68	0.963562753	0.979423868	0.948207171
FLT3 AND NOT-AGPAT6	0.940936864	0.9625	0.920318725
FUT4 AND NOT-CR2	0.936416185	0.906716418	0.96812749
P2RY8 AND NOT-CR2	0.92481203	0.87544484	0.980079681
NRROS AND NOT-CR2	0.943319838	0.958847737	0.928286853
NOT-PFN2 AND NOT-CR2	0.96124031	0.935849057	0.988047809
FLT3 AND NOT-SELM	0.948453608	0.982905983	0.916334661
NOT-SELM AND CXCL8	0.928030303	0.884476534	0.976095618
NOT-SELM AND SMIM3	0.962524655	0.953125	0.972111554
FUT4 AND NOT-TMEM37	0.964285714	0.960474308	0.96812749
SMIM3 AND NOT-TMEM37	0.945736434	0.920754717	0.972111554
FUT4 AND NOT-TRPM6	0.958415842	0.952755906	0.964143426
NOT-FAM69A AND FAM76B	0.945179584	0.899280576	0.996015936
NOT-VAPB AND FAM76B	0.925093633	0.872791519	0.984063745
FLT3 AND NOT-LAYN	0.94214876	0.978540773	0.908366534
NOT-FAM69A AND CSF3R	0.964	0.967871486	0.960159363
NOT-PTPRK AND CSF3R	0.936090226	0.886120996	0.992031873
FLT3 AND NOT-SLC38A6	0.952182952	0.995652174	0.912350598
FLT3 AND NOT-CMTM4	0.940936864	0.9625	0.920318725
FUT4 AND NOT-CMTM4	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-TMED6	0.958415842	0.952755906	0.964143426
FLT3 AND NOT-CD300LG	0.942857143	0.966527197	0.920318725
NOT-FAM69A AND TMC8	0.961538462	0.92936803	0.996015936
NOT-PHLDB2 AND TMC8	0.930320151	0.882142857	0.984063745
FLT3 AND NOT-TMEM99	0.952380952	0.99137931	0.916334661
FUT4 AND NOT-TMEM99	0.949019608	0.934362934	0.964143426
FUT4 AND NOT-APCDD1	0.93877551	0.962343096	0.916334661
FLT3 AND NOT-SMIM17	0.93877551	0.962343096	0.916334661
FUT4 AND NOT-TMC4	0.952191235	0.952191235	0.952191235
FUT4 AND NOT-MFSD4	0.956692913	0.945525292	0.96812749

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
FDPS AND NOT-EDAR	1	1	1
MLANA AND NOT-EDAR	1	1	1
AUP1 AND NOT-EDAR	1	1	1
PARL AND NOT-EDAR	1	1	1
GPR137B AND NOT-EDAR	1	1	1
TMEM187 AND NOT-EDAR	1	1	1
CLCN5 AND NOT-BTNL3	1	1	1
FDPS AND NOT-BTNL3	1	1	1
MLANA AND NOT-BTNL3	1	1	1
NPC1 AND NOT-BTNL3	1	1	1
FDPS AND NOT-GPR75	1	1	1
MLANA AND NOT-GPR75	1	1	1
MLANA AND AFG3L2	1	1	1
NOT-MLANA AND KDELR1	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND OS9	1	1	1
NOT-OS9 AND LRFN1	1	1	1
NOT-OS9 AND NOT-NPFFR1	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND LMAN2	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND MLANA	1	1	1
CLCN5 AND NOT-SLC38A3	1	1	1
FDPS AND NOT-SLC38A3	1	1	1
MLANA AND NOT-SLC38A3	1	1	1
NPC1 AND NOT-SLC38A3	1	1	1
SLC35E3 AND NOT-SLC38A3	1	1	1
TYRP1 AND NOT-SLC38A3	1	1	1
FDPS AND ILVBL	1	1	1
MLANA AND ILVBL	1	1	1
MLANA AND SLC27A5	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND NOT-NAT2	1	1	1
NPC1 AND NOT-NAT2	1	1	1
FDPS AND SLC27A3	1	1	1
MLANA AND MLANA	1	1	1
FDPS AND NOT-SLC27A2	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND LILRB4	1	1	1
MLANA AND MLANA	1	1	1
AUP1 AND NOT-GLIPR1	1	1	1
PARL AND NOT-GLIPR1	1	1	1
MLANA AND KDELR2	1	1	1
MLANA AND KDELR3	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND TMED1	1	1	1
MLANA AND TDRKH	1	1	1
MLANA AND LILRB3	1	1	1
MLANA AND B4GAT1	1	1	1
AUP1 AND B4GAT1	1	1	1
MLANA AND NOT-UPK1A	1	1	1
MLANA AND SLC35D2	1	1	1
CLCN5 AND NOT-ZPBP	1	1	1
FDPS AND NOT-ZPBP	1	1	1
MLANA AND NOT-ZPBP	1	1	1
NPC1 AND NOT-ZPBP	1	1	1
TYRP1 AND NOT-ZPBP	1	1	1
MLANA AND NOT-LECT1	1	1	1
NPC1 AND NOT-LECT1	1	1	1
MLANA AND CYB561D2	1	1	1
MLANA AND TMEM115	1	1	1
FDPS AND RER1	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND NOT-ADAM30	1	1	1
NPC1 AND NOT-ADAM30	1	1	1
FDPS AND NOT-ADAM29	1	1	1
MLANA AND NOT-ADAM29	1	1	1
NOT-ADAM29 AND GPR19	1	1	1
NPC1 AND NOT-ADAM29	1	1	1
NOT-ADAM29 AND NOT-EDDM3B	1	1	1
NOT-ADAM29 AND CD63	1	1	1

Antigen Pair Logic	F1	Precision	Recall
FUT4 AND NOT-C1orf210	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-PROM2	0.958579882	0.94921875	0.96812749
FUT4 AND NOT-ADRA2A	0.956349206	0.95256917	0.960159363
FLT3 AND NOT-ARL6IP6	0.944785276	0.970588235	0.920318725
NOT-FAM69A AND METTL21A	0.925373134	0.870175439	0.988047809
FLT3 AND NOT-RMDN2	0.948453608	0.982905983	0.916334661
FUT4 AND NOT-RMDN2	0.962075848	0.964	0.960159363
FLT3 AND NOT-SLC16A14	0.926624738	0.977876106	0.880478088
ADGRE1 AND NOT-GPR155	0.940206186	0.974358974	0.908366534
FLT3 AND NOT-GPR155	0.944785276	0.970588235	0.920318725
HBD AND NOT-GPR155	0.939805825	0.916666667	0.964143426
P2RX1 AND NOT-GPR155	0.958415842	0.952755906	0.964143426
NOT-GPR155 AND P2RX4	0.93852459	0.966244726	0.912350598
NOT-PFN2 AND NOT-GPR155	0.931818182	0.888086643	0.980079681
NOT-MTUS1 AND NOT-GPR155	0.935849057	0.888888889	0.988047809
NOT-PFN2 AND NOT-BTLA	0.957364341	0.932075472	0.984063745
FLT3 AND NOT-CMTM8	0.938271605	0.970212766	0.908366534
FUT4 AND NOT-CMTM8	0.957575758	0.971311475	0.944223108
HBD AND NOT-CMTM8	0.929460581	0.96969697	0.892430279
FLT3 AND NOT-NFXL1	0.939203354	0.991150442	0.892430279
FLT3 AND NOT-CXADR	0.935550936	0.97826087	0.896414343
FUT4 AND NOT-CXADR	0.9498998	0.955645161	0.944223108
FLT3 AND NOT-THAP6	0.954166667	1	0.912350598
FUT4 AND NOT-THAP6	0.957055215	0.983193277	0.932270916
FLT3 AND NOT-CYB5A	0.918454936	0.995348837	0.852589641
FLT3 AND NOT-SLC38A9	0.939958592	0.978448276	0.90438247
FLT3 AND NOT-TMEM161B	0.950413223	0.987124464	0.916334661
FUT4 AND NOT-CYB561	0.96031746	0.956521739	0.964143426
FUT4 AND NOT-MARVELD2	0.960474308	0.952941176	0.96812749
FLT3 AND NOT-NKAIN2	0.93877551	0.962343096	0.916334661
SMIM3 AND NOT-VKORC1L1	0.934156379	0.965957447	0.90438247
FLT3 AND NOT-CYP2J2	0.93877551	0.962343096	0.916334661
FUT4 AND NOT-CYP2J2	0.958415842	0.952755906	0.964143426
FUT4 AND NOT-RDH10	0.9500998	0.952	0.948207171
FLT3 AND NOT-ANKRD46	0.950413223	0.987124464	0.916334661
FUT4 AND NOT-ANKRD46	0.959183673	0.983263598	0.93625498
FUT4 AND NOT-CYP3A5	0.960474308	0.952941176	0.96812749
FLT3 AND NOT-FREM1	0.940936864	0.9625	0.920318725
FUT4 AND NOT-FAAH2	0.938223938	0.91011236	0.96812749
FUT4 AND NOT-SLC35G1	0.960474308	0.952941176	0.96812749
NOT-FAM69A AND CLEC12A	0.94368932	0.920454545	0.96812749
FLT3 AND NOT-DAD1	0.920770878	0.99537037	0.856573705
FLT3 AND NOT-TMTC3	0.948665298	0.978813559	0.920318725
FUT4 AND NOT-TMTC3	0.971428571	0.9958159	0.948207171
NOT-TMTC3 AND SUSD1	0.934615385	0.903345725	0.96812749
NOT-TMTC3 AND SLC30A1	0.938697318	0.904059041	0.976095618
FLT3 AND NOT-DENND5B	0.948665298	0.978813559	0.920318725
FLT3 AND NOT-DAG1	0.948665298	0.978813559	0.920318725
FUT4 AND NOT-DAG1	0.962376238	0.956692913	0.96812749
FLT3 AND NOT-GPR180	0.940936864	0.9625	0.920318725
FLT3 AND NOT-FITM1	0.940936864	0.9625	0.920318725
FUT4 AND NOT-TMEM30B	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-SYNE4	0.947368421	0.927480916	0.96812749
FUT4 AND NOT-TOR1AIP2	0.940206186	0.974358974	0.908366534
FUT4 AND NOT-IFNLR1	0.964143426	0.964143426	0.964143426
P2RY8 AND NOT-IFNLR1	0.940726577	0.904411765	0.980079681
NRROS AND NOT-IFNLR1	0.947154472	0.966804979	0.928286853
P2RX1 AND NOT-IFNLR1	0.954990215	0.938461538	0.972111554
NOT-IFNLR1 AND UBE2J1	0.927643785	0.868055556	0.996015936
HBD AND NOT-SPTSSB	0.932821497	0.9	0.96812749
P2RX1 AND NOT-SPTSSB	0.933078394	0.897058824	0.972111554
FLT3 AND NOT-GIMAP1	0.958506224	1	0.920318725
NOT-PFN2 AND NOT-GIMAP1	0.918819188	0.855670103	0.992031873
FLT3 AND NOT-CLYBL	0.94214876	0.978540773	0.908366534
FUT4 AND NOT-CLYBL	0.952380952	0.948616601	0.956175299
FLT3 AND NOT-SPTSSA	0.952380952	0.99137931	0.916334661
FUT4 AND NOT-SPTSSA	0.957230143	0.979166667	0.93625498
FLT3 AND NOT-DHCR24	0.948665298	0.978813559	0.920318725
FUT4 AND NOT-DHCR24	0.954455446	0.948818898	0.960159363
FUT4 AND NOT-DPP4	0.967741935	0.979591837	0.956175299
HBD AND NOT-DPP4	0.941176471	0.926640927	0.956175299
P2RX1 AND NOT-DPP4	0.937984496	0.913207547	0.964143426
FUT4 AND NOT-SLC26A3	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-DSC2	0.95481336	0.941860465	0.96812749
FUT4 AND NOT-DSG2	0.960474308	0.952941176	0.96812749
FLT3 AND NOT-SLC26A2	0.942622951	0.970464135	0.916334661

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
MLANA AND MLANA	1	1	1
FDPS AND NOT-FDPS	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND MLANA	1	1	1
AUP1 AND NOT-BTN3A2	1	1	1
PARL AND NOT-BTN3A2	1	1	1
AUP1 AND NOT-BTN3A1	1	1	1
PARL AND NOT-BTN3A1	1	1	1
MLANA AND BTN2A1	1	1	1
MLANA AND MLANA	1	1	1
FDPS AND NOT-IL1RAPL1	1	1	1
MLANA AND NOT-IL1RAPL1	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND BVES	1	1	1
MLANA AND MLANA	1	1	1
NPC1 AND NOT-FICD	1	1	1
MLANA AND C14orf1	1	1	1
MLANA AND MLANA	1	1	1
NPC1 AND CHI3L2	1	1	1
MLANA AND SLC2A6	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND NOT-ADCY5	1	1	1
FDPS AND NOT-KLK8	1	1	1
MLANA AND NOT-KLK8	1	1	1
MLANA AND NOT-FZD10	1	1	1
FDPS AND NOT-GALNT6	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND PRAF2	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND MLANA	1	1	1
CLCN5 AND NOT-GPR45	1	1	1
FDPS AND NOT-GPR45	1	1	1
MLANA AND NOT-GPR45	1	1	1
NPC1 AND NOT-GPR45	1	1	1
TMEM187 AND NOT-GPR45	1	1	1
NOT-CD300LB AND PTGDR2	1	1	1
FDPS AND PTGDR2	1	1	1
MLANA AND NOT-PTGDR2	1	1	1
NOT-ITGA2B AND PTGDR2	1	1	1
FDPS AND MAN1B1	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND MLANA	1	1	1
CLCN5 AND NOT-HRH3	1	1	1
FDPS AND NOT-HRH3	1	1	1
MLANA AND NOT-HRH3	1	1	1
NPC1 AND NOT-HRH3	1	1	1
TMEM187 AND NOT-HRH3	1	1	1
MLANA AND NOT-MRAP2	1	1	1
MLANA AND CMTM7	1	1	1
CLCN5 AND NOT-STX1B	1	1	1
FDPS AND NOT-STX1B	1	1	1
MLANA AND NOT-STX1B	1	1	1
GALNS AND NOT-STX1B	1	1	1
NPC1 AND NOT-STX1B	1	1	1
SLC35E3 AND NOT-STX1B	1	1	1
TMEM187 AND NOT-STX1B	1	1	1
MLANA AND KLHL2	1	1	1
MLANA AND GLMP	1	1	1
MLANA AND NOT-SCN11A	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND MGAT4B	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND MLANA	1	1	1
CLCN5 AND NOT-CHRM2	1	1	1
FDPS AND NOT-CHRM2	1	1	1
MLANA AND NOT-CHRM2	1	1	1
NOT-CHRNA2 AND ADCY6	1	1	1
NOT-CLDN19 AND ADCY6	1	1	1
FDPS AND ADCY6	1	1	1
MLANA AND MLANA	1	1	1
NOT-KCNS2 AND ADCY6	1	1	1
AUP1 AND ADCY6	1	1	1

Antigen Pair Logic	F1	Precision	Recall
FUT4 AND NOT-SLC26A2	0.96222664	0.96031746	0.964143426
FLT3 AND NOT-AGTR1	0.948665298	0.978813559	0.920318725
SMIM3 AND NOT-APLNR	0.943907157	0.917293233	0.972111554
FLT3 AND NOT-S1PR1	0.956340956	1	0.916334661
NRROS AND NOT-S1PR1	0.931392931	0.973913043	0.892430279
FUT4 AND NOT-LPAR1	0.979757085	0.995884774	0.964143426
NOT-LPAR1 AND CXCL8	0.944550669	0.908088235	0.984063745
P2RX1 AND NOT-LPAR1	0.9348659	0.900369004	0.972111554
NOT-LPAR1 AND B3GNT5	0.92936803	0.871080139	0.996015936
NOT-LPAR1 AND SMIM3	0.954198473	0.915750916	0.996015936
FLT3 AND NOT-EDNRB	0.948665298	0.978813559	0.920318725
FUT4 AND NOT-EDNRB	0.957915832	0.963709677	0.952191235
NOT-EDNRB AND SMIM3	0.962671906	0.949612403	0.976095618
FLT3 AND NOT-EFNB2	0.939958592	0.978448276	0.90438247
FUT4 AND NOT-EFNB2	0.964	0.967871486	0.960159363
NOT-FAM69A AND NOT-EFNB2	0.935361217	0.894545455	0.980079681
FLT3 AND NOT-MEGF8	0.940936864	0.9625	0.920318725
FLT3 AND NOT-MPZL3	0.936082474	0.97008547	0.90438247
FUT4 AND NOT-MPZL3	0.935010482	0.986725664	0.888446215
ELANE AND NOT-FAM69A	0.938016529	0.974248927	0.90438247
ELANE AND NOT-PHLDB2	0.931392931	0.973913043	0.892430279
NOT-MTUS1 AND TMEM201	0.904761905	0.990521327	0.832669323
FLT3 AND NOT-PLD5	0.940936864	0.9625	0.920318725
FLT3 AND NOT-KRTCAP3	0.93877551	0.962343096	0.916334661
FUT4 AND NOT-KRTCAP3	0.958415842	0.952755906	0.964143426
FUT4 AND NOT-EMP2	0.960474308	0.952941176	0.96812749
NOT-FAM69A AND NOT-EMP2	0.919413919	0.850847458	1
NOT-PTPRK AND EMP3	0.930841121	0.876760563	0.992031873
NOT-FAM69A AND ADGRE1	0.936660269	0.903703704	0.972111554
NOT-HEG1 AND ADGRE1	0.919148936	0.98630137	0.860557769
FLT3 AND NOT-TIGIT	0.952380952	0.99137931	0.916334661
HBD AND NOT-TIGIT	0.936416185	0.906716418	0.96812749
NOT-PFN2 AND NOT-TIGIT	0.930581614	0.879432624	0.988047809
NOT-PTPRK AND NOT-TIGIT	0.922222222	0.861591696	0.992031873
FLT3 AND NOT-TMEM154	0.921775899	0.981981982	0.868525896
FUT4 AND NOT-SMIM14	0.966202783	0.964285714	0.96812749
ADGRE2 AND NOT-SMIM14	0.909090909	0.995260664	0.836653386
NRROS AND NOT-SMIM14	0.943089431	0.962655602	0.924302789
P2RX1 AND NOT-SMIM14	0.948207171	0.948207171	0.948207171
FXYD5 AND NOT-SMIM14	0.940944882	0.929961089	0.952191235
FLT3 AND NOT-C9orf91	0.951983299	1	0.908366534
FLT3 AND NOT-CCDC107	0.943632568	0.99122807	0.900398406
FUT4 AND NOT-CCDC107	0.933884298	0.969957082	0.900398406
FUT4 AND NOT-SLC25A43	0.941414141	0.954918033	0.928286853
NOT-LRIG1 AND VMA21	0.923364486	0.86971831	0.984063745
NOT-MINOS1 AND VMA21	0.952941176	0.938223938	0.96812749
NOT-SLC37A3 AND VMA21	0.938223938	0.91011236	0.96812749
NOT-VAPB AND VMA21	0.924214418	0.862068966	0.996015936
NOT-ARMCX2 AND VMA21	0.92393321	0.864583333	0.992031873
FLT3 AND NOT-CERS3	0.940936864	0.9625	0.920318725
FUT4 AND NOT-EPHA4	0.952755906	0.941634241	0.964143426
HBD AND NOT-EPHA4	0.943469786	0.923664122	0.964143426
P2RX1 AND NOT-EPHA4	0.953307393	0.931558935	0.976095618
FUT4 AND NOT-EPHB2	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-ERBB2	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-ERBB3	0.960474308	0.952941176	0.96812749
FLT3 AND NOT-ETFDH	0.9375	0.982532751	0.896414343
FUT4 AND NOT-ETFDH	0.953907816	0.959677419	0.948207171
NOT-LRIG1 AND ETV6	0.946322068	0.944444444	0.948207171
NOT-FAM69A AND ETV6	0.952380952	0.912408759	0.996015936
NOT-MINOS1 AND ETV6	0.940944882	0.929961089	0.952191235
NOT-HEG1 AND ETV6	0.919831224	0.977578475	0.868525896
NOT-SLC37A3 AND ETV6	0.958415842	0.952755906	0.964143426
NOT-PHLDB2 AND ETV6	0.954990215	0.938461538	0.972111554
NOT-VAPB AND ETV6	0.941176471	0.926640927	0.956175299
NOT-FAM69A AND EVI2A	0.92962963	0.868512111	1
NOT-FAM69A AND EVI2B	0.967117988	0.939849624	0.996015936
FLT3 AND NOT-EXT1	0.956521739	0.995689655	0.920318725
FUT4 AND NOT-EXT1	0.97188755	0.979757085	0.964143426
NOT-EXT1 AND SMIM3	0.951267057	0.93129771	0.972111554
FUT4 AND NOT-F2RL1	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-FAT1	0.958415842	0.952755906	0.964143426
FLT3 AND NOT-TMEM136	0.940936864	0.9625	0.920318725
FLT3 AND NOT-ABCA3	0.940936864	0.9625	0.920318725
FLT3 AND NOT-CYB561A3	0.911447084	0.995283019	0.84063745
FUT4 AND NOT-SLC16A9	0.954635108	0.9453125	0.964143426

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
PARL AND ADCY6	1	1	1
NOT-CACNG6 AND ADCY6	1	1	1
NOT-SLC5A5 AND ADCY6	1	1	1
NOT-TM4SF5 AND ADCY6	1	1	1
NOT-SLC22A8 AND ADCY6	1	1	1
NOT-PTH2 AND TLCD1	1	1	1
NOT-PTH2 AND CLCN5	1	1	1
NOT-PTH2 AND SLC44A3	1	1	1
NOT-PTH2 AND LRRC15	1	1	1
NOT-PTH2 AND MANEAL	1	1	1
NOT-PTH2 AND PROM2	1	1	1
NOT-PTH2 AND CCDC80	1	1	1
NOT-PTH2 AND DSC3	1	1	1
NOT-PTH2 AND ELN	1	1	1
NOT-PTH2 AND EMP2	1	1	1
NOT-PTH2 AND ENG	1	1	1
FDPS AND NOT-PTH2	1	1	1
MLANA AND NOT-PTH2	1	1	1
NOT-PTH2 AND INAFM1	1	1	1
NOT-PTH2 AND SUSD5	1	1	1
NOT-PTH2 AND GPR19	1	1	1
NOT-PTH2 AND SLC2A8	1	1	1
NOT-PTH2 AND IL1R1	1	1	1
NOT-PTH2 AND JAG2	1	1	1
NOT-PTH2 AND KIT	1	1	1
NOT-PTH2 AND TMEM220	1	1	1
NOT-PTH2 AND TSPAN11	1	1	1
NOT-PTH2 AND ATP1A1	1	1	1
NPC1 AND NOT-PTH2	1	1	1
NOT-PTH2 AND CDON	1	1	1
NOT-PTH2 AND ENPP2	1	1	1
NOT-PTH2 AND PMP22	1	1	1
NOT-PTH2 AND IL20RB	1	1	1
NOT-PTH2 AND DNAJC25	1	1	1
NOT-PTH2 AND SLC47A1	1	1	1
NOT-PTH2 AND LGR4	1	1	1
NOT-PTH2 AND TNFRSF19	1	1	1
NOT-PTH2 AND SLC35E3	1	1	1
NOT-PTH2 AND PCDHB14	1	1	1
NOT-PTH2 AND IGDCC4	1	1	1
NOT-PTH2 AND SORT1	1	1	1
NOT-PTH2 AND SDC1	1	1	1
NOT-PTH2 AND PERP	1	1	1
NOT-PTH2 AND TPBG	1	1	1
NOT-PTH2 AND TYRP1	1	1	1
NOT-PTH2 AND COL14A1	1	1	1
NOT-PTH2 AND VCAM1	1	1	1
NOT-PTH2 AND DHRS11	1	1	1
NOT-PTH2 AND WLS	1	1	1
NOT-PTH2 AND PIGZ	1	1	1
TMEM187 AND NOT-PTH2	1	1	1
NOT-PTH2 AND FZD4	1	1	1
NOT-PTH2 AND PLVAP	1	1	1
NOT-PTH2 AND ZNRF3	1	1	1
NOT-PTH2 AND C15orf48	1	1	1
NOT-PTH2 AND CNTNAP1	1	1	1
NOT-PTH2 AND LY6D	1	1	1
NOT-PTH2 AND OSMR	1	1	1
NOT-PTH2 AND REEP6	1	1	1
AUP1 AND NOT-CD300A	1	1	1
MLANA AND NOT-SCAMP4	1	1	1
CLCN5 AND NOT-GPR182	1	1	1
FDPS AND NOT-GPR182	1	1	1
MLANA AND NOT-GPR182	1	1	1
NPC1 AND NOT-GPR182	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND MGAT4A	1	1	1
MLANA AND MLANA	1	1	1
AUP1 AND NOT-TMC6	1	1	1
PARL AND NOT-TMC6	1	1	1
MLANA AND SLC46A1	1	1	1
MLANA AND NOT-GLCCI1	1	1	1
AUP1 AND NOT-GLCCI1	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND TMEM106A	1	1	1
FDPS AND NOT-SLC52A3	1	1	1

Antigen Pair Logic	F1	Precision	Recall
FUT4 AND NOT-REEP3	0.944	0.947791165	0.940239044
NOT-FAM171A1 AND SMIM3	0.953156823	0.975	0.932270916
NOT-FAM69A AND NOT-ADGRF5	0.945386064	0.896428571	1
FLT3 AND NOT-FKTN	0.940936864	0.9625	0.920318725
FLT3 AND NOT-SLC35F1	0.940936864	0.9625	0.920318725
FLT3 AND NOT-ALDH3A2	0.948665298	0.978813559	0.920318725
FUT4 AND NOT-ALDH3A2	0.935010482	0.986725664	0.888446215
FUT4 AND NOT-FGFR3	0.949494949	0.963114754	0.93625498
FUT4 AND NOT-FGFR2	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-CLCA4	0.960474308	0.952941176	0.96812749
FLT3 AND NOT-DOLK	0.944099379	0.982758621	0.908366534
FUT4 AND NOT-DOLK	0.961460446	0.979338843	0.944223108
FUT4 AND NOT-LMTK2	0.953535354	0.967213115	0.940239044
FLT3 AND NOT-ADGRL1	0.940936864	0.9625	0.920318725
NOT-DIP2C AND FUT4	0.952015355	0.918518519	0.988047809
NOT-DIP2C AND NOT-SIT1	0.918819188	0.855670103	0.992031873
NOT-DIP2C AND NOT-HLA-DOB	0.917431193	0.850340136	0.996015936
NOT-DIP2C AND CXCL8	0.930320151	0.882142857	0.984063745
NOT-FAM69A AND NOT-DIP2C	0.922509225	0.859106529	0.996015936
NOT-DIP2C AND SCCPDH	0.94368932	0.920454545	0.96812749
NOT-DIP2C AND SMIM3	0.97415507	0.972222222	0.976095618
NOT-DIP2C AND NOT-PHLDB2	0.927102804	0.873239437	0.988047809
NOT-DIP2C AND NOT-FCMR	0.920810313	0.856164384	0.996015936
FUT4 AND NOT-KLHDC10	0.958083832	0.96	0.956175299
FLT3 AND NOT-TMCC1	0.945378151	1	0.896414343
FUT4 AND NOT-TMCC1	0.947807933	0.995614035	0.90438247
P2RX1 AND NOT-TMCC1	0.952	0.955823293	0.948207171
NOT-PTPRK AND NOT-TMCC1	0.934615385	0.903345725	0.96812749
SLC22A16 AND NOT-TMCC1	0.926931106	0.973684211	0.884462151
FUT4 AND NOT-ENDOD1	0.956521739	0.995689655	0.920318725
FUT4 AND NOT-ATP10B	0.960474308	0.952941176	0.96812749
FLT3 AND NOT-LRCH1	0.948665298	0.978813559	0.920318725
FUT4 AND NOT-CLCC1	0.963709677	0.975510204	0.952191235
FUT4 AND NOT-SNX13	0.911447084	0.995283019	0.84063745
FUT4 AND NOT-SLC35D1	0.954274354	0.952380952	0.956175299
FLT3 AND NOT-SYT11	0.936708861	0.995515695	0.884462151
NOT-FAM69A AND NUP210	0.956349206	0.95256917	0.960159363
FLT3 AND NOT-ABCA5	0.935817805	0.974137931	0.900398406
FLT3 AND NOT-ICMT	0.938271605	0.970212766	0.908366534
FLT3 AND NOT-FOLH1	0.944785276	0.970588235	0.920318725
FLT3 AND NOT-NNT	0.940936864	0.9625	0.920318725
FLT3 AND NOT-MMD	0.906318083	1	0.828685259
FLT3 AND NOT-PIGN	0.940936864	0.9625	0.920318725
FLT3 AND NOT-LEMD3	0.945606695	0.995594714	0.900398406
FLT3 AND NOT-RABGAP1	0.939958592	0.978448276	0.90438247
FLT3 AND NOT-TMEM2	0.950413223	0.987124464	0.916334661
FLT3 AND NOT-STX12	0.951983299	1	0.908366534
FLT3 AND NOT-APOL2	0.942622951	0.970464135	0.916334661
FLT3 AND NOT-FJX1	0.948665298	0.978813559	0.920318725
FLT3 AND NOT-LCLAT1	0.942857143	0.966527197	0.920318725
FLT3 AND NOT-MFSD8	0.948665298	0.978813559	0.920318725
FLT3 AND NOT-BACE2	0.948665298	0.978813559	0.920318725
FLT3 AND NOT-TMEM184B	0.940936864	0.9625	0.920318725
FLT3 AND NOT-TMEM186	0.954166667	1	0.912350598
FLT3 AND NOT-ABI3BP	0.948665298	0.978813559	0.920318725
FLT3 AND NOT-RNF19A	0.948665298	0.978813559	0.920318725
FLT3 AND NOT-C2CD2	0.946502058	0.978723404	0.916334661
FLT3 AND NOT-UNC50	0.925053533	1	0.860557769
FLT3 AND NOT-DHRS7B	0.913419913	1	0.84063745
FLT3 AND NOT-LRIG1	0.954545455	0.991416309	0.920318725
FLT3 AND NOT-TMEM98	0.933333333	0.978165939	0.892430279
FLT3 AND NOT-GLCE	0.940936864	0.9625	0.920318725
FLT3 AND NOT-LRRTM2	0.940936864	0.9625	0.920318725
FLT3 AND NOT-SEZ6L2	0.940936864	0.9625	0.920318725
FLT3 AND NOT-OR7A5	0.940936864	0.9625	0.920318725
FLT3 AND NOT-GHR	0.948665298	0.978813559	0.920318725
FLT3 AND NOT-LAMP3	0.944785276	0.970588235	0.920318725
FLT3 AND NOT-GLG1	0.939203354	0.991150442	0.892430279
FLT3 AND NOT-SLC46A3	0.944785276	0.970588235	0.920318725
FLT3 AND NOT-ANG	0.940695297	0.966386555	0.916334661
FLT3 AND NOT-GPR18	0.947589099	1	0.900398406
FLT3 AND NOT-CYB561D1	0.950617284	0.982978723	0.920318725
FLT3 AND NOT-SLC9A9	0.954545455	0.991416309	0.920318725
FLT3 AND NOT-PRRT3	0.940936864	0.9625	0.920318725
FLT3 AND NOT-CYP4V2	0.941908714	0.982683983	0.90438247
FLT3 AND NOT-XKR6	0.940936864	0.9625	0.920318725

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
MLANA AND NOT-SLC52A3	1	1	1
FDPS AND NOT-CHRM4	1	1	1
MLANA AND NOT-CHRM4	1	1	1
NPC1 AND NOT-CHRM4	1	1	1
TYRP1 AND NOT-CHRM4	1	1	1
TMEM187 AND NOT-CHRM4	1	1	1
FDPS AND NOT-CHRM5	1	1	1
MLANA AND NOT-CHRM5	1	1	1
NPC1 AND NOT-CHRM5	1	1	1
MLANA AND SFT2D1	1	1	1
MLANA AND TEX261	1	1	1
MLANA AND NOT-SCRG1	1	1	1
MLANA AND RNF13	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND CMTM1	1	1	1
NOT-CHRNA2 AND ABHD15	1	1	1
NOT-CHRNA2 AND TLCD1	1	1	1
NOT-CHRNA2 AND CLCN5	1	1	1
NOT-CHRNA2 AND CPXM2	1	1	1
NOT-CHRNA2 AND SLC44A3	1	1	1
NOT-CHRNA2 AND TMEM178A	1	1	1
NOT-CHRNA2 AND LRRC15	1	1	1
NOT-CHRNA2 AND TRAM1L1	1	1	1
NOT-CHRNA2 AND SLC47A2	1	1	1
NOT-CHRNA2 AND APCDD1	1	1	1
NOT-CHRNA2 AND MANEAL	1	1	1
NOT-CHRNA2 AND PROM2	1	1	1
NOT-CHRNA2 AND ADRA2A	1	1	1
NOT-CHRNA2 AND CCDC80	1	1	1
NOT-CHRNA2 AND ADRA2C	1	1	1
NOT-CHRNA2 AND ADGRA3	1	1	1
NOT-CHRNA2 AND DSC3	1	1	1
NOT-CHRNA2 AND APLNR	1	1	1
NOT-CHRNA2 AND EFNB2	1	1	1
NOT-CHRNA2 AND EMP1	1	1	1
NOT-CHRNA2 AND EMP2	1	1	1
NOT-CHRNA2 AND ENG	1	1	1
NOT-CHRNA2 AND FAT2	1	1	1
FDPS AND NOT-CHRNA2	1	1	1
MLANA AND NOT-CHRNA2	1	1	1
NOT-CHRNA2 AND INAFM1	1	1	1
NOT-CHRNA2 AND GABRE	1	1	1
NOT-CHRNA2 AND KIAA1549L	1	1	1
NOT-CHRNA2 AND ATL3	1	1	1
NOT-CHRNA2 AND TENM4	1	1	1
NOT-CHRNA2 AND SUSD5	1	1	1
NOT-CHRNA2 AND ABCA12	1	1	1
NOT-CHRNA2 AND GREM1	1	1	1
NOT-CHRNA2 AND GJA1	1	1	1
NOT-CHRNA2 AND GJA4	1	1	1
NOT-CHRNA2 AND GPR19	1	1	1
NOT-CHRNA2 AND DLL1	1	1	1
NOT-CHRNA2 AND GUSB	1	1	1
NOT-CHRNA2 AND SLC2A8	1	1	1
NOT-CHRNA2 AND HRH1	1	1	1
NOT-CHRNA2 AND GLDN	1	1	1
NOT-CHRNA2 AND IFI27	1	1	1
NOT-CHRNA2 AND IL1R1	1	1	1
NOT-CHRNA2 AND AQP1	1	1	1
NOT-CHRNA2 AND IL12B	1	1	1
NOT-CHRNA2 AND JAG2	1	1	1
NOT-CHRNA2 AND KCNJ8	1	1	1
NOT-CHRNA2 AND KDR	1	1	1
NOT-CHRNA2 AND KIT	1	1	1
NOT-CHRNA2 AND SERP2	1	1	1
NOT-CHRNA2 AND TMEM220	1	1	1
NOT-CHRNA2 AND SAMD5	1	1	1
NOT-CHRNA2 AND FADS3	1	1	1
NOT-CHRNA2 AND TSPAN11	1	1	1
NOT-CHRNA2 AND ATP1A1	1	1	1
NPC1 AND NOT-CHRNA2	1	1	1
NOT-CHRNA2 AND ROR1	1	1	1
NOT-CHRNA2 AND NSDHL	1	1	1
NOT-CHRNA2 AND NTM	1	1	1
NOT-CHRNA2 AND CDON	1	1	1

Antigen Pair Logic	F1	Precision	Recall
FLT3 AND NOT-MICU3	0.937759336	0.978354978	0.900398406
FLT3 AND NOT-SLC6A16	0.940936864	0.9625	0.920318725
FLT3 AND NOT-MAGEH1	0.929637527	1	0.868525896
FLT3 AND NOT-TIMM21	0.950207469	0.991341991	0.912350598
FLT3 AND NOT-DNAJC15	0.933054393	0.982378855	0.888446215
FLT3 AND NOT-GRM5	0.940936864	0.9625	0.920318725
FLT3 AND NOT-SLC25A4	0.942857143	0.966527197	0.920318725
FLT3 AND NOT-GNL2	0.901531729	1	0.820717131
FLT3 AND NOT-ALG6	0.931677019	0.969827586	0.896414343
FLT3 AND NOT-SLC2A8	0.944558522	0.974576271	0.916334661
FLT3 AND NOT-GZMK	0.945378151	1	0.896414343
FLT3 AND NOT-HLA-DQB2	0.942857143	0.966527197	0.920318725
FLT3 AND NOT-HMGCR	0.956340956	1	0.916334661
FLT3 AND NOT-HMOX2	0.954545455	0.991416309	0.920318725
FLT3 AND NOT-ACACB	0.938271605	0.970212766	0.908366534
FLT3 AND NOT-UBAC2	0.947589099	1	0.900398406
FLT3 AND NOT-ICAM2	0.922746781	1	0.856573705
FLT3 AND NOT-C2orf74	0.95257732	0.987179487	0.920318725
FLT3 AND NOT-FAM174A	0.90712743	0.990566038	0.836653386
FLT3 AND NOT-AMIGO2	0.956521739	0.995689655	0.920318725
FLT3 AND NOT-ZACN	0.940936864	0.9625	0.920318725
FLT3 AND NOT-IL1R1	0.940206186	0.974358974	0.908366534
FLT3 AND NOT-IL11RA	0.940936864	0.9625	0.920318725
FLT3 AND NOT-AQP7	0.948453608	0.982905983	0.916334661
FLT3 AND NOT-ITGAV	0.943866944	0.986956522	0.90438247
FLT3 AND NOT-ITGB5	0.948665298	0.978813559	0.920318725
FLT3 AND NOT-SFT2D2	0.93220339	0.995475113	0.876494024
FLT3 AND NOT-KCNMA1	0.948665298	0.978813559	0.920318725
FLT3 AND NOT-KDR	0.948665298	0.978813559	0.920318725
FLT3 AND NOT-SERP2	0.940936864	0.9625	0.920318725
FLT3 AND NOT-SHISA6	0.940936864	0.9625	0.920318725
FLT3 AND NOT-FAM69A	0.958506224	1	0.920318725
FLT3 AND NOT-OR7A10	0.940936864	0.9625	0.920318725
FLT3 AND NOT-C4orf3	0.923728814	0.986425339	0.868525896
FLT3 AND NOT-PPAPDC2	0.954545455	0.991416309	0.920318725
FLT3 AND NOT-LRMP	0.928571429	0.982222222	0.880478088
FLT3 AND NOT-SMAD2	0.949790795	1	0.90438247
FLT3 AND CD46	0.931677019	0.969827586	0.896414343
FLT3 AND NOT-MME	0.93852459	0.966244726	0.912350598
FLT3 AND NOT-TMEM41B	0.929637527	1	0.868525896
FLT3 AND NOT-MINOS1	0.950617284	0.982978723	0.920318725
FLT3 AND NOT-TMEM150C	0.948665298	0.978813559	0.920318725
FLT3 AND NOT-NDUFC2	0.94214876	0.978540773	0.908366534
FLT3 AND NOT-NPY	0.940936864	0.9625	0.920318725
FLT3 AND NOT-NPC1	0.951983299	1	0.908366534
FLT3 AND NOT-OCA2	0.940936864	0.9625	0.920318725
FLT3 AND NOT-NSDHL	0.937759336	0.978354978	0.900398406
FLT3 AND NOT-SLC35B3	0.945606695	0.995594714	0.900398406
FLT3 AND NOT-NDUFA13	0.901960784	0.995192308	0.824701195
FLT3 AND NOT-KLHL5	0.923728814	0.986425339	0.868525896
FLT3 AND NOT-SIDT2	0.933333333	0.978165939	0.892430279
FLT3 AND NOT-RDH11	0.927966102	0.990950226	0.87250996
FLT3 AND NOT-INSIG2	0.956340956	1	0.916334661
FLT3 AND NOT-C3orf18	0.931392931	0.973913043	0.892430279
FLT3 AND NOT-TUBD1	0.958506224	1	0.920318725
FLT3 AND NOT-CRIM1	0.948665298	0.978813559	0.920318725
FLT3 AND NOT-TMEM69	0.918803419	0.99078341	0.856573705
FLT3 AND NOT-GOLM1	0.948665298	0.978813559	0.920318725
FLT3 AND NOT-BFAR	0.93220339	0.995475113	0.876494024
FLT3 AND NOT-TIMMDC1	0.950413223	0.987124464	0.916334661
FLT3 AND NOT-ARMCX1	0.926624738	0.977876106	0.880478088
FLT3 AND NOT-SLC22A17	0.940936864	0.9625	0.920318725
FLT3 AND NOT-TEX264	0.922105263	0.977678571	0.87250996
FLT3 AND NOT-WNT16	0.944785276	0.970588235	0.920318725
FLT3 AND NOT-TMEM138	0.940936864	0.9625	0.920318725
FLT3 AND NOT-FKBP7	0.93877551	0.962343096	0.916334661
FLT3 AND NOT-ENPP2	0.948665298	0.978813559	0.920318725
FLT3 AND NOT-ACSL5	0.948024948	0.991304348	0.908366534
FLT3 AND NOT-EMCN	0.948665298	0.978813559	0.920318725
FLT3 AND NOT-ATP8B1	0.948665298	0.978813559	0.920318725
FLT3 AND NOT-PFN2	0.948665298	0.978813559	0.920318725
FLT3 AND NOT-PLA2G2A	0.93877551	0.962343096	0.916334661
FLT3 AND NOT-IL17D	0.940936864	0.9625	0.920318725
FLT3 AND NOT-FXYD1	0.942857143	0.966527197	0.920318725
FLT3 AND NOT-PLOD2	0.944329897	0.978632479	0.912350598
FLT3 AND NOT-PLP1	0.940936864	0.9625	0.920318725

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
NOT-CHRNA2 AND ARMCX1	1	1	1
NOT-CHRNA2 AND PDGFRA	1	1	1
NOT-CHRNA2 AND FKBP7	1	1	1
NOT-CHRNA2 AND SERPINB13	1	1	1
NOT-CHRNA2 AND PMP22	1	1	1
NOT-CHRNA2 AND IL20RB	1	1	1
NOT-CHRNA2 AND PCDH18	1	1	1
NOT-CHRNA2 AND ANO10	1	1	1
NOT-CHRNA2 AND RMDN3	1	1	1
NOT-CHRNA2 AND SLC47A1	1	1	1
NOT-CHRNA2 AND TMEM63B	1	1	1
NOT-CHRNA2 AND LGR4	1	1	1
NOT-CHRNA2 AND SLC35E3	1	1	1
NOT-CHRNA2 AND ACSS2	1	1	1
NOT-CHRNA2 AND PCDHB14	1	1	1
NOT-CHRNA2 AND GKN1	1	1	1
NOT-CHRNA2 AND TMEM27	1	1	1
NOT-CHRNA2 AND IGDCC4	1	1	1
NOT-CHRNA2 AND ROBO2	1	1	1
NOT-CHRNA2 AND SORT1	1	1	1
NOT-CHRNA2 AND BDNF	1	1	1
NOT-CHRNA2 AND SCN4B	1	1	1
NOT-CHRNA2 AND SDC1	1	1	1
NOT-CHRNA2 AND PERP	1	1	1
NOT-CHRNA2 AND ADGRL4	1	1	1
NOT-CHRNA2 AND CDH24	1	1	1
NOT-CHRNA2 AND SLC7A2	1	1	1
NOT-CHRNA2 AND SLC15A1	1	1	1
NOT-CHRNA2 AND TIE1	1	1	1
NOT-CHRNA2 AND TSPAN7	1	1	1
NOT-CHRNA2 AND TPBG	1	1	1
NOT-CHRNA2 AND TYRO3	1	1	1
NOT-CHRNA2 AND TYRP1	1	1	1
NOT-CHRNA2 AND COL14A1	1	1	1
NOT-CHRNA2 AND VCAM1	1	1	1
NOT-CHRNA2 AND XG	1	1	1
NOT-CHRNA2 AND PTTG1IP	1	1	1
NOT-CHRNA2 AND FZD5	1	1	1
NOT-CHRNA2 AND SEMA3B	1	1	1
NOT-CHRNA2 AND DHRS11	1	1	1
NOT-CHRNA2 AND TMEM231	1	1	1
NOT-CHRNA2 AND EPHX3	1	1	1
NOT-CHRNA2 AND THSD4	1	1	1
NOT-CHRNA2 AND WLS	1	1	1
NOT-CHRNA2 AND CWH43	1	1	1
NOT-CHRNA2 AND PIGZ	1	1	1
NOT-CHRNA2 AND COLEC12	1	1	1
TMEM187 AND NOT-CHRNA2	1	1	1
NOT-CHRNA2 AND FZD4	1	1	1
NOT-CHRNA2 AND FZD6	1	1	1
NOT-CHRNA2 AND FZD7	1	1	1
NOT-CHRNA2 AND PLVAP	1	1	1
NOT-CHRNA2 AND ZNRF3	1	1	1
NOT-CHRNA2 AND C15orf48	1	1	1
NOT-CHRNA2 AND CNTNAP1	1	1	1
NOT-CHRNA2 AND CAV1	1	1	1
NOT-CHRNA2 AND LY6D	1	1	1
NOT-CHRNA2 AND SEMA5A	1	1	1
NOT-CHRNA2 AND DUOXA1	1	1	1
NOT-CHRNA2 AND KCNQ4	1	1	1
NOT-CHRNA2 AND OSMR	1	1	1
NOT-CHRNA2 AND DSEL	1	1	1
NOT-CHRNA2 AND CDHR1	1	1	1
NOT-CHRNA2 AND G6PC3	1	1	1
NOT-CHRNA2 AND REEP6	1	1	1
NOT-CHRNA2 AND CLCA2	1	1	1
NOT-CHRNA2 AND CD63	1	1	1
MLANA AND CYP2U1	1	1	1
FDPS AND NOT-FDPS	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND SDSL	1	1	1
MLANA AND NOT-CHRNA3	1	1	1
CLCN5 AND NOT-CHRNA4	1	1	1
FDPS AND NOT-CHRNA4	1	1	1
MLANA AND NOT-CHRNA4	1	1	1
NPC1 AND NOT-CHRNA4	1	1	1

Antigen Pair Logic	F1	Precision	Recall
FLT3 AND NOT-PLXNA1	0.940936864	0.9625	0.920318725
FLT3 AND NOT-PLXNA2	0.940936864	0.9625	0.920318725
FLT3 AND NOT-PMP22	0.946502058	0.978723404	0.916334661
FLT3 AND NOT-FXYD6	0.937759336	0.978354978	0.900398406
FLT3 AND NOT-ACP2	0.933884298	0.969957082	0.900398406
FLT3 AND NOT-PODXL	0.948665298	0.978813559	0.920318725
FLT3 AND NOT-GDAP1	0.948665298	0.978813559	0.920318725
FLT3 AND NOT-YIPF1	0.901960784	0.995192308	0.824701195
FLT3 AND NOT-TMX3	0.947807933	0.995614035	0.90438247
FLT3 AND NOT-ZDHHC13	0.940451745	0.970338983	0.912350598
FLT3 AND NOT-CRLS1	0.944099379	0.982758621	0.908366534
FLT3 AND NOT-WBP1L	0.913793103	0.995305164	0.844621514
FLT3 AND NOT-RNF43	0.940936864	0.9625	0.920318725
FLT3 AND NOT-SEMA4C	0.928870293	0.977973568	0.884462151
FLT3 AND NOT-OCIAD1	0.93877551	0.962343096	0.916334661
FLT3 AND NOT-BANP	0.943157895	1	0.892430279
FLT3 AND NOT-SLC35A5	0.918103448	1	0.848605578
FLT3 AND NOT-TMEM248	0.936440678	1	0.880478088
FLT3 AND NOT-PRPF38B	0.937759336	0.978354978	0.900398406
FLT3 AND NOT-LRRC8D	0.935010482	0.986725664	0.888446215
FLT3 AND NOT-SLC38A7	0.940936864	0.9625	0.920318725
FLT3 AND NOT-SIRPG	0.952182952	0.995652174	0.912350598
FLT3 AND NOT-SLC35E3	0.927659574	0.99543379	0.868525896
FLT3 AND NOT-VEZT	0.935010482	0.986725664	0.888446215
FLT3 AND NOT-CARKD	0.938689218	1	0.884462151
FLT3 AND NOT-COA1	0.918803419	0.99078341	0.856573705
FLT3 AND NOT-SAYSD1	0.946721311	0.974683544	0.920318725
FLT3 AND NOT-ERMARD	0.944558522	0.974576271	0.916334661
FLT3 AND NOT-PAG1	0.948240166	0.987068966	0.912350598
FLT3 AND NOT-ECHDC1	0.925053533	1	0.860557769
FLT3 AND NOT-AXL	0.944329897	0.978632479	0.912350598
FLT3 AND NOT-ALG1	0.944558522	0.974576271	0.916334661
FLT3 AND NOT-ANKH	0.940936864	0.9625	0.920318725
FLT3 AND NOT-PROS1	0.944099379	0.982758621	0.908366534
FLT3 AND NOT-DIABLO	0.901960784	0.995192308	0.824701195
FLT3 AND NOT-CLDND1	0.95257732	0.987179487	0.920318725
FLT3 AND NOT-MDM1	0.93852459	0.966244726	0.912350598
FLT3 AND NOT-PMEPA1	0.944558522	0.974576271	0.916334661
FLT3 AND NOT-CEACAM19	0.940936864	0.9625	0.920318725
FLT3 AND NOT-MAN1C1	0.95257732	0.987179487	0.920318725
FLT3 AND NOT-SMIM8	0.942857143	0.966527197	0.920318725
FLT3 AND NOT-NIPAL3	0.942857143	0.966527197	0.920318725
FLT3 AND NOT-ATP10D	0.93418259	1	0.876494024
FLT3 AND NOT-SLC45A4	0.935550936	0.97826087	0.896414343
FLT3 AND NOT-MRS2	0.944558522	0.974576271	0.916334661
FLT3 AND NOT-ABHD6	0.948665298	0.978813559	0.920318725
FLT3 AND NOT-RNF150	0.946502058	0.978723404	0.916334661
FLT3 AND NOT-KIAA1467	0.940936864	0.9625	0.920318725
FLT3 AND NOT-PTN	0.948665298	0.978813559	0.920318725
FLT3 AND NOT-GPR107	0.940936864	0.9625	0.920318725
FLT3 AND NOT-WDR19	0.940936864	0.9625	0.920318725
FLT3 AND NOT-SCUBE2	0.948665298	0.978813559	0.920318725
FLT3 AND NOT-PTPRM	0.948665298	0.978813559	0.920318725
FLT3 AND NOT-PTPRS	0.940936864	0.9625	0.920318725
FLT3 AND NOT-PTPRZ1	0.946721311	0.974683544	0.920318725
FLT3 AND NOT-SRPRB	0.94214876	0.978540773	0.908366534
FLT3 AND NOT-JAM2	0.940936864	0.9625	0.920318725
FLT3 AND NOT-BBS4	0.927966102	0.990950226	0.87250996
FLT3 AND NOT-RARRES3	0.949790795	1	0.90438247
FLT3 AND NOT-EVA1C	0.947807933	0.995614035	0.90438247
FLT3 AND NOT-BCL2L2	0.940936864	0.9625	0.920318725
NOT-FLT3 AND RHAG	0.940952381	0.901459854	0.984063745
NOT-FLT3 AND RHCE	0.939805825	0.916666667	0.964143426
FLT3 AND NOT-RNASE1	0.937759336	0.978354978	0.900398406
FLT3 AND NOT-RTN1	0.954545455	0.991416309	0.920318725
FLT3 AND NOT-RYR1	0.940936864	0.9625	0.920318725
FLT3 AND NOT-SCNN1G	0.940936864	0.9625	0.920318725
FLT3 AND NOT-CX3CL1	0.940936864	0.9625	0.920318725
FLT3 AND NOT-THADA	0.944785276	0.970588235	0.920318725
FLT3 AND NOT-MANBAL	0.945378151	1	0.896414343
FLT3 AND NOT-GALNT11	0.934736842	0.991071429	0.884462151
FLT3 AND NOT-TMBIM1	0.914893617	0.98173516	0.856573705
FLT3 AND NOT-ADGRL4	0.937759336	0.978354978	0.900398406
FLT3 AND NOT-XYLT2	0.940936864	0.9625	0.920318725
FLT3 AND NOT-SMIM15	0.925690021	0.990909091	0.868525896
FLT3 AND NOT-CDH22	0.940936864	0.9625	0.920318725

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
MLANA AND MLANA	1	1	1
MLANA AND NOT-ADCY7	1	1	1
FDPS AND CHRNB1	1	1	1
MLANA AND MLANA	1	1	1
CLCN5 AND NOT-SIGLEC11	1	1	1
MLANA AND NOT-SIGLEC11	1	1	1
NPC1 AND NOT-SIGLEC11	1	1	1
SLC35E3 AND NOT-SIGLEC11	1	1	1
TMEM187 AND NOT-SIGLEC11	1	1	1
MLANA AND SLC2A13	1	1	1
MLANA AND NOT-CHRNB2	1	1	1
MLANA AND NOT-CHRNB3	1	1	1
NPC1 AND NOT-CHRNB3	1	1	1
FDPS AND CHRNB4	1	1	1
MLANA AND NOT-CHRNB4	1	1	1
FDPS AND NOT-CHRND	1	1	1
MLANA AND NOT-CHRND	1	1	1
MLANA AND NOT-MAL2	1	1	1
NPC1 AND MAL2	1	1	1
AUP1 AND MAL2	1	1	1
PARL AND MAL2	1	1	1
MLANA AND NOT-SLC22A9	1	1	1
FDPS AND NOT-CHRNE	1	1	1
MLANA AND NOT-CHRNE	1	1	1
MLANA AND NOT-ERMAP	1	1	1
MLANA AND MLANA	1	1	1
FDPS AND NOT-CHRNG	1	1	1
MLANA AND NOT-CHRNG	1	1	1
FDPS AND PKD1L2	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND NOT-LMTK3	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND NOT-XKR4	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND NOT-ELFN2	1	1	1
FDPS AND NOT-SLITRK1	1	1	1
MLANA AND NOT-SLITRK1	1	1	1
MLANA AND NOT-TMEM200A	1	1	1
CLCN5 AND NOT-GALNT13	1	1	1
FDPS AND NOT-GALNT13	1	1	1
MLANA AND NOT-GALNT13	1	1	1
NPC1 AND NOT-GALNT13	1	1	1
TYRP1 AND NOT-GALNT13	1	1	1
TMEM187 AND NOT-GALNT13	1	1	1
MLANA AND SORCS1	1	1	1
MLANA AND MLANA	1	1	1
FDPS AND NOT-FDPS	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND NOT-VASN	1	1	1
MLANA AND NOT-SLC26A9	1	1	1
FDPS AND NOT-SLC26A7	1	1	1
MLANA AND NOT-SLC26A7	1	1	1
MLANA AND SLC25A26	1	1	1
NOT-PMEL AND SLC25A26	1	1	1
MLANA AND MLANA	1	1	1
PARL AND NOT-FCRL1	1	1	1
MLANA AND FCRL3	1	1	1
FDPS AND NOT-SLC5A11	1	1	1
MLANA AND NOT-SLC5A11	1	1	1
NPC1 AND NOT-SLC5A11	1	1	1
FDPS AND NOT-TNFRSF13C	1	1	1
MLANA AND NOT-TNFRSF13C	1	1	1
NPC1 AND NOT-TNFRSF13C	1	1	1
TMEM187 AND NOT-TNFRSF13C	1	1	1
FDPS AND NOT-EVI5L	1	1	1
MLANA AND NOT-EVI5L	1	1	1
NPC1 AND NOT-EVI5L	1	1	1
MLANA AND MLANA	1	1	1
MLANA AND LYSMD3	1	1	1
FDPS AND NOT-SLC22A12	1	1	1
MLANA AND NOT-SLC22A12	1	1	1
MLANA AND FAM210B	1	1	1
MLANA AND NOT-CYYR1	1	1	1

Antigen Pair Logic	F1	Precision	Recall
FLT3 AND NOT-ZDHHC6	0.927966102	0.990950226	0.87250996
FLT3 AND NOT-LMF1	0.940936864	0.9625	0.920318725
FLT3 AND NOT-CASD1	0.95257732	0.987179487	0.920318725
FLT3 AND NOT-TMEM237	0.940936864	0.9625	0.920318725
FLT3 AND NOT-SLC1A6	0.940936864	0.9625	0.920318725
FLT3 AND NOT-SLC3A1	0.940451745	0.970338983	0.912350598
FLT3 AND NOT-SLC8A3	0.940936864	0.9625	0.920318725
FLT3 AND NOT-SLC22A5	0.940936864	0.9625	0.920318725
FLT3 AND NOT-SLN	0.940936864	0.9625	0.920318725
FLT3 AND NOT-GRAMD3	0.946502058	0.978723404	0.916334661
FLT3 AND NOT-BMPR2	0.950413223	0.987124464	0.916334661
FLT3 AND NOT-DST	0.946502058	0.978723404	0.916334661
FLT3 AND NOT-SRD5A1	0.935281837	0.98245614	0.892430279
FLT3 AND NOT-SSR1	0.922746781	1	0.856573705
FLT3 AND NOT-BST2	0.911062907	1	0.836653386
FLT3 AND NOT-TEK	0.948665298	0.978813559	0.920318725
FLT3 AND NOT-TGFB3	0.940936864	0.9625	0.920318725
FLT3 AND NOT-SEC62	0.956340956	1	0.916334661
FLT3 AND NOT-TSPAN6	0.948665298	0.978813559	0.920318725
FLT3 AND NOT-GPR137B	0.906318083	1	0.828685259
FLT3 AND NOT-CLDN5	0.948665298	0.978813559	0.920318725
FLT3 AND NOT-TNFRSF1A	0.923076923	0.995391705	0.860557769
FLT3 AND NOT-TRPC1	0.944329897	0.978632479	0.912350598
FLT3 AND NOT-TYRP1	0.940936864	0.9625	0.920318725
FLT3 AND NOT-TRPV1	0.940936864	0.9625	0.920318725
FLT3 AND NOT-VRK2	0.926315789	0.982142857	0.876494024
FLT3 AND NOT-ZNF7	0.946502058	0.978723404	0.916334661
FLT3 AND NOT-LRP8	0.940936864	0.9625	0.920318725
FLT3 AND NOT-PVRIG	0.948453608	0.982905983	0.916334661
FLT3 AND NOT-ELOVL6	0.93877551	0.962343096	0.916334661
FLT3 AND NOT-TMEM109	0.947807933	0.995614035	0.90438247
FLT3 AND NOT-SMIM7	0.936440678	1	0.880478088
FLT3 AND NOT-FAM134A	0.935817805	0.974137931	0.900398406
FLT3 AND NOT-FA2H	0.940936864	0.9625	0.920318725
FLT3 AND NOT-SLC39A7	0.938271605	0.970212766	0.908366534
FLT3 AND NOT-TTC13	0.956340956	1	0.916334661
FLT3 AND NOT-TMEM231	0.946721311	0.974683544	0.920318725
FLT3 AND NOT-RNF128	0.940936864	0.9625	0.920318725
FLT3 AND NOT-FAT4	0.948665298	0.978813559	0.920318725
FLT3 AND NOT-C3orf52	0.942857143	0.966527197	0.920318725
FLT3 AND NOT-CCDC51	0.94092827	1	0.888446215
FLT3 AND NOT-FZD3	0.940936864	0.9625	0.920318725
FLT3 AND NOT-TXNDC15	0.936440678	1	0.880478088
FLT3 AND NOT-C12orf49	0.946721311	0.974683544	0.920318725
FLT3 AND NOT-UBA5	0.924369748	0.977777778	0.876494024
FLT3 AND NOT-PAQR6	0.940936864	0.9625	0.920318725
FLT3 AND NOT-ELOVL7	0.939958592	0.978448276	0.90438247
FLT3 AND NOT-MYCT1	0.94214876	0.978540773	0.908366534
FLT3 AND NOT-TMEM134	0.929936306	0.995454545	0.87250996
FLT3 AND NOT-ALPK1	0.940936864	0.9625	0.920318725
FLT3 AND NOT-PAAF1	0.935817805	0.974137931	0.900398406
FLT3 AND NOT-PIGZ	0.940936864	0.9625	0.920318725
FLT3 AND NOT-SLC35F5	0.937759336	0.978354978	0.900398406
FLT3 AND NOT-ADAM33	0.940936864	0.9625	0.920318725
FLT3 AND NOT-LY6G6C	0.940936864	0.9625	0.920318725
FLT3 AND NOT-SSPN	0.935550936	0.97826087	0.896414343
FLT3 AND NOT-FXR1	0.942857143	0.966527197	0.920318725
FLT3 AND NOT-CLPTM1L	0.941422594	0.991189427	0.896414343
FLT3 AND NOT-ITFG1	0.946502058	0.978723404	0.916334661
FLT3 AND NOT-VWA9	0.948240166	0.987068966	0.912350598
FLT3 AND NOT-APOLD1	0.944329897	0.978632479	0.912350598
FLT3 AND NOT-NIPA2	0.936440678	1	0.880478088
FLT3 AND NOT-TMEM163	0.940206186	0.974358974	0.908366534
FLT3 AND NOT-CANX	0.924686192	0.973568282	0.880478088
FLT3 AND NOT-FZD7	0.940936864	0.9625	0.920318725
FLT3 AND NOT-SLC10A7	0.956521739	0.995689655	0.920318725
FLT3 AND NOT-ZNRF3	0.940936864	0.9625	0.920318725
FLT3 AND NOT-STX7	0.954166667	1	0.912350598
FLT3 AND NOT-SLC37A3	0.946721311	0.974683544	0.920318725
FLT3 AND NOT-DNAJC30	0.946280992	0.982832618	0.912350598
FLT3 AND NOT-FAM213A	0.937759336	0.978354978	0.900398406
FLT3 AND NOT-TMEM246	0.940936864	0.9625	0.920318725
FLT3 AND NOT-TMEM101	0.948453608	0.982905983	0.916334661
FLT3 AND NOT-ZNF559	0.948024948	0.991304348	0.908366534
FLT3 AND NOT-ACSS1	0.919831224	0.977578475	0.868525896
FLT3 AND NOT-ST6GAL2	0.940936864	0.9625	0.920318725

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
MLANA AND NOT-CMTM5	1	1	1
NOT-CD300LB AND ABHD15	1	1	1
NOT-FCN2 AND ABHD15	1	1	1
FDPS AND ABHD15	1	1	1
MLANA AND ABHD15	1	1	1
NOT-TMEM105 AND ABHD15	1	1	1
NOT-TMEM150B AND ABHD15	1	1	1
NPC1 AND ABHD15	1	1	1
NOT-SLC4A1 AND ABHD15	1	1	1
NOT-ACKR2 AND TLCD1	1	1	1
NOT-CD300LB AND TLCD1	1	1	1
NOT-SYT2 AND TLCD1	1	1	1
NOT-PROKR2 AND TLCD1	1	1	1
NOT-CLDN7 AND TLCD1	1	1	1
NOT-SLC34A3 AND TLCD1	1	1	1
NOT-ISM2 AND TLCD1	1	1	1
NOT-ODF4 AND TLCD1	1	1	1
NOT-CLDN19 AND TLCD1	1	1	1
NOT-CYP2A6 AND TLCD1	1	1	1
NOT-MFSD6L AND TLCD1	1	1	1
NOT-DNAH10 AND TLCD1	1	1	1
NOT-FCN2 AND TLCD1	1	1	1
MLANA AND MLANA	1	1	1
NOT-FSHR AND TLCD1	1	1	1
NOT-FUT6 AND TLCD1	1	1	1
NOT-CALHM1 AND TLCD1	1	1	1
NOT-OR2F1 AND TLCD1	1	1	1
NOT-OR10J1 AND TLCD1	1	1	1
NOT-GIPR AND TLCD1	1	1	1
NOT-AMHR2 AND TLCD1	1	1	1
NOT-NOX1 AND TLCD1	1	1	1
NOT-CACNG5 AND TLCD1	1	1	1
NOT-CACNG4 AND TLCD1	1	1	1
NOT-GLP1R AND TLCD1	1	1	1
NOT-GLRA2 AND TLCD1	1	1	1
NOT-GP2 AND TLCD1	1	1	1
NOT-B4GALNT3 AND TLCD1	1	1	1
NOT-TMEM105 AND TLCD1	1	1	1
NOT-TMEM150B AND TLCD1	1	1	1
NOT-FFAR2 AND TLCD1	1	1	1
Lymphoma, Large-Cell, Anaplastic (Anaplastic Lymphoma)			
TNFRSF8 AND NOT-GPC3	0.928571429	1	0.866666667
TNFRSF8 AND NOT-L1CAM	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SDC1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TNFRSF10A	0.965517241	1	0.933333333
TNFRSF8 AND NOT-ERBB2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-IL11RA	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SYT15	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CYP4F2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KCNK13	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-NTN1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GGCX	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ANO4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CGA	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-HYAL4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMEM231	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-AGPAT4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FGFR4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ITGA11	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-OR7C2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-DNASE1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CD151	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-NPC1L1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CMTM3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-LRRTM2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMEM169	0.965517241	1	0.933333333
TNFRSF8 AND NOT-MMP24	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-LRTOMT	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PAQR6	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PTPRH	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FAM69A	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-DKKL1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-OR7A17	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ITIH1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PTAFR	0.965517241	1	0.933333333
TNFRSF8 AND NOT-KIR2DL3	0.933333333	0.93333333	0.933333333

Antigen Pair Logic	F1	Precision	Recall
FLT3 AND NOT-HIATL1	0.950207469	0.991341991	0.912350598
FLT3 AND NOT-COX4I2	0.942857143	0.966527197	0.920318725
FLT3 AND NOT-SFT2D3	0.950413223	0.987124464	0.916334661
FLT3 AND NOT-LRCH3	0.944785276	0.970588235	0.920318725
FLT3 AND NOT-TMEM128	0.947589099	1	0.900398406
FLT3 AND NOT-CNTNAP1	0.940936864	0.9625	0.920318725
FLT3 AND NOT-PAQR8	0.944329897	0.978632479	0.912350598
FLT3 AND NOT-CDC14B	0.940695297	0.966386555	0.916334661
FLT3 AND NOT-CAV1	0.948665298	0.978813559	0.920318725
FLT3 AND NOT-PPAP2A	0.937759336	0.978354978	0.900398406
FLT3 AND NOT-PPAP2B	0.944329897	0.978632479	0.912350598
FLT3 AND NOT-S1PR4	0.914893617	0.98173516	0.856573705
FLT3 AND NOT-ST3GAL5	0.958506224	1	0.920318725
FLT3 AND NOT-SGPL1	0.958506224	1	0.920318725
FLT3 AND NOT-P4HA2	0.948665298	0.978813559	0.920318725
FLT3 AND NOT-ABCC10	0.944558522	0.974576271	0.916334661
FLT3 AND NOT-TMEM116	0.927350427	1	0.864541833
FLT3 AND NOT-PHLDB2	0.954166667	1	0.912350598
FLT3 AND NOT-SPAG9	0.9375	0.982532751	0.896414343
FLT3 AND NOT-CLDN12	0.940695297	0.966386555	0.916334661
FLT3 AND NOT-AIFM1	0.909871245	0.986046512	0.844621514
FLT3 AND NOT-VAPB	0.944785276	0.970588235	0.920318725
FLT3 AND NOT-ATP6AP1L	0.940936864	0.9625	0.920318725
FLT3 AND NOT-PXYLP1	0.943866944	0.986956522	0.90438247
FLT3 AND NOT-ACVR2A	0.944329897	0.978632479	0.912350598
FLT3 AND NOT-KL	0.948453608	0.982905983	0.916334661
FLT3 AND NOT-STOML1	0.940936864	0.9625	0.920318725
FLT3 AND NOT-SFXN1	0.935550936	0.97826087	0.896414343
FLT3 AND NOT-ORMDL1	0.929460581	0.96969697	0.892430279
FLT3 AND NOT-KCNK6	0.944785276	0.970588235	0.920318725
FLT3 AND NOT-CHST10	0.93877551	0.962343096	0.916334661
FLT3 AND NOT-ABCG1	0.946502058	0.978723404	0.916334661
FLT3 AND NOT-EDEM1	0.9375	0.982532751	0.896414343
FLT3 AND NOT-CD79B	0.909090909	0.995260664	0.836653386
FLT3 AND NOT-RNF144A	0.946502058	0.978723404	0.916334661
FLT3 AND NOT-ARMCX2	0.946502058	0.978723404	0.916334661
FLT3 AND NOT-HEPH	0.948665298	0.978813559	0.920318725
FLT3 AND NOT-C2CD5	0.932489451	0.99103139	0.880478088
FLT3 AND NOT-AREL1	0.933884298	0.969957082	0.900398406
FLT3 AND NOT-SLC35E2	0.940936864	0.9625	0.920318725
FLT3 AND NOT-GOLGA5	0.925053533	1	0.860557769
FLT3 AND NOT-HS3ST2	0.940936864	0.9625	0.920318725
FUT4 AND DENND5A	0.948616601	0.941176471	0.956175299
FUT4 AND NOT-ADGRL2	0.964	0.967871486	0.960159363
FUT4 AND NOT-FMO5	0.964143426	0.964143426	0.964143426
FUT4 AND NOT-DNAJC16	0.9453125	0.927203065	0.964143426
FUT4 AND NOT-SLC35A3	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-ICMT	0.952380952	0.948616601	0.956175299
FUT4 AND NOT-SLC39A14	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-TSPAN12	0.960159363	0.960159363	0.960159363
FUT4 AND NOT-TSPAN15	0.960474308	0.952941176	0.96812749
CXCL8 AND NOT-TSPAN15	0.927102804	0.873239437	0.988047809
SMIM3 AND NOT-TSPAN15	0.947572816	0.924242424	0.972111554
FUT4 AND NOT-VSIG2	0.958579882	0.94921875	0.96812749
FUT4 AND NOT-TMEM2	0.972	0.975903614	0.96812749
FUT4 AND NOT-STX12	0.951219512	0.970954357	0.932270916
FUT4 AND NOT-CADM1	0.940451745	0.970338983	0.912350598
FUT4 AND NOT-EML2	0.951076321	0.934615385	0.96812749
FUT4 AND NOT-FUT2	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-FUT3	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-FUT6	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-HEPACAM2	0.958415842	0.952755906	0.964143426
FUT4 AND NOT-LCLAT1	0.944	0.947791165	0.940239044
FUT4 AND NOT-SLC37A4	0.9498998	0.955645161	0.944223108
FUT4 AND NOT-LPCAT4	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-ZDHHC23	0.948	0.951807229	0.944223108
FUT4 AND NOT-MCOLN2	0.943248532	0.926923077	0.960159363
FUT4 AND NOT-MFSD8	0.964	0.967871486	0.960159363
FUT4 AND NOT-SERINC5	0.967871486	0.975708502	0.960159363
FUT4 AND NOT-TPSG1	0.958415842	0.952755906	0.964143426
FUT4 AND NOT-BACE2	0.9498998	0.955645161	0.944223108
FUT4 AND NOT-PARM1	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-TMEM186	0.95473251	0.987234043	0.924302789
FUT4 AND NOT-ABI3BP	0.966067864	0.968	0.964143426
FUT4 AND NOT-PVRL3	0.958415842	0.952755906	0.964143426
FUT4 AND NOT-HIGD1A	0.966202783	0.964285714	0.96812749

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
TNFRSF8 AND NOT-PCDHB14	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC7A1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-C1orf233	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PTPRT	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PIGB	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-VNN3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-UCN2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-HHLA2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KCNH7	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CACNA1G	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMEM97	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-C2orf83	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GAL3ST4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TNFRSF13C	0.928571429	1	0.866666667
TNFRSF8 AND NOT-NPY4R	0.928571429	1	0.866666667
TNFRSF8 AND NOT-TMEM2	0.928571429	1	0.866666667
TNFRSF8 AND NOT-ERMP1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CHST8	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CCL22	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMEM132B	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-WNT4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMEM62	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-RGS9BP	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ADGRE3	0.965517241	1	0.933333333
TNFRSF8 AND NOT-RMDN3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-B4GALNT4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MLC1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-WNT5A	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PPAPDC1A	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PCDHB11	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-COLEC11	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-VIPR2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SLC30A1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-TMEM257	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ANGPTL1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PRADC1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KLHDC7A	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PCDHB12	0.928571429	1	0.866666667
TNFRSF8 AND NOT-ANG	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC18A2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC35F5	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PLD6	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SDK2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SLC39A8	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MFSD6L	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PI3	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SLC44A4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PTCHD2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GRIA1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CA14	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-C1orf186	0.928571429	1	0.866666667
TNFRSF8 AND NOT-KCNA3	0.928571429	1	0.866666667
TNFRSF8 AND NOT-MAVS	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CLEC2D	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-NDUFC2	0.928571429	1	0.866666667
TNFRSF8 AND NOT-ZNRF3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-LY6K	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KCNJ1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ADRB1	0.928571429	1	0.866666667
TNFRSF8 AND NOT-CACFD1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GLP1R	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-HS3ST3B1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TAS2R9	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TAAR1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-YIPF3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ADSSL1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-LRCH3	0.928571429	1	0.866666667
TNFRSF8 AND NOT-TMEM130	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-XG	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GYPA	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GAL	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FITM1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SMIM6	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC39A1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SYNGR1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-RNF121	0.928571429	1	0.866666667

Antigen Pair Logic	F1	Precision	Recall
FUT4 AND NOT-LRIG1	0.97983871	0.991836735	0.96812749
FUT4 AND NOT-TMEM98	0.951807229	0.95951417	0.944223108
FUT4 AND NOT-GLCE	0.96812749	0.96812749	0.96812749
FUT4 AND NOT-ABHD12	0.946107784	0.948	0.944223108
FUT4 AND NOT-STEAP2	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-SLC17A5	0.968	0.97188755	0.964143426
FUT4 AND NOT-CNNM4	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-GCNT2	0.95951417	0.975308642	0.944223108
FUT4 AND NOT-TBL2	0.955823293	0.963562753	0.948207171
FUT4 AND NOT-STEAP1	0.954274354	0.952380952	0.956175299
FUT4 AND NOT-GHR	0.961923848	0.967741935	0.956175299
FUT4 AND NOT-GJB1	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-GJB2	0.956349206	0.95256917	0.960159363
FUT4 AND NOT-LAMP3	0.934615385	0.903345725	0.96812749
FUT4 AND NOT-GPD2	0.945098039	0.930501931	0.960159363
FUT4 AND NOT-AQP11	0.956692913	0.945525292	0.96812749
FUT4 AND NOT-SLC39A5	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-SLC46A3	0.967741935	0.979591837	0.956175299
FUT4 AND C16orf54	0.937743191	0.91634981	0.960159363
FUT4 AND NOT-ANG	0.967871486	0.975708502	0.960159363
FUT4 AND NOT-CYB561D1	0.93877551	0.962343096	0.916334661
FUT4 AND NOT-SLC25A34	0.954274354	0.952380952	0.956175299
FUT4 AND NOT-CYP4V2	0.953535354	0.967213115	0.940239044
FUT4 AND NOT-DCBLD1	0.962376238	0.956692913	0.96812749
FUT4 AND NOT-YIPF6	0.961923848	0.967741935	0.956175299
FUT4 AND P2RY8	0.937743191	0.91634981	0.960159363
FUT4 AND NOT-TIMM21	0.949698189	0.959349593	0.940239044
FUT4 AND NOT-ORMDL2	0.914529915	0.986175115	0.852589641
FUT4 AND NOT-DNAJC15	0.93877551	0.962343096	0.916334661
FUT4 AND NOT-SLC25A4	0.96812749	0.96812749	0.96812749
FUT4 AND NOT-GUCA2B	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-GUCY2C	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-SLC2A8	0.956349206	0.95256917	0.960159363
FUT4 AND NOT-HAS3	0.934615385	0.903345725	0.96812749
FUT4 AND HBD	0.937984496	0.913207547	0.964143426
FUT4 AND NOT-ST6GALNAC6	0.954635108	0.9453125	0.964143426
FUT4 AND NOT-HMGCR	0.965656566	0.979508197	0.952191235
FUT4 AND NOT-HMOX2	0.961770624	0.971544715	0.952191235
FUT4 AND NOT-HRH1	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-HSD17B2	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-ACACB	0.956	0.959839357	0.952191235
FUT4 AND NOT-ANO9	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-LRRC66	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-SERINC2	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-IHH	0.960474308	0.952941176	0.96812749
FUT4 AND CXCL8	0.958083832	0.96	0.956175299
FUT4 AND NOT-ITGA2	0.958415842	0.952755906	0.964143426
FUT4 AND NOT-ITGA3	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-ITGAV	0.918454936	0.995348837	0.852589641
FUT4 AND NOT-ITGB5	0.95951417	0.975308642	0.944223108
FUT4 AND NOT-ITGB6	0.964285714	0.960474308	0.96812749
FUT4 AND NOT-ITPR3	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-FAM73A	0.962075848	0.964	0.960159363
FUT4 AND NOT-MIA3	0.950690335	0.94140625	0.960159363
FUT4 AND NOT-KCNJ2	0.950207469	0.991341991	0.912350598
FUT4 AND NOT-KCNK1	0.952191235	0.952191235	0.952191235
FUT4 AND NOT-KCNMA1	0.96812749	0.96812749	0.96812749
FUT4 AND NOT-C10orf99	0.960474308	0.952941176	0.96812749
NOT-FAM69A AND FUT4	0.990138067	0.98046875	1
FUT4 AND NOT-SAMD5	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-IYD	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-LETM1	0.9500998	0.952	0.948207171
FUT4 AND NOT-SNX19	0.948412698	0.944664032	0.952191235
FUT4 AND NOT-LMO7	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-PPAPDC2	0.968	0.97188755	0.964143426
FUT4 AND NOT-EPCAM	0.952755906	0.941634241	0.964143426
FUT4 AND NOT-ARSE	0.958579882	0.94921875	0.96812749
FUT4 AND NOT-MEP1A	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-MET	0.952191235	0.952191235	0.952191235
FUT4 AND NOT-MFAP3	0.948616601	0.941176471	0.956175299
FUT4 AND NOT-MGAT5	0.965931864	0.971774194	0.960159363
FUT4 AND NOT-KITLG	0.961923848	0.967741935	0.956175299
FUT4 AND NOT-MID1	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-ALDH6A1	0.944223108	0.944223108	0.944223108
FUT4 AND NOT-TMEM41B	0.940451745	0.970338983	0.912350598
FUT4 AND NOT-SMIM22	0.960474308	0.952941176	0.96812749

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
TNFRSF8 AND NOT-GPER1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KCNH1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MYADML2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ABHD3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ALG1	0.928571429	1	0.866666667
TNFRSF8 AND NOT-SORCS3	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SYS1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-FAM187B	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TAS2R39	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TNMD	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-HSD3B2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ADRB2	0.928571429	1	0.866666667
TNFRSF8 AND NOT-FAM209B	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CTAGE1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FAR1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ARSE	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-LRTM1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KCNE5	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CHRNA4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PCDHGA11	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ATRN	0.965517241	1	0.933333333
TNFRSF8 AND NOT-PANX3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FGFR3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-IL13	0.928571429	1	0.866666667
TNFRSF8 AND NOT-TLR7	0.965517241	1	0.933333333
TNFRSF8 AND NOT-GTF3C3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GALNT15	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FCGR3B	0.928571429	1	0.866666667
TNFRSF8 AND NOT-KLK5	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-OR52A1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-THSD7B	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC2A1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KCNH6	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TTYH2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TSPAN2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-LRRTM1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC30A5	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MPPE1	0.928571429	1	0.866666667
TNFRSF8 AND NOT-GHSR	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-LNPEP	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GABRR1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CLDN15	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-XPR1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PTPRJ	0.965517241	1	0.933333333
TNFRSF8 AND NOT-CYP2C8	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GABBR2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GPRC5D	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-IGSF6	0.965517241	1	0.933333333
TNFRSF8 AND NOT-ACVR2A	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-B4GALNT2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GABRG1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-LY6G6D	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-OR10A3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC3A1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-OR8D2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FAM205A	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GDAP1L1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC29A4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-UGT1A8	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CD207	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ANKH	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC27A3	0.965517241	1	0.933333333
TNFRSF8 AND NOT-MS4A3	0.965517241	1	0.933333333
TNFRSF8 AND NOT-CERS4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MGAT3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MEGF9	0.965517241	1	0.933333333
TNFRSF8 AND NOT-FXYD7	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC38A11	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GPA33	0.965517241	1	0.933333333
TNFRSF8 AND NOT-GALR1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CACNA1B	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC13A2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PARM1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMPRSS12	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC5A3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-RECK	0.965517241	1	0.933333333

Antigen Pair Logic	F1	Precision	Recall
FUT4 AND NOT-MINOS1	0.971544715	0.991701245	0.952191235
FUT4 AND NOT-TMEM150C	0.964143426	0.964143426	0.964143426
FUT4 AND NOT-ASPH	0.960159363	0.960159363	0.960159363
FUT4 AND NOT-MST1R	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-MUC1	0.941883768	0.947580645	0.93625498
FUT4 AND NOT-MUC4	0.958579882	0.94921875	0.96812749
FUT4 AND NOT-NEO1	0.954274354	0.952380952	0.956175299
FUT4 AND NOT-NPY1R	0.962075848	0.964	0.960159363
FUT4 AND NOT-NT5E	0.958250497	0.956349206	0.960159363
P2RX1 AND FUT4	0.936660269	0.903703704	0.972111554
FUT4 AND NOT-NSDHL	0.931392931	0.973913043	0.892430279
FUT4 AND NOT-SLC35B3	0.963562753	0.979423868	0.948207171
FUT4 AND NOT-ZDHHC9	0.958415842	0.952755906	0.964143426
FUT4 AND NOT-INSIG2	0.936170213	0.909774436	0.964143426
FUT4 AND NOT-CRIM1	0.958415842	0.952755906	0.964143426
FUT4 AND NOT-GOLM1	0.955823293	0.963562753	0.948207171
FUT4 AND NOT-BFAR	0.923728814	0.986425339	0.868525896
FUT4 AND NOT-CD320	0.948207171	0.948207171	0.948207171
FUT4 AND NOT-ZDHHC3	0.947368421	0.962962963	0.932270916
FUT4 AND NOT-PDE3A	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-GULP1	0.949494949	0.963114754	0.93625498
FUT4 AND NOT-PDGFRA	0.966067864	0.968	0.964143426
FUT4 AND NOT-LSR	0.956349206	0.95256917	0.960159363
FUT4 AND NOT-ENPP3	0.958415842	0.952755906	0.964143426
FUT4 AND NOT-ACSL5	0.958083832	0.96	0.956175299
FUT4 AND NOT-GALNT7	0.937743191	0.91634981	0.960159363
FUT4 AND NOT-ATP8B1	0.958415842	0.952755906	0.964143426
NOT-PFN2 AND FUT4	0.968627451	0.953667954	0.984063745
FUT4 AND NOT-PIGR	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-PLA2G2A	0.958579882	0.94921875	0.96812749
FUT4 AND NOT-FXYD3	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-PLOD2	0.957746479	0.967479675	0.948207171
FUT4 AND NOT-PLXNA2	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-PMP22	0.960159363	0.960159363	0.960159363
FUT4 AND NOT-C11orf24	0.96	0.963855422	0.956175299
FUT4 AND NOT-CDHR5	0.960474308	0.952941176	0.96812749
NOT-PODXL AND FUT4	0.957528958	0.928838951	0.988047809
FUT4 AND NOT-PON2	0.954274354	0.952380952	0.956175299
FUT4 AND NOT-FAM134B	0.969818913	0.979674797	0.960159363
FUT4 AND NOT-FAM105A	0.965656566	0.979508197	0.952191235
FUT4 AND NOT-ZDHHC13	0.955102041	0.979079498	0.932270916
FUT4 AND NOT-RNF186	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-NDFIP2	0.958415842	0.952755906	0.964143426
FUT4 AND NOT-VSIG10	0.952191235	0.952191235	0.952191235
FUT4 AND NOT-TMEM106B	0.961770624	0.971544715	0.952191235
FUT4 AND NOT-CRLS1	0.95	0.995633188	0.908366534
FUT4 AND NOT-MANSC1	0.958415842	0.952755906	0.964143426
FUT4 AND NOT-GRAMD1C	0.956349206	0.95256917	0.960159363
FUT4 AND NOT-PPIC	0.964143426	0.964143426	0.964143426
FUT4 AND NOT-PAQR5	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-RETSAT	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-RNF43	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-TMEM45A	0.946938776	0.970711297	0.924302789
FUT4 AND NOT-TMEM51	0.952191235	0.952191235	0.952191235
FUT4 AND NOT-ANO10	0.946322068	0.944444444	0.948207171
FUT4 AND NOT-TMEM19	0.965931864	0.971774194	0.960159363
FUT4 AND NOT-TMEM144	0.948	0.951807229	0.944223108
FUT4 AND NOT-ACER3	0.954091816	0.956	0.952191235
FUT4 AND NOT-SYNJ2BP	0.967741935	0.979591837	0.956175299
FUT4 AND NOT-SLC35C1	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-STYK1	0.958415842	0.952755906	0.964143426
FUT4 AND NOT-LGR4	0.954274354	0.952380952	0.956175299
FUT4 AND NOT-SMPD3	0.96031746	0.956521739	0.964143426
FUT4 AND NOT-IL17RB	0.958579882	0.94921875	0.96812749
FUT4 AND NOT-SLC30A6	0.965931864	0.971774194	0.960159363
FUT4 AND NOT-FAR2	0.926931106	0.973684211	0.884462151
FUT4 AND NOT-SAYSD1	0.94921875	0.931034483	0.96812749
FUT4 AND NOT-ST6GALNAC1	0.956349206	0.95256917	0.960159363
FUT4 AND NOT-PAG1	0.956	0.959839357	0.952191235
FUT4 AND NOT-ACSS2	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-ANKH	0.956701031	0.991452991	0.924302789
FUT4 AND NOT-PROS1	0.961616162	0.975409836	0.948207171
FUT4 AND NOT-PRRG1	0.964285714	0.960474308	0.96812749
FUT4 AND NOT-PSEN1	0.94921875	0.931034483	0.96812749
FUT4 AND NOT-TMPRSS4	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-MUC13	0.960474308	0.952941176	0.96812749

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
TNFRSF8 AND NOT-TNFRSF11A	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-S1PR2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GSG1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-P2RY2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-NLGN3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-RNF182	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC38A3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMEM88	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMEM33	0.965517241	1	0.933333333
TNFRSF8 AND NOT-HLA-DOB	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-LRCH1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-XKR6	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KIAA1161	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-RHOT2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TCIRG1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-UPK3B	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-DCST2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GLT8D1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-NDST4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TAS2R13	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TNFRSF4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GOLT1A	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CYP4A11	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CHRNG	0.965517241	1	0.933333333
TNFRSF8 AND NOT-MUC4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ITGB2	0.928571429	1	0.866666667
TNFRSF8 AND NOT-CEP95	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TAAR5	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TXNDC15	0.928571429	1	0.866666667
TNFRSF8 AND NOT-ADTRP	0.928571429	1	0.866666667
TNFRSF8 AND NOT-TNFSF9	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ABCC3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PCDHB5	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CLDN23	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ASIC5	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GABBR1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MUC12	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GABRA6	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-STOML3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PLXND1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PCDHA6	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-LECT1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-LRRN4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-COL17A1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ST8SIA5	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ELOVL2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PAQR8	0.965517241	1	0.933333333
TNFRSF8 AND NOT-NFXL1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KCNC2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-LRRC3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ABCA7	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SPN	0.928571429	1	0.866666667
TNFRSF8 AND NOT-SLC26A11	0.928571429	1	0.866666667
TNFRSF8 AND NOT-TPSG1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-EVC	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SPINT1	0.928571429	1	0.866666667
TNFRSF8 AND NOT-KCNK2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-HCN3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SPATA3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GPR146	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-OR2B6	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC18A1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SPTSSB	0.965517241	1	0.933333333
TNFRSF8 AND NOT-MFSD6	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-RHAG	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SPPL2C	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-NAAA	0.965517241	1	0.933333333
TNFRSF8 AND NOT-RFT1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ADRA1D	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PIRT	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CCR3	0.965517241	1	0.933333333
TNFRSF8 AND NOT-CES5A	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLCO6A1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-COX10	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SEMA3D	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FBXW7	0.965517241	1	0.933333333

Antigen Pair Logic	F1	Precision	Recall
FUT4 AND NOT-AGPAT3	0.93970894	0.982608696	0.900398406
FUT4 AND NOT-PMEPA1	0.964	0.967871486	0.960159363
FUT4 AND NOT-ACKR3	0.91416309	0.990697674	0.848605578
FUT4 AND NOT-LPAR5	0.97188755	0.979757085	0.964143426
FUT4 AND NOT-SLC44A2	0.969325153	0.995798319	0.944223108
FUT4 AND NOT-DOLPP1	0.95049505	0.94488189	0.956175299
FUT4 AND NOT-SLC45A4	0.947580645	0.959183673	0.93625498
FUT4 AND NOT-SMAGP	0.956	0.959839357	0.952191235
FUT4 AND NOT-PTCH1	0.967741935	0.979591837	0.956175299
FUT4 AND NOT-PTGDR	0.968	0.97188755	0.964143426
FUT4 AND NOT-MRS2	0.928571429	0.982222222	0.880478088
FUT4 AND NOT-PTGFRN	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-CYP20A1	0.94302554	0.930232558	0.956175299
FUT4 AND NOT-ABHD6	0.973947896	0.97983871	0.96812749
FUT4 AND NOT-NDRG2	0.964	0.967871486	0.960159363
FUT4 AND NOT-TMCC3	0.950617284	0.982978723	0.920318725
FUT4 AND NOT-HEG1	0.948024948	0.991304348	0.908366534
FUT4 AND NOT-MTUS1	0.97005988	0.972	0.96812749
FUT4 AND NOT-KIAA1324	0.958415842	0.952755906	0.964143426
FUT4 AND NOT-SEMA6A	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-SYT13	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-PTN	0.96812749	0.96812749	0.96812749
FUT4 AND NOT-SLAMF7	0.971544715	0.991701245	0.952191235
FUT4 AND NOT-PTPRD	0.955823293	0.963562753	0.948207171
FUT4 AND NOT-PTPRF	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-PTPRH	0.960474308	0.952941176	0.96812749
NOT-PTPRK AND FUT4	0.952380952	0.912408759	0.996015936
FUT4 AND NOT-PTPRM	0.948240166	0.987068966	0.912350598
FUT4 AND NOT-PTPRO	0.938223938	0.91011236	0.96812749
FUT4 AND NOT-PTPRR	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-IL22RA1	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-ENPP5	0.967871486	0.975708502	0.960159363
FUT4 AND NOT-EVA1C	0.951417004	0.967078189	0.93625498
FUT4 AND NOT-ACE2	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-NTN4	0.97005988	0.972	0.96812749
FUT4 AND NOT-RNASE1	0.955645161	0.967346939	0.944223108
FUT4 AND NOT-RNASE4	0.959016393	0.987341772	0.932270916
FUT4 AND NOT-TNFRSF17	0.965376782	0.9875	0.944223108
FUT4 AND NOT-RRBP1	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-BDKRB2	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-SLC22A23	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-TSPAN31	0.944444444	0.940711462	0.948207171
FUT4 AND NOT-SCNN1A	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-SCNN1G	0.945525292	0.923954373	0.96812749
FUT4 AND NOT-CEACAM1	0.951219512	0.970954357	0.932270916
FUT4 AND NOT-SDC1	0.964285714	0.960474308	0.96812749
FUT4 AND NOT-MCUR1	0.9500998	0.952	0.948207171
FUT4 AND NOT-PERP	0.965794769	0.975609756	0.956175299
FUT4 AND NOT-SMOC2	0.962376238	0.956692913	0.96812749
FUT4 AND NOT-ATL2	0.934615385	0.903345725	0.96812749
FUT4 AND NOT-TMEM168	0.956349206	0.95256917	0.960159363
FUT4 AND NOT-SGCB	0.966067864	0.968	0.964143426
FUT4 AND NOT-ITSN1	0.957403651	0.975206612	0.940239044
FUT4 AND NOT-HRCT1	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-SI	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-CDCP1	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-CASD1	0.952755906	0.941634241	0.964143426
FUT4 AND NOT-LRRC19	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-SLC30A5	0.946322068	0.944444444	0.948207171
FUT4 AND NOT-SLC26A6	0.954274354	0.952380952	0.956175299
FUT4 AND NOT-SLC1A1	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-SLC3A1	0.956349206	0.95256917	0.960159363
FUT4 AND NOT-SLC12A2	0.952755906	0.941634241	0.964143426
FUT4 AND NOT-SLC16A1	0.954274354	0.952380952	0.956175299
FUT4 AND NOT-SLC20A2	0.962376238	0.956692913	0.96812749
FUT4 AND NOT-SLCO2A1	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-BMPR1A	0.950617284	0.982978723	0.920318725
FUT4 AND NOT-SLC22A5	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-GRAMD3	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-BMPR2	0.960824742	0.995726496	0.928286853
FUT4 AND NOT-CYP4F12	0.958415842	0.952755906	0.964143426
FUT4 AND NOT-SMPD1	0.958250497	0.956349206	0.960159363
FUT4 AND NOT-SPINT1	0.954455446	0.948818898	0.960159363
FUT4 AND NOT-SSTR1	0.958579882	0.94921875	0.96812749
FUT4 AND NOT-ST14	0.956	0.959839357	0.952191235
FUT4 AND NOT-TGFA	0.956349206	0.95256917	0.960159363

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
TNFRSF8 AND NOT-C1orf27	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-HCRT	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-NDUFA4L2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ECM1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC6A12	0.965517241	1	0.933333333
TNFRSF8 AND NOT-TMEM189	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-DSCAM	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GPR135	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-VSIG10	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMEM79	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CA9	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GOSR2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-RTN2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMEM121	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GPR63	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-NPBWR2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMEM52B	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KCNJ5	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-UNC93B1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MAS1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MARCH8	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-P2RY6	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PEBP4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SCAMP5	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC12A9	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-REEP2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-MTUS1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-G6PC2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ST8SIA3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ARSB	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MRGPRX4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-UTY	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC36A4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GPR137	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KCNJ12	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-HHATL	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ADGRG7	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-UPK1A	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC25A2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-DCBLD2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-OPRK1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC15A3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GPR31	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SSTR5	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FAM9C	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-LINGO3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-AASDH	0.965517241	1	0.933333333
TNFRSF8 AND NOT-GPR12	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FIBCD1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CD40LG	0.965517241	1	0.933333333
TNFRSF8 AND NOT-TIMM50	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TRAF3IP3	0.965517241	1	0.933333333
TNFRSF8 AND NOT-TAS1R2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-C9orf91	0.965517241	1	0.933333333
TNFRSF8 AND NOT-NDST3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MIEF2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-EPHB6	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KCNA7	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-B3GALT2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ADAM20	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SMIM8	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MGAT4A	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-IL1RAPL2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CACHD1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMBIM1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-BEST1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MUC15	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MALRD1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-AMIGO2	0.928571429	1	0.866666667
TNFRSF8 AND NOT-PKD1L2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GHRHR	0.928571429	1	0.866666667
TNFRSF8 AND NOT-CD5	0.965517241	1	0.933333333
TNFRSF8 AND NOT-KCNK7	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SMIM2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SELP	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMEM125	0.933333333	0.93333333	0.933333333

Antigen Pair Logic	F1	Precision	Recall
FUT4 AND NOT-TLR3	0.954274354	0.952380952	0.956175299
FUT4 AND NOT-TSPAN8	0.962376238	0.956692913	0.96812749
FUT4 AND NOT-TSPAN6	0.966202783	0.964285714	0.96812749
FUT4 AND NOT-TMPRSS2	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-TLCD2	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-UGT2B15	0.94368932	0.920454545	0.96812749
FUT4 AND NOT-UGT2B17	0.94368932	0.920454545	0.96812749
FUT4 AND NOT-UGT8	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-VIPR1	0.953907816	0.959677419	0.948207171
FUT4 AND NOT-WNT5A	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-CA4	0.955645161	0.967346939	0.944223108
FUT4 AND NOT-CA12	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-DDR1	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-FZD5	0.956175299	0.956175299	0.956175299
FUT4 AND NOT-SLMAP	0.9375	0.982532751	0.896414343
FUT4 AND NOT-ELOVL6	0.956349206	0.95256917	0.960159363
FUT4 AND NOT-FA2H	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-GDPD3	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-DHRS11	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-MFSD11	0.94214876	0.978540773	0.908366534
FUT4 AND NOT-RNF128	0.962376238	0.956692913	0.96812749
FUT4 AND NOT-C3orf52	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-GAL3ST4	0.934156379	0.965957447	0.90438247
FUT4 AND NOT-GALNT12	0.943548387	0.955102041	0.932270916
FUT4 AND NOT-CCDC51	0.948665298	0.978813559	0.920318725
FUT4 AND NOT-C1orf115	0.964143426	0.964143426	0.964143426
FUT4 AND NOT-TXNDC15	0.921775899	0.981981982	0.868525896
FUT4 AND NOT-CLMN	0.953346856	0.97107438	0.93625498
FUT4 AND NOT-C12orf49	0.96993988	0.975806452	0.964143426
FUT4 AND NOT-UGT2A3	0.956349206	0.95256917	0.960159363
FUT4 AND NOT-TMC5	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-THSD4	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-BTNL8	0.952755906	0.941634241	0.964143426
FUT4 AND NOT-DNAJC22	0.960474308	0.952941176	0.96812749
NOT-WLS AND FUT4	0.96222664	0.96031746	0.964143426
FUT4 AND NOT-CPED1	0.943632568	0.99122807	0.900398406
FUT4 AND NOT-ELOVL7	0.961770624	0.971544715	0.952191235
FUT4 AND NOT-TMEM62	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-SEMA6D	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-CWH43	0.958579882	0.94921875	0.96812749
FUT4 AND NOT-TMEM254	0.954274354	0.952380952	0.956175299
FUT4 AND NOT-PIGZ	0.956521739	0.949019608	0.964143426
FUT4 AND NOT-SLC35F5	0.957230143	0.979166667	0.93625498
FUT4 AND NOT-SLC44A4	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-CYB5B	0.964	0.967871486	0.960159363
FUT4 AND NOT-SSPN	0.953722334	0.963414634	0.944223108
FUT4 AND NOT-SLC2A10	0.962376238	0.956692913	0.96812749
FUT4 AND NOT-SGPP1	0.901960784	0.995192308	0.824701195
FUT4 AND NOT-YIPF5	0.973843058	0.983739837	0.964143426
FUT4 AND NOT-VWA9	0.945674044	0.955284553	0.93625498
FUT4 AND NOT-AKAP1	0.96031746	0.956521739	0.964143426
FUT4 AND NOT-RNF170	0.942386831	0.974468085	0.912350598
FUT4 AND NOT-GLT8D2	0.96812749	0.96812749	0.96812749
FUT4 AND NOT-TMEM47	0.96812749	0.96812749	0.96812749
FUT4 AND NOT-MS4A8	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-TMEM133	0.96812749	0.96812749	0.96812749
FUT4 AND NOT-PLA2G10	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-SLC41A2	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-ZNRF3	0.956349206	0.95256917	0.960159363
FUT4 AND NOT-TMEM117	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-RHBDD1	0.936416185	0.906716418	0.96812749
FUT4 AND NOT-SLC37A3	0.963414634	0.98340249	0.944223108
FUT4 AND NOT-YIPF4	0.968	0.97188755	0.964143426
FUT4 AND NOT-FAM213A	0.959839357	0.967611336	0.952191235
FUT4 AND NOT-TMEM246	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-SLC12A8	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-HIATL1	0.946322068	0.944444444	0.948207171
FUT4 AND NOT-MFSD9	0.958415842	0.952755906	0.964143426
FUT4 AND NOT-ADTRP	0.968	0.97188755	0.964143426
FUT4 AND NOT-AIFM2	0.958415842	0.952755906	0.964143426
FUT4 AND NOT-TMEM87B	0.942857143	0.966527197	0.920318725
FUT4 AND NOT-LRP11	0.9437751	0.951417004	0.93625498
FUT4 AND NOT-PAQR8	0.961460446	0.979338843	0.944223108
FUT4 AND NOT-CDC14B	0.975806452	0.987755102	0.964143426
FUT4 AND NOT-PPAP2A	0.941176471	0.958677686	0.924302789
FUT4 AND NOT-PPAP2C	0.960474308	0.952941176	0.96812749

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
TNFRSF8 AND NOT-LY6G6F	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-RPS23	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CD1C	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SLC22A11	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CHRNB4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-EPHA10	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GABRA5	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PCDHGA9	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC6A1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-COL25A1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC41A1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-HMGCR	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ZDHHC21	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ADAM8	0.965517241	1	0.933333333
TNFRSF8 AND NOT-VANGL2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-P2RY12	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CDRT15L2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-NSG1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TSPAN10	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-NIPA1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-S1PR5	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SCN3B	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC29A1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SUSD3	0.928571429	1	0.866666667
TNFRSF8 AND NOT-GLRA3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ITPR3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ACHE	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-OR4N4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CD300LB	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SIGLEC7	0.928571429	1	0.866666667
TNFRSF8 AND NOT-RIC3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ERMAP	0.965517241	1	0.933333333
TNFRSF8 AND NOT-ADGRG4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PCDH11X	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KRT5	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-HTR2A	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FREM1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-LVRN	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GLTPD2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FXYD1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-TMCC1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TNFRSF1B	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SLC2A12	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CAMKMT	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PGAP3	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SLC22A25	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-LRP8	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ST7L	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-NUP210	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMPRSS13	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PCDHA2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TRPC3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-IL22RA1	0.928571429	1	0.866666667
TNFRSF8 AND NOT-PHTF1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PSEN1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-NEMP2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC13A5	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PTPRC	0.965517241	1	0.933333333
TNFRSF8 AND NOT-OS9	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-OCLM	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ABCA3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-AIFM1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-RARRES3	0.965517241	1	0.933333333
TNFRSF8 AND NOT-TMEM86B	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-NELL2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-RNF167	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PCDHA10	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SV2C	0.965517241	1	0.933333333
TNFRSF8 AND NOT-CACNG8	0.928571429	1	0.866666667
TNFRSF8 AND NOT-PCDHA9	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-HCRTR1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CLPTM1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-DPM2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-RPRML	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KIAA1549L	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CYP3A7	0.933333333	0.93333333	0.933333333

Antigen Pair Logic	F1	Precision	Recall
FUT4 AND NOT-AOC3	0.966067864	0.968	0.964143426
FUT4 AND NOT-SLC4A4	0.956349206	0.95256917	0.960159363
FUT4 AND NOT-B4GALT4	0.957915832	0.963709677	0.952191235
FUT4 AND NOT-B3GALT4	0.948665298	0.978813559	0.920318725
FUT4 AND NOT-ABCC3	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-TNFSF10	0.937759336	0.978354978	0.900398406
FUT4 AND NOT-TNFRSF11A	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-PEX11B	0.965931864	0.971774194	0.960159363
FUT4 AND NOT-PEX11A	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-SGPL1	0.97188755	0.979757085	0.964143426
FUT4 AND NOT-SGCE	0.962075848	0.964	0.960159363
FUT4 AND NOT-P4HA2	0.966202783	0.964285714	0.96812749
FUT4 AND NOT-GLB1L2	0.954274354	0.952380952	0.956175299
FUT4 AND NOT-PHLDB2	0.975708502	0.991769547	0.960159363
FUT4 AND NOT-MPZL1	0.958415842	0.952755906	0.964143426
FUT4 AND NOT-TM4SF5	0.958579882	0.94921875	0.96812749
FUT4 AND NOT-SEMA5A	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-SPAG9	0.949290061	0.966942149	0.932270916
FUT4 AND NOT-TMEM263	0.935817805	0.974137931	0.900398406
FUT4 AND NOT-GPRC5A	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-MCU	0.954091816	0.956	0.952191235
FUT4 AND NOT-SLC13A2	0.940038685	0.913533835	0.96812749
FUT4 AND NOT-ANGPTL1	0.964285714	0.960474308	0.96812749
FUT4 AND NOT-CLDN8	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-PKDCC	0.96031746	0.956521739	0.964143426
FUT4 AND NOT-SLC28A2	0.938223938	0.91011236	0.96812749
FUT4 AND NOT-PCSK7	0.967741935	0.979591837	0.956175299
FUT4 AND NOT-MARVELD3	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-SLC33A1	0.95481336	0.941860465	0.96812749
FUT4 AND NOT-ACVR1B	0.959839357	0.967611336	0.952191235
FUT4 AND NOT-LARGE	0.938223938	0.91011236	0.96812749
FUT4 AND NOT-VAPB	0.973947896	0.97983871	0.96812749
FUT4 AND NOT-TMEM129	0.959839357	0.967611336	0.952191235
FUT4 AND NOT-CRB3	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-PXYLP1	0.948240166	0.987068966	0.912350598
FUT4 AND NOT-GCNT3	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-ACVR2A	0.967741935	0.979591837	0.956175299
FUT4 AND NOT-PGAP3	0.964	0.967871486	0.960159363
FUT4 AND NOT-SDR42E1	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-KL	0.959677419	0.971428571	0.948207171
FUT4 AND NOT-CD27	0.929166667	0.973799127	0.888446215
FUT4 AND NOT-ORMDL3	0.939958592	0.978448276	0.90438247
FUT4 AND NOT-KCNK6	0.959016393	0.987341772	0.932270916
FUT4 AND NOT-ACVRL1	0.958579882	0.94921875	0.96812749
FUT4 AND NOT-SCARB2	0.941649899	0.951219512	0.932270916
FUT4 AND NOT-GOSR1	0.946280992	0.982832618	0.912350598
FUT4 AND NOT-CXCL14	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-ENTPD5	0.960474308	0.952941176	0.96812749
FUT4 AND NOT-CD151	0.948	0.951807229	0.944223108
FUT4 AND NOT-HEPH	0.954274354	0.952380952	0.956175299
FUT4 AND NOT-ATP2C2	0.954091816	0.956	0.952191235
FUT4 AND NOT-MFN2	0.954918033	0.983122363	0.928286853
FUT4 AND NOT-CDH1	0.960474308	0.952941176	0.96812749
P2RX1 AND NOT-MFSD8	0.929791271	0.887681159	0.976095618
SMIM3 AND NOT-PARM1	0.960629921	0.949416342	0.972111554
NOT-FAM69A AND NOT-ABI3BP	0.956022945	0.919117647	0.996015936
NOT-PFN2 AND NOT-NCR3	0.937853107	0.889285714	0.992031873
NOT-LRIG1 AND CXCL8	0.946360153	0.911439114	0.984063745
NRROS AND NOT-LRIG1	0.940936864	0.9625	0.920318725
P2RX1 AND NOT-LRIG1	0.962524655	0.953125	0.972111554
NOT-LRIG1 AND UBE2J1	0.921933086	0.864111498	0.988047809
NOT-LRIG1 AND FXYD5	0.930581614	0.879432624	0.988047809
NOT-LRIG1 AND SUSD1	0.946322068	0.944444444	0.948207171
NOT-LRIG1 AND SMIM3	0.953846154	0.921933086	0.988047809
SMIM3 AND NOT-STEAP1	0.946954813	0.934108527	0.960159363
NOT-GHR AND SMIM3	0.961089494	0.939163498	0.984063745
P2RY8 AND NOT-SIT1	0.940726577	0.904411765	0.980079681
HBD AND NOT-SIT1	0.941860465	0.916981132	0.96812749
NOT-PFN2 AND NOT-SIT1	0.966861598	0.946564885	0.988047809
NOT-PFN2 AND NOT-P2RY10	0.953125	0.9348659	0.972111554
NOT-MINOS1 AND TOR2A	0.93957115	0.919847328	0.960159363
NOT-VAPB AND TOR2A	0.935849057	0.888888889	0.988047809
NOT-PFN2 AND NOT-SLC46A3	0.9348659	0.900369004	0.972111554
NOT-FAM69A AND C16orf54	0.972762646	0.950570342	0.996015936
SMIM3 AND NOT-ANG	0.936170213	0.909774436	0.964143426
NOT-MTUS1 AND NOT-CYB561D1	0.936660269	0.903703704	0.972111554

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
TNFRSF8 AND NOT-SLC43A2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-KCNN2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMEM35	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC9B1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-HAS3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PTPN5	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GJD4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PIGN	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CHIC1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MEGF6	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-HSD17B2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-OR2B3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ACSL6	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SPAST	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GPR37	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMEM63A	0.928571429	1	0.866666667
TNFRSF8 AND NOT-GPR180	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PLA2G2E	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CACNG3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ROMO1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PDZD8	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC18B1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-TGFBR1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-FAIM2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ACSS3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLITRK5	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-XKRX	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TNFRSF10C	0.928571429	1	0.866666667
TNFRSF8 AND NOT-MARCH3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GPATCH2L	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-WNT6	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-IFNE	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KCNN3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMUB2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MSR1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CLCN4	0.965517241	1	0.933333333
TNFRSF8 AND NOT-LRIT1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CLMN	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SEMA4B	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-HCN4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-IGSF3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-IL18R1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-DAGLB	0.928571429	1	0.866666667
TNFRSF8 AND NOT-KIAA1549	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC12A7	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TIMM21	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FCRLA	0.928571429	1	0.866666667
TNFRSF8 AND NOT-SYNDIG1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-DRD4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SEC61A2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-NOX1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-VAMP5	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KCNH5	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CLDN2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ALG10	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ARSF	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC22A12	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ACSS1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ICAM4	0.928571429	1	0.866666667
TNFRSF8 AND NOT-PTDSS2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SPTLC2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-CDH7	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-APOB	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CRY2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TSPO	0.928571429	1	0.866666667
TNFRSF8 AND NOT-BMP10	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SCNN1B	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CD180	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-C12orf73	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CDHR1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TSPAN16	0.965517241	1	0.933333333
TNFRSF8 AND NOT-ADGRB1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SMO	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FAM210A	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SCAMP4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GPR3	0.933333333	0.93333333	0.933333333

Antigen Pair Logic	F1	Precision	Recall
NOT-DCBLD1 AND IL1RAP	0.919148936	0.98630137	0.860557769
SMIM3 AND NOT-SCARA5	0.964426877	0.956862745	0.972111554
P2RY8 AND NOT-HLA-DOB	0.944550669	0.908088235	0.984063745
P2RY8 AND NOT-ITPR3	0.933837429	0.888489209	0.984063745
NOT-FAM69A AND P2RY8	0.968627451	0.953667954	0.984063745
P2RY8 AND NOT-SLC44A2	0.941176471	0.926640927	0.956175299
P2RY8 AND NOT-ADAM19	0.93129771	0.893772894	0.972111554
NOT-PHLDB2 AND P2RY8	0.933078394	0.897058824	0.972111554
P2RY8 AND NOT-FCMR	0.930320151	0.882142857	0.984063745
NOT-PFN2 AND NOT-CLEC2D	0.936090226	0.886120996	0.992031873
NOT-SLC25A4 AND CXCL8	0.921933086	0.864111498	0.988047809
NOT-SLC25A4 AND SMIM3	0.933586338	0.891304348	0.980079681
NOT-PHLDB2 AND SEC61A1	0.946564885	0.908424908	0.988047809
HBD AND NOT-HLA-DOB	0.938223938	0.91011236	0.96812749
HBD AND NOT-INPP4A	0.934615385	0.903345725	0.96812749
HBD AND NOT-KLRD1	0.934615385	0.903345725	0.96812749
HBD AND NOT-LY9	0.936416185	0.906716418	0.96812749
HBD AND NOT-MGAT5	0.942574257	0.937007874	0.948207171
HBD AND NOT-INSIG2	0.936170213	0.909774436	0.964143426
HBD AND NOT-TUBD1	0.94368932	0.920454545	0.96812749
HBD AND NOT-TMEM140	0.942800789	0.93359375	0.952191235
HBD AND NOT-SLC44A2	0.948207171	0.948207171	0.948207171
HBD AND NOT-PTCH1	0.937743191	0.91634981	0.960159363
HBD AND NOT-PTGDR	0.941634241	0.920152091	0.964143426
HBD AND NOT-PTPRO	0.936416185	0.906716418	0.96812749
HBD AND NOT-PVRIG	0.939805825	0.916666667	0.964143426
HBD AND NOT-RHBDD1	0.934362934	0.906367041	0.964143426
HBD AND NOT-ADTRP	0.936170213	0.909774436	0.964143426
HBD AND NOT-KMO	0.934615385	0.903345725	0.96812749
HBD AND NOT-SLC16A7	0.937743191	0.91634981	0.960159363
HBD AND NOT-FCMR	0.941860465	0.916981132	0.96812749
HBD AND NOT-CD27	0.937743191	0.91634981	0.960159363
SMIM3 AND NOT-ST6GALNAC6	0.947368421	0.927480916	0.96812749
NOT-FAM69A AND ADGRE2	0.967479675	0.987551867	0.948207171
NOT-FAM134B AND ADGRE2	0.940206186	0.974358974	0.908366534
ADGRE2 AND NOT-MAN1C1	0.907526882	0.985981308	0.84063745
ADGRE2 AND NOT-SLC44A2	0.924686192	0.973568282	0.880478088
NOT-HEG1 AND ADGRE2	0.918454936	0.995348837	0.852589641
NOT-TGFBR3 AND ADGRE2	0.91416309	0.990697674	0.848605578
NOT-SLC37A3 AND ADGRE2	0.945454545	0.959016393	0.932270916
NOT-PHLDB2 AND ADGRE2	0.9500998	0.952	0.948207171
NOT-PFN2 AND NOT-HLA-DOB	0.966990291	0.943181818	0.992031873
NOT-MTUS1 AND NOT-HLA-DOB	0.970873786	0.946969697	0.996015936
SMIM3 AND NOT-ACACB	0.947162427	0.930769231	0.964143426
NOT-FAM69A AND ICAM3	0.947169811	0.899641577	1
NOT-PHLDB2 AND ICAM3	0.932330827	0.882562278	0.988047809
SMIM3 AND NOT-TMEM119	0.94368932	0.920454545	0.96812749
NRROS AND NOT-AMIGO2	0.944558522	0.974576271	0.916334661
P2RX1 AND NOT-AMIGO2	0.940451745	0.970338983	0.912350598
NOT-HEG1 AND IL1RAP	0.911447084	0.995283019	0.84063745
NOT-PHLDB2 AND IL1RAP	0.943319838	0.958847737	0.928286853
NRROS AND NOT-IL7R	0.941176471	0.958677686	0.924302789
NOT-PFN2 AND NOT-IL7R	0.935361217	0.894545455	0.980079681
CXCL8 AND NOT-ITPR3	0.927102804	0.873239437	0.988047809
CXCL8 AND NOT-KDR	0.923364486	0.86971831	0.984063745
P2RX1 AND CXCL8	0.936170213	0.909774436	0.964143426
CXCL8 AND NOT-FXYD1	0.923649907	0.867132867	0.988047809
CXCL8 AND NOT-TMTC1	0.938223938	0.91011236	0.96812749
SMIM3 AND CXCL8	0.93957115	0.919847328	0.960159363
CXCL8 AND NOT-CDC14B	0.930056711	0.884892086	0.980079681
CXCL8 AND NOT-ST3GAL5	0.938223938	0.91011236	0.96812749
CXCL8 AND NOT-SLC16A4	0.937618147	0.892086331	0.988047809
NOT-ARMCX2 AND CXCL8	0.942748092	0.904761905	0.984063745
NOT-PFN2 AND NOT-INPP4A	0.922222222	0.861591696	0.992031873
NOT-FAM69A AND ITGA4	0.948240166	0.987068966	0.912350598
ITGA4 AND NOT-SLC44A2	0.926931106	0.973684211	0.884462151
NOT-FAM69A AND ITGB2	0.91773309	0.847972973	1
NOT-ITGB5 AND MGST1	0.929166667	0.973799127	0.888446215
NOT-ITGB5 AND SMIM3	0.947368421	0.962962963	0.932270916
P2RX1 AND NOT-ITPR3	0.945945946	0.917602996	0.976095618
NOT-PFN2 AND NOT-ITPR3	0.941398866	0.895683453	0.992031873
FXYD5 AND NOT-ITPR3	0.926199262	0.862542955	1
P2RX1 AND NOT-FAM73A	0.935114504	0.897435897	0.976095618
NRROS AND NOT-FAM69A	0.959016393	0.987341772	0.932270916
NRROS AND NOT-LY9	0.941649899	0.951219512	0.932270916
NRROS AND NOT-MINOS1	0.93877551	0.962343096	0.916334661

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
TNFRSF8 AND NOT-ABCC6	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CLEC3A	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC9B2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC13A3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-POMK	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-NPHS2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PEX2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-LEPROT	0.965517241	1	0.933333333
TNFRSF8 AND NOT-CYP2U1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC1A2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SPTSSA	0.965517241	1	0.933333333
TNFRSF8 AND NOT-ITGA9	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLITRK1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-BNIP2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-CHST12	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC39A11	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMEM8B	0.965517241	1	0.933333333
TNFRSF8 AND NOT-FAT2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PARP16	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC1A1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SFTPC	0.965517241	1	0.933333333
TNFRSF8 AND NOT-EFNB3	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SYT2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CMTM1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GGT7	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-HERPUD2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-TOR1AIP2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-BSND	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CYSLTR2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC12A1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PIANP	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-OR2L2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-APOL2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-IFNL2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GUCY2D	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-IFITM10	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FANCM	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-STX1B	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PCNXL2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FAM73A	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-OR51I1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-PMEL	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-EI24	0.965517241	1	0.933333333
TNFRSF8 AND NOT-TMEM25	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-C16orf92	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMEM255B	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GPD2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-OR6B1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-OR14J1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC9A8	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMEM74	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GPR50	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GJA10	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-LDLRAD4	0.965517241	1	0.933333333
TNFRSF8 AND NOT-NAALADL1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PIGR	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-DNAJC14	0.928571429	1	0.866666667
TNFRSF8 AND NOT-GPRC6A	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-HCAR1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-IGSF1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SHISA2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CX3CR1	0.928571429	1	0.866666667
TNFRSF8 AND NOT-TAAR2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-IL20RB	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-DPY19L3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ITGB7	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC12A6	0.965517241	1	0.933333333
TNFRSF8 AND NOT-C2CD2L	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMEM170A	0.965517241	1	0.933333333
TNFRSF8 AND NOT-TREM1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-TAPT1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-NGFR	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-DRD1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MS4A6E	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CXCL8	0.928571429	1	0.866666667
TNFRSF8 AND NOT-LHFPL1	0.933333333	0.93333333	0.933333333

Antigen Pair Logic	F1	Precision	Recall
NRROS AND NOT-IL21R	0.941176471	0.958677686	0.924302789
NRROS AND NOT-FAM134B	0.931392931	0.973913043	0.892430279
NRROS AND NOT-HEG1	0.948240166	0.987068966	0.912350598
NRROS AND NOT-SLAMF7	0.940936864	0.9625	0.920318725
NRROS AND NOT-PVRIG	0.945233266	0.962809917	0.928286853
NRROS AND NOT-ST3GAL5	0.944329897	0.978632479	0.912350598
NRROS AND NOT-PHLDB2	0.941176471	0.958677686	0.924302789
NRROS AND NOT-KIAA0040	0.945454545	0.959016393	0.932270916
NRROS AND NOT-CD72	0.943089431	0.962655602	0.924302789
SMIM3 AND NOT-KDR	0.96812749	0.96812749	0.96812749
NOT-PFN2 AND KIT	0.941176471	0.995555556	0.892430279
NOT-PFN2 AND NOT-KLRD1	0.934333959	0.882978723	0.992031873
NOT-FAM69A AND LAIR1	0.949698189	0.959349593	0.940239044
NOT-FAM69A AND NOT-LIFR	0.925650558	0.867595819	0.992031873
NOT-FAM69A AND HACD4	0.925925926	0.865051903	0.996015936
NOT-FAM69A AND LRMP	0.944761905	0.905109489	0.988047809
NOT-FAM69A AND NOT-MAOB	0.943396226	0.896057348	0.996015936
NOT-FAM69A AND NOT-MGP	0.96484375	0.946360153	0.984063745
NOT-FAM69A AND NOT-MID1	0.943609023	0.893238434	1
NOT-FAM69A AND NKG7	0.946534653	0.940944882	0.952191235
NOT-FAM69A AND P2RX1	0.986138614	0.980314961	0.992031873
NOT-FAM69A AND CHST11	0.952	0.955823293	0.948207171
NOT-FAM69A AND TMED5	0.916058394	0.845117845	1
NOT-FAM69A AND UBE2J1	0.921100917	0.853741497	1
NOT-FAM69A AND NOT-PDGFRA	0.959692898	0.925925926	0.996015936
NOT-FAM69A AND NOT-PFN2	0.97265625	0.954022989	0.992031873
NOT-FAM69A AND FXYD5	0.940074906	0.886925795	1
NOT-FAM69A AND ADPRM	0.961089494	0.939163498	0.984063745
NOT-FAM69A AND PTGER2	0.946058091	0.987012987	0.908366534
NOT-FAM69A AND NOT-NDRG2	0.943396226	0.896057348	0.996015936
NOT-FAM69A AND NOT-MTUS1	0.961538462	0.92936803	0.996015936
NOT-FAM69A AND NOT-PTN	0.947169811	0.899641577	1
NOT-FAM69A AND PTPRC	0.942622951	0.970464135	0.916334661
NOT-FAM69A AND NOT-PTPRK	0.940074906	0.886925795	1
NOT-FAM69A AND NOT-NTN4	0.922794118	0.85665529	1
NOT-FAM69A AND SELL	0.94980695	0.921348315	0.980079681
NOT-FAM69A AND NOT-PERP	0.934086629	0.885714286	0.988047809
NOT-FAM69A AND SLC39A8	0.95481336	0.941860465	0.96812749
NOT-FAM69A AND SUSD1	0.96	0.963855422	0.956175299
NOT-FAM69A AND VAMP7	0.919413919	0.850847458	1
NOT-FAM69A AND NOT-TSPAN6	0.934579439	0.88028169	0.996015936
NOT-FAM69A AND VRK1	0.936567164	0.880701754	1
NOT-FAM69A AND SLC30A1	0.936567164	0.880701754	1
NOT-FAM69A AND LAPTM5	0.947169811	0.899641577	1
NOT-FAM69A AND NOT-DDR1	0.953488372	0.928301887	0.980079681
NOT-FAM69A AND DERL1	0.921100917	0.853741497	1
NOT-FAM69A AND NOT-RNF128	0.92962963	0.868512111	1
NOT-FAM69A AND NOT-C1orf115	0.925650558	0.867595819	0.992031873
NOT-FAM69A AND ATP8B4	0.95481336	0.941860465	0.96812749
NOT-FAM69A AND NOT-SSPN	0.929791271	0.887681159	0.976095618
NOT-FAM69A AND NOT-GLT8D2	0.933085502	0.87456446	1
NOT-FAM69A AND NOT-TMEM47	0.974757282	0.950757576	1
NOT-FAM69A AND USP48	0.938317757	0.883802817	1
NOT-FAM69A AND C9orf89	0.932835821	0.877192982	0.996015936
NOT-FAM69A AND SMIM3	0.934823091	0.877622378	1
NOT-FAM69A AND SLC22A16	0.953125	0.9348659	0.972111554
NOT-FAM69A AND NOT-SGCE	0.936090226	0.886120996	0.992031873
NOT-FAM69A AND NOT-PHLDB2	0.942965779	0.901818182	0.988047809
NOT-FAM69A AND MYADM	0.933586338	0.891304348	0.980079681
NOT-FAM69A AND CD33	0.96031746	0.956521739	0.964143426
NOT-FAM69A AND CD37	0.95703125	0.938697318	0.976095618
NOT-FAM69A AND CD44	0.944971537	0.902173913	0.992031873
NOT-FAM69A AND CD53	0.943181818	0.898916968	0.992031873
NOT-FAM69A AND CD69	0.934333959	0.882978723	0.992031873
NOT-FAM69A AND ADGRE5	0.947169811	0.899641577	1
NOT-FAM69A AND PTDSS1	0.937618147	0.892086331	0.988047809
NOT-MINOS1 AND HACD4	0.95049505	0.94488189	0.956175299
NOT-SLC37A3 AND HACD4	0.939334638	0.923076923	0.956175299
NOT-VAPB AND HACD4	0.946153846	0.914498141	0.980079681
SMIM3 AND NOT-LRP1	0.949698189	0.959349593	0.940239044
NOT-PFN2 AND NOT-LTB	0.941649899	0.951219512	0.932270916
NOT-PHLDB2 AND LTBR	0.937743191	0.91634981	0.960159363
NOT-MAOB AND HACD1	0.942574257	0.937007874	0.948207171
SMIM3 AND NOT-MCAM	0.95481336	0.941860465	0.96812749
NOT-PFN2 AND CD46	0.926553672	0.878571429	0.980079681
SUSD1 AND NOT-MGAT5	0.919831224	0.977578475	0.868525896

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
TNFRSF8 AND NOT-OR5I1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ITGAM	0.965517241	1	0.933333333
TNFRSF8 AND NOT-ANO10	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC7A4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-VIPR1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-TMPRSS6	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-OR10H3	0.928571429	1	0.866666667
TNFRSF8 AND NOT-PTCHD4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-C6orf89	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-STX7	0.965517241	1	0.933333333
TNFRSF8 AND NOT-APLF	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FBN2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-GPR17	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GDAP1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CEPT1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-ERLIN1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-DRAXIN	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GALNT3	0.928571429	1	0.866666667
TNFRSF8 AND NOT-TLR6	0.965517241	1	0.933333333
TNFRSF8 AND NOT-CLEC1B	0.965517241	1	0.933333333
TNFRSF8 AND NOT-DNAJC16	0.965517241	1	0.933333333
TNFRSF8 AND NOT-RHBG	0.965517241	1	0.933333333
TNFRSF8 AND NOT-FAM105A	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CYP4V2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-FRRS1L	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CATSPER1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-GABRQ	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC7A10	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KCNA5	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SFXN3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-B4GALT3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-RNF43	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CLDND1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CFAP61	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SHISA4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CATSPERB	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GAPT	0.965517241	1	0.933333333
TNFRSF8 AND NOT-PPAPDC2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PNPLA7	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GNPTAB	0.965517241	1	0.933333333
TNFRSF8 AND NOT-CAV3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-RHBDL3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-101805491?	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GRID1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PLIN1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MCHR1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC22A1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MOSPD1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-LTB	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SLC6A2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ARMCX2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ST6GALNAC5	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC25A45	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ITGA2B	0.965517241	1	0.933333333
TNFRSF8 AND NOT-TMEM159	0.965517241	1	0.933333333
TNFRSF8 AND NOT-CTXN3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GRM7	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CFTR	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SPINK6	0.928571429	1	0.866666667
TNFRSF8 AND NOT-TLR9	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SYT1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CD101	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MFSD11	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-WBSCR17	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SYT6	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-LRMP	0.965517241	1	0.933333333
TNFRSF8 AND NOT-CLN5	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KLK4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CEACAM3	0.965517241	1	0.933333333
TNFRSF8 AND NOT-CHRNA1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-OR51B4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PCDHGA1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-COQ7	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TRPV6	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CHRNB1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-NMBR	0.933333333	0.93333333	0.933333333

Antigen Pair Logic	F1	Precision	Recall
CD33 AND NOT-MGAT5	0.929166667	0.973799127	0.888446215
P2RX1 AND NOT-MINOS1	0.954455446	0.948818898	0.960159363
NOT-MINOS1 AND PTGER4	0.94302554	0.930232558	0.956175299
NOT-MINOS1 AND SUSD1	0.955823293	0.963562753	0.948207171
NOT-MINOS1 AND TMEM183A	0.947368421	0.927480916	0.96812749
SMIM3 AND NOT-ASPH	0.960474308	0.952941176	0.96812749
SMIM3 AND NOT-NPY1R	0.954635108	0.9453125	0.964143426
P2RX1 AND NOT-NT5E	0.950884086	0.937984496	0.964143426
SMIM3 AND NOT-DDR2	0.966336634	0.960629921	0.972111554
SMIM3 AND NOT-GPX8	0.947572816	0.924242424	0.972111554
P2RX1 AND NOT-INSIG2	0.9348659	0.900369004	0.972111554
P2RX1 AND NOT-TUBD1	0.936902486	0.900735294	0.976095618
NOT-PFN2 AND P2RX1	0.923364486	0.86971831	0.984063745
P2RX1 AND NOT-FAM134B	0.947791165	0.955465587	0.940239044
P2RX1 AND NOT-SYNJ2BP	0.94302554	0.930232558	0.956175299
P2RX1 AND NOT-ANKH	0.942800789	0.93359375	0.952191235
P2RX1 AND NOT-LPAR5	0.92952381	0.890510949	0.972111554
P2RX1 AND NOT-PLXDC1	0.931558935	0.890909091	0.976095618
P2RX1 AND NOT-SLC44A2	0.947368421	0.927480916	0.96812749
P2RX1 AND NOT-PTCH1	0.941634241	0.920152091	0.964143426
P2RX1 AND NOT-HEG1	0.973737374	0.987704918	0.960159363
NOT-PTPRK AND P2RX1	0.948374761	0.911764706	0.988047809
P2RX1 AND NOT-SCN3A	0.949019608	0.934362934	0.964143426
P2RX1 AND NOT-BMPR2	0.947791165	0.955465587	0.940239044
P2RX1 AND NOT-SP4	0.942800789	0.93359375	0.952191235
P2RX1 AND SPINK2	0.929460581	0.96969697	0.892430279
P2RX1 AND NOT-TMEM204	0.931558935	0.890909091	0.976095618
P2RX1 AND NOT-STEAP4	0.941883768	0.947580645	0.93625498
P2RX1 AND NOT-SLC37A3	0.961616162	0.975409836	0.948207171
P2RX1 AND NOT-ADTRP	0.944123314	0.914179104	0.976095618
P2RX1 AND NOT-LRCH3	0.931558935	0.890909091	0.976095618
P2RX1 AND NOT-CDC14B	0.936660269	0.903703704	0.972111554
P2RX1 AND NOT-AOC3	0.92952381	0.890510949	0.972111554
P2RX1 AND NOT-ST3GAL5	0.946745562	0.9375	0.956175299
P2RX1 AND NOT-PHLDB2	0.962376238	0.956692913	0.96812749
P2RX1 AND NOT-VAPB	0.947775629	0.921052632	0.976095618
P2RX1 AND NOT-ARMCX2	0.95481336	0.941860465	0.96812749
P2RX1 AND NOT-MFAP3L	0.934615385	0.903345725	0.96812749
P2RX1 AND NOT-MFN2	0.934615385	0.903345725	0.96812749
NOT-SLC37A3 AND P2RX4	0.942345924	0.94047619	0.944223108
NOT-PHLDB2 AND P2RX4	0.936416185	0.906716418	0.96812749
NOT-SLC37A3 AND CHST11	0.941176471	0.958677686	0.924302789
NOT-PFN2 AND NOT-IL21R	0.953488372	0.928301887	0.980079681
NOT-MTUS1 AND NOT-IL21R	0.959223301	0.935606061	0.984063745
NOT-PTPRK AND NOT-IL21R	0.928571429	0.879003559	0.984063745
NOT-PHLDB2 AND TMED5	0.939393939	0.895306859	0.988047809
NOT-PFN2 AND SCCPDH	0.943248532	0.926923077	0.960159363
NOT-MTUS1 AND SCCPDH	0.958415842	0.952755906	0.964143426
NOT-PFN2 AND NOT-TUBD1	0.929791271	0.887681159	0.976095618
SMIM3 AND NOT-PDE3A	0.9348659	0.900369004	0.972111554
NOT-VAPB AND UBE2J1	0.917431193	0.850340136	0.996015936
NOT-PFN2 AND GDE1	0.927374302	0.870629371	0.992031873
SMIM3 AND NOT-PDGFRB	0.92952381	0.890510949	0.972111554
SMIM3 AND NOT-EMCN	0.964285714	0.960474308	0.96812749
NOT-ATP8B1 AND SMIM3	0.941176471	0.898550725	0.988047809
NOT-PFN2 AND NOT-SIRPG	0.937381404	0.894927536	0.984063745
NOT-PFN2 AND RNF130	0.92051756	0.85862069	0.992031873
NOT-PFN2 AND NOT-LPAR5	0.939393939	0.895306859	0.988047809
NOT-PFN2 AND NOT-PTGDR	0.930581614	0.879432624	0.988047809
NOT-PFN2 AND NOT-HEG1	0.93129771	0.893772894	0.972111554
NOT-PFN2 AND NOT-STIM2	0.923649907	0.867132867	0.988047809
NOT-PFN2 AND NOT-SLAMF7	0.938461538	0.907063197	0.972111554
NOT-PFN2 AND SUSD1	0.923364486	0.86971831	0.984063745
NOT-PFN2 AND SLC30A1	0.922222222	0.861591696	0.992031873
NOT-PFN2 AND NOT-PVRIG	0.941649899	0.951219512	0.932270916
NOT-PFN2 AND NOT-TTC13	0.935361217	0.894545455	0.980079681
NOT-PFN2 AND NOT-TRAF3IP3	0.935849057	0.888888889	0.988047809
NOT-PFN2 AND NOT-TLR10	0.941398866	0.895683453	0.992031873
NOT-PFN2 AND NOT-FCRL5	0.957364341	0.932075472	0.984063745
NOT-PFN2 AND B3GNT5	0.925650558	0.867595819	0.992031873
NOT-PFN2 AND NOT-STX7	0.926829268	0.875886525	0.984063745
NOT-PFN2 AND NOT-FCRLA	0.959381044	0.932330827	0.988047809
NOT-PFN2 AND NOT-ADTRP	0.928838951	0.876325088	0.988047809
NOT-PFN2 AND SMIM3	0.927374302	0.870629371	0.992031873
NOT-PFN2 AND NOT-KMO	0.944761905	0.905109489	0.988047809
NOT-PFN2 AND NOT-PHLDB2	0.926553672	0.878571429	0.980079681

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
TNFRSF8 AND NOT-SMAD2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC9A6	0.965517241	1	0.933333333
TNFRSF8 AND NOT-ART4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-LRRC8A	0.928571429	1	0.866666667
TNFRSF8 AND NOT-DCUN1D5	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-AMICA1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-FXYD6	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-HVCN1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-GRID2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ZNF7	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FPR1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-MGST2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-CA4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-STEAP4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-AMN	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ADGRF2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MYCT1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CD1E	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-NRG4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TTC13	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-LRP1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SLC18A3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-UBXN8	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MGAM	0.965517241	1	0.933333333
TNFRSF8 AND NOT-C10orf54	0.965517241	1	0.933333333
TNFRSF8 AND NOT-MAG	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TLR4	0.965517241	1	0.933333333
TNFRSF8 AND NOT-DSCAML1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-C5AR1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-CD300LF	0.965517241	1	0.933333333
TNFRSF8 AND NOT-PRPF38B	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GPR35	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ADAMTS13	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC14A1	0.928571429	1	0.866666667
TNFRSF8 AND NOT-SLC13A4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ADPRM	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FAM19A5	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FAM171A2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-LMF1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-YIPF1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CXCR1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-ASGR1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-LEMD3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PCSK1N	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMTC4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC6A5	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KIR2DS4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SCN11A	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ROS1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-HTR1D	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-OPRM1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ZMYM6	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SLC29A3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SGMS2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-TAOK2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ADAM22	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-BBS4	0.965517241	1	0.933333333
TNFRSF8 AND NOT-CADM4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-IFNAR2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMEM39A	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ACVR1C	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SIRPG	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-RGR	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GUCY2F	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FCRL4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KCNK4	0.965517241	1	0.933333333
TNFRSF8 AND NOT-AQP4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CDH8	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FCRL1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-S1PR3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ELOVL5	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GRIK4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CATSPER2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ITGA4	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SIGIRR	0.965517241	1	0.933333333
TNFRSF8 AND NOT-MANEA	0.933333333	0.93333333	0.933333333

Antigen Pair Logic	F1	Precision	Recall
NOT-PFN2 AND NOT-PCSK7	0.936170213	0.909774436	0.964143426
NOT-PFN2 AND HACD1	0.954274354	0.952380952	0.956175299
NOT-PFN2 AND NOT-FCMR	0.952198853	0.915441176	0.992031873
NOT-PFN2 AND NOT-CD8B	0.930320151	0.882142857	0.984063745
NOT-PFN2 AND NOT-MS4A1	0.95703125	0.938697318	0.976095618
NOT-PFN2 AND NOT-CD22	0.959223301	0.935606061	0.984063745
NOT-PFN2 AND NOT-CD27	0.948	0.951807229	0.944223108
NOT-PFN2 AND NOT-CD40LG	0.925373134	0.870175439	0.988047809
NOT-PFN2 AND NOT-KIAA0040	0.937853107	0.889285714	0.992031873
SMIM3 AND NOT-PLA2G2A	0.94368932	0.920454545	0.96812749
SMIM3 AND NOT-PLN	0.933078394	0.897058824	0.972111554
NOT-PHLDB2 AND PLP2	0.925373134	0.870175439	0.988047809
NOT-PHLDB2 AND FXYD5	0.941176471	0.898550725	0.988047809
NOT-VAPB AND FXYD5	0.924214418	0.862068966	0.996015936
NOT-FAM134B AND SLC39A8	0.941414141	0.954918033	0.928286853
SMIM3 AND NOT-PCDH18	0.954990215	0.938461538	0.972111554
SMIM3 AND NOT-RETSAT	0.94026975	0.910447761	0.972111554
NOT-TMEM45A AND SMIM3	0.942084942	0.913857678	0.972111554
SMIM3 AND NOT-TMEM100	0.951267057	0.93129771	0.972111554
NOT-MTUS1 AND NOT-TMEM140	0.933837429	0.888489209	0.984063745
TLR2 AND NOT-TMEM140	0.926315789	0.982142857	0.876494024
SMIM3 AND NOT-TMEM140	0.949494949	0.963114754	0.93625498
CD33 AND NOT-TMEM140	0.938016529	0.974248927	0.90438247
NOT-MTUS1 AND NOT-SIRPG	0.934086629	0.885714286	0.988047809
NOT-PTPRK AND NOT-SIRPG	0.928838951	0.876325088	0.988047809
NOT-PTPRK AND RNF130	0.941619586	0.892857143	0.996015936
SMIM3 AND NOT-AXL	0.964143426	0.964143426	0.964143426
NOT-PROS1 AND SMIM3	0.962671906	0.949612403	0.976095618
SMIM3 AND NOT-PRRG1	0.938461538	0.907063197	0.972111554
NOT-PHLDB2 AND ADPRM	0.94368932	0.920454545	0.96812749
NOT-VAPB AND ADPRM	0.930056711	0.884892086	0.980079681
CD33 AND NOT-MAN1C1	0.927966102	0.990950226	0.87250996
CD33 AND NOT-SLC44A2	0.934156379	0.965957447	0.90438247
NOT-MTUS1 AND NOT-PTGDR	0.930841121	0.876760563	0.992031873
NOT-SLC37A3 AND PTGER2	0.924686192	0.973568282	0.880478088
NOT-VAPB AND PTGER4	0.941398866	0.895683453	0.992031873
NOT-HEG1 AND ESYT2	0.923404255	0.99086758	0.864541833
NOT-HEG1 AND QSOX1	0.924050633	0.98206278	0.87250996
NOT-HEG1 AND SUSD1	0.923076923	0.995391705	0.860557769
NOT-HEG1 AND SPNS1	0.913419913	1	0.84063745
NOT-HEG1 AND SMIM3	0.93697479	0.991111111	0.888446215
NOT-HEG1 AND MYADM	0.921108742	0.990825688	0.860557769
MS4A3 AND NOT-HEG1	0.941414141	0.954918033	0.928286853
NOT-MTUS1 AND NOT-RARRES3	0.926553672	0.878571429	0.980079681
NOT-MTUS1 AND NOT-STX7	0.952015355	0.918518519	0.988047809
NOT-PHLDB2 AND NOT-MTUS1	0.928838951	0.876325088	0.988047809
NOT-MTUS1 AND HACD1	0.953907816	0.959677419	0.948207171
NOT-MTUS1 AND NOT-CD8B	0.926829268	0.875886525	0.984063745
NOT-MTUS1 AND NOT-MS4A1	0.953488372	0.928301887	0.980079681
NOT-MTUS1 AND NOT-CD27	0.953907816	0.959677419	0.948207171
NOT-PHLDB2 AND NOT-PTN	0.928838951	0.876325088	0.988047809
NOT-PTPRK AND QSOX1	0.928301887	0.88172043	0.980079681
SMIM3 AND NOT-PTPRG	0.943907157	0.917293233	0.972111554
NOT-PTPRK AND SPINK2	0.941176471	0.926640927	0.956175299
NOT-PTPRK AND MLLT10	0.938931298	0.901098901	0.980079681
NOT-PTPRK AND B3GNT5	0.943396226	0.896057348	0.996015936
NOT-PTPRK AND NOT-PHLDB2	0.939163498	0.898181818	0.984063745
NOT-PTPRK AND SPTLC2	0.935606061	0.891696751	0.984063745
NOT-PTPRM AND SMIM3	0.947791165	0.955465587	0.940239044
SMIM3 AND NOT-JAM2	0.949416342	0.927756654	0.972111554
NOT-EVA1C AND SMIM3	0.957575758	0.971311475	0.944223108
NOT-PERP AND SLC30A1	0.921933086	0.864111498	0.988047809
SMIM3 AND NOT-SMOC2	0.954990215	0.938461538	0.972111554
NOT-SLC37A3 AND SLC39A8	0.953907816	0.959677419	0.948207171
NOT-PHLDB2 AND SLC39A8	0.935281837	0.98245614	0.892430279
SMIM3 AND NOT-ADGRL4	0.956	0.959839357	0.952191235
SMIM3 AND NOT-ECSCR	0.934615385	0.903345725	0.96812749
SMIM3 AND NOT-SFRP1	0.93129771	0.893772894	0.972111554
NOT-SLC37A3 AND SUSD1	0.957746479	0.967479675	0.948207171
NOT-VAPB AND SUSD1	0.951456311	0.928030303	0.976095618
NOT-SGCB AND SMIM3	0.95703125	0.938697318	0.976095618
SMIM3 AND NOT-ITSN1	0.957403651	0.975206612	0.940239044
NOT-PHLDB2 AND FNDC3B	0.94026975	0.910447761	0.972111554
CD33 AND NOT-BMPR2	0.929460581	0.96969697	0.892430279
SMIM3 AND NOT-TGFBI	0.921443737	0.986363636	0.864541833
NOT-TGFBR3 AND SMIM3	0.928571429	0.982222222	0.880478088

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
TNFRSF8 AND NOT-PIGW	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MIP	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMPRSS11E	0.928571429	1	0.866666667
TNFRSF8 AND NOT-AVPR2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CLEC4G	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KCNQ1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-TRPC4	0.928571429	1	0.866666667
TNFRSF8 AND NOT-TRPM2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ADGRV1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TSPAN32	0.928571429	1	0.866666667
TNFRSF8 AND NOT-VSIG10L	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-DUOX2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FZD2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PYY	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PPAP2C	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FLT3LG	0.965517241	1	0.933333333
TNFRSF8 AND NOT-HFE2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-DNAJC18	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC35F4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC34A1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CLEC4A	0.965517241	1	0.933333333
TNFRSF8 AND NOT-MDM1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CLEC12B	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-OR1G1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CSF2RA	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CNTNAP2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-KCNG3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-IFNB1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-XK	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ERN1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-B3GNT3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MMP15	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMEM59L	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ADGRG6	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PTPRO	0.965517241	1	0.933333333
TNFRSF8 AND NOT-FUT7	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FRMD3	0.965517241	1	0.933333333
TNFRSF8 AND NOT-NPHS1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMEM86A	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC44A1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FAM19A1	0.928571429	1	0.866666667
TNFRSF8 AND NOT-CHRNA9	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ATP10D	0.928571429	1	0.866666667
TNFRSF8 AND NOT-NKAIN2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-HFE	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-B3GALNT1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-AMELX	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMEM156	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CLN8	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-EPDR1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TRPV4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CEACAM7	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CASR	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KCNQ2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ITPR2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PM20D1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-LRRC4C	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-LPAR4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC12A4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GPR20	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MPZL2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-MFAP3L	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SCN1A	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KCND3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-B4GALT1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SLC51B	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KCNQ4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PTGER4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-NIPAL3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ACSL5	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLITRK2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SUSD4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-JPH3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMEM237	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-EQTN	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CCR10	0.933333333	0.93333333	0.933333333

Antigen Pair Logic	F1	Precision	Recall
NOT-TGFBR3 AND MYADM	0.922105263	0.977678571	0.87250996
SMIM3 AND NOT-TLCD2	0.945736434	0.920754717	0.972111554
SMIM3 AND NOT-WFS1	0.938461538	0.907063197	0.972111554
NOT-SLC37A3 AND SLC30A1	0.938223938	0.91011236	0.96812749
NOT-PHLDB2 AND LAPTM5	0.932330827	0.882562278	0.988047809
NOT-DDR1 AND SMIM3	0.941176471	0.926640927	0.956175299
NOT-PHLDB2 AND CXCR4	0.932075472	0.885304659	0.984063745
SMIM3 AND NOT-FAT4	0.943907157	0.917293233	0.972111554
NOT-C1orf115 AND SMIM3	0.974358974	0.96484375	0.984063745
NOT-WLS AND SMIM3	0.949019608	0.934362934	0.964143426
SMIM3 AND NOT-CPED1	0.93697479	0.991111111	0.888446215
SMIM3 AND NOT-SLC2A10	0.945525292	0.923954373	0.96812749
SMIM3 AND NOT-COLEC12	0.941860465	0.916981132	0.96812749
SMIM3 AND NOT-GLT8D2	0.964426877	0.956862745	0.972111554
SMIM3 AND NOT-PLVAP	0.958742633	0.945736434	0.972111554
NOT-PHLDB2 AND NOT-TMEM47	0.934086629	0.885714286	0.988047809
NOT-TMEM47 AND HACD1	0.946534653	0.940944882	0.952191235
SMIM3 AND NOT-C19orf12	0.947368421	0.927480916	0.96812749
SMIM3 AND NOT-TMTC1	0.96	0.963855422	0.956175299
NOT-ANTXR1 AND HACD1	0.946534653	0.940944882	0.952191235
NOT-VAPB AND USP48	0.9348659	0.900369004	0.972111554
NOT-SLC37A3 AND SMIM3	0.947368421	0.927480916	0.96812749
NOT-FAM213A AND SMIM3	0.952755906	0.941634241	0.964143426
NOT-VAPB AND ABHD13	0.942965779	0.901818182	0.988047809
SMIM3 AND NOT-AOC3	0.972	0.975903614	0.96812749
SMIM3 AND NOT-NRP1	0.945736434	0.920754717	0.972111554
NOT-PHLDB2 AND SMIM3	0.966861598	0.946564885	0.988047809
SMIM3 AND NOT-ANGPTL1	0.966336634	0.960629921	0.972111554
SMIM3 AND NOT-PKDCC	0.966067864	0.968	0.964143426
SMIM3 AND NOT-BOC	0.933078394	0.897058824	0.972111554
SMIM3 AND NOT-OSMR	0.960629921	0.949416342	0.972111554
NOT-VAPB AND SMIM3	0.945525292	0.923954373	0.96812749
SMIM3 AND NOT-KL	0.94140625	0.923371648	0.960159363
NOT-ARMCX2 AND SMIM3	0.96875	0.950191571	0.988047809
NOT-PHLDB2 AND MYADM	0.954274354	0.952380952	0.956175299
NOT-PHLDB2 AND CD44	0.947775629	0.921052632	0.976095618
NOT-PHLDB2 AND CD53	0.928301887	0.88172043	0.980079681
NOT-PHLDB2 AND CLEC2B	0.942307692	0.910780669	0.976095618
NOT-VAPB AND TMEM183A	0.950381679	0.912087912	0.992031873
NOT-VAPB AND PTDSS1	0.936902486	0.900735294	0.976095618
CD33 AND NOT-PXYLP1	0.93081761	0.982300885	0.884462151
NOT-ARMCX2 AND ADGRE5	0.931818182	0.888086643	0.980079681
Stomach Neoplasms|Adenocarcinoma (Stomach A)			
MUC13 AND NOT-SLC30A10	0.942528736	0.931818182	0.953488372
MUC13 AND NOT-SCNN1B	0.976744186	0.976744186	0.976744186
MUC13 AND APOLD1	0.911111111	0.872340426	0.953488372
GREM1 AND MUC13	0.926829268	0.974358974	0.88372093
MUC13 AND NOT-DHRS9	0.921348315	0.891304348	0.953488372
MUC13 AND NOT-TRPM6	0.9	0.972972973	0.837209302
MUC13 AND NOT-CLDN8	0.952380952	0.975609756	0.930232558
MUC13 AND NOT-SLC26A2	0.953488372	0.953488372	0.953488372
MUC13 AND NOT-CWH43	0.930232558	0.930232558	0.930232558
GREM1 AND TNFRSF10B	0.915662651	0.95	0.88372093
GREM1 AND MUC1	0.913580247	0.973684211	0.860465116
GREM1 AND KCNE3	0.938271605	1	0.88372093
GREM1 AND ATP9A	0.9	0.972972973	0.837209302
GREM1 AND TSPAN1	0.915662651	0.95	0.88372093
TMC5 AND NOT-TSPAN1	0.904761905	0.926829268	0.88372093
GREM1 AND PDZK1IP1	0.904761905	0.926829268	0.88372093
MUC13 AND NOT-DHRS9	0.921348315	0.891304348	0.953488372
GREM1 AND ADAM8	0.915662651	0.95	0.88372093
GREM1 AND ZMPSTE24	0.938271605	1	0.88372093
GREM1 AND PRSS16	0.904761905	0.926829268	0.88372093
GREM1 AND LILRB2	0.926829268	0.974358974	0.88372093
GREM1 AND B3GNT3	0.926829268	0.974358974	0.88372093
B3GNT3 AND NOT-SCNN1B	0.915662651	0.95	0.88372093
B3GNT3 AND NOT-CLDN8	0.9	0.972972973	0.837209302
GREM1 AND LYPLA1	0.925	1	0.860465116
GREM1 AND HTATIP2	0.911392405	1	0.837209302
GREM1 AND AGPAT2	0.938271605	1	0.88372093
GREM1 AND TNFSF13B	0.938271605	1	0.88372093
GREM1 AND SLC12A7	0.911392405	1	0.837209302
GREM1 AND ADAM28	0.902439024	0.948717949	0.860465116
GRIN2D AND JTB	0.909090909	0.888888889	0.930232558
GREM1 AND AFG3L2	0.938271605	1	0.88372093
GREM1 AND SLC35D2	0.938271605	1	0.88372093

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
TNFRSF8 AND NOT-FAM57A	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-STX3	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SLC25A34	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-OR10C1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC44A2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MGAT4D	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-IL1RL2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SV2A	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CLDN6	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PTGER3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMEM198	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-IZUMO1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-LTK	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PILRA	0.965517241	1	0.933333333
TNFRSF8 AND NOT-VSTM4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-JPH1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PLXDC1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMEM200C	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ALG9	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ANTXR2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-LMAN2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-ABCC11	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMEM199	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MS4A14	0.965517241	1	0.933333333
TNFRSF8 AND NOT-GLMP	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-BEST4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-M6PR	0.928571429	1	0.866666667
TNFRSF8 AND NOT-CD36	0.965517241	1	0.933333333
TNFRSF8 AND NOT-WDR33	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-UXS1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SLC8A1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SLC39A14	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ISLR2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-UNC5C	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KCNJ11	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-HTR3A	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ODF4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-NIPAL4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC16A8	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC7A6	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC7A9	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MFSD9	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-VSIG1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-P2RY4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ECE2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ELFN2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-THADA	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SCNN1G	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-DERL2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC27A1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-HTR3B	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC16A1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CDH15	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SAMD8	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMEM251	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SOX1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-STX18	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SGCD	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PPAPDC3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ECEL1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-CDH24	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GP5	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-OPN5	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PF4	0.965517241	1	0.933333333
TNFRSF8 AND NOT-PCDHGC4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMEM27	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ADAM11	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-NCR2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ADGRG3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-EPHA5	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MGAT1	0.928571429	1	0.866666667
TNFRSF8 AND NOT-SLMAP	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FAM19A2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-NCAM1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-CHST13	0.965517241	1	0.933333333
TNFRSF8 AND NOT-PVRL1	0.933333333	0.93333333	0.933333333

Antigen Pair Logic	F1	Precision	Recall
GREM1 AND GALNT6	0.925	1	0.860465116
GREM1 AND GLMP	0.925	1	0.860465116
GREM1 AND MGAT4B	0.938271605	1	0.88372093
GREM1 AND MGAT4A	0.911392405	1	0.837209302
GREM1 AND TMC6	0.904761905	0.926829268	0.88372093
GREM1 AND MAL2	0.915662651	0.95	0.88372093
GREM1 AND NOT-GALNT13	0.938271605	1	0.88372093
GREM1 AND SLC18B1	0.938271605	1	0.88372093
GREM1 AND CLCN5	0.904761905	0.926829268	0.88372093
GREM1 AND ZFYVE27	0.926829268	0.974358974	0.88372093
GREM1 AND SFXN4	0.925	1	0.860465116
GREM1 AND TMEM45B	0.915662651	0.95	0.88372093
GREM1 AND CLPTM1	0.904761905	0.926829268	0.88372093
GREM1 AND NIPA1	0.904761905	0.926829268	0.88372093
GREM1 AND TMEM219	0.926829268	0.974358974	0.88372093
GREM1 AND CANT1	0.938271605	1	0.88372093
GREM1 AND SLC44A3	0.926829268	0.974358974	0.88372093
GREM1 AND TMEM125	0.915662651	0.95	0.88372093
GREM1 AND NOT-TMEM207	0.9	0.972972973	0.837209302
GREM1 AND COX8A	0.926829268	0.974358974	0.88372093
GREM1 AND CLDN4	0.915662651	0.95	0.88372093
MUC13 AND NOT-TRPM6	0.9	0.972972973	0.837209302
GREM1 AND CMTM4	0.926829268	0.974358974	0.88372093
GREM1 AND TMC4	0.926829268	0.974358974	0.88372093
GREM1 AND CXADR	0.904761905	0.926829268	0.88372093
GREM1 AND TMEM161B	0.926829268	0.974358974	0.88372093
GREM1 AND RDH10	0.926829268	0.974358974	0.88372093
GREM1 AND CYP3A5	0.915662651	0.95	0.88372093
GREM1 AND ABHD3	0.925	1	0.860465116
GREM1 AND DPAGT1	0.938271605	1	0.88372093
GREM1 AND DSG2	0.938271605	1	0.88372093
DSG2 AND NOT-CLDN8	0.942528736	0.931818182	0.953488372
MUC13 AND NOT-SLC26A2	0.953488372	0.953488372	0.953488372
GREM1 AND NOT-PLD5	0.915662651	0.95	0.88372093
GREM1 AND KRTCAP3	0.925	1	0.860465116
GREM1 AND TIGIT	0.904761905	0.926829268	0.88372093
GREM1 AND EPHA1	0.904761905	0.926829268	0.88372093
GREM1 AND ERN1	0.9	0.972972973	0.837209302
GREM1 AND EVI2A	0.915662651	0.95	0.88372093
GREM1 AND EVI2B	0.904761905	0.926829268	0.88372093
GREM1 AND F2RL1	0.926829268	0.974358974	0.88372093
GREM1 AND MPEG1	0.926829268	0.974358974	0.88372093
GREM1 AND FCER1G	0.925	1	0.860465116
CYP2S1 AND NOT-SLC16A9	0.921348315	0.891304348	0.953488372
GREM1 AND C6orf136	0.926829268	0.974358974	0.88372093
GREM1 AND FCGR3B	0.911392405	1	0.837209302
GREM1 AND TMED4	0.911392405	1	0.837209302
GREM1 AND UNC5CL	0.915662651	0.95	0.88372093
GREM1 AND CD93	0.913580247	0.973684211	0.860465116
GREM1 AND NUP210	0.926829268	0.974358974	0.88372093
GREM1 AND SEL1L3	0.938271605	1	0.88372093
GREM1 AND ATP11A	0.902439024	0.948717949	0.860465116
GREM1 AND FMO5	0.9	0.972972973	0.837209302
GREM1 AND SLC9A8	0.904761905	0.926829268	0.88372093
GREM1 AND SLC35A3	0.926829268	0.974358974	0.88372093
GREM1 AND TRAM1	0.911392405	1	0.837209302
GREM1 AND SEC61G	0.926829268	0.974358974	0.88372093
GREM1 AND VSIG2	0.9	0.972972973	0.837209302
GREM1 AND LEMD3	0.925	1	0.860465116
GREM1 AND PLD3	0.926829268	0.974358974	0.88372093
GREM1 AND TMEM2	0.938271605	1	0.88372093
GREM1 AND ALOX5AP	0.925	1	0.860465116
GREM1 AND FUT2	0.904761905	0.926829268	0.88372093
GREM1 AND FUT4	0.938271605	1	0.88372093
GREM1 AND FUT8	0.913580247	0.973684211	0.860465116
GREM1 AND CERS6	0.926829268	0.974358974	0.88372093
GREM1 AND IFI6	0.911392405	1	0.837209302
GREM1 AND ZDHHC23	0.926829268	0.974358974	0.88372093
GREM1 AND GALNT3	0.938271605	1	0.88372093
GREM1 AND IL4I1	0.904761905	0.926829268	0.88372093
GREM1 AND PVRL3	0.9	0.972972973	0.837209302
GREM1 AND PNKD	0.911392405	1	0.837209302
GREM1 AND STEAP2	0.915662651	0.95	0.88372093
GREM1 AND KCNG2	0.904761905	0.926829268	0.88372093
GREM1 AND CLEC4E	0.902439024	0.948717949	0.860465116
GREM1 AND CNNM4	0.915662651	0.95	0.88372093

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
TNFRSF8 AND NOT-RYR1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMEM151B	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GABRA4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-NTSR1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CNNM2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-B3GLCT	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CLEC1A	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GABRD	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-VKORC1L1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MUC3A	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-OPN1SW	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GCNT2	0.928571429	1	0.866666667
TNFRSF8 AND NOT-BCL2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SPI1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-EGF	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PTPRS	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-DISP1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-COL11A2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SMIM10	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMEM19	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SNRNP40	0.965517241	1	0.933333333
TNFRSF8 AND NOT-CGRRF1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-POMT1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ST6GAL2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PTCH1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MUC17	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CFAP47	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMEM221	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ATP10A	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-HMP19	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-BTC	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KCNA6	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMEM141	0.965517241	1	0.933333333
TNFRSF8 AND NOT-ERVMER34-1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ESYT3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-HTR1B	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-STX17	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMCC3	0.965517241	1	0.933333333
TNFRSF8 AND NOT-CCL11	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PIGA	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ELANE	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GPR6	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-C11orf87	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-LGR6	0.965517241	1	0.933333333
TNFRSF8 AND NOT-CD3G	0.928571429	1	0.866666667
TNFRSF8 AND NOT-KCNH8	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ENPP5	0.965517241	1	0.933333333
TNFRSF8 AND NOT-NRXN2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-NPY5R	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ABCG4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC25A48	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GPR155	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SNX19	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KLRF1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-GRIK3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ATP2A3	0.965517241	1	0.933333333
TNFRSF8 AND NOT-PCDH19	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PCDHB16	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PCDHGC5	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ZFPL1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SPCS2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ENPP1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-COL24A1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-RTN1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-GPR107	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC30A3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC4A9	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MTNR1B	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-DPCR1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MARCH4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC35D3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-LRRN1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC47A2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KLHL2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SLC23A1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MPZL3	0.933333333	0.93333333	0.933333333

Antigen Pair Logic	F1	Precision	Recall
GREM1 AND STEAP1	0.915662651	0.95	0.88372093
GREM1 AND GPR160	0.938271605	1	0.88372093
GREM1 AND CNPPD1	0.926829268	0.974358974	0.88372093
GREM1 AND GJB1	0.926829268	0.974358974	0.88372093
GREM1 AND NOT-GLRA2	0.9	0.972972973	0.837209302
GREM1 AND GOLGB1	0.911392405	1	0.837209302
GREM1 AND NOT-GPM6B	0.902439024	0.948717949	0.860465116
GREM1 AND B4GALNT3	0.904761905	0.926829268	0.88372093
GREM1 AND TMEM176B	0.915662651	0.95	0.88372093
GREM1 AND CYP2S1	0.915662651	0.95	0.88372093
GREM1 AND DPP7	0.925	1	0.860465116
GREM1 AND HLA-DOA	0.904761905	0.926829268	0.88372093
GREM1 AND HLA-DRA	0.911392405	1	0.837209302
GREM1 AND HLA-F	0.926829268	0.974358974	0.88372093
GREM1 AND HSD17B2	0.915662651	0.95	0.88372093
GREM1 AND ANO9	0.904761905	0.926829268	0.88372093
GREM1 AND SERINC2	0.938271605	1	0.88372093
GREM1 AND IL7R	0.938271605	1	0.88372093
GREM1 AND IL13	0.904761905	0.926829268	0.88372093
GREM1 AND INPP4A	0.926829268	0.974358974	0.88372093
GREM1 AND INSR	0.926829268	0.974358974	0.88372093
GREM1 AND ITGA6	0.925	1	0.860465116
GREM1 AND ITGAM	0.911392405	1	0.837209302
GREM1 AND ITGAX	0.904761905	0.926829268	0.88372093
GREM1 AND ITGB2	0.938271605	1	0.88372093
GREM1 AND ITPR3	0.938271605	1	0.88372093
GREM1 AND TMEM205	0.926829268	0.974358974	0.88372093
GREM1 AND KCNK1	0.911392405	1	0.837209302
GREM1 AND KCNN4	0.913580247	0.973684211	0.860465116
GREM1 AND KCNQ1	0.926829268	0.974358974	0.88372093
GREM1 AND TMEM189	0.915662651	0.95	0.88372093
GREM1 AND NOT-SPATA31D1	0.915662651	0.95	0.88372093
GREM1 AND LAMP1	0.913580247	0.973684211	0.860465116
GREM1 AND EPCAM	0.938271605	1	0.88372093
GREM1 AND MGST2	0.926829268	0.974358974	0.88372093
GREM1 AND ABCC1	0.915662651	0.95	0.88372093
GREM1 AND NOT-SMCO3	0.925	1	0.860465116
GREM1 AND SMIM22	0.925	1	0.860465116
GREM1 AND MST1R	0.926829268	0.974358974	0.88372093
GREM1 AND MTHFD1	0.938271605	1	0.88372093
GREM1 AND MUC1	0.913580247	0.973684211	0.860465116
GREM1 AND MUC3A	0.904761905	0.926829268	0.88372093
GREM1 AND ATP2A3	0.904761905	0.926829268	0.88372093
GREM1 AND NTRK1	0.913580247	0.973684211	0.860465116
GREM1 AND NUCB1	0.904761905	0.926829268	0.88372093
GREM1 AND P2RX4	0.926829268	0.974358974	0.88372093
GREM1 AND F11R	0.926829268	0.974358974	0.88372093
GREM1 AND TMX2	0.925	1	0.860465116
GREM1 AND SLC15A3	0.904761905	0.926829268	0.88372093
GREM1 AND LSR	0.926829268	0.974358974	0.88372093
GREM1 AND PIGT	0.925	1	0.860465116
GREM1 AND TMBIM4	0.911392405	1	0.837209302
GREM1 AND ACSL5	0.911392405	1	0.837209302
GREM1 AND GALNT7	0.938271605	1	0.88372093
GREM1 AND CECR1	0.913580247	0.973684211	0.860465116
GREM1 AND SERPINA1	0.926829268	0.974358974	0.88372093
GREM1 AND NOT-FXYD1	0.913580247	0.973684211	0.860465116
GREM1 AND PLXNA1	0.904761905	0.926829268	0.88372093
GREM1 AND ATP6AP1	0.938271605	1	0.88372093
GREM1 AND PODXL	0.9	0.972972973	0.837209302
GREM1 AND POR	0.938271605	1	0.88372093
GREM1 AND VSIG10	0.904761905	0.926829268	0.88372093
GREM1 AND P4HTM	0.926829268	0.974358974	0.88372093
GREM1 AND TRPM4	0.926829268	0.974358974	0.88372093
GREM1 AND CDHR2	0.938271605	1	0.88372093
GREM1 AND RETSAT	0.915662651	0.95	0.88372093
GREM1 AND TMEM248	0.938271605	1	0.88372093
GREM1 AND TAPBPL	0.904761905	0.926829268	0.88372093
GREM1 AND TMEM51	0.915662651	0.95	0.88372093
GREM1 AND ZDHHC4	0.913580247	0.973684211	0.860465116
GREM1 AND TMEM57	0.9	0.972972973	0.837209302
GREM1 AND NOT-SLC30A10	0.913580247	0.973684211	0.860465116
GREM1 AND IL17RB	0.902439024	0.948717949	0.860465116
GREM1 AND SLC39A4	0.926829268	0.974358974	0.88372093
GREM1 AND CARKD	0.902439024	0.948717949	0.860465116
GREM1 AND ST6GALNAC1	0.938271605	1	0.88372093

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
TNFRSF8 AND NOT-DGAT1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-LPPR3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-NRG3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KIDINS220	0.965517241	1	0.933333333
TNFRSF8 AND NOT-PRRG3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CALHM1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC24A2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ZDHHC22	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ZDHHC18	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMED3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-OPRD1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CLEC5A	0.965517241	1	0.933333333
TNFRSF8 AND NOT-LCN10	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-C6orf25	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-UGT2B17	0.928571429	1	0.866666667
TNFRSF8 AND NOT-DAGLA	0.965517241	1	0.933333333
TNFRSF8 AND NOT-TMEM265	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-BTN1A1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-WLS	0.965517241	1	0.933333333
TNFRSF8 AND NOT-ABCC4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-NIPAL2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-HRH3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ATCAY	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KCNK6	0.965517241	1	0.933333333
TNFRSF8 AND NOT-ATP13A4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FMNL1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ATP6V0A4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-LRRN3	0.965517241	1	0.933333333
TNFRSF8 AND NOT-GDNF	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-XCR1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-DDX59	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-LMBR1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CHST10	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ADAM30	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GALNTL5	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC5A1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-C17orf74	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CSF3R	0.965517241	1	0.933333333
TNFRSF8 AND NOT-FAM132A	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ERBB4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CYP4F8	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC5A2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MCU	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CCPG1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-USP27X	0.965517241	1	0.933333333
TNFRSF8 AND NOT-CCDC109B	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KMO	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PCDHGA10	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-LRFN4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PTPN2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-BTNL2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ADRA1B	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMEM61	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-LAMP2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GXYLT1	0.928571429	1	0.866666667
TNFRSF8 AND NOT-PTGS1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-KCNMB1	0.928571429	1	0.866666667
TNFRSF8 AND NOT-ADAM18	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GDF9	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CDCP1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-DERL1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMEM132D	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ZAN	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MRAP	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MUC16	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLCO4C1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-CDH26	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CACNA1F	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMEM144	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ADGRA3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TAS2R7	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ADCY10	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-LTB4R	0.928571429	1	0.866666667
TNFRSF8 AND NOT-RYR3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLAMF9	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-OR7C1	0.933333333	0.93333333	0.933333333

Antigen Pair Logic	F1	Precision	Recall
GREM1 AND LRRC8A	0.904761905	0.926829268	0.88372093
GREM1 AND TMPRSS4	0.904761905	0.926829268	0.88372093
GREM1 AND MUC13	0.926829268	0.974358974	0.88372093
GREM1 AND SLAMF8	0.904761905	0.926829268	0.88372093
GREM1 AND C1GALT1	0.911392405	1	0.837209302
GREM1 AND NOT-OTOR	0.915662651	0.95	0.88372093
GREM1 AND ATP13A1	0.926829268	0.974358974	0.88372093
GREM1 AND SLC45A4	0.925	1	0.860465116
GREM1 AND SYT13	0.913580247	0.973684211	0.860465116
GREM1 AND SEMA4G	0.926829268	0.974358974	0.88372093
GREM1 AND PTPRCAP	0.904761905	0.926829268	0.88372093
GREM1 AND PTPRE	0.926829268	0.974358974	0.88372093
GREM1 AND PTPRH	0.925	1	0.860465116
GREM1 AND PTPRK	0.938271605	1	0.88372093
GREM1 AND CXCL16	0.925	1	0.860465116
GREM1 AND PVRL2	0.904761905	0.926829268	0.88372093
GREM1 AND BCL2L1	0.915662651	0.95	0.88372093
GREM1 AND RNASE6	0.926829268	0.974358974	0.88372093
GREM1 AND RPN2	0.911392405	1	0.837209302
GREM1 AND CEACAM1	0.902439024	0.948717949	0.860465116
GREM1 AND SDC4	0.911392405	1	0.837209302
GREM1 AND BIK	0.938271605	1	0.88372093
GREM1 AND ABCG4	0.902439024	0.948717949	0.860465116
GREM1 AND EDDM3B	0.926829268	0.974358974	0.88372093
GREM1 AND DNAJC1	0.926829268	0.974358974	0.88372093
GREM1 AND NOT-SFRP1	0.913580247	0.973684211	0.860465116
GREM1 AND CYP3A43	0.904761905	0.926829268	0.88372093
GREM1 AND SLC4A2	0.926829268	0.974358974	0.88372093
GREM1 AND NOT-SLC10A1	0.926829268	0.974358974	0.88372093
GREM1 AND SLC12A2	0.911392405	1	0.837209302
GREM1 AND SLC16A1	0.913580247	0.973684211	0.860465116
GREM1 AND SLC20A1	0.911392405	1	0.837209302
GREM1 AND SPAG4	0.915662651	0.95	0.88372093
GREM1 AND SPI1	0.915662651	0.95	0.88372093
GREM1 AND SPINT1	0.904761905	0.926829268	0.88372093
GREM1 AND ST14	0.913580247	0.973684211	0.860465116
GREM1 AND STX3	0.925	1	0.860465116
GREM1 AND TAPBP	0.915662651	0.95	0.88372093
GREM1 AND TLR2	0.911392405	1	0.837209302
GREM1 AND TSPAN8	0.915662651	0.95	0.88372093
GREM1 AND TMPRSS2	0.915662651	0.95	0.88372093
GREM1 AND TNFRSF1B	0.904761905	0.926829268	0.88372093
GREM1 AND C1QB	0.915662651	0.95	0.88372093
GREM1 AND TYROBP	0.915662651	0.95	0.88372093
GREM1 AND UGT8	0.904761905	0.926829268	0.88372093
GREM1 AND XK	0.902439024	0.948717949	0.860465116
GREM1 AND FZD5	0.913580247	0.973684211	0.860465116
GREM1 AND FA2H	0.926829268	0.974358974	0.88372093
GREM1 AND ABHD16A	0.915662651	0.95	0.88372093
GREM1 AND LST1	0.904761905	0.926829268	0.88372093
GREM1 AND TTC13	0.925	1	0.860465116
GREM1 AND RNF128	0.926829268	0.974358974	0.88372093
GREM1 AND RHBDF2	0.902439024	0.948717949	0.860465116
GREM1 AND C3orf52	0.925	1	0.860465116
GREM1 AND GALNT12	0.913580247	0.973684211	0.860465116
GREM1 AND TMC5	0.938271605	1	0.88372093
GREM1 AND LPCAT1	0.911392405	1	0.837209302
GREM1 AND DNAJC22	0.904761905	0.926829268	0.88372093
GREM1 AND ALPK1	0.926829268	0.974358974	0.88372093
GREM1 AND HSD3B7	0.904761905	0.926829268	0.88372093
GREM1 AND TTYH3	0.915662651	0.95	0.88372093
GREM1 AND SLC44A4	0.926829268	0.974358974	0.88372093
GREM1 AND NOT-SPX	0.913580247	0.973684211	0.860465116
GREM1 AND HM13	0.926829268	0.974358974	0.88372093
GREM1 AND AKAP1	0.926829268	0.974358974	0.88372093
GREM1 AND CRISPLD2	0.911392405	1	0.837209302
GREM1 AND ARMC10	0.911392405	1	0.837209302
GREM1 AND SPNS1	0.938271605	1	0.88372093
GREM1 AND PLA2G10	0.925	1	0.860465116
GREM1 AND SLC41A2	0.9	0.972972973	0.837209302
GREM1 AND FAR1	0.913580247	0.973684211	0.860465116
GREM1 AND SLC12A8	0.938271605	1	0.88372093
GREM1 AND MFSD9	0.915662651	0.95	0.88372093
GREM1 AND TMEM87B	0.911392405	1	0.837209302
GREM1 AND SLC43A1	0.913580247	0.973684211	0.860465116
GREM1 AND PAQR8	0.911392405	1	0.837209302

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
TNFRSF8 AND NOT-VAMP2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-RNASE7	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SMCO3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ABCG8	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC28A2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMEM239	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-C19orf18	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ABCB4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FAM69C	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-100132596?	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ABCB11	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KCNA1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CISD2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CACNA1D	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MXRA7	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SUMF2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-BTN2A1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ABCD4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GDPD5	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC35E2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-HRH2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ITGB3	0.928571429	1	0.866666667
TNFRSF8 AND NOT-FLRT1	0.928571429	1	0.866666667
TNFRSF8 AND NOT-MSLN	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-NAALAD2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-C5AR2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMEM55B	0.928571429	1	0.866666667
TNFRSF8 AND NOT-SLC16A6	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-HRH4	0.965517241	1	0.933333333
TNFRSF8 AND NOT-MFSD8	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-LST1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SPPL2B	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KCNK10	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ATP1A4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SEZ6L	0.928571429	1	0.866666667
TNFRSF8 AND NOT-KCNA4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-NCR1	0.928571429	1	0.866666667
TNFRSF8 AND NOT-SSTR4	0.965517241	1	0.933333333
TNFRSF8 AND NOT-CYP2A6	0.928571429	1	0.866666667
TNFRSF8 AND NOT-AGPAT9	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SIRPA	0.965517241	1	0.933333333
TNFRSF8 AND NOT-GCNT1	0.928571429	1	0.866666667
TNFRSF8 AND NOT-DNAJC4	0.965517241	1	0.933333333
TNFRSF8 AND NOT-CLEC4E	0.965517241	1	0.933333333
TNFRSF8 AND NOT-USP30	0.928571429	1	0.866666667
TNFRSF8 AND NOT-ADGRD1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-XXYLT1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-KLRC3	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SNX13	0.965517241	1	0.933333333
TNFRSF8 AND NOT-ABCB1	0.928571429	1	0.866666667
TNFRSF8 AND NOT-SIGLEC9	0.928571429	1	0.866666667
TNFRSF8 AND NOT-CCR2	0.928571429	1	0.866666667
TNFRSF8 AND NOT-ORAI3	0.965517241	1	0.933333333
TNFRSF8 AND NOT-TMCC2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-CLCN3	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SLC36A1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-FCGR2B	0.965517241	1	0.933333333
TNFRSF8 AND NOT-CD244	0.965517241	1	0.933333333
TNFRSF8 AND NOT-CUX1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-MAN2A2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-CKLF	0.965517241	1	0.933333333
TNFRSF8 AND NOT-LFNG	0.965517241	1	0.933333333
TNFRSF8 AND NOT-CLEC12A	0.965517241	1	0.933333333
TNFRSF8 AND NOT-CYSLTR1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-S1PR1	0.928571429	1	0.866666667
TNFRSF8 AND NOT-CLEC10A	0.928571429	1	0.866666667
TNFRSF8 AND NOT-CKAP4	0.965517241	1	0.933333333
TNFRSF8 AND NOT-ADCY5	0.928571429	1	0.866666667
TNFRSF8 AND NOT-VSTM1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-UPK3A	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SIRPB2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-P2RX7	0.928571429	1	0.866666667
TNFRSF8 AND NOT-RHO	0.965517241	1	0.933333333
TNFRSF8 AND NOT-BPI	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SLC39A7	0.965517241	1	0.933333333
TNFRSF8 AND NOT-TMEM175	0.965517241	1	0.933333333

Antigen Pair Logic	F1	Precision	Recall
GREM1 AND PPAP2C	0.926829268	0.974358974	0.88372093
GREM1 AND NOT-PPAP2B	0.902439024	0.948717949	0.860465116
GREM1 AND DGAT1	0.938271605	1	0.88372093
GREM1 AND CD164	0.911392405	1	0.837209302
GREM1 AND TNFRSF14	0.938271605	1	0.88372093
GREM1 AND TNFRSF10B	0.915662651	0.95	0.88372093
GREM1 AND GPRC5A	0.926829268	0.974358974	0.88372093
GREM1 AND MCU	0.911392405	1	0.837209302
GREM1 AND SLC7A7	0.925	1	0.860465116
GREM1 AND CLDN12	0.938271605	1	0.88372093
GREM1 AND SYNGR2	0.938271605	1	0.88372093
GREM1 AND MARVELD3	0.925	1	0.860465116
GREM1 AND CRB3	0.904761905	0.926829268	0.88372093
GREM1 AND GCNT3	0.904761905	0.926829268	0.88372093
GREM1 AND CD7	0.904761905	0.926829268	0.88372093
GREM1 AND CD27	0.913580247	0.973684211	0.860465116
GREM1 AND PIGS	0.915662651	0.95	0.88372093
GREM1 AND EPSTI1	0.925	1	0.860465116
GREM1 AND CD86	0.904761905	0.926829268	0.88372093
GREM1 AND ENTPD6	0.938271605	1	0.88372093
GREM1 AND ENTPD4	0.938271605	1	0.88372093
GREM1 AND MLEC	0.925	1	0.860465116
GREM1 AND ADGRE5	0.904761905	0.926829268	0.88372093
GREM1 AND PIEZO1	0.904761905	0.926829268	0.88372093
GREM1 AND EFCAB14	0.915662651	0.95	0.88372093
GREM1 AND AREL1	0.911392405	1	0.837209302
CYP2S1 AND NOT-SCNN1B	0.954545455	0.933333333	0.976744186
CYP2S1 AND NOT-CWH43	0.909090909	0.888888889	0.930232558
MUC13 AND NOT-SLC30A10	0.942528736	0.931818182	0.953488372
MUC13 AND NOT-SCNN1B	0.976744186	0.976744186	0.976744186
MUC13 AND NOT-CWH43	0.930232558	0.930232558	0.930232558
MUC13 AND APOLD1	0.911111111	0.872340426	0.953488372
MUC13 AND NOT-CLDN8	0.952380952	0.975609756	0.930232558
SMAGP AND NOT-SCNN1B	0.903225806	0.84	0.976744186
TMC5 AND NOT-SCNN1B	0.904761905	0.926829268	0.88372093
BIK AND NOT-CLDN8	0.917647059	0.928571429	0.906976744
Lymphoma, Large B-Cell, Diffuse (B-Cell Diffuse)			
PLA2G2D AND NOT-GABRD	0.916666667	0.942857143	0.891891892
PLA2G2D AND NOT-BMPR1B	0.931506849	0.944444444	0.918918919
PLA2G2D AND NOT-CRYGC	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-HCN4	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-SLC17A4	0.918918919	0.918918919	0.918918919
PLA2G2D AND NOT-CDH7	0.918918919	0.918918919	0.918918919
PLA2G2D AND NOT-CDH12	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-CLEC3A	0.918918919	0.918918919	0.918918919
PLA2G2D AND NOT-PDZK1IP1	0.942857143	1	0.891891892
PLA2G2D AND NOT-TENM1	0.916666667	0.942857143	0.891891892
PLA2G2D AND NOT-DHRS2	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-ANGPTL7	0.918918919	0.918918919	0.918918919
PLA2G2D AND NOT-GPA33	0.904109589	0.916666667	0.891891892
PLA2G2D AND NOT-MSLN	0.918918919	0.918918919	0.918918919
PLA2G2D AND NOT-NMUR1	0.929577465	0.970588235	0.891891892
PLA2G2D AND NOT-ABCA8	0.944444444	0.971428571	0.918918919
PLA2G2D AND NOT-CACNG2	0.931506849	0.944444444	0.918918919
PLA2G2D AND NOT-WFDC2	0.904109589	0.916666667	0.891891892
PLA2G2D AND NOT-LRRN2	0.929577465	0.970588235	0.891891892
PLA2G2D AND NOT-SEMA6B	0.914285714	0.96969697	0.864864865
PLA2G2D AND NOT-ST6GALNAC2	0.914285714	0.96969697	0.864864865
PLA2G2D AND NOT-CERS1	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-NRG3	0.904109589	0.916666667	0.891891892
PLA2G2D AND NOT-CCR9	0.942857143	1	0.891891892
PLA2G2D AND NOT-FAXDC2	0.916666667	0.942857143	0.891891892
PLA2G2D AND NOT-SLC26A1	0.931506849	0.944444444	0.918918919
PLA2G2D AND NOT-NAT2	0.914285714	0.96969697	0.864864865
PLA2G2D AND NOT-SLC7A9	0.944444444	0.971428571	0.918918919
PLA2G2D AND NOT-GALNT5	0.904109589	0.916666667	0.891891892
PLA2G2D AND NOT-MRAP2	0.942857143	1	0.891891892
PLA2G2D AND NOT-SCN11A	0.901408451	0.941176471	0.864864865
PLA2G2D AND NOT-CHRM2	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-PTH2	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-SCRG1	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-MFSD3	0.904109589	0.916666667	0.891891892
PLA2G2D AND NOT-SIGLEC11	0.904109589	0.916666667	0.891891892
PLA2G2D AND NOT-CHRNE	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-CSMD3	0.918918919	0.918918919	0.918918919
PLA2G2D AND NOT-GALNT13	0.918918919	0.918918919	0.918918919

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
TNFRSF8 AND NOT-SEMA4D	0.965517241	1	0.933333333
TNFRSF8 AND NOT-ARSD	0.965517241	1	0.933333333
TNFRSF8 AND NOT-IL10RA	0.965517241	1	0.933333333
TNFRSF8 AND NOT-MARCH9	0.965517241	1	0.933333333
TNFRSF8 AND NOT-KCNE3	0.965517241	1	0.933333333
TNFRSF8 AND NOT-LPAR5	0.965517241	1	0.933333333
TNFRSF8 AND NOT-NXPE3	0.965517241	1	0.933333333
TNFRSF8 AND NOT-PPBP	0.965517241	1	0.933333333
TNFRSF8 AND NOT-MFSD2A	0.965517241	1	0.933333333
TNFRSF8 AND NOT-LPGAT1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-CHST11	0.965517241	1	0.933333333
TNFRSF8 AND NOT-DRAM2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-ATG9A	0.965517241	1	0.933333333
TNFRSF8 AND NOT-KLRC4	0.965517241	1	0.933333333
TNFRSF8 AND NOT-FMO5	0.928571429	1	0.866666667
TNFRSF8 AND NOT-SLC5A11	0.928571429	1	0.866666667
TNFRSF8 AND NOT-SSR1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SLC25A20	0.965517241	1	0.933333333
TNFRSF8 AND NOT-DPEP2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SLC22A15	0.928571429	1	0.866666667
TNFRSF8 AND NOT-CLEC4D	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SLC16A7	0.965517241	1	0.933333333
TNFRSF8 AND NOT-MBOAT1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-NFAM1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-TM6SF1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-CYBB	0.928571429	1	0.866666667
TNFRSF8 AND NOT-PTGDR2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-FUT4	0.928571429	1	0.866666667
TNFRSF8 AND NOT-ALDH6A1	0.928571429	1	0.866666667
TNFRSF8 AND NOT-CECR6	0.965517241	1	0.933333333
TNFRSF8 AND NOT-RHBDF2	0.928571429	1	0.866666667
TNFRSF8 AND NOT-BAX	0.965517241	1	0.933333333
TNFRSF8 AND NOT-MYADM	0.965517241	1	0.933333333
TNFRSF8 AND NOT-DNAJB12	0.928571429	1	0.866666667
TNFRSF8 AND NOT-LRP3	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SLC46A2	0.928571429	1	0.866666667
TNFRSF8 AND NOT-GPR82	0.965517241	1	0.933333333
TNFRSF8 AND NOT-NOTCH2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-TPST2	0.928571429	1	0.866666667
TNFRSF8 AND NOT-SMPD3	0.965517241	1	0.933333333
TNFRSF8 AND NOT-TMEM129	0.965517241	1	0.933333333
TNFRSF8 AND NOT-CEACAM4	0.965517241	1	0.933333333
TNFRSF8 AND NOT-ANPEP	0.965517241	1	0.933333333
TNFRSF8 AND NOT-BTN3A3	0.928571429	1	0.866666667
TNFRSF8 AND NOT-KLRB1	0.928571429	1	0.866666667
TNFRSF8 AND NOT-TGFBR3	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SLC35B3	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SUSD1	0.928571429	1	0.866666667
TNFRSF8 AND NOT-EXTL3	0.965517241	1	0.933333333
TNFRSF8 AND NOT-EPHA4	0.965517241	1	0.933333333
TNFRSF8 AND NOT-CTAGE5	0.965517241	1	0.933333333
TNFRSF8 AND NOT-TLR2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-PLXDC2	0.928571429	1	0.866666667
TNFRSF8 AND NOT-KCNA10	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SFT2D3	0.965517241	1	0.933333333
TNFRSF8 AND NOT-XYLT1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-CMTM5	0.965517241	1	0.933333333
TNFRSF8 AND NOT-TNFSF8	0.928571429	1	0.866666667
TNFRSF8 AND NOT-TLR5	0.965517241	1	0.933333333
TNFRSF8 AND NOT-ICAM2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-P2RY14	0.928571429	1	0.866666667
TNFRSF8 AND NOT-CLCN5	0.965517241	1	0.933333333
TNFRSF8 AND NOT-COQ2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-GRM2	0.928571429	1	0.866666667
TNFRSF8 AND NOT-MARC1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-FBXL17	0.928571429	1	0.866666667
TNFRSF8 AND NOT-SUN2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-CAMP	0.965517241	1	0.933333333
TNFRSF8 AND NOT-NRG1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-TMEM242	0.965517241	1	0.933333333
TNFRSF8 AND NOT-KLRD1	0.928571429	1	0.866666667
TNFRSF8 AND NOT-STT3A	0.928571429	1	0.866666667
TNFRSF8 AND NOT-SFXN5	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SLC11A1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-ACPP	0.965517241	1	0.933333333
TNFRSF8 AND NOT-PRR7	0.965517241	1	0.933333333

Antigen Pair Logic	F1	Precision	Recall
PLA2G2D AND NOT-GRIN3B	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-SLC16A10	0.916666667	0.942857143	0.891891892
PLA2G2D AND NOT-CLRN3	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-NXPE1	0.918918919	0.918918919	0.918918919
PLA2G2D AND NOT-LRIG3	0.914285714	0.96969697	0.864864865
PLA2G2D AND NOT-TMEM132D	0.916666667	0.942857143	0.891891892
PLA2G2D AND NOT-GPHB5	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-ABCC2	0.904109589	0.916666667	0.891891892
PLA2G2D AND NOT-WFIKKN2	0.944444444	0.971428571	0.918918919
PLA2G2D AND NOT-SEZ6	0.929577465	0.970588235	0.891891892
PLA2G2D AND NOT-CNGA1	0.927536232	1	0.864864865
PLA2G2D AND NOT-CYP4F22	0.918918919	0.918918919	0.918918919
PLA2G2D AND NOT-SLC44A3	0.904109589	0.916666667	0.891891892
PLA2G2D AND NOT-CNTN1	0.911764706	1	0.837837838
PLA2G2D AND NOT-DCST2	0.914285714	0.96969697	0.864864865
PLA2G2D AND NOT-KLHDC7A	0.914285714	0.96969697	0.864864865
PLA2G2D AND NOT-FAM3D	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-IL17RE	0.929577465	0.970588235	0.891891892
PLA2G2D AND NOT-LSMEM2	0.927536232	1	0.864864865
PLA2G2D AND NOT-ADORA1	0.929577465	0.970588235	0.891891892
PLA2G2D AND NOT-RAET1E	0.918918919	0.918918919	0.918918919
PLA2G2D AND NOT-WBSCR28	0.916666667	0.942857143	0.891891892
PLA2G2D AND NOT-TMEM139	0.931506849	0.944444444	0.918918919
PLA2G2D AND NOT-CLDN3	0.904109589	0.916666667	0.891891892
PLA2G2D AND NOT-TRPM6	0.904109589	0.916666667	0.891891892
PLA2G2D AND NOT-CST11	0.901408451	0.941176471	0.864864865
PLA2G2D AND NOT-CRY2	0.914285714	0.96969697	0.864864865
PLA2G2D AND NOT-SYT9	0.914285714	0.96969697	0.864864865
PLA2G2D AND NOT-PRIMA1	0.916666667	0.942857143	0.891891892
PLA2G2D AND NOT-CSPG4	0.901408451	0.941176471	0.864864865
PLA2G2D AND NOT-MGAT5B	0.918918919	0.918918919	0.918918919
PLA2G2D AND NOT-CD300LG	0.904109589	0.916666667	0.891891892
PLA2G2D AND NOT-TMEM190	0.929577465	0.970588235	0.891891892
PLA2G2D AND NOT-ADRA1B	0.931506849	0.944444444	0.918918919
PLA2G2D AND NOT-ATP8B3	0.918918919	0.918918919	0.918918919
PLA2G2D AND NOT-SYT6	0.904109589	0.916666667	0.891891892
PLA2G2D AND NOT-HFE2	0.901408451	0.941176471	0.864864865
PLA2G2D AND NOT-PROM2	0.904109589	0.916666667	0.891891892
PLA2G2D AND NOT-PLB1	0.916666667	0.942857143	0.891891892
PLA2G2D AND NOT-SLC38A11	0.931506849	0.944444444	0.918918919
PLA2G2D AND NOT-SLC2A12	0.901408451	0.941176471	0.864864865
PLA2G2D AND NOT-CYP2A7	0.918918919	0.918918919	0.918918919
PLA2G2D AND NOT-TMEM213	0.901408451	0.941176471	0.864864865
PLA2G2D AND NOT-CYP2E1	0.944444444	0.971428571	0.918918919
PLA2G2D AND NOT-PRUNE2	0.901408451	0.941176471	0.864864865
PLA2G2D AND NOT-SLC5A8	0.929577465	0.970588235	0.891891892
PLA2G2D AND NOT-DBH	0.931506849	0.944444444	0.918918919
PLA2G2D AND NOT-SLC16A11	0.931506849	0.944444444	0.918918919
PLA2G2D AND NOT-SPPL2C	0.944444444	0.971428571	0.918918919
PLA2G2D AND NOT-CABP7	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-CLEC4F	0.916666667	0.942857143	0.891891892
PLA2G2D AND NOT-SLC30A8	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-KCNG3	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-PTCRA	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-AGER	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-DPP6	0.916666667	0.942857143	0.891891892
PLA2G2D AND NOT-DRD4	0.914285714	0.96969697	0.864864865
PLA2G2D AND NOT-EGFR	0.929577465	0.970588235	0.891891892
PLA2G2D AND NOT-VSTM4	0.914285714	0.96969697	0.864864865
PLA2G2D AND NOT-CADM4	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-SLC5A9	0.931506849	0.944444444	0.918918919
PLA2G2D AND NOT-SPNS3	0.931506849	0.944444444	0.918918919
PLA2G2D AND NOT-ANO5	0.931506849	0.944444444	0.918918919
PLA2G2D AND NOT-EPHA5	0.914285714	0.96969697	0.864864865
PLA2G2D AND NOT-EPHA8	0.918918919	0.918918919	0.918918919
PLA2G2D AND NOT-ABCA2	0.904109589	0.916666667	0.891891892
PLA2G2D AND NOT-EXTL1	0.914285714	0.96969697	0.864864865
PLA2G2D AND NOT-F2RL1	0.929577465	0.970588235	0.891891892
PLA2G2D AND NOT-LRRC55	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-FCER1A	0.929577465	0.970588235	0.891891892
PLA2G2D AND NOT-LRRN4CL	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-SEMA3D	0.929577465	0.970588235	0.891891892
PLA2G2D AND NOT-FGFR3	0.942857143	1	0.891891892
PLA2G2D AND NOT-PLA2R1	0.914285714	0.96969697	0.864864865
PLA2G2D AND NOT-ATP10B	0.916666667	0.942857143	0.891891892
PLA2G2D AND NOT-STAB1	0.901408451	0.941176471	0.864864865

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
TNFRSF8 AND NOT-CD1D	0.965517241	1	0.933333333
TNFRSF8 AND NOT-TLR1	0.928571429	1	0.866666667
TNFRSF8 AND NOT-HGSNAT	0.965517241	1	0.933333333
TNFRSF8 AND NOT-MS4A2	0.928571429	1	0.866666667
TNFRSF8 AND NOT-LIM2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-HCST	0.965517241	1	0.933333333
TNFRSF8 AND NOT-PTGER2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-ST3GAL3	0.965517241	1	0.933333333
TNFRSF8 AND NOT-FKRP	0.928571429	1	0.866666667
TNFRSF8 AND NOT-TMEM154	0.965517241	1	0.933333333
TNFRSF8 AND NOT-CHIC2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-FAM73B	0.928571429	1	0.866666667
TNFRSF8 AND NOT-EPHA8	0.928571429	1	0.866666667
TNFRSF8 AND NOT-PCDH9	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SLC4A4	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SLC31A2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-TMEM170B	0.965517241	1	0.933333333
TNFRSF8 AND NOT-LMTK2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-BST1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-OSTM1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-STX10	0.965517241	1	0.933333333
TNFRSF8 AND NOT-UGGT1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-PTGDR	0.965517241	1	0.933333333
TNFRSF8 AND NOT-LINGO2	0.928571429	1	0.866666667
TNFRSF8 AND NOT-MCTP1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SLC2A6	0.965517241	1	0.933333333
TNFRSF8 AND NOT-NTSR2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-ZACN	0.928571429	1	0.866666667
TNFRSF8 AND NOT-VNN2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-MCHR2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-IFNAR1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-NAALAD2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-NAALADL1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KCNE3	0.965517241	1	0.933333333
TNFRSF8 AND NOT-TMEM170B	0.965517241	1	0.933333333
TNFRSF8 AND NOT-TMEM221	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-LRRC70	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SMIM6	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-NEMP2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-100132596?	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-HCN4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ERVMER34-1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMEM239	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CDH6	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MRLN	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-100507547?	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC17A4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ABCB6	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CDH7	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SCAMP3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CDH8	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MUC12	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TSPAN32	0.928571429	1	0.866666667
TNFRSF8 AND NOT-TMEM265	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KCNK7	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TSPAN2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CDH15	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CLEC3A	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CDH16	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PDZK1IP1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CHST4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-101805491?	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-LHFPL2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ADAM8	0.965517241	1	0.933333333
TNFRSF8 AND NOT-MPZL2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-GPA33	0.965517241	1	0.933333333
TNFRSF8 AND NOT-COQ7	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MSLN	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KCNMB2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ABCC4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-IGSF6	0.965517241	1	0.933333333
TNFRSF8 AND NOT-MARCH6	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TCIRG1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-B3GALT5	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-B3GNT3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TLR6	0.965517241	1	0.933333333

Antigen Pair Logic	F1	Precision	Recall
PLA2G2D AND NOT-FLT4	0.911764706	1	0.837837838
PLA2G2D AND NOT-WSCD1	0.944444444	0.971428571	0.918918919
PLA2G2D AND NOT-KCNH4	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-CRB1	0.904109589	0.916666667	0.891891892
PLA2G2D AND NOT-NPTXR	0.914285714	0.96969697	0.864864865
PLA2G2D AND NOT-SEZ6L	0.931506849	0.944444444	0.918918919
PLA2G2D AND NOT-LPAR3	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-FLRT3	0.914285714	0.96969697	0.864864865
PLA2G2D AND NOT-FLRT1	0.918918919	0.918918919	0.918918919
PLA2G2D AND NOT-FUT1	0.911764706	1	0.837837838
PLA2G2D AND NOT-FUT2	0.904109589	0.916666667	0.891891892
PLA2G2D AND NOT-FUT3	0.904109589	0.916666667	0.891891892
PLA2G2D AND NOT-FUT6	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-C19orf26	0.931506849	0.944444444	0.918918919
PLA2G2D AND NOT-MYADML2	0.914285714	0.96969697	0.864864865
PLA2G2D AND NOT-GABRD	0.916666667	0.942857143	0.891891892
PLA2G2D AND NOT-FAM19A5	0.904109589	0.916666667	0.891891892
PLA2G2D AND NOT-KLK5	0.901408451	0.941176471	0.864864865
PLA2G2D AND NOT-MRGPRX1	0.914285714	0.96969697	0.864864865
PLA2G2D AND NOT-CYP4X1	0.931506849	0.944444444	0.918918919
PLA2G2D AND NOT-PCDHB5	0.918918919	0.918918919	0.918918919
PLA2G2D AND NOT-OR1A2	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-OR2L2	0.911764706	1	0.837837838
PLA2G2D AND NOT-GCGR	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-OR10H2	0.931506849	0.944444444	0.918918919
PLA2G2D AND NOT-GDNF	0.916666667	0.942857143	0.891891892
PLA2G2D AND NOT-ADGRF1	0.916666667	0.942857143	0.891891892
PLA2G2D AND NOT-GHRHR	0.918918919	0.918918919	0.918918919
PLA2G2D AND NOT-GIPR	0.929577465	0.970588235	0.891891892
PLA2G2D AND NOT-AMHR2	0.929577465	0.970588235	0.891891892
PLA2G2D AND NOT-GJA3	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-NSG1	0.929577465	0.970588235	0.891891892
PLA2G2D AND NOT-GJB5	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-GPR162	0.918918919	0.918918919	0.918918919
PLA2G2D AND NOT-BMP10	0.904109589	0.916666667	0.891891892
PLA2G2D AND NOT-GLP1R	0.918918919	0.918918919	0.918918919
PLA2G2D AND NOT-GP2	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-GP9	0.914285714	0.96969697	0.864864865
PLA2G2D AND NOT-SLC13A5	0.918918919	0.918918919	0.918918919
PLA2G2D AND NOT-IZUMO1	0.916666667	0.942857143	0.891891892
PLA2G2D AND NOT-GPR20	0.931506849	0.944444444	0.918918919
PLA2G2D AND NOT-GPR22	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-GPR25	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-DNAJC5G	0.931506849	0.944444444	0.918918919
PLA2G2D AND NOT-CDH19	0.942857143	1	0.891891892
PLA2G2D AND NOT-CYP4V2	0.916666667	0.942857143	0.891891892
PLA2G2D AND NOT-GPR35	0.914285714	0.96969697	0.864864865
PLA2G2D AND NOT-GPR39	0.916666667	0.942857143	0.891891892
PLA2G2D AND NOT-TUSC5	0.916666667	0.942857143	0.891891892
PLA2G2D AND NOT-GRID2	0.918918919	0.918918919	0.918918919
PLA2G2D AND NOT-ABO	0.931506849	0.944444444	0.918918919
PLA2G2D AND NOT-GRIN2C	0.916666667	0.942857143	0.891891892
PLA2G2D AND NOT-GRM3	0.904109589	0.916666667	0.891891892
PLA2G2D AND NOT-ZDHHC1	0.944444444	0.971428571	0.918918919
PLA2G2D AND NOT-GUCA2A	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-GYPE	0.918918919	0.918918919	0.918918919
PLA2G2D AND NOT-GUCY2D	0.916666667	0.942857143	0.891891892
PLA2G2D AND NOT-HCRTR1	0.957746479	1	0.918918919
PLA2G2D AND NOT-KCNIP3	0.929577465	0.970588235	0.891891892
PLA2G2D AND NOT-HPN	0.904109589	0.916666667	0.891891892
PLA2G2D AND NOT-HSD17B2	0.918918919	0.918918919	0.918918919
PLA2G2D AND NOT-ACACB	0.931506849	0.944444444	0.918918919
PLA2G2D AND NOT-HTR1A	0.904109589	0.916666667	0.891891892
PLA2G2D AND NOT-HTR1B	0.904109589	0.916666667	0.891891892
PLA2G2D AND NOT-HTR6	0.957746479	1	0.918918919
PLA2G2D AND NOT-FFAR4	0.931506849	0.944444444	0.918918919
PLA2G2D AND NOT-LRRC66	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-MOGAT3	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-TMEM255B	0.914285714	0.96969697	0.864864865
PLA2G2D AND NOT-IHH	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-IL13	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-AQP2	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-AQP5	0.944444444	0.971428571	0.918918919
PLA2G2D AND NOT-ITGA2B	0.931506849	0.944444444	0.918918919
PLA2G2D AND NOT-ITGA9	0.914285714	0.96969697	0.864864865
PLA2G2D AND NOT-ABCC6	0.929577465	0.970588235	0.891891892

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
TNFRSF8 AND NOT-ABCA7	0.965517241	1	0.933333333
TNFRSF8 AND NOT-CACNG3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CACNG2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-BTN3A3	0.928571429	1	0.866666667
TNFRSF8 AND NOT-CEPT1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-WFDC2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CLEC10A	0.928571429	1	0.866666667
TNFRSF8 AND NOT-SLC25A17	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC9A6	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SEMA6B	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SEMA4D	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SEMA4B	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-POMT1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ERLIN1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CGRRF1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-B3GNT2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CLDN16	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-RRH	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CORIN	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CERS1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-NRG3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC12A7	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PHTF1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC17A3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-VAMP5	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-OR5I1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CYSLTR1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-CFTR	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CGA	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CEACAM3	0.965517241	1	0.933333333
TNFRSF8 AND NOT-HCST	0.965517241	1	0.933333333
TNFRSF8 AND NOT-CEACAM7	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GPR83	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MMP24	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-OCLM	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CEACAM4	0.965517241	1	0.933333333
TNFRSF8 AND NOT-ADCY2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MAN1A2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GPR75	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-OS9	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-LMAN2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-CKAP4	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SLC38A3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC27A4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC27A3	0.965517241	1	0.933333333
TNFRSF8 AND NOT-LILRB4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ATF7	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TDRKH	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-UPK1A	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC35D2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-LECT1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMEM115	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ADAM30	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ADAMTS13	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CACFD1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-BTN2A1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PTPRT	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC7A9	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-HHLA2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-C14orf1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC2A6	0.965517241	1	0.933333333
TNFRSF8 AND NOT-CHIT1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ADCY5	0.928571429	1	0.866666667
TNFRSF8 AND NOT-PRRT2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PTGDR2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-HRH3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MRAP2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-STX1B	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KLHL2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-GLMP	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SCN11A	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CYP4F8	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CHRM1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CHRM2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SCAMP4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GPR182	0.933333333	0.93333333	0.933333333

Antigen Pair Logic	F1	Precision	Recall
PLA2G2D AND NOT-KCNA5	0.944444444	0.971428571	0.918918919
PLA2G2D AND NOT-KCNA6	0.931506849	0.944444444	0.918918919
PLA2G2D AND NOT-KCNC2	0.904109589	0.916666667	0.891891892
PLA2G2D AND NOT-TEX38	0.901408451	0.941176471	0.864864865
PLA2G2D AND NOT-KCNF1	0.929577465	0.970588235	0.891891892
PLA2G2D AND NOT-ENHO	0.929577465	0.970588235	0.891891892
PLA2G2D AND NOT-KCNJ3	0.918918919	0.918918919	0.918918919
PLA2G2D AND NOT-KCNJ4	0.929577465	0.970588235	0.891891892
PLA2G2D AND NOT-KCNJ12	0.929577465	0.970588235	0.891891892
PLA2G2D AND NOT-KCNK3	0.931506849	0.944444444	0.918918919
PLA2G2D AND NOT-KIR2DL1	0.904109589	0.916666667	0.891891892
PLA2G2D AND NOT-C10orf99	0.918918919	0.918918919	0.918918919
PLA2G2D AND NOT-GLTPD2	0.929577465	0.970588235	0.891891892
PLA2G2D AND NOT-SCIMP	0.911764706	1	0.837837838
PLA2G2D AND NOT-RPRML	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-FAM132A	0.916666667	0.942857143	0.891891892
PLA2G2D AND NOT-OR51I2	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-OR52D1	0.916666667	0.942857143	0.891891892
PLA2G2D AND NOT-C14orf180	0.931506849	0.944444444	0.918918919
PLA2G2D AND NOT-KLHL31	0.904109589	0.916666667	0.891891892
PLA2G2D AND NOT-LRP4	0.944444444	0.971428571	0.918918919
PLA2G2D AND NOT-LTK	0.914285714	0.96969697	0.864864865
PLA2G2D AND NOT-BCAM	0.901408451	0.941176471	0.864864865
PLA2G2D AND NOT-TACSTD2	0.904109589	0.916666667	0.891891892
PLA2G2D AND NOT-MAG	0.914285714	0.96969697	0.864864865
PLA2G2D AND NOT-ARSE	0.914285714	0.96969697	0.864864865
PLA2G2D AND NOT-MC5R	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-ADAM11	0.916666667	0.942857143	0.891891892
PLA2G2D AND NOT-MIP	0.918918919	0.918918919	0.918918919
PLA2G2D AND NOT-MLN	0.901408451	0.941176471	0.864864865
PLA2G2D AND NOT-CTRB2	0.904109589	0.916666667	0.891891892
PLA2G2D AND NOT-TMEM151B	0.904109589	0.916666667	0.891891892
PLA2G2D AND NOT-OR10C1	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-MUC3A	0.929577465	0.970588235	0.891891892
PLA2G2D AND NOT-MUC4	0.929577465	0.970588235	0.891891892
PLA2G2D AND NOT-ASTN1	0.904109589	0.916666667	0.891891892
PLA2G2D AND NOT-ATP1B2	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-NPR3	0.904109589	0.916666667	0.891891892
PLA2G2D AND NOT-NPY5R	0.904109589	0.916666667	0.891891892
PLA2G2D AND NOT-NTF3	0.904109589	0.916666667	0.891891892
PLA2G2D AND NOT-NTRK1	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-NTRK2	0.944444444	0.971428571	0.918918919
PLA2G2D AND NOT-NTRK3	0.918918919	0.918918919	0.918918919
PLA2G2D AND NOT-ATP4A	0.929577465	0.970588235	0.891891892
PLA2G2D AND NOT-OMG	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-OPCML	0.927536232	1	0.864864865
PLA2G2D AND NOT-OPRD1	0.914285714	0.96969697	0.864864865
PLA2G2D AND NOT-OPRL1	0.931506849	0.944444444	0.918918919
PLA2G2D AND NOT-P2RY2	0.916666667	0.942857143	0.891891892
PLA2G2D AND NOT-P2RY4	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-CD207	0.944444444	0.971428571	0.918918919
PLA2G2D AND NOT-PODXL2	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-GALNT9	0.944444444	0.971428571	0.918918919
PLA2G2D AND NOT-CALY	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-KCNK4	0.931506849	0.944444444	0.918918919
PLA2G2D AND NOT-PCDH1	0.916666667	0.942857143	0.891891892
PLA2G2D AND NOT-CLDN18	0.918918919	0.918918919	0.918918919
PLA2G2D AND NOT-CRIM1	0.929577465	0.970588235	0.891891892
PLA2G2D AND NOT-EMCN	0.904109589	0.916666667	0.891891892
PLA2G2D AND NOT-FXYD3	0.904109589	0.916666667	0.891891892
PLA2G2D AND NOT-PLP1	0.944444444	0.971428571	0.918918919
PLA2G2D AND NOT-S1PR5	0.918918919	0.918918919	0.918918919
PLA2G2D AND NOT-FXYD7	0.929577465	0.970588235	0.891891892
PLA2G2D AND NOT-FGFRL1	0.901408451	0.941176471	0.864864865
PLA2G2D AND NOT-ATP7B	0.904109589	0.916666667	0.891891892
PLA2G2D AND NOT-GPR88	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-KCNK10	0.914285714	0.96969697	0.864864865
PLA2G2D AND NOT-GPR173	0.929577465	0.970588235	0.891891892
PLA2G2D AND NOT-WNT4	0.918918919	0.918918919	0.918918919
PLA2G2D AND NOT-PON3	0.916666667	0.942857143	0.891891892
PLA2G2D AND NOT-RNF186	0.904109589	0.916666667	0.891891892
PLA2G2D AND NOT-SDK2	0.914285714	0.96969697	0.864864865
PLA2G2D AND NOT-NXPE4	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-BEST2	0.927536232	1	0.864864865
PLA2G2D AND NOT-GDPD2	0.944444444	0.971428571	0.918918919
PLA2G2D AND NOT-MS4A12	0.906666667	0.894736842	0.918918919

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
TNFRSF8 AND NOT-CHRM3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MGAT4A	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CHRM4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CHRM5	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CHRNA1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CMTM1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CYP2U1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MFSD3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CHRNA3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CHRNA4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CHRNB1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SIGLEC11	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CHRNB3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CHRNB4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ERMAP	0.965517241	1	0.933333333
TNFRSF8 AND NOT-LRRC37B	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CHRNG	0.965517241	1	0.933333333
TNFRSF8 AND NOT-PKD1L2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC25A25	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ELFN2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMEM132B	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLITRK1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-C1QTNF6	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ADCY8	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MARCH3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GPR146	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FCRL1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FCRL3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC5A11	0.928571429	1	0.866666667
TNFRSF8 AND NOT-TNFRSF13C	0.928571429	1	0.866666667
TNFRSF8 AND NOT-SLC22A12	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CMTM5	0.965517241	1	0.933333333
TNFRSF8 AND NOT-ABHD15	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PANX3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GRIN3B	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MAS1L	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC18B1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-CATSPER1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-CATSPER2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MRGPRX2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MRGPRX4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GALNT15	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CLCN3	0.965517241	1	0.933333333
TNFRSF8 AND NOT-CLCN4	0.965517241	1	0.933333333
TNFRSF8 AND NOT-ANTXR2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GPR62	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CLCN5	0.965517241	1	0.933333333
TNFRSF8 AND NOT-CLCN6	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FAM24A	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ZFYVE27	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PDZD8	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC36A4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CYP2R1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CLN5	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-NXPE1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-AMICA1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-TMEM52B	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CLPTM1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMEM132D	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ANO4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PLD4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ADSSL1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC51B	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CCR3	0.965517241	1	0.933333333
TNFRSF8 AND NOT-NIPA1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CMTM3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-LTB4R	0.928571429	1	0.866666667
TNFRSF8 AND NOT-TMEM170A	0.965517241	1	0.933333333
TNFRSF8 AND NOT-CD300LB	0.965517241	1	0.933333333
TNFRSF8 AND NOT-B4GALNT2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC43A2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-MIEF2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SLC5A10	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FAM210A	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FAM69C	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CNGA3	0.933333333	0.93333333	0.933333333

Antigen Pair Logic	F1	Precision	Recall
PLA2G2D AND NOT-ST7L	0.916666667	0.942857143	0.891891892
PLA2G2D AND NOT-RHBDL2	0.916666667	0.942857143	0.891891892
PLA2G2D AND NOT-KIRREL	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-TMEM40	0.918918919	0.918918919	0.918918919
PLA2G2D AND NOT-SPTLC3	0.929577465	0.970588235	0.891891892
PLA2G2D AND NOT-VNN3	0.914285714	0.96969697	0.864864865
PLA2G2D AND NOT-NPY4R	0.931506849	0.944444444	0.918918919
PLA2G2D AND NOT-AVPR2	0.944444444	0.971428571	0.918918919
PLA2G2D AND NOT-TNFRSF19	0.916666667	0.942857143	0.891891892
PLA2G2D AND NOT-SVOP	0.918918919	0.918918919	0.918918919
PLA2G2D AND NOT-SLC30A10	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-PCDHGC5	0.931506849	0.944444444	0.918918919
PLA2G2D AND NOT-PCDHGA11	0.931506849	0.944444444	0.918918919
PLA2G2D AND NOT-PCDHB10	0.901408451	0.941176471	0.864864865
PLA2G2D AND NOT-PCDHB9	0.918918919	0.918918919	0.918918919
PLA2G2D AND NOT-PCDHB7	0.916666667	0.942857143	0.891891892
PLA2G2D AND NOT-PCDHB6	0.904109589	0.916666667	0.891891892
PLA2G2D AND NOT-PCDHA10	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-PCDHA5	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-PROS1	0.929577465	0.970588235	0.891891892
PLA2G2D AND NOT-PRRG1	0.914285714	0.96969697	0.864864865
PLA2G2D AND NOT-LTB4R2	0.931506849	0.944444444	0.918918919
PLA2G2D AND NOT-RPRM	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-MUC13	0.918918919	0.918918919	0.918918919
PLA2G2D AND NOT-NMUR2	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-PYY	0.944444444	0.971428571	0.918918919
PLA2G2D AND NOT-CHRNA10	0.914285714	0.96969697	0.864864865
PLA2G2D AND NOT-ANO2	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-CYSLTR2	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-JPH2	0.904109589	0.916666667	0.891891892
PLA2G2D AND NOT-GJC2	0.944444444	0.971428571	0.918918919
PLA2G2D AND NOT-ADGRG6	0.942857143	1	0.891891892
PLA2G2D AND NOT-PTGER1	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-TTYH1	0.901408451	0.941176471	0.864864865
PLA2G2D AND NOT-GJD2	0.931506849	0.944444444	0.918918919
PLA2G2D AND NOT-KIAA1161	0.914285714	0.96969697	0.864864865
PLA2G2D AND NOT-PCDH19	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-PTK7	0.901408451	0.941176471	0.864864865
PLA2G2D AND NOT-SYT13	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-SLC7A14	0.904109589	0.916666667	0.891891892
PLA2G2D AND NOT-SEMA4G	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-SCUBE2	0.916666667	0.942857143	0.891891892
PLA2G2D AND NOT-ADGRB3	0.918918919	0.918918919	0.918918919
PLA2G2D AND NOT-PTPRZ1	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-PVRL1	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-BCHE	0.929577465	0.970588235	0.891891892
PLA2G2D AND NOT-NTN4	0.929577465	0.970588235	0.891891892
PLA2G2D AND NOT-CACNG7	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-PRPH2	0.929577465	0.970588235	0.891891892
PLA2G2D AND NOT-RHAG	0.918918919	0.918918919	0.918918919
PLA2G2D AND NOT-NYX	0.929577465	0.970588235	0.891891892
PLA2G2D AND NOT-ROM1	0.901408451	0.941176471	0.864864865
PLA2G2D AND NOT-BDKRB1	0.904109589	0.916666667	0.891891892
PLA2G2D AND NOT-SCN7A	0.929577465	0.970588235	0.891891892
PLA2G2D AND NOT-SCNN1A	0.904109589	0.916666667	0.891891892
PLA2G2D AND NOT-SCNN1B	0.929577465	0.970588235	0.891891892
PLA2G2D AND NOT-SCTR	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-SDC1	0.927536232	1	0.864864865
PLA2G2D AND NOT-SFRP1	0.944444444	0.971428571	0.918918919
PLA2G2D AND NOT-CHST8	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-SGCD	0.929577465	0.970588235	0.891891892
PLA2G2D AND NOT-ITSN1	0.914285714	0.96969697	0.864864865
PLA2G2D AND NOT-LPPR2	0.901408451	0.941176471	0.864864865
PLA2G2D AND NOT-IL25	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-SLC13A3	0.929577465	0.970588235	0.891891892
PLA2G2D AND NOT-SLC1A7	0.914285714	0.96969697	0.864864865
PLA2G2D AND NOT-SLC5A1	0.927536232	1	0.864864865
PLA2G2D AND NOT-SLC6A11	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-SLC6A13	0.918918919	0.918918919	0.918918919
PLA2G2D AND NOT-SLC7A4	0.914285714	0.96969697	0.864864865
PLA2G2D AND NOT-SLC16A2	0.911764706	1	0.837837838
PLA2G2D AND NOT-SLC18A2	0.918918919	0.918918919	0.918918919
PLA2G2D AND NOT-BMPR1B	0.931506849	0.944444444	0.918918919
PLA2G2D AND NOT-SOX1	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-SYT5	0.914285714	0.96969697	0.864864865
PLA2G2D AND NOT-TEK	0.916666667	0.942857143	0.891891892

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
TNFRSF8 AND NOT-CNGA4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-OR1I1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMCO2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GJB4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-DCST2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KLHDC7A	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SYT2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GOLT1A	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMEM125	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-DRAM2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-NKAIN4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FITM2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PROKR2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CST9L	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CST9	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MBOAT2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CNTNAP5	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-COL11A2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SGPP2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ACVR1C	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SPATA3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMEM198	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMEM182	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-COL17A1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KCNH8	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-DNAJC19	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-DCBLD2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC31A2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-RTP1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-LSMEM2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-AASDH	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SLC9B2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLCO6A1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-UGT3A1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FAM173B	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TAAR1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-COX10	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-COX15	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-DPCR1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-OR6B1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CLDN3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CLDN23	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CR1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-HGSNAT	0.965517241	1	0.933333333
TNFRSF8 AND NOT-FAM69B	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ADGRG4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MUC17	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ASB11	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TRPM6	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ROMO1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SIRPA	0.965517241	1	0.933333333
TNFRSF8 AND NOT-CRY2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SAMD8	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-VTI1A	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MUC15	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FAM76B	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CSF2RA	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMEM86A	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CSF3R	0.965517241	1	0.933333333
TNFRSF8 AND NOT-ALG10B	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-BEST3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-B3GLCT	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-LRFN5	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-NRG4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-C16orf92	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GSG1L	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMED6	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CD300LF	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SLC47A2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ODF4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ADRA1D	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMEM199	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-VSIG10L	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-C19orf18	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-DPY19L3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ADRA1B	0.933333333	0.93333333	0.933333333

Antigen Pair Logic	F1	Precision	Recall
PLA2G2D AND NOT-TPO	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-TLCD2	0.916666667	0.942857143	0.891891892
PLA2G2D AND NOT-FAM198A	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-VIP	0.911764706	1	0.837837838
PLA2G2D AND NOT-VIPR2	0.904109589	0.916666667	0.891891892
PLA2G2D AND NOT-CACNA1S	0.931506849	0.944444444	0.918918919
PLA2G2D AND NOT-SEMA3B	0.901408451	0.941176471	0.864864865
PLA2G2D AND NOT-NOX5	0.916666667	0.942857143	0.891891892
PLA2G2D AND NOT-LRFN3	0.916666667	0.942857143	0.891891892
PLA2G2D AND NOT-EPHX3	0.904109589	0.916666667	0.891891892
PLA2G2D AND NOT-BTNL8	0.916666667	0.942857143	0.891891892
PLA2G2D AND NOT-SEMA6D	0.914285714	0.96969697	0.864864865
PLA2G2D AND NOT-FRAS1	0.944444444	0.971428571	0.918918919
PLA2G2D AND NOT-BPIFB2	0.901408451	0.941176471	0.864864865
PLA2G2D AND NOT-PRRT1	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-LRRC3	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-PVRL4	0.904109589	0.916666667	0.891891892
PLA2G2D AND NOT-AMN	0.918918919	0.918918919	0.918918919
PLA2G2D AND NOT-RTP3	0.918918919	0.918918919	0.918918919
PLA2G2D AND NOT-MS4A8	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-SLC4A9	0.914285714	0.96969697	0.864864865
PLA2G2D AND NOT-KCNK16	0.918918919	0.918918919	0.918918919
PLA2G2D AND NOT-KIAA1109	0.918918919	0.918918919	0.918918919
PLA2G2D AND NOT-TMEM246	0.944444444	0.971428571	0.918918919
PLA2G2D AND NOT-SOAT2	0.929577465	0.970588235	0.891891892
PLA2G2D AND NOT-ADGRA1	0.904109589	0.916666667	0.891891892
PLA2G2D AND NOT-ST6GAL2	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-KIRREL3	0.931506849	0.944444444	0.918918919
PLA2G2D AND NOT-ABHD1	0.914285714	0.96969697	0.864864865
PLA2G2D AND NOT-GALR3	0.918918919	0.918918919	0.918918919
PLA2G2D AND NOT-PTPN5	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-ABCC11	0.931506849	0.944444444	0.918918919
PLA2G2D AND NOT-LY6D	0.904109589	0.916666667	0.891891892
PLA2G2D AND NOT-CAV3	0.942857143	1	0.891891892
PLA2G2D AND NOT-UNC5C	0.911764706	1	0.837837838
PLA2G2D AND NOT-SLC4A4	0.927536232	1	0.864864865
PLA2G2D AND NOT-B3GALT1	0.911764706	1	0.837837838
PLA2G2D AND NOT-ADAM23	0.916666667	0.942857143	0.891891892
PLA2G2D AND NOT-GALR2	0.901408451	0.941176471	0.864864865
PLA2G2D AND NOT-PROM1	0.916666667	0.942857143	0.891891892
PLA2G2D AND NOT-CCKAR	0.901408451	0.941176471	0.864864865
PLA2G2D AND NOT-CACNA1I	0.904109589	0.916666667	0.891891892
PLA2G2D AND NOT-TSPAN18	0.914285714	0.96969697	0.864864865
PLA2G2D AND NOT-TM4SF5	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-SLC13A2	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-ANGPTL1	0.931506849	0.944444444	0.918918919
PLA2G2D AND NOT-CLDN10	0.904109589	0.916666667	0.891891892
PLA2G2D AND NOT-CLDN1	0.918918919	0.918918919	0.918918919
PLA2G2D AND NOT-TMEM257	0.904109589	0.916666667	0.891891892
PLA2G2D AND NOT-MYADM	0.901408451	0.941176471	0.864864865
PLA2G2D AND NOT-IL1RL1	0.916666667	0.942857143	0.891891892
PLA2G2D AND NOT-DSEL	0.929577465	0.970588235	0.891891892
PLA2G2D AND NOT-LARGE	0.916666667	0.942857143	0.891891892
PLA2G2D AND NOT-CDHR1	0.918918919	0.918918919	0.918918919
PLA2G2D AND NOT-DHRS3	0.901408451	0.941176471	0.864864865
PLA2G2D AND NOT-SLC22A6	0.918918919	0.918918919	0.918918919
PLA2G2D AND NOT-NRXN3	0.914285714	0.96969697	0.864864865
PLA2G2D AND NOT-LRRC4B	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-OPN4	0.918918919	0.918918919	0.918918919
PLA2G2D AND NOT-NTN1	0.918918919	0.918918919	0.918918919
PLA2G2D AND NOT-NCR1	0.931506849	0.944444444	0.918918919
PLA2G2D AND NOT-ITM2B	0.906666667	0.894736842	0.918918919
PLA2G2D AND NOT-ENTPD2	0.918918919	0.918918919	0.918918919
PLA2G2D AND NOT-ENTPD3	0.931506849	0.944444444	0.918918919
PLA2G2D AND NOT-MFAP3L	0.957746479	1	0.918918919
Adenocarcinoma|Lung Neoplasms (Lung A)			
SFTPD AND NOT-IL3RA	0.918918919	0.894736842	0.944444444
SFTPD AND NOT-CDH5	0.941176471	1	0.888888889
SFTPC AND NOT-RAMP2	0.909090909	1	0.833333333
SFTPD AND NOT-RAMP2	0.941176471	1	0.888888889
SFTPD AND NOT-SIRPB1	0.918918919	0.894736842	0.944444444
SFTPD AND NOT-ABCA8	0.941176471	1	0.888888889
SLC34A2 AND NOT-DLL1	0.914285714	0.941176471	0.888888889
SFTPC AND NOT-LYVE1	0.909090909	1	0.833333333
SFTPD AND NOT-LYVE1	0.971428571	1	0.944444444
SFTPD AND NOT-PNPLA6	0.918918919	0.894736842	0.944444444

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
TNFRSF8 AND NOT-FAM187B	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ATP8B3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SYT6	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-HFE2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FAM163A	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MFSD4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PM20D1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC30A7	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KNCN	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-DCST1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SHISA4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC9B1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-NFAM1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-KANSL1L	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FAM132B	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ARL6IP6	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC38A11	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GPR155	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FAM19A4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-XXYLT1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-CUX1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-IGSF11	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CX3CR1	0.928571429	1	0.866666667
TNFRSF8 AND NOT-NFXL1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PAQR3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MGAT4D	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CXADR	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-THAP6	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC25A48	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MARVELD2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CYBB	0.928571429	1	0.866666667
TNFRSF8 AND NOT-ADRB1	0.928571429	1	0.866666667
TNFRSF8 AND NOT-SLC2A12	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MBOAT1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-RNF217	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-NKAIN2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CYP1A1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-VKORC1L1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CYP2A6	0.928571429	1	0.866666667
TNFRSF8 AND NOT-ADRB2	0.928571429	1	0.866666667
TNFRSF8 AND NOT-CYP3A7	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CYP2A13	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CYP2C8	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CYP2E1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PEBP4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMEM65	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMEM74	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CYP4A11	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-LINGO2	0.928571429	1	0.866666667
TNFRSF8 AND NOT-FREM1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FMR1NB	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FAAH2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ZDHHC15	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CLEC12A	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SLC5A8	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GPR180	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-STOML3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FITM1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMEM30B	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MFSD6L	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMEM92	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-RHBDL3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SPPL2C	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-DCC	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SYNE4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TOR1AIP2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-DCT	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PAQR7	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-LRRN4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMPRSS6	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ADGRF3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-OXER1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SPTSSB	0.965517241	1	0.933333333
TNFRSF8 AND NOT-CHST13	0.965517241	1	0.933333333
TNFRSF8 AND NOT-ADGRA3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SGMS2	0.965517241	1	0.933333333
SFTPD AND NOT-SLC25A25	0.944444444	0.944444444	0.944444444
SFTPD AND NOT-CYYR1	0.914285714	0.941176471	0.888888889
SFTPD AND NOT-BTNL9	0.971428571	1	0.944444444
SFTPC AND NOT-CLEC14A	0.909090909	1	0.833333333
SFTPD AND NOT-CLEC14A	0.971428571	1	0.944444444
SFTPC AND NOT-AGER	0.909090909	1	0.833333333
SFTPD AND NOT-AGER	0.944444444	0.944444444	0.944444444
SFTPD AND NOT-HBEGF	0.944444444	0.944444444	0.944444444
SFTPD AND NOT-AGTR1	0.918918919	0.894736842	0.944444444
SFTPD AND NOT-S1PR1	0.914285714	0.941176471	0.888888889
SFTPD AND NOT-EDNRB	0.918918919	0.894736842	0.944444444
SFTPD AND NOT-ADCY4	0.941176471	1	0.888888889
SFTPD AND NOT-MCEMP1	0.914285714	0.941176471	0.888888889
SFTPD AND NOT-EMP2	0.941176471	1	0.888888889
SFTPD AND NOT-ENG	0.941176471	1	0.888888889
BAX AND NOT-ABCA2	0.914285714	0.941176471	0.888888889
SFTPD AND NOT-RNF182	0.914285714	0.941176471	0.888888889
SFTPD AND NOT-FGFR4	0.971428571	1	0.944444444
SFTPD AND NOT-ADGRL2	0.909090909	1	0.833333333
SFTPD AND NOT-FUT1	0.914285714	0.941176471	0.888888889
SFTPD AND NOT-ACKR1	0.914285714	0.941176471	0.888888889
SFTPD AND NOT-ABI3BP	0.909090909	1	0.833333333
SFTPD AND NOT-LRRC32	0.971428571	1	0.944444444
SFTPD AND NOT-GPM6A	0.944444444	0.944444444	0.944444444
SFTPD AND NOT-IGSF10	0.971428571	1	0.944444444
SFTPD AND NOT-GRIA1	0.914285714	0.941176471	0.888888889
SFTPD AND NOT-ACACB	0.909090909	1	0.833333333
SFTPD AND NOT-KCNT2	0.918918919	0.894736842	0.944444444
SFTPD AND NOT-IL3RA	0.918918919	0.894736842	0.944444444
SFTPD AND NOT-CXCR2	0.914285714	0.941176471	0.888888889
SFTPD AND NOT-PEAR1	0.941176471	1	0.888888889
SFTPC AND NOT-KCNK3	0.909090909	1	0.833333333
SFTPD AND NOT-KCNK3	0.914285714	0.941176471	0.888888889
SFTPD AND NOT-SERTM1	0.914285714	0.941176471	0.888888889
SFTPD AND NOT-MME	0.971428571	1	0.944444444
SFTPC AND NOT-MARCH11	0.909090909	1	0.833333333
SFTPD AND NOT-NOTCH4	0.941176471	1	0.888888889
SFTPD AND NOT-CLEC1A	0.914285714	0.941176471	0.888888889
SFTPD AND NOT-PCDH12	0.909090909	1	0.833333333
SFTPD AND NOT-PECAM1	0.971428571	1	0.944444444
BAX AND NOT-ABCB1	0.918918919	0.894736842	0.944444444
SFTPD AND NOT-FXYD6	0.909090909	1	0.833333333
SFTPD AND NOT-LRRN3	0.941176471	1	0.888888889
BAX AND NOT-PRPF38B	0.909090909	1	0.833333333
SFTPC AND NOT-TMEM74B	0.909090909	1	0.833333333
SFTPD AND NOT-TMEM74B	0.918918919	0.894736842	0.944444444
SFTPD AND NOT-PRRG1	0.941176471	1	0.888888889
SFTPD AND NOT-HEG1	0.941176471	1	0.888888889
SFTPD AND NOT-SEMA6A	0.909090909	1	0.833333333
SFTPD AND NOT-PTPRM	0.944444444	0.944444444	0.944444444
SFTPD AND NOT-JAM2	0.941176471	1	0.888888889
SFTPD AND NOT-RXFP1	0.918918919	0.894736842	0.944444444
SFTPD AND NOT-SELE	0.909090909	1	0.833333333
SFTPC AND NOT-ECSCR	0.909090909	1	0.833333333
SFTPD AND NOT-ECSCR	0.971428571	1	0.944444444
SFTPD AND NOT-SGCG	0.971428571	1	0.944444444
SFTPD AND NOT-TEK	0.914285714	0.941176471	0.888888889
SFTPD AND NOT-TGFBR3	0.971428571	1	0.944444444
SFTPD AND NOT-THBD	0.909090909	1	0.833333333
SFTPD AND NOT-TIE1	0.941176471	1	0.888888889
SFTPD AND NOT-CLDN5	0.909090909	1	0.833333333
SFTPD AND TMEM258	0.944444444	0.944444444	0.944444444
SFTPD AND NOT-CA4	0.944444444	0.944444444	0.944444444
SFTPD AND NOT-C1orf115	0.971428571	1	0.944444444
SFTPD AND NOT-MYCT1	0.941176471	1	0.888888889
SFTPD AND NOT-ADTRP	0.918918919	0.894736842	0.944444444
SFTPD AND NOT-AOC3	0.941176471	1	0.888888889
SFTPD AND NOT-IL18R1	0.914285714	0.941176471	0.888888889
SFTPD AND NOT-KL	0.909090909	1	0.833333333
SFTPD AND NOT-FAM189A2	0.909090909	1	0.833333333
SFTPD AND NOT-ITM2A	0.941176471	1	0.888888889
SFTPD AND NOT-CD36	0.918918919	0.894736842	0.944444444
SFTPD AND NOT-ACVRL1	0.941176471	1	0.888888889
TMEM258 AND NOT-TGFBR3	0.914285714	0.941176471	0.888888889
Oligodendroglioma (Oligodendroglioma)			
NLGN3 AND NOT-FOLR2	0.933333333	0.933333333	0.933333333

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
TNFRSF8 AND NOT-GALNTL5	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PKD1L1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC30A8	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KCNV2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CLEC4C	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-HTR3C	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KCNG3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CLYBL	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FAM9C	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SPTSSA	0.965517241	1	0.933333333
TNFRSF8 AND NOT-ABHD3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-DHCR24	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-DIO1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-DIO3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-DNASE1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-DPP6	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-DRD1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-DRD3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-DRD4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-DSC3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-DSCAM	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-DTNA	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-AGTR2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-E2F5	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ECM1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-S1PR1	0.928571429	1	0.866666667
TNFRSF8 AND NOT-S1PR3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SCAMP5	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-EFNA5	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-EFNB3	0.965517241	1	0.933333333
TNFRSF8 AND NOT-EGF	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CELSR2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MEGF6	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MEGF9	0.965517241	1	0.933333333
TNFRSF8 AND NOT-PPAPDC1A	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MPZL3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PIANP	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-VSTM4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GRAMD2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ELANE	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CADM4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMEM61	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FAM209A	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-APLF	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMEM17	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GABRR3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PLD6	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-C17orf74	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC39A11	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMEM154	0.965517241	1	0.933333333
TNFRSF8 AND NOT-TAPT1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-DNAJC18	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GAPT	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SLC29A1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-HTRA4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-C9orf91	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SUSD3	0.928571429	1	0.866666667
TNFRSF8 AND NOT-ANO5	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-EPHA3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-EPHA4	0.965517241	1	0.933333333
TNFRSF8 AND NOT-EPHA5	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-EPHA8	0.928571429	1	0.866666667
TNFRSF8 AND NOT-EPHB4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-EPHB6	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CLN8	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-LVRN	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC36A1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-ERBB2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-ERBB3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ERBB4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-EREG	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ERN1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-ETFDH	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-EVC	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-EXTL3	0.965517241	1	0.933333333
TNFRSF8 AND NOT-F10	0.933333333	0.93333333	0.933333333

Antigen Pair Logic	F1	Precision	Recall
NLGN3 AND NOT-SLAMF7	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SDC1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TNFRSF8	0.903225806	0.875	0.933333333
NLGN3 AND NOT-IL11RA	0.903225806	0.875	0.933333333
NLGN3 AND NOT-MUC1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-KDR	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-IGF1R	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ALK	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ERBB3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TNFRSF10B	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-B4GALNT1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-GPA33	0.903225806	0.875	0.933333333
NLGN3 AND NOT-MUC13	0.903225806	0.875	0.933333333
NLGN3 AND NOT-EPHA3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-MAGEA1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-IL3RA	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ROR1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TNFSF11	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-MUC17	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ITGB3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-FOLH1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-EPHA2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-CA9	0.903225806	0.875	0.933333333
NLGN3 AND NOT-MS4A1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TNFRSF10A	0.903225806	0.875	0.933333333
NLGN3 AND NOT-MUC16	0.903225806	0.875	0.933333333
NLGN3 AND NOT-MLANA	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CD160	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NAALADL1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-KCNE3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TMEM221	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-LRRC70	0.903225806	0.875	0.933333333
NLGN3 AND NOT-SMIM6	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-NEMP2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ARMCX4	0.903225806	0.875	0.933333333
NLGN3 AND NOT-100132596?	0.903225806	0.875	0.933333333
NLGN3 AND NOT-BCL2L10	0.903225806	0.875	0.933333333
NLGN3 AND NOT-PIGK	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ERVMER34-1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TMEM239	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CDH4	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-APELA	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ZMYM6NB	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-MRLN	0.903225806	0.875	0.933333333
NLGN3 AND NOT-SMLR1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-MICA	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SLC17A4	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ABCC5	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ABCB6	0.903225806	0.875	0.933333333
NLGN3 AND NOT-SCAMP3	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-MUC12	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TMEM265	0.903225806	0.875	0.933333333
NLGN3 AND NOT-KCNK7	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TSPAN5	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TSPAN2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TSPAN1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CDH12	0.903225806	0.875	0.933333333
NLGN3 AND NOT-PREB	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-CLEC3A	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ADGRG2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-PDZK1IP1	0.903225806	0.875	0.933333333
SLC4A4 AND NOT-PDZK1IP1	0.928571429	1	0.866666667
NLGN3 AND NOT-CDH17	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-LPCAT3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CHST4	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CDH18	0.903225806	0.875	0.933333333
NLGN3 AND NOT-DHRS9	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TENM1	0.903225806	0.875	0.933333333

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
TNFRSF8 AND NOT-LRRC55	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMEM26	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FAT2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-C10orf35	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GJD4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ABCA3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-LRTOMT	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FBN2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-FCAR	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MS4A2	0.928571429	1	0.866666667
TNFRSF8 AND NOT-FCER2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MARCH8	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-LRRN4CL	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CES5A	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-OPN5	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FCGR2B	0.965517241	1	0.933333333
TNFRSF8 AND NOT-PI16	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-C6orf89	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FCGR3B	0.928571429	1	0.866666667
TNFRSF8 AND NOT-RNF182	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-DAGLB	0.928571429	1	0.866666667
TNFRSF8 AND NOT-ADGRG3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GPRC6A	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ADGRF2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-UNC5CL	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMEM130	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC29A4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SEMA3D	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FGF4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FGFR3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FGFR2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FGFR4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ITGA11	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FKBP1A	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-LMTK2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-ZP1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SORCS3	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SV2C	0.965517241	1	0.933333333
TNFRSF8 AND NOT-FAIM2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMCC1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PLXND1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-LRCH1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SNX13	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SLC35D1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ATP11B	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MLC1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-NUP210	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SEL1L3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FLT3LG	0.965517241	1	0.933333333
TNFRSF8 AND NOT-KIAA1024	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ACSL6	0.965517241	1	0.933333333
TNFRSF8 AND NOT-FMO5	0.928571429	1	0.866666667
TNFRSF8 AND NOT-SLC9A8	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-DNAJC16	0.965517241	1	0.933333333
TNFRSF8 AND NOT-FAM189A1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KCNH4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMED3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ATP1B4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC44A1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ABCA5	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FOLR1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-LRRC8B	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC39A14	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-NNT	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-OR52A1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC16A8	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SEZ6L	0.928571429	1	0.866666667
TNFRSF8 AND NOT-SYNGR4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-HYAL4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PIGN	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FPR1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-VSIG2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-LEMD3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CLEC5A	0.965517241	1	0.933333333
TNFRSF8 AND NOT-NTSR2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-KCNE5	0.933333333	0.93333333	0.933333333

Antigen Pair Logic	F1	Precision	Recall
NLGN3 AND NOT-LHFPL2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ADAM8	0.903225806	0.875	0.933333333
NLGN3 AND NOT-DHRS2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-MPZL2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TRIM13	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ANGPTL7	0.903225806	0.875	0.933333333
NLGN3 AND NOT-KLRG1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-GPA33	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CD96	0.903225806	0.875	0.933333333
NLGN3 AND NOT-STX6	0.903225806	0.875	0.933333333
NLGN3 AND NOT-SLC35B1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-IGSF6	0.903225806	0.875	0.933333333
NLGN3 AND NOT-RAMP2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-RAMP3	0.928571429	1	0.866666667
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-SIGMAR1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-BET1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-LILRB2	0.965517241	1	0.933333333
NLGN3 AND NOT-MARCH6	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TCIRG1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-NMUR1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-B3GALT5	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CRISP3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-SIRPB1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-B3GNT3	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-CLEC4M	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TLR6	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-PCGF3	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-PKDREJ	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-BTN2A2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-CEPT1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ATP8A1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NDRG1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ST3GAL6	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TIMM17A	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-MERTK	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-CLEC10A	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SLC9A6	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-VTI1B	0.903225806	0.875	0.933333333
CSPG5 AND CRTAP	0.909090909	0.833333333	1
NLGN3 AND NOT-CRTAP	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-DEAF1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-HTATIP2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-AGPAT2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SLC19A2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SLC34A2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-SLCO1B1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ST6GALNAC2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-SPINT2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-DMRT2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CXCR6	0.903225806	0.875	0.933333333
NLGN3 AND NOT-C6orf10	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CD226	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
CSPG5 AND TMEM115	0.909090909	0.833333333	1
CSPG5 AND TSPAN11	0.909090909	0.833333333	1
CSPG5 AND P2RX4	0.909090909	0.833333333	1
CSPG5 AND SOAT1	0.909090909	0.833333333	1
CSPG5 AND JAGN1	0.909090909	0.833333333	1
CSPG5 AND GPAA1	0.909090909	0.833333333	1
NLGN3 AND NOT-CLDN16	0.903225806	0.875	0.933333333
NLGN3 AND NOT-RRH	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CORIN	0.903225806	0.875	0.933333333
NLGN3 AND NOT-FAM189B	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NRG3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-SLC12A7	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-PHTF1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
TNFRSF8 AND NOT-CA14	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMEM2	0.928571429	1	0.866666667
TNFRSF8 AND NOT-STX12	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FRRS1L	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FLRT1	0.928571429	1	0.866666667
TNFRSF8 AND NOT-FKBP8	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-APOL2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-BCL2L13	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-EML2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CLDN15	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MS4A6E	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FUT4	0.928571429	1	0.866666667
TNFRSF8 AND NOT-FUT7	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FZD2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CNIH2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC41A1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-NAALADL2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CALHM1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMEM86B	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-C19orf26	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GABBR1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MYADML2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-COL24A1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GABRA4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GABRA5	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GABRA6	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMEM196	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CDRT15L2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-STEAP1B	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GABRB3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GABRD	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ST6GALNAC3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MFSD8	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GABRG1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GABRG2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GABRP	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GABRR1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FRMD3	0.965517241	1	0.933333333
TNFRSF8 AND NOT-GABRR2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-UTS2B	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KIAA1549L	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC24A2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SUN2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-TMEM59L	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FAM19A5	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KLK5	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GALC	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TPSG1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-YIPF3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PARM1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SUMF2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GALR1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GALNT3	0.928571429	1	0.866666667
TNFRSF8 AND NOT-LY6G6F	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TAS2R39	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TAS2R40	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TAS2R41	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FAM205A	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-RNF167	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ATRNL1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GLCE	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-LRRTM2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CNTNAP2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SLITRK5	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CFAP61	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-HERC4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-LRIT1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PCDHB5	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-NPHP4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMEM251	0.965517241	1	0.933333333
TNFRSF8 AND NOT-OR2F1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-OR2B6	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-OR2L2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CLEC4E	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SLC13A4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-IL1RAPL2	0.933333333	0.93333333	0.933333333

Antigen Pair Logic	F1	Precision	Recall
NLGN3 AND NOT-ZMYND11	0.903225806	0.875	0.933333333
NLGN3 AND NOT-SLC17A3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-OR5I1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CYSLTR1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-CFTR	0.903225806	0.875	0.933333333
NLGN3 AND NOT-FAXDC2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-LILRB1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-SLC26A1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ADAM28	0.903225806	0.875	0.933333333
NLGN3 AND NOT-SLC22A7	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TSPAN9	0.903225806	0.875	0.933333333
NLGN3 AND NOT-USP19	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CD300C	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-EDDM3A	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CEACAM7	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NPFFR2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-LYVE1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-OCLM	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CEACAM4	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ADCY2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-BLCAP	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-EDAR	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-BTNL3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-KDELR1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-STARD3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ERP29	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TMED10	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-LILRB5	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SLC27A5	0.903225806	0.875	0.933333333
NLGN3 AND NOT-SLC27A4	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NAT2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-SLC27A3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-LILRB4	0.903225806	0.875	0.933333333
NLGN3 AND NOT-GLIPR1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-KDELR3	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ATF7	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TDRKH	0.903225806	0.875	0.933333333
NLGN3 AND NOT-LILRB3	0.965517241	1	0.933333333
NLGN3 AND NOT-UPK1A	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ZPBP	0.903225806	0.875	0.933333333
NLGN3 AND NOT-LECT1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CYB561D2	0.928571429	1	0.866666667
SORCS3 AND TMEM115	0.909090909	0.833333333	1
NLGN3 AND NOT-TMEM115	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ADAM30	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ADAM29	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CACFD1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-BTN3A1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-BTN2A1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-PTPRT	0.903225806	0.875	0.933333333
NLGN3 AND NOT-IL1RAPL1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-PLA2G16	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-HHLA2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ABCB8	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ADCY5	0.903225806	0.875	0.933333333
NLGN3 AND NOT-KLK8	0.903225806	0.875	0.933333333
NLGN3 AND NOT-FZD10	0.903225806	0.875	0.933333333
NLGN3 AND NOT-GALNT5	0.903225806	0.875	0.933333333
NLGN3 AND NOT-PRAF2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SEC63	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-GPR45	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-MAN1B1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-SLC6A14	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CMTM7	0.965517241	1	0.933333333
NLGN3 AND NOT-STX1B	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-SCN11A	0.903225806	0.875	0.933333333
NLGN3 AND NOT-MGAT4B	0.933333333	0.933333333	0.933333333

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
TNFRSF8 AND NOT-GALNT8	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-OR7A17	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-OR8B8	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-OR10A3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GCNT1	0.928571429	1	0.866666667
TNFRSF8 AND NOT-CHIC2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-GCNT2	0.928571429	1	0.866666667
TNFRSF8 AND NOT-TSPAN16	0.965517241	1	0.933333333
TNFRSF8 AND NOT-OR10H3	0.928571429	1	0.866666667
TNFRSF8 AND NOT-AMELX	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GDF9	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-OR7C2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-OR7C1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-BEST4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-HS6ST3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GDNF	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ADGRF1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-OR2H1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-OR1J2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GGCX	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-B4GALT1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-GGT7	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GHRHR	0.928571429	1	0.866666667
TNFRSF8 AND NOT-GHSR	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SEC22A	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-NOX1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SIGLEC7	0.928571429	1	0.866666667
TNFRSF8 AND NOT-PKD2L2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-NSG1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GJB3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ST6GALNAC4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-DKKL1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KCNH5	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-NAAA	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SLC39A1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SIGLEC9	0.928571429	1	0.866666667
TNFRSF8 AND NOT-SIGLEC8	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GPR82	0.965517241	1	0.933333333
TNFRSF8 AND NOT-HCAR1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-C5AR2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-COQ2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-GPR162	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-BMP10	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PCDH11X	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PCSK1N	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMEM97	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TOR1B	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GLP1R	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CECR6	0.965517241	1	0.933333333
TNFRSF8 AND NOT-GNRHR	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-WFDC10B	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GP2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GP5	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GPD2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GPM6B	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-IFNL2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CCR10	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GPR3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GPR4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-XCR1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GPR6	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC25A45	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-OR8D2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CDH20	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-OR10A4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-NPBWR2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ADGRD1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-GXYLT1	0.928571429	1	0.866666667
TNFRSF8 AND NOT-TMPRSS12	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ZDHHC22	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GPR12	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-OR4N4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-UTS2R	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMEM235	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ANG	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FAM171A2	0.933333333	0.93333333	0.933333333

Antigen Pair Logic	F1	Precision	Recall
NLGN3 AND NOT-B4GALT7	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-CHRM1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CHRM2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-SLCO2B1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CD300A	0.903225806	0.875	0.933333333
NLGN3 AND NOT-SCAMP4	0.903225806	0.875	0.933333333
NLGN3 AND NOT-GPR182	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CASC4	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-SLC46A1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-VSIG4	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TMEM106A	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SLC52A3	0.928571429	1	0.866666667
NLGN3 AND NOT-CHRM5	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-RNF13	0.928571429	1	0.866666667
NLGN3 AND NOT-SMIM12	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CHRNA1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CMTM1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CHRNA2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-MFSD3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-SDSL	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CHRNA3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-PIK3IP1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-SLC35A4	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-CHRNA5	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ADCY7	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CHRNB1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-CHRNB3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CHRNB4	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CHRND	0.903225806	0.875	0.933333333
NLGN3 AND NOT-SLC22A9	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ERMAP	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CHRNG	0.903225806	0.875	0.933333333
NLGN3 AND NOT-PKD1L2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CSMD2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-XKR4	0.903225806	0.875	0.933333333
NLGN3 AND NOT-SLC25A25	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-SLITRK1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TMEM200A	0.933333333	0.933333333	0.933333333
GRIK3 AND NOT-SORCS1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-SORCS1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-OSBPL8	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-C1QTNF6	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ADCY8	0.903225806	0.875	0.933333333
NLGN3 AND NOT-SLC26A9	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-OMA1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-FCRL1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-FCRL3	0.903225806	0.875	0.933333333
HEPACAM AND NOT-SLC5A11	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SLC5A11	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TNFRSF13C	0.903225806	0.875	0.933333333
NLGN3 AND NOT-EVI5L	0.903225806	0.875	0.933333333
NLGN3 AND NOT-LYSMD3	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SLC22A12	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CYYR1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ABHD15	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TLCD1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-PANX3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-GRIN3A	0.903225806	0.875	0.933333333
NLGN3 AND NOT-GRIN3B	0.903225806	0.875	0.933333333
NLGN3 AND NOT-MRGPRF	0.965517241	1	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CATSPER1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CATSPER2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-MRGPRX2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-MRGPRX4	0.903225806	0.875	0.933333333
NLGN3 AND NOT-SLC16A10	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-GALNT15	0.903225806	0.875	0.933333333

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
TNFRSF8 AND NOT-PIGW	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-C17orf78	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GPR17	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC13A5	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC26A11	0.928571429	1	0.866666667
TNFRSF8 AND NOT-IZUMO1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GPR20	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-VSTM1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-GPR22	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CYB561D1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-EPHA10	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-LPAR4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC25A34	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SIRPB2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-MCHR1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-DNAJC5G	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC9A9	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GPER1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GPR31	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CYP4V2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-GPR32	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-RELL2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-DCBLD1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ATG9B	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GPR35	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-LSMEM1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-XKR6	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GPR37	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CRB2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CFAP47	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FFAR1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ANK1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GRIA1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GRIA3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GRIA4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GRID1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GRID2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-OSTM1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SLC6A16	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMPRSS11E	0.928571429	1	0.866666667
TNFRSF8 AND NOT-GRIK3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GRIK4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GRIN1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GRIN2B	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TIMM21	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ORMDL2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ANPEP	0.965517241	1	0.933333333
TNFRSF8 AND NOT-GRM1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CLEC2D	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GRM2	0.928571429	1	0.866666667
TNFRSF8 AND NOT-GRM5	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GRM7	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ZDHHC1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ZDHHC8	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GRHL1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ICOS	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GUCY2F	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-NPC1L1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ST8SIA5	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PCDHB1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GYPA	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC39A3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC39A2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PILRA	0.965517241	1	0.933333333
TNFRSF8 AND NOT-GUCY2D	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-HAS1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-HAS3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-HCRT	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-HCRTR1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-HFE	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PLA2G2E	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KCNIP3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-NRG1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-HLA-DOB	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-HMGCR	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC29A2	0.933333333	0.93333333	0.933333333

Antigen Pair Logic	F1	Precision	Recall
NLGN3 AND NOT-TMC1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TMC2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CLCN1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CLCN2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-UBE2J2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ANTXR2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-GPR62	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-CLCN5	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CLCN6	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CLCN7	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-CLCNKB	0.903225806	0.875	0.933333333
NLGN3 AND NOT-PDZD8	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CALHM3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CLRN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CPXM2	0.965517241	1	0.933333333
NLGN3 AND NOT-OR5P2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-GYLTL1B	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TMEM45B	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CLN5	0.903225806	0.875	0.933333333
NLGN3 AND NOT-AMICA1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-CLPS	0.903225806	0.875	0.933333333
NLGN3 AND NOT-LRIG3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TEX29	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-DEGS2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CCR3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CCR4	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CCR5	0.903225806	0.875	0.933333333
NLGN3 AND NOT-C15orf27	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CCR6	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CCR7	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CCR8	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ACKR2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CMTM3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CMKLR1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-LTB4R	0.903225806	0.875	0.933333333
NLGN3 AND NOT-PAQR4	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TMEM219	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TMEM170A	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ABCC2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-SLC38A10	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-CD300LB	0.903225806	0.875	0.933333333
NLGN3 AND NOT-B4GALNT2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-GJD3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-SLC5A10	0.903225806	0.875	0.933333333
NLGN3 AND NOT-FAM210A	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-CLDND2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ZNF816	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CNGA1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-IZUMO2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-OR1I1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-CYP4F22	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ANKLE1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-B3GALT6	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CNR1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-SLC44A3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CNR2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TMCO2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-GJB4	0.903225806	0.875	0.933333333
NLGN3 AND NOT-RNF19B	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-DCST2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TEDDM1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-KLHDC7A	0.903225806	0.875	0.933333333
NLGN3 AND NOT-GOLT1A	0.903225806	0.875	0.933333333
NLGN3 AND NOT-C1orf162	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-FITM2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TBC1D20	0.903225806	0.875	0.933333333
NLGN3 AND NOT-PROKR2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CST9L	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CST9	0.903225806	0.875	0.933333333
NLGN3 AND NOT-EMID1	0.933333333	0.933333333	0.933333333

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
TNFRSF8 AND NOT-HRH2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-HSD3B2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-HSD17B3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-HSD17B2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ACACB	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-IGSF3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-DNAJC4	0.965517241	1	0.933333333
TNFRSF8 AND NOT-HTR1B	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-HTR1D	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-HTR2A	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-HTR3A	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-HTR5A	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-HTR6	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-HTR7	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FAM151A	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CLEC4D	0.965517241	1	0.933333333
TNFRSF8 AND NOT-IFNE	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ANO9	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ICAM2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-TMEM225	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ICAM4	0.928571429	1	0.866666667
TNFRSF8 AND NOT-B4GALNT4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FAM19A2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-APOB	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ZNF546	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CLEC4G	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-C2orf74	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-LRRC66	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC35D3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TREML1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ZDHHC21	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-VSIG1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-LHFPL1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MALRD1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC35F4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GLDN	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-IFNAR1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-IFNAR2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-IFNB1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MOGAT3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-LRRTM1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-AMIGO2	0.928571429	1	0.866666667
TNFRSF8 AND NOT-TMEM255B	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC6A18	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-NIPAL4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-HCN1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ZACN	0.928571429	1	0.866666667
TNFRSF8 AND NOT-SLCO4C1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-RGSL1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-IGSF1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-IL3RA	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-IL5RA	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-IL6ST	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CXCL8	0.928571429	1	0.866666667
TNFRSF8 AND NOT-CXCR1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-IL10RA	0.965517241	1	0.933333333
TNFRSF8 AND NOT-IL11RA	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-IL12RB1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-IL13	0.928571429	1	0.866666667
TNFRSF8 AND NOT-AQP4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-AQP5	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-INPP4A	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-AQP6	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-INSRR	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-AQP7	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ITGA2B	0.965517241	1	0.933333333
TNFRSF8 AND NOT-ITGA4	0.965517241	1	0.933333333
TNFRSF8 AND NOT-ITGA9	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ITGAM	0.965517241	1	0.933333333
TNFRSF8 AND NOT-ITGB2	0.928571429	1	0.866666667
TNFRSF8 AND NOT-ABCC6	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ITGB3	0.928571429	1	0.866666667
TNFRSF8 AND NOT-ITGB6	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ITGB7	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ITIH1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-STT3A	0.928571429	1	0.866666667

Antigen Pair Logic	F1	Precision	Recall
NLGN3 AND NOT-TMEM150A	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-CNTNAP5	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ACMSD	0.903225806	0.875	0.933333333
NLGN3 AND NOT-COL11A2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-SPATA3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-LYPD6B	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TMEM198	0.903225806	0.875	0.933333333
NLGN3 AND NOT-PQLC3	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TMEM182	0.903225806	0.875	0.933333333
NLGN3 AND NOT-COL17A1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-KCNH8	0.903225806	0.875	0.933333333
NLGN3 AND NOT-DNAJC19	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-FAM3D	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ZPLD1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CD200R1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ZDHHC19	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-DCBLD2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-LRRC15	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TPRA1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SLC31A2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TMEM207	0.903225806	0.875	0.933333333
NLGN3 AND NOT-IL17RE	0.903225806	0.875	0.933333333
NLGN3 AND NOT-RTP1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-EVC2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-SLC9B2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-IL31RA	0.903225806	0.875	0.933333333
NLGN3 AND NOT-EGFLAM	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-UGT3A1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CCDC127	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TMEM171	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TMEM174	0.903225806	0.875	0.933333333
NLGN3 AND NOT-PACRG	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-COX11	0.903225806	0.875	0.933333333
NLGN3 AND NOT-DPCR1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-WBSCR28	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TMEM139	0.903225806	0.875	0.933333333
NLGN3 AND NOT-OR6B1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-OR2F2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CPD	0.903225806	0.875	0.933333333
NLGN3 AND NOT-SVOPL	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CLDN4	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CLDN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ADORA2B	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CLDN23	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CPT1A	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TMEM71	0.903225806	0.875	0.933333333
NLGN3 AND NOT-SGCZ	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CR1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-LETM2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-HGSNAT	0.903225806	0.875	0.933333333
NLGN3 AND NOT-RNF183	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CR2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-C9orf57	0.903225806	0.875	0.933333333
NLGN3 AND NOT-FAM69B	0.903225806	0.875	0.933333333
NLGN3 AND NOT-SLITRK4	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ADGRG4	0.903225806	0.875	0.933333333
NLGN3 AND NOT-PTCHD1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CRHR1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-AADAC	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-MUC17	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ASB11	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CHODL	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TMEM37	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TRPM6	0.903225806	0.875	0.933333333
NLGN3 AND NOT-WFDC10A	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-C20orf173	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CST11	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CRY2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ADORA3	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SLC34A3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-VTI1A	0.903225806	0.875	0.933333333

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
TNFRSF8 AND NOT-ITPR2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ITPR3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CD82	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KCNA1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KCNA2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KCNA3	0.928571429	1	0.866666667
TNFRSF8 AND NOT-KCNA4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KCNA5	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KCNA6	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KCNA7	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KCNA10	0.965517241	1	0.933333333
TNFRSF8 AND NOT-KCNC1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KCNC2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-LRRC37A3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-DRAXIN	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FAM73A	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KCNC4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MIA3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-C1orf95	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KCND3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KCNH1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ENHO	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PNPLA7	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KCNJ1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KCNJ5	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KCNJ6	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC27A1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-KCNJ11	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KCNJ12	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KCNJ14	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KCNJ15	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KCNK2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KCNMB1	0.928571429	1	0.866666667
TNFRSF8 AND NOT-KCNN2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KCNN3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KCNQ1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-KCNQ2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KIR2DL3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KIR2DS4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KLRB1	0.928571429	1	0.866666667
TNFRSF8 AND NOT-KLRC3	0.965517241	1	0.933333333
TNFRSF8 AND NOT-KLRD1	0.928571429	1	0.866666667
TNFRSF8 AND NOT-KRT5	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-AMIGO3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GPR153	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMEM189	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC22A25	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CLEC12B	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SHISA2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GLTPD2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-RPRML	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-RGS9BP	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FAM132A	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FAM69A	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FAM209B	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-C5orf46	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-IYD	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-L1CAM	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-USP27X	0.965517241	1	0.933333333
TNFRSF8 AND NOT-OR51I1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-OR52D1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-LAMP2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-LFNG	0.965517241	1	0.933333333
TNFRSF8 AND NOT-LIM2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-C11orf87	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SNX19	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-LNPEP	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-XKRX	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-IFITM10	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PPAPDC2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-LRMP	0.965517241	1	0.933333333
TNFRSF8 AND NOT-LRP1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-LRP3	0.965517241	1	0.933333333
TNFRSF8 AND NOT-HAPLN4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SPINK6	0.928571429	1	0.866666667
TNFRSF8 AND NOT-LTB	0.965517241	1	0.933333333

Antigen Pair Logic	F1	Precision	Recall
NLGN3 AND NOT-SYT9	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CSF1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-MUC15	0.903225806	0.875	0.933333333
NLGN3 AND NOT-FAM76B	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CSF1R	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-CSF2RA	0.903225806	0.875	0.933333333
NLGN3 AND NOT-LAYN	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TMEM86A	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CSF3R	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ALG10B	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TMEM120B	0.933333333	0.933333333	0.933333333
BEST3 AND RPRM	0.909090909	0.833333333	1
BEST3 AND NOT-CD36	0.967741935	0.9375	1
NLGN3 AND NOT-B3GLCT	0.903225806	0.875	0.933333333
NLGN3 AND NOT-SLC38A6	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-C14orf37	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ISM2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-LRFN5	0.903225806	0.875	0.933333333
NLGN3 AND NOT-LYSMD4	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NRG4	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CMTM4	0.903225806	0.875	0.933333333
NLGN3 AND NOT-C16orf92	0.903225806	0.875	0.933333333
NLGN3 AND NOT-GSG1L	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SLC22A31	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TMED6	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CD300LF	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CD300LG	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TMEM199	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TMC8	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TMEM99	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-VSIG10L	0.903225806	0.875	0.933333333
NLGN3 AND NOT-C19orf18	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TMEM190	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TMC4	0.903225806	0.875	0.933333333
SLC4A4 AND NOT-TMC4	0.928571429	1	0.866666667
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ADRA1B	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ZNRF4	0.903225806	0.875	0.933333333
NLGN3 AND NOT-FAM187B	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ATP8B3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-HFE2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-FAM163A	0.903225806	0.875	0.933333333
NLGN3 AND NOT-PM20D1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-SLC30A7	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-KNCN	0.903225806	0.875	0.933333333
NLGN3 AND NOT-DCST1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-IL23R	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-CTLA4	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CLDN19	0.903225806	0.875	0.933333333
NLGN3 AND NOT-C1orf210	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ADIG	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CTNS	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SLC9B1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NFAM1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-PROM2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ADRA2A	0.903225806	0.875	0.933333333
NLGN3 AND NOT-PLB1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-SLC23A3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-GPBAR1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-RMDN2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-FAM19A4	0.965517241	1	0.933333333
NLGN3 AND NOT-CCDC80	0.903225806	0.875	0.933333333
NLGN3 AND NOT-BTLA	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ADRA2B	0.903225806	0.875	0.933333333
NLGN3 AND NOT-XXYLT1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CTSZ	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-CUX1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-IGSF11	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-PAQR3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-SHISA3	0.928571429	1	0.866666667
NLGN3 AND NOT-THAP6	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-C4orf26	0.903225806	0.875	0.933333333

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
TNFRSF8 AND NOT-LTBP3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ERVFRD-1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-LTK	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CD180	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TACSTD2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-M6PR	0.928571429	1	0.866666667
TNFRSF8 AND NOT-FAM19A1	0.928571429	1	0.866666667
TNFRSF8 AND NOT-SMAD2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MAG	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ARSB	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MAN1A1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MAN2A2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-MAS1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-LCN10	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ARSD	0.965517241	1	0.933333333
TNFRSF8 AND NOT-ARSE	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MC5R	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ARSF	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ADAM11	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ART3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ART4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MGAT1	0.928571429	1	0.866666667
TNFRSF8 AND NOT-MGAT3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CTAGE5	0.965517241	1	0.933333333
TNFRSF8 AND NOT-MGST2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-MIP	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TRPM1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MMP15	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MMP16	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ALDH6A1	0.928571429	1	0.866666667
TNFRSF8 AND NOT-ASGR1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-MOG	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MPL	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MXRA7	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ACHE	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SMCO3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-LRRC52	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-C1orf186	0.928571429	1	0.866666667
TNFRSF8 AND NOT-SHISA8	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SMIM4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMEM151B	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-OR2B3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-OR14J1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-OR10C1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PTCHD4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MSR1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MTNR1B	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MUC3A	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-MUC4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-NCAM1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-NDUFC2	0.928571429	1	0.866666667
TNFRSF8 AND NOT-NELL2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-ATP1A3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-NGFR	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ATP1A4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-NMBR	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-NOTCH2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-NPHS1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-NPPA	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-NPY5R	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC11A2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ATP2A3	0.965517241	1	0.933333333
TNFRSF8 AND NOT-NTRK3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ROR2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-NTSR1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CISD2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ATP4B	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-OPRD1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-OPRK1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-OPRL1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-OPRM1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-OR2C1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-OR3A1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-P2RX1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-P2RX7	0.928571429	1	0.866666667
TNFRSF8 AND NOT-P2RY1	0.933333333	0.93333333	0.933333333

Antigen Pair Logic	F1	Precision	Recall
NLGN3 AND NOT-KLB	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CYB5A	0.933333333	0.93333333	0.933333333
NLGN3 AND NOT-SPINK13	0.933333333	0.93333333	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TMEM161B	0.903225806	0.875	0.933333333
NLGN3 AND NOT-MARVELD2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-BTNL9	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CYBA	0.933333333	0.93333333	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CCDC112	0.903225806	0.875	0.933333333
NLGN3 AND NOT-RNF145	0.933333333	0.93333333	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-SLC2A12	0.903225806	0.875	0.933333333
NLGN3 AND NOT-MBOAT1	0.933333333	0.93333333	0.933333333
NLGN3 AND NOT-RNF217	0.933333333	0.93333333	0.933333333
NLGN3 AND NOT-CYP1A1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CYP1A2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ABCA13	0.933333333	0.93333333	0.933333333
NLGN3 AND NOT-VKORC1L1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CYP2A6	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ADRB2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TMEM213	0.903225806	0.875	0.933333333
NLGN3 AND ATP6V0E2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CYP3A7	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CYP2C8	0.933333333	0.93333333	0.933333333
NLGN3 AND NOT-ADRB3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-PEBP4	0.933333333	0.93333333	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CYP3A4	0.933333333	0.93333333	0.933333333
NLGN3 AND NOT-CYP3A5	0.933333333	0.93333333	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CYP4B1	0.965517241	1	0.933333333
NLGN3 AND NOT-CYP8B1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-FREM1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-PRUNE2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-FMR1NB	0.965517241	1	0.933333333
NLGN3 AND NOT-FAAH2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-MOSPD2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-AWAT1	0.903225806	0.875	0.933333333
TSPAN7 AND ZDHHC15	0.933333333	0.93333333	0.933333333
NLGN3 AND NOT-CYP19A1	0.933333333	0.93333333	0.933333333
NLGN3 AND NOT-SLC35G1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-SLC5A12	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CCDC67	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TMTC2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CLEC12A	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TMTC3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-DAG1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-SLC5A8	0.903225806	0.875	0.933333333
NLGN3 AND NOT-GPR180	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TMEM229B	0.903225806	0.875	0.933333333
NLGN3 AND NOT-SYNE3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CLEC14A	0.903225806	0.875	0.933333333
NLGN3 AND NOT-FITM1	0.933333333	0.93333333	0.933333333
NLGN3 AND NOT-TMEM30B	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ITPRIPL2	0.933333333	0.93333333	0.933333333
NLGN3 AND NOT-DBH	0.903225806	0.875	0.933333333
NLGN3 AND NOT-MFSD6L	0.903225806	0.875	0.933333333
NLGN3 AND NOT-FAM134C	0.933333333	0.93333333	0.933333333
NLGN3 AND NOT-TMEM92	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TRPV3	0.933333333	0.93333333	0.933333333
NLGN3 AND NOT-SPPL2C	0.903225806	0.875	0.933333333
NLGN3 AND NOT-OR7D2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-LPPR5	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TOR1AIP2	0.933333333	0.93333333	0.933333333
NLGN3 AND NOT-ACE	0.903225806	0.875	0.933333333
NLGN3 AND NOT-IFNLR1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-DCT	0.903225806	0.875	0.933333333
NLGN3 AND NOT-PAQR7	0.933333333	0.93333333	0.933333333
NLGN3 AND NOT-LRRN4	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TMPRSS6	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ADGRF3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-DDOST	0.933333333	0.93333333	0.933333333

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
TNFRSF8 AND NOT-P2RY2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-P2RY4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-P2RY6	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CD207	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-DUOX2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CHST11	0.965517241	1	0.933333333
TNFRSF8 AND NOT-ATP6V0A4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CYHR1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KCNK4	0.965517241	1	0.933333333
TNFRSF8 AND NOT-TAS2R1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TAS2R9	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TAS2R7	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TAS2R13	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CLEC4A	0.965517241	1	0.933333333
TNFRSF8 AND NOT-PCDH1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PCDHGC3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC35B3	0.965517241	1	0.933333333
TNFRSF8 AND NOT-DERL2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PCDH9	0.965517241	1	0.933333333
TNFRSF8 AND NOT-ST8SIA3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ATL1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GAL	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-A4GNT	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC45A2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CKLF	0.965517241	1	0.933333333
TNFRSF8 AND NOT-DUSP13	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CLDN18	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CLEC1B	0.965517241	1	0.933333333
TNFRSF8 AND NOT-CLEC1A	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PIPOX	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TLR7	0.965517241	1	0.933333333
TNFRSF8 AND NOT-CEND1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-RXFP3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC15A3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-REEP2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-PDCD1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-KLRF1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-MBTPS2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-WNT16	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SCARA3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GULP1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMEM138	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PDE6B	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-HMP19	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FKBP7	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ENPP1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ENPP3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ACSL5	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CD244	0.965517241	1	0.933333333
TNFRSF8 AND NOT-HIGD1B	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TMEM8B	0.965517241	1	0.933333333
TNFRSF8 AND NOT-ASIC5	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TUBA8	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PF4	0.965517241	1	0.933333333
TNFRSF8 AND NOT-ABCB1	0.928571429	1	0.866666667
TNFRSF8 AND NOT-ABCB4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PI3	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SERPINB13	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PIGA	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PIGR	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PKHD1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-IL17D	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-CHIC1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TM6SF1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-STX18	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PLIN1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FXYD1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-FXYD3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PLP1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ADAM22	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-S1PR5	0.965517241	1	0.933333333
TNFRSF8 AND NOT-FXYD7	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FXYD6	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FXYD4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-IL20RB	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-GPR87	0.933333333	0.93333333	0.933333333

Antigen Pair Logic	F1	Precision	Recall
NLGN3 AND NOT-CLEC4F	0.903225806	0.875	0.933333333
NLGN3 AND NOT-GPR156	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CHST13	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ADGRA3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-SGMS2	0.903225806	0.875	0.933333333
SLC4A4 AND NOT-SGMS2	0.928571429	1	0.866666667
NLGN3 AND NOT-GALNTL5	0.903225806	0.875	0.933333333
NLGN3 AND NOT-RNF133	0.903225806	0.875	0.933333333
NLGN3 AND NOT-PKD1L1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-SLC30A8	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TMEM64	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-KCNV2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TMEM252	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-GIMAP1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-KCNG3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CLYBL	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-FAM9C	0.903225806	0.875	0.933333333
NLGN3 AND NOT-SPTSSA	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ABHD3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-DHCR24	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-DIO1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-DNASE1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-AGER	0.903225806	0.875	0.933333333
NLGN3 AND NOT-DPAGT1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-DPP4	0.903225806	0.875	0.933333333
NLGN3 AND NOT-DPP6	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-DRD2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-DRD3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-DRD4	0.903225806	0.875	0.933333333
NLGN3 AND NOT-DSC1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-DSC2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-DSC3	0.933333333	0.933333333	0.933333333
NOT-MMP16 AND DSCAM	0.909090909	0.833333333	1
NLGN3 AND NOT-DSG2	0.903225806	0.875	0.933333333
SLC4A4 AND NOT-DSG2	0.928571429	1	0.866666667
NLGN3 AND NOT-SLC26A2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-DTNA	0.903225806	0.875	0.933333333
NLGN3 AND NOT-AGTR1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-AGTR2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-GPR183	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ECE1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-EDA	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-LPAR1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-EDNRA	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-PGAM5	0.903225806	0.875	0.933333333
NLGN3 AND NOT-EFNB1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-EFNB3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-EGF	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-MPZL3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-DNAH10	0.903225806	0.875	0.933333333
NLGN3 AND NOT-PLBD2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ANO6	0.903225806	0.875	0.933333333
NLGN3 AND NOT-VSTM4	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-FAM24B	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ADCY4	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-EPHA2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-LCTL	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ELANE	0.903225806	0.875	0.933333333
NLGN3 AND NOT-MCEMP1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TMEM201	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TMEM61	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-SLC5A9	0.903225806	0.875	0.933333333
NLGN3 AND NOT-FAM209A	0.903225806	0.875	0.933333333
NLGN3 AND NOT-KRTCAP3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-SLC51A	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TVP23C	0.928571429	1	0.866666667
NLGN3 AND NOT-SLC39A11	0.903225806	0.875	0.933333333

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
TNFRSF8 AND NOT-DUOX1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLCO1C1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FAM3B	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TLR9	0.965517241	1	0.933333333
TNFRSF8 AND NOT-KCNK10	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TREM2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-TREM1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-GDAP1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-WNT4	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-NLGN3	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-DNAJC10	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-YIPF1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-FAM105A	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-ZDHHC13	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SDK2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-UGT1A8	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-EQTN	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-NDFIP2	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-VSIG10	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-LRRN3	0.965517241	1	0.933333333
TNFRSF8 AND NOT-CRLS1	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-SLC6A20	0.933333333	0.93333333	0.933333333
TNFRSF8 AND NOT-PPBP	0.965517241	1	0.933333333
TNFRSF8 AND NOT-LEPROT	0.965517241	1	0.933333333
TNFRSF8 AND NOT-DNAJB12	0.928571429	1	0.866666667
TNFRSF8 AND NOT-TMEM33	0.965517241	1	0.933333333
TNFRSF8 AND NOT-FBXW7	0.965517241	1	0.933333333
TNFRSF8 AND NOT-RNF121	0.928571429	1	0.866666667
TNFRSF8 AND NOT-SLC22A15	0.928571429	1	0.866666667
TNFRSF8 AND NOT-NPY4R	0.928571429	1	0.866666667
TNFRSF8 AND NOT-SMPD3	0.965517241	1	0.933333333
TNFRSF8 AND NOT-ACPP	0.965517241	1	0.933333333
TNFRSF8 AND NOT-ALG1	0.928571429	1	0.866666667
TNFRSF8 AND NOT-PCDHB12	0.928571429	1	0.866666667
TNFRSF8 AND NOT-LRRC8A	0.928571429	1	0.866666667
TNFRSF8 AND NOT-UGGT1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-LPAR5	0.965517241	1	0.933333333
TNFRSF8 AND NOT-RHBG	0.965517241	1	0.933333333
TNFRSF8 AND NOT-TMEM159	0.965517241	1	0.933333333
TNFRSF8 AND NOT-ATP10D	0.928571429	1	0.866666667
TNFRSF8 AND NOT-PTAFR	0.965517241	1	0.933333333
TNFRSF8 AND NOT-PTGDR	0.965517241	1	0.933333333
TNFRSF8 AND NOT-PTGER2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-PTGS1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-TMCC3	0.965517241	1	0.933333333
TNFRSF8 AND NOT-KIDINS220	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SLC46A2	0.928571429	1	0.866666667
TNFRSF8 AND NOT-PTPRC	0.965517241	1	0.933333333
TNFRSF8 AND NOT-PTPRJ	0.965517241	1	0.933333333
TNFRSF8 AND NOT-PTPRO	0.965517241	1	0.933333333
TNFRSF8 AND NOT-BAX	0.965517241	1	0.933333333
TNFRSF8 AND NOT-BBS4	0.965517241	1	0.933333333
TNFRSF8 AND NOT-IL22RA1	0.928571429	1	0.866666667
TNFRSF8 AND NOT-ENPP5	0.965517241	1	0.933333333
TNFRSF8 AND NOT-RARRES3	0.965517241	1	0.933333333
TNFRSF8 AND NOT-CACNG8	0.928571429	1	0.866666667
TNFRSF8 AND NOT-SIGIRR	0.965517241	1	0.933333333
TNFRSF8 AND NOT-HRH4	0.965517241	1	0.933333333
TNFRSF8 AND NOT-LGR6	0.965517241	1	0.933333333
TNFRSF8 AND NOT-RHO	0.965517241	1	0.933333333
TNFRSF8 AND NOT-RTN1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-TMBIM1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-C10orf54	0.965517241	1	0.933333333
TNFRSF8 AND NOT-XYLT1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-DPEP2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-HERPUD2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SFTPC	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SUSD1	0.928571429	1	0.866666667
TNFRSF8 AND NOT-MARC1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-FBXL17	0.928571429	1	0.866666667
TNFRSF8 AND NOT-ST3GAL3	0.965517241	1	0.933333333
TNFRSF8 AND NOT-MPPE1	0.928571429	1	0.866666667
TNFRSF8 AND NOT-SLC6A12	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SLC8A1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SLC11A1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SLC14A1	0.928571429	1	0.866666667

Antigen Pair Logic	F1	Precision	Recall
NLGN3 AND NOT-RDM1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ADGRE1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TIGIT	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TMEM192	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TAPT1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ENG	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ENPEP	0.903225806	0.875	0.933333333
NLGN3 AND NOT-SLC29A1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-HTRA4	0.903225806	0.875	0.933333333
NLGN3 AND NOT-C9orf91	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NRK	0.965517241	1	0.933333333
NLGN3 AND NOT-VMA21	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TMEM31	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ANO5	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-EPHA1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CERS3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-EPHA3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-EPHA4	0.903225806	0.875	0.933333333
NLGN3 AND NOT-EPHA8	0.903225806	0.875	0.933333333
NLGN3 AND NOT-SLC44A5	0.903225806	0.875	0.933333333
NLGN3 AND NOT-EPHB4	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-EPHB6	0.903225806	0.875	0.933333333
NLGN3 AND NOT-STX2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-EPO	0.903225806	0.875	0.933333333
NLGN3 AND NOT-LVRN	0.903225806	0.875	0.933333333
NLGN3 AND NOT-SLC36A1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ERBB3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-EREG	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ERN1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ETV6	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-EVC	0.903225806	0.875	0.933333333
NLGN3 AND NOT-EXT1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-F2R	0.903225806	0.875	0.933333333
NLGN3 AND NOT-F2RL1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-F2RL2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-F10	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ACSL1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-LRRC55	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-UNC5B	0.903225806	0.875	0.933333333
NLGN3 AND NOT-FAT2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TMEM218	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TMEM136	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-MPEG1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-CYB561A3	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-LRTOMT	0.903225806	0.875	0.933333333
NLGN3 AND NOT-FBN2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-RNF152	0.903225806	0.875	0.933333333
NLGN3 AND NOT-FCAR	0.903225806	0.875	0.933333333
NLGN3 AND NOT-FCER1A	0.903225806	0.875	0.933333333
NLGN3 AND NOT-MS4A2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-FCER2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-MARCH8	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CES5A	0.903225806	0.875	0.933333333
NLGN3 AND NOT-FAM26D	0.903225806	0.875	0.933333333
NLGN3 AND NOT-FAM162B	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-OPN5	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ADGRF4	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ADGRF5	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TMEM217	0.903225806	0.875	0.933333333
NLGN3 AND NOT-PI16	0.903225806	0.875	0.933333333
NLGN3 AND NOT-C6orf89	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-LEMD2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-FAM200A	0.903225806	0.875	0.933333333
NLGN3 AND NOT-FCGRT	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
TNFRSF8 AND NOT-BNIP2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SPINT1	0.928571429	1	0.866666667
TNFRSF8 AND NOT-SPN	0.928571429	1	0.866666667
TNFRSF8 AND NOT-BPI	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SSR1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SSTR4	0.965517241	1	0.933333333
TNFRSF8 AND NOT-STX3	0.965517241	1	0.933333333
TNFRSF8 AND NOT-BST1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-TGFBR1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-TGFBR3	0.965517241	1	0.933333333
TNFRSF8 AND NOT-TSPO	0.928571429	1	0.866666667
TNFRSF8 AND NOT-TLR1	0.928571429	1	0.866666667
TNFRSF8 AND NOT-TLR2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-TLR4	0.965517241	1	0.933333333
TNFRSF8 AND NOT-TLR5	0.965517241	1	0.933333333
TNFRSF8 AND NOT-TNFRSF1B	0.965517241	1	0.933333333
TNFRSF8 AND NOT-TRPC4	0.928571429	1	0.866666667
TNFRSF8 AND NOT-C5AR1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-CCR2	0.928571429	1	0.866666667
TNFRSF8 AND NOT-TMEM242	0.965517241	1	0.933333333
TNFRSF8 AND NOT-UGT2B17	0.928571429	1	0.866666667
TNFRSF8 AND NOT-UPK3A	0.965517241	1	0.933333333
TNFRSF8 AND NOT-VIPR1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-VIPR2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-DAGLA	0.965517241	1	0.933333333
TNFRSF8 AND NOT-LDLRAD4	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SLC30A1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SLC25A20	0.965517241	1	0.933333333
TNFRSF8 AND NOT-ATG9A	0.965517241	1	0.933333333
TNFRSF8 AND NOT-FKRP	0.928571429	1	0.866666667
TNFRSF8 AND NOT-GNPTAB	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SLC39A7	0.965517241	1	0.933333333
TNFRSF8 AND NOT-LST1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-RHBDF2	0.928571429	1	0.866666667
TNFRSF8 AND NOT-TXNDC15	0.928571429	1	0.866666667
TNFRSF8 AND NOT-MCTP1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-CLMN	0.965517241	1	0.933333333
TNFRSF8 AND NOT-WLS	0.965517241	1	0.933333333
TNFRSF8 AND NOT-UXS1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-TRAF3IP3	0.965517241	1	0.933333333
TNFRSF8 AND NOT-PRR7	0.965517241	1	0.933333333
TNFRSF8 AND NOT-CAMP	0.965517241	1	0.933333333
TNFRSF8 AND NOT-KLRC4	0.965517241	1	0.933333333
TNFRSF8 AND NOT-STX7	0.965517241	1	0.933333333
TNFRSF8 AND NOT-TMEM175	0.965517241	1	0.933333333
TNFRSF8 AND NOT-HVCN1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-RECK	0.965517241	1	0.933333333
TNFRSF8 AND NOT-MCHR2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-ATRN	0.965517241	1	0.933333333
TNFRSF8 AND NOT-TPST2	0.928571429	1	0.866666667
TNFRSF8 AND NOT-ADGRE3	0.965517241	1	0.933333333
TNFRSF8 AND NOT-MS4A14	0.965517241	1	0.933333333
TNFRSF8 AND NOT-USP30	0.928571429	1	0.866666667
TNFRSF8 AND NOT-AGPAT9	0.965517241	1	0.933333333
TNFRSF8 AND NOT-FCRLA	0.928571429	1	0.866666667
TNFRSF8 AND NOT-SFT2D3	0.965517241	1	0.933333333
TNFRSF8 AND NOT-ADTRP	0.928571429	1	0.866666667
TNFRSF8 AND NOT-LRCH3	0.928571429	1	0.866666667
TNFRSF8 AND NOT-MFSD2A	0.965517241	1	0.933333333
TNFRSF8 AND NOT-FAM73B	0.928571429	1	0.866666667
TNFRSF8 AND NOT-PLXDC2	0.928571429	1	0.866666667
TNFRSF8 AND NOT-TMEM141	0.965517241	1	0.933333333
TNFRSF8 AND NOT-PAQR8	0.965517241	1	0.933333333
TNFRSF8 AND NOT-DNAJC14	0.928571429	1	0.866666667
TNFRSF8 AND NOT-SLC4A4	0.965517241	1	0.933333333
TNFRSF8 AND NOT-STX10	0.965517241	1	0.933333333
TNFRSF8 AND NOT-DGAT1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-TNFRSF10C	0.928571429	1	0.866666667
TNFRSF8 AND NOT-TNFRSF10A	0.965517241	1	0.933333333
TNFRSF8 AND NOT-VNN2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-MGAM	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SYS1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-TMEM55B	0.928571429	1	0.866666667
TNFRSF8 AND NOT-CD1C	0.965517241	1	0.933333333
TNFRSF8 AND NOT-CD1D	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SYNGR1	0.965517241	1	0.933333333

Antigen Pair Logic	F1	Precision	Recall
NLGN3 AND NOT-DAGLB	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TMED4	0.903225806	0.875	0.933333333
NLGN3 AND NOT-FCN2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-KIAA1324L	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ADGRG3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-GPRC6A	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ADGRF2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TSPO2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-UNC5CL	0.903225806	0.875	0.933333333
NLGN3 AND NOT-LHFPL5	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ALDH3A2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-FGF6	0.903225806	0.875	0.933333333
NLGN3 AND NOT-FGF7	0.903225806	0.875	0.933333333
NLGN3 AND NOT-FGF10	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-FGFR3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-FGFR2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-FGFR4	0.903225806	0.875	0.933333333
NLGN3 AND NOT-FGG	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ITGA11	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-CLCA4	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN4Y	0.903225806	0.875	0.933333333
NLGN3 AND NOT-DOLK	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-LMTK2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CHSY1	0.903225806	0.875	0.933333333
KCNQ2 AND NOT-ADGRL1	0.928571429	1	0.866666667
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-SLITRK3	0.903225806	0.875	0.933333333
NRCAM AND NOT-CLSTN1	0.928571429	1	0.866666667
NLGN3 AND NOT-ZP1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CD93	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-PLA2R1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SCAP	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
SORCS3 AND TSPAN11	0.909090909	0.833333333	1
SORCS3 AND P2RX4	0.909090909	0.833333333	1
SORCS3 AND OGFOD3	0.909090909	0.833333333	1
SORCS3 AND GPAA1	0.909090909	0.833333333	1
SORCS3 AND MPZL1	0.909090909	0.833333333	1
NLGN3 AND NOT-SV2C	0.903225806	0.875	0.933333333
NLGN3 AND NOT-PCNX	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TMCC1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TBC1D2B	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-NFASC	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ATP10B	0.903225806	0.875	0.933333333
NLGN3 AND NOT-PLXND1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ANKLE2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-LRCH1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-GRAMD4	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-MLANA	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-STAB1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SLC35D1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ATP11B	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NUP210	0.903225806	0.875	0.933333333
NLGN3 AND NOT-FLT3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ATP11A	0.903225806	0.875	0.933333333
NLGN3 AND NOT-KIAA1024	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ADGRL2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-FMO3	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-FMO4	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ICOSLG	0.903225806	0.875	0.933333333
NLGN3 AND NOT-FMO5	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SLC9A8	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-DNAJC16	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ESYT1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-FAM189A1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ZDHHC17	0.903225806	0.875	0.933333333

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
TNFRSF8 AND NOT-MYADM	0.965517241	1	0.933333333
TNFRSF8 AND NOT-NXPE3	0.965517241	1	0.933333333
TNFRSF8 AND NOT-CD3G	0.928571429	1	0.866666667
TNFRSF8 AND NOT-SLC16A7	0.965517241	1	0.933333333
TNFRSF8 AND NOT-ZMYM6	0.965517241	1	0.933333333
TNFRSF8 AND NOT-CD5	0.965517241	1	0.933333333
TNFRSF8 AND NOT-TMEM129	0.965517241	1	0.933333333
TNFRSF8 AND NOT-TMEM169	0.965517241	1	0.933333333
TNFRSF8 AND NOT-MARCH9	0.965517241	1	0.933333333
TNFRSF8 AND NOT-ORAI3	0.965517241	1	0.933333333
TNFRSF8 AND NOT-PGAP3	0.965517241	1	0.933333333
TNFRSF8 AND NOT-MS4A3	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SFXN5	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SNRNP40	0.965517241	1	0.933333333
TNFRSF8 AND NOT-KCNK6	0.965517241	1	0.933333333
TNFRSF8 AND NOT-ECEL1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-NCR1	0.928571429	1	0.866666667
TNFRSF8 AND NOT-TNFSF8	0.928571429	1	0.866666667
TNFRSF8 AND NOT-CD36	0.965517241	1	0.933333333
TNFRSF8 AND NOT-SPTLC2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-EI24	0.965517241	1	0.933333333
TNFRSF8 AND NOT-CD40LG	0.965517241	1	0.933333333
TNFRSF8 AND NOT-TMEM63A	0.928571429	1	0.866666667
TNFRSF8 AND NOT-TMCC2	0.965517241	1	0.933333333
TNFRSF8 AND NOT-LPGAT1	0.965517241	1	0.933333333
TNFRSF8 AND NOT-P2RY14	0.928571429	1	0.866666667
TNFRSF8 AND NOT-SLC12A6	0.965517241	1	0.933333333
Leiomyosarcoma (Leiomyosarcoma)			
NOT-SGPP2 AND VANGL1	1	1	1
NOT-KRTCAP3 AND VANGL1	0.923076923	0.92307692	0.923076923
NOT-SMIM5 AND NEMP1	0.916666667	1	0.846153846
NOT-CRB3 AND ATP2A2	0.916666667	1	0.846153846
NOT-F11R AND VANGL1	0.96	1	0.923076923
NOT-RNF43 AND VANGL1	0.96	1	0.923076923
NOT-SCNN1A AND VANGL1	0.96	1	0.923076923
NOT-SMIM5 AND VANGL1	0.923076923	0.92307692	0.923076923
NOT-ERMP1 AND VANGL1	0.962962963	0.92857143	1
NOT-CRB3 AND VANGL1	0.96	1	0.923076923
Astrocytoma (Astrocytoma)			
PTPRZ1 AND CLCN5	0.911111111	0.93181818	0.891304348
PTPRZ1 AND ADGRE1	0.917647059	1	0.847826087
PTPRZ1 AND F2R	0.930232558	1	0.869565217
PTPRZ1 AND CYB561A3	0.901098901	0.91111111	0.891304348
PTPRZ1 AND IL27	0.901098901	0.91111111	0.891304348
PTPRZ1 AND NOT-GHITM	0.921348315	0.95348837	0.891304348
PTPRZ1 AND NOT-PRRT3	0.901098901	0.91111111	0.891304348
PTPRZ1 AND HAS2	0.917647059	1	0.847826087
PTPRZ1 AND LCAT	0.91954023	0.97560976	0.869565217
PTPRZ1 AND NOT-EPCAM	0.901098901	0.91111111	0.891304348
PTPRZ1 AND NOT-CYB5R2	0.909090909	0.95238095	0.869565217
PTPRZ1 AND NOT-ATP6V0C	0.911111111	0.93181818	0.891304348
PTPRZ1 AND ATP13A1	0.909090909	0.95238095	0.869565217
PTPRZ1 AND NOT-LRP11	0.901098901	0.91111111	0.891304348
PTPRZ1 AND NOT-INPP4B	0.906976744	0.975	0.847826087
PTPRZ1 AND NOT-C14orf2	0.901098901	0.91111111	0.891304348
Ovarian Neoplasms|Cystadenocarcinoma, Serous|Adenocarcinoma, Papillary (Ovarian)			
SVOPL AND NOT-ARL6IP5	0.923076923	0.85714286	1
NOT-ARL6IP5 AND FAAH2	0.916666667	0.91666667	0.916666667
NOT-ARL6IP5 AND GAL3ST4	0.916666667	0.91666667	0.916666667
SYNE4 AND NOT-FAXDC2	0.916666667	0.91666667	0.916666667
SYNE4 AND NOT-OS9	0.916666667	0.91666667	0.916666667
SVOPL AND NOT-ANTXR2	0.923076923	0.85714286	1
SVOPL AND NOT-COMT	0.923076923	0.85714286	1
SVOPL AND NOT-SLC31A2	0.96	0.92307692	1
SVOPL AND NOT-MFSD4	0.923076923	0.85714286	1
SVOPL AND NOT-FMO5	0.96	0.92307692	1
SVOPL AND NOT-TMEM131	0.923076923	0.85714286	1
SVOPL AND NOT-CHIC2	0.923076923	0.85714286	1
SVOPL AND NOT-C4orf3	0.923076923	0.85714286	1
NOT-NDUFA1 AND SVOPL	0.923076923	0.85714286	1
SVOPL AND NOT-TAS2R16	0.956521739	1	0.916666667
SVOPL AND NOT-DHRS7	0.923076923	0.85714286	1
SVOPL AND NOT-CYB5R1	0.923076923	0.85714286	1
SVOPL AND NOT-PKD2	0.923076923	0.85714286	1
SVOPL AND NOT-OCIAD1	0.923076923	0.85714286	1

Antigen Pair Logic	F1	Precision	Recall
NLGN3 AND NOT-KCNH4	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-SLC44A1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ABCB10	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ABCB9	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ABCA6	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ICMT	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-FOLH1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TRAM1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-LEPROTL1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-FOLR1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TNFRSF13B	0.903225806	0.875	0.933333333
NLGN3 AND NOT-MACF1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TMEM131	0.903225806	0.875	0.933333333
NLGN3 AND NOT-LRRC8B	0.903225806	0.875	0.933333333
NLGN3 AND NOT-SLC39A14	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NNT	0.903225806	0.875	0.933333333
NLGN3 AND NOT-OR52A1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-SLC16A8	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-SYNGR4	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TSPAN12	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TSPAN15	0.903225806	0.875	0.933333333
NLGN3 AND NOT-PIGN	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-CLDN14	0.903225806	0.875	0.933333333
NLGN3 AND NOT-LPAR3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-VSIG2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-FPR2	0.928571429	1	0.866666667
NLGN3 AND NOT-CLEC5A	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-PRND	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CA14	0.903225806	0.875	0.933333333
NLGN3 AND NOT-RABGAP1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TMEM2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TMEFF2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-STX12	0.903225806	0.875	0.933333333
NLGN3 AND NOT-KCNE4	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TMEM245	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-IL17RA	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-FLRT2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-MTCH1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-LAMP5	0.903225806	0.875	0.933333333
NLGN3 AND NOT-PANX1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CLDN15	0.903225806	0.875	0.933333333
GRIK3 AND FJX1	0.909090909	0.833333333	1
NLGN3 AND NOT-MS4A6E	0.903225806	0.875	0.933333333
NLGN3 AND NOT-SLC17A8	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ADAM2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-FUT2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-FUT3	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-FUT4	0.903225806	0.875	0.933333333
NLGN3 AND NOT-FUT5	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TMEM9	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-FUT6	0.903225806	0.875	0.933333333
NLGN3 AND NOT-FUT7	0.903225806	0.875	0.933333333
NLGN3 AND NOT-HEPACAM2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TECRL	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-FUT8	0.903225806	0.875	0.933333333
NLGN3 AND NOT-KDSR	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ACKR1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-FZD2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-IFI6	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ZDHHC20	0.903225806	0.875	0.933333333
NLGN3 AND NOT-G6PC	0.903225806	0.875	0.933333333
NLGN3 AND NOT-FAM26E	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SLC37A4	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TMEM256	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ZDHHC23	0.903225806	0.875	0.933333333
NLGN3 AND NOT-GAA	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CALHM1	0.903225806	0.875	0.933333333

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
SVOPL AND NOT-ZDHHC4	0.923076923	0.85714286	1
SVOPL AND NOT-ZDHHC7	0.923076923	0.85714286	1
SVOPL AND NOT-NDRG2	0.923076923	0.85714286	1
SVOPL AND NOT-TMEM181	0.923076923	0.85714286	1
SVOPL AND NOT-RNASE4	0.923076923	0.85714286	1
SVOPL AND NOT-DNAJC1	0.923076923	0.85714286	1
SVOPL AND NOT-PCDH20	0.909090909	1	0.833333333
SVOPL AND NOT-SLC20A1	0.923076923	0.85714286	1
SVOPL AND NOT-BNIP3L	0.923076923	0.85714286	1
SVOPL AND NOT-SPG7	0.923076923	0.85714286	1
SVOPL AND NOT-TSPO	0.923076923	0.85714286	1
SVOPL AND NOT-AGPAT9	0.923076923	0.85714286	1
SVOPL AND NOT-SEC11C	0.923076923	0.85714286	1
SVOPL AND NOT-CLDN8	0.916666667	0.91666667	0.916666667
SVOPL AND NOT-SMDT1	0.923076923	0.85714286	1
SVOPL AND NOT-TMEM59	0.923076923	0.85714286	1
IL18RAP AND NOT-TMEM71	0.909090909	1	0.833333333
SYNE4 AND NOT-SELM	0.916666667	0.91666667	0.916666667
NOT-TMED6 AND OR10H2	0.916666667	0.91666667	0.916666667
NOT-NDRG2 AND FAAH2	0.909090909	1	0.833333333
NOT-TMEM59 AND FAAH2	0.916666667	0.91666667	0.916666667
SYNE4 AND NOT-FAM134C	0.956521739	1	0.916666667
SYNE4 AND NOT-TMED3	0.916666667	0.91666667	0.916666667
SYNE4 AND NOT-SUN2	0.916666667	0.91666667	0.916666667
SYNE4 AND NOT-NDUFA1	0.916666667	0.91666667	0.916666667
SYNE4 AND NOT-NDUFB8	0.956521739	1	0.916666667
SYNE4 AND NOT-MAN1C1	0.916666667	0.91666667	0.916666667
SYNE4 AND NOT-C16orf58	0.909090909	1	0.833333333
SYNE4 AND NOT-SPG7	0.916666667	0.91666667	0.916666667
SYNE4 AND NOT-SSR3	0.909090909	1	0.833333333
SYNE4 AND NOT-VKORC1	0.909090909	1	0.833333333
NOT-MAN2A1 AND SFT2D2	0.916666667	0.91666667	0.916666667
NOT-CYB5R1 AND LRRC8E	0.923076923	0.85714286	1
NOT-NDRG2 AND DAPL1	0.909090909	1	0.833333333
GAL3ST4 AND NOT-SMPD1	0.909090909	1	0.833333333
NOT-TMEM59 AND GAL3ST4	0.909090909	1	0.833333333
NOT-TMEM59 AND LRRC8E	0.916666667	0.91666667	0.916666667
Neuroblastoma (Neuroblastoma)			
MARCH11	0.924855491	0.8988764	0.952380952
CHRNA3	0.927710843	0.93902439	0.916666667
ST8SIA2	0.922155689	0.92771084	0.916666667
MARCH11 AND SDC1	0.905882353	0.89534884	0.916666667
SLC10A4 AND NOT-SDC1	0.922222222	0.86458333	0.988095238
ST8SIA2 AND NOT-SDC1	0.944785276	0.97468354	0.916666667
VANGL2 AND NOT-SDC1	0.918238994	0.97333333	0.869047619
CHRNA3 AND CHRNA3	0.913580247	0.94871795	0.880952381
DIABLO AND NOT-SDC1	0.902439024	0.925	0.880952381
SLC10A4 AND NOT-WT1	0.917127072	0.8556701	0.988095238
ST8SIA2 AND NOT-WT1	0.922155689	0.92771084	0.916666667
CHRNA3 AND CHRNA3	0.913580247	0.94871795	0.880952381
MARCH11 AND MARCH11	0.924855491	0.8988764	0.952380952
ST8SIA2 AND NOT-FAP	0.938271605	0.97435897	0.904761905
CHRNA3 AND CHRNA3	0.913580247	0.94871795	0.880952381
SLC10A4 AND NOT-MAGEA1	0.917127072	0.8556701	0.988095238
ST8SIA2 AND NOT-ST8SIA2	0.904458599	0.97260274	0.845238095
CHRNA3 AND CHRNA3	0.913580247	0.94871795	0.880952381
MARCH11 AND NOT-MET	0.94047619	0.94047619	0.94047619
SLC10A4 AND NOT-MET	0.917127072	0.8556701	0.988095238
ST8SIA2 AND NOT-MET	0.938271605	0.97435897	0.904761905
CHRNA3 AND CHRNA3	0.913580247	0.94871795	0.880952381
MARCH11 AND MARCH11	0.924855491	0.8988764	0.952380952
SLC10A4 AND NOT-CD70	0.917127072	0.8556701	0.988095238
ST8SIA2 AND CD70	0.922155689	0.92771084	0.916666667
CHRNA3 AND CHRNA3	0.913580247	0.94871795	0.880952381
MARCH11 AND CD44	0.912280702	0.89655172	0.928571429
SLC10A4 AND NOT-CD44	0.912087912	0.84693878	0.988095238
ST8SIA2 AND NOT-ST8SIA2	0.904458599	0.97260274	0.845238095
CHRNA3 AND CHRNA3	0.913580247	0.94871795	0.880952381
SLC10A4 AND NOT-MS4A1	0.912087912	0.84693878	0.988095238
ST8SIA2 AND NOT-ST8SIA2	0.904458599	0.97260274	0.845238095
CHRNA3 AND CHRNA3	0.913580247	0.94871795	0.880952381
MARCH11 AND MARCH11	0.924855491	0.8988764	0.952380952
SLC10A4 AND NOT-CD160	0.917127072	0.8556701	0.988095238
ST8SIA2 AND NOT-CD160	0.922155689	0.92771084	0.916666667
CHRNA3 AND CHRNA3	0.913580247	0.94871795	0.880952381
SLC10A4 AND NOT-TNFSF11	0.917127072	0.8556701	0.988095238

Antigen Pair Logic	F1	Precision	Recall
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-C19orf26	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CCDC108	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TMCO4	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-MCOLN2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-RNF144B	0.903225806	0.875	0.933333333
NLGN3 AND NOT-GABRA2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-GABRA4	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CNEP1R1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SYT14	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CDRT15L2	0.965517241	1	0.933333333
NLGN3 AND NOT-STEAP1B	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ST6GALNAC3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-MFSD8	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-GABRE	0.903225806	0.875	0.933333333
NLGN3 AND NOT-GABRG1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-GABRP	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NPB	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-GABRR1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-FRMD3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-C1orf101	0.903225806	0.875	0.933333333
NLGN3 AND NOT-GABRR2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-KIAA1549L	0.903225806	0.875	0.933333333
NLGN3 AND NOT-SUN2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-BRI3	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-RHBDD3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-KLK5	0.903225806	0.875	0.933333333
NLGN3 AND NOT-BACE2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-B4GALNT1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-YIPF3	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SUMF2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-GALNS	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ABI3BP	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-RNF19A	0.903225806	0.875	0.933333333
NLGN3 AND NOT-GALNT1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-GALNT3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-SGMS1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-WFDC11	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ATL3	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-WFDC9	0.903225806	0.875	0.933333333
NLGN3 AND NOT-MRGPRX1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TAS2R39	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TAS2R40	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TAS2R41	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TAS2R50	0.903225806	0.875	0.933333333
NLGN3 AND NOT-IL4I1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-FAM205A	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-PNKD	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TMEM87A	0.903225806	0.875	0.933333333
NLGN3 AND NOT-C2CD2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TKFC	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-LRP10	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TMEM98	0.965517241	1	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-FDCSP	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CFAP61	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ABHD12	0.903225806	0.875	0.933333333
NLGN3 AND NOT-HERC4	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-LRIT1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TCTN3	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ABCA12	0.903225806	0.875	0.933333333
NLGN3 AND NOT-GIMAP2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-PCDHB5	0.903225806	0.875	0.933333333
NLGN3 AND NOT-STEAP2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NPHP4	0.903225806	0.875	0.933333333
NLGN3 AND NOT-OR1A2	0.903225806	0.875	0.933333333

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
CHRNA3 AND CHRNA3	0.913580247	0.94871795	0.880952381
MARCH11 AND MARCH11	0.924855491	0.8988764	0.952380952
FAM163A AND NOT-CA9	0.911111111	0.85416667	0.976190476
SLC10A4 AND NOT-CA9	0.921348315	0.87234043	0.976190476
ST8SIA2 AND NOT-CA9	0.922155689	0.92771084	0.916666667
CHRNA3 AND NOT-CA9	0.944099379	0.98701299	0.904761905
SLC10A4 AND NOT-MUC17	0.917127072	0.8556701	0.988095238
CHRNA3 AND NOT-MUC17	0.924050633	0.98648649	0.869047619
MARCH11 AND MARCH11	0.924855491	0.8988764	0.952380952
SLC10A4 AND NOT-FOLR2	0.917127072	0.8556701	0.988095238
ST8SIA2 AND FOLR2	0.922155689	0.92771084	0.916666667
CHRNA3 AND CHRNA3	0.913580247	0.94871795	0.880952381
MARCH11 AND NOT-FOLH1	0.927710843	0.93902439	0.916666667
FAM163A AND NOT-FOLH1	0.907103825	0.83838384	0.988095238
SLC10A4 AND NOT-FOLH1	0.954022989	0.92222222	0.988095238
CHRNA3 AND NOT-FOLH1	0.950617284	0.98717949	0.916666667
MARCH11 AND MARCH11	0.924855491	0.8988764	0.952380952
SLC10A4 AND NOT-ST8SIA1	0.916201117	0.86315789	0.976190476
CHRNA3 AND CHRNA3	0.913580247	0.94871795	0.880952381
SLC10A4 AND NOT-SLAMF7	0.917127072	0.8556701	0.988095238
ST8SIA2 AND SLAMF7	0.922155689	0.92771084	0.916666667
CHRNA3 AND CHRNA3	0.913580247	0.94871795	0.880952381
MARCH11 AND MARCH11	0.924855491	0.8988764	0.952380952
SLC10A4 AND NOT-CD19	0.911111111	0.85416667	0.976190476
ST8SIA2 AND NOT-ST8SIA2	0.904458599	0.97260274	0.845238095
CHRNA3 AND CHRNA3	0.913580247	0.94871795	0.880952381
FAM163A AND NOT-NCAM1	0.901098901	0.83673469	0.976190476
SLC10A4 AND NOT-NCAM1	0.937142857	0.9010989	0.976190476
CHRNA3 AND NOT-NCAM1	0.909090909	1	0.833333333
DIABLO AND NCAM1	0.925	0.97368421	0.880952381
SLC10A4 AND NOT-MLANA	0.917127072	0.8556701	0.988095238
CHRNA3 AND NOT-MLANA	0.93081761	0.98666667	0.880952381
MARCH11 AND MARCH11	0.924855491	0.8988764	0.952380952
SLC10A4 AND NOT-MUC1	0.917127072	0.8556701	0.988095238
ST8SIA2 AND NOT-MUC1	0.929936306	1	0.869047619
CHRNA3 AND NOT-MUC1	0.950617284	0.98717949	0.916666667
SLC10A4 AND NOT-TNFRSF17	0.917127072	0.8556701	0.988095238
CHRNA3 AND CHRNA3	0.913580247	0.94871795	0.880952381
SLC10A4 AND NOT-ROR1	0.917127072	0.8556701	0.988095238
CHRNA3 AND CHRNA3	0.913580247	0.94871795	0.880952381
SLC10A4 AND NOT-PSCA	0.917127072	0.8556701	0.988095238
ST8SIA2 AND NOT-PSCA	0.944099379	0.98701299	0.904761905
VANGL2 AND NOT-PSCA	0.9	0.94736842	0.857142857
CHRNA3 AND NOT-PSCA	0.950617284	0.98717949	0.916666667
MARCH11 AND NOT-ERBB4	0.946745562	0.94117647	0.952380952
SLC10A4 AND NOT-ERBB4	0.927374302	0.87368421	0.988095238
ST8SIA2 AND NOT-ERBB4	0.931677019	0.97402597	0.892857143
CHRNA3 AND NOT-ERBB4	0.9375	0.98684211	0.892857143
FAM163A AND NOT-MSLN	0.907103825	0.83838384	0.988095238
SLC10A4 AND NOT-MSLN	0.927374302	0.87368421	0.988095238
ST8SIA2 AND NOT-MSLN	0.944785276	0.97468354	0.916666667
CHRNA3 AND NOT-MSLN	0.944099379	0.98701299	0.904761905
MARCH11 AND MARCH11	0.924855491	0.8988764	0.952380952
SLC10A4 AND NOT-ERBB3	0.917127072	0.8556701	0.988095238
CHRNA3 AND CHRNA3	0.913580247	0.94871795	0.880952381
MARCH11 AND MARCH11	0.924855491	0.8988764	0.952380952
SLC10A4 AND NOT-CD33	0.922222222	0.86458333	0.988095238
ST8SIA2 AND NOT-CD33	0.938271605	0.97435897	0.904761905
CHRNA3 AND CHRNA3	0.913580247	0.94871795	0.880952381
MARCH11 AND MARCH11	0.924855491	0.8988764	0.952380952
SLC10A4 AND NOT-IGF1R	0.911111111	0.85416667	0.976190476
ST8SIA2 AND NOT-IGF1R	0.922155689	0.92771084	0.916666667
CHRNA3 AND CHRNA3	0.913580247	0.94871795	0.880952381
ALK AND NOT-KCNJ16	0.901960784	1	0.821428571
NOT-MARCH11 AND ALK	0.937142857	0.9010989	0.976190476
NOT-GLCCI1 AND ALK	0.909090909	0.92592593	0.892857143
NOT-ALK AND PODXL2	0.906976744	0.88636364	0.928571429
SLC10A4 AND NOT-ALK	0.906077348	0.84536082	0.976190476
NOT-ALK AND LRRN3	0.91954023	0.88888889	0.952380952
NOT-ST8SIA2 AND ALK	0.941176471	0.93023256	0.952380952
NOT-SLC6A2 AND ALK	0.917647059	0.90697674	0.928571429
ALK AND TMEM258	0.901960784	1	0.821428571
NOT-GDAP1L1 AND ALK	0.946745562	0.94117647	0.952380952
NOT-PCDHB6 AND ALK	0.933333333	0.95061728	0.916666667
NOT-ALK AND PTCHD1	0.925714286	0.89010989	0.964285714
NOT-DBH AND ALK	0.905027933	0.85263158	0.964285714

Antigen Pair Logic	F1	Precision	Recall
NLGN3 AND NOT-OR2F1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-OR1J4	0.903225806	0.875	0.933333333
NLGN3 AND NOT-B3GAT3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-KCNG2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CLEC4E	0.903225806	0.875	0.933333333
NLGN3 AND NOT-IL1RAPL2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CLDN17	0.903225806	0.875	0.933333333
NLGN3 AND NOT-GALNT8	0.903225806	0.875	0.933333333
NLGN3 AND NOT-OR7A17	0.903225806	0.875	0.933333333
NLGN3 AND NOT-OR5K1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-GCGR	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-OR1OJ1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-OR8B8	0.903225806	0.875	0.933333333
NLGN3 AND NOT-OR8G1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-OR10A3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-CNNM4	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-MYOF	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-GCNT1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-GCNT2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TSPAN16	0.903225806	0.875	0.933333333
NLGN3 AND NOT-OR12D2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-OR11A1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-OR10H3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-OR10H2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-AMELX	0.903225806	0.875	0.933333333
NLGN3 AND NOT-TBL2	0.903225806	0.875	0.933333333
SLC1A2 AND TBL2	0.909090909	0.833333333	1
NLGN3 AND NOT-GDF9	0.903225806	0.875	0.933333333
NLGN3 AND NOT-OR7C2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-OR7A5	0.903225806	0.875	0.933333333
NLGN3 AND NOT-OR7C1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-GDNF	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-ADGRF1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-OR2J2	0.903225806	0.875	0.933333333
NLGN3 AND NOT-OR2H1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-AMFR	0.903225806	0.875	0.933333333
NLGN3 AND NOT-B4GALT1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-NLGN3	0.903225806	0.875	0.933333333
NLGN3 AND NOT-STEAP1	0.903225806	0.875	0.933333333
NLGN3 AND NOT-GGT5	0.933333333	0.933333333	0.933333333
SLC4A4 AND NOT-GPR160	0.928571429	1	0.866666667
NLGN3 AND NOT-AMHR2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-CNPPD1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-GJA4	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SIGLEC7	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-DKKL1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SLC39A1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SERP1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-PGAP2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-KCNMB4	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-GNS	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-GOLGB1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SLCO4A1	0.933333333	0.933333333	0.933333333
GPM6A AND DOLPP1	0.909090909	0.833333333	1
NLGN3 AND NOT-GPR3	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-OR8D2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-CD163L1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-B4GALNT3	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-DPY19L2	0.933333333	0.933333333	0.933333333
ZDHHC22 AND NOT-CCDC136	0.928571429	1	0.866666667
NLGN3 AND NOT-ANG	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SLC26A11	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SYPL2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SLC25A34	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SIRPB2	0.965517241	1	0.933333333
NLGN3 AND NOT-RNF149	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-IGSF10	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-GPR31	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-CYP4V2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-COX18	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SCARA5	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ATP11C	0.933333333	0.933333333	0.933333333

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
ALK AND NOT-TMC4	0.901960784	1	0.821428571
CHRNA3 AND CHRNA3	0.913580247	0.94871795	0.880952381
NOT-PCDHB10 AND ALK	0.934911243	0.92941176	0.94047619
DIABLO AND ALK	0.9	0.94736842	0.857142857
MARCH11 AND MARCH11	0.924855491	0.8988764	0.952380952
SLC10A4 AND NOT-TNFRSF10B	0.911111111	0.85416667	0.976190476
CHRNA3 AND NOT-TNFRSF10B	0.903225806	0.98591549	0.833333333
ST8SIA2 AND NOT-IL13RA2	0.922155689	0.92771084	0.916666667
CHRNA3 AND CHRNA3	0.913580247	0.94871795	0.880952381
SLC10A4 AND NOT-KDR	0.917127072	0.8556701	0.988095238
PCSK1N AND CD276	0.909090909	0.92592593	0.892857143
MARCH11 AND MARCH11	0.924855491	0.8988764	0.952380952
ST8SIA3 AND CD276	0.928571429	0.92857143	0.928571429
NOT-EGFR AND CD276	0.935672515	0.91954023	0.952380952
SYT11 AND CD276	0.904761905	0.9047619	0.904761905
SVOP AND CD276	0.9	0.94736842	0.857142857
SLC10A4 AND NOT-CD276	0.917127072	0.8556701	0.988095238
ST8SIA2 AND NOT-ST8SIA2	0.904458599	0.97260274	0.845238095
SCN3B AND CD276	0.914634146	0.9375	0.892857143
NRSN1 AND CD276	0.9	0.94736842	0.857142857
HMP19 AND CD276	0.906832298	0.94805195	0.869047619
GPR19 AND CD276	0.920245399	0.94936709	0.892857143
RTN1 AND CD276	0.944785276	0.97468354	0.916666667
FAIM2 AND CD276	0.901734104	0.87640449	0.928571429
LHFPL4 AND CD276	0.9	0.94736842	0.857142857
CHRNA3 AND CHRNA3	0.913580247	0.94871795	0.880952381
NOT-ALPK1 AND CD276	0.90797546	0.93670886	0.880952381
MARCH11 AND MARCH11	0.924855491	0.8988764	0.952380952
SLC10A4 AND NOT-IL11RA	0.917127072	0.8556701	0.988095238
ST8SIA2 AND NOT-ST8SIA2	0.904458599	0.97260274	0.845238095
CHRNA3 AND CHRNA3	0.913580247	0.94871795	0.880952381
SLC10A4 AND NOT-EPHA2	0.917127072	0.8556701	0.988095238
ST8SIA2 AND NOT-EPHA2	0.944785276	0.97468354	0.916666667
VANGL2 AND NOT-EPHA2	0.911392405	0.97297297	0.857142857
CHRNA3 AND NOT-EPHA2	0.93081761	0.98666667	0.880952381
SLC10A4 AND NOT-IL3RA	0.917127072	0.8556701	0.988095238
ST8SIA2 AND NOT-IL3RA	0.911392405	0.97297297	0.857142857
CHRNA3 AND CHRNA3	0.913580247	0.94871795	0.880952381
SLC10A4 AND NOT-GPC3	0.917127072	0.8556701	0.988095238
CHRNA3 AND NOT-GPC3	0.927710843	0.93902439	0.916666667
SLC10A4 AND NOT-MUC16	0.917127072	0.8556701	0.988095238
CHRNA3 AND NOT-MUC16	0.950617284	0.98717949	0.916666667
MARCH11 AND MARCH11	0.924855491	0.8988764	0.952380952
SLC10A4 AND NOT-EPHA3	0.917127072	0.8556701	0.988095238
ST8SIA2 AND NOT-EPHA3	0.938271605	0.97435897	0.904761905
CHRNA3 AND CHRNA3	0.913580247	0.94871795	0.880952381
PCDHB10 AND NOT-EPHA3	0.909090909	1	0.833333333
SLC10A4 AND NOT-CD38	0.916201117	0.86315789	0.976190476
ST8SIA2 AND NOT-ST8SIA2	0.904458599	0.97260274	0.845238095
CHRNA3 AND CHRNA3	0.913580247	0.94871795	0.880952381
MARCH11 AND ERBB2	0.912280702	0.89655172	0.928571429
SLC10A4 AND NOT-ERBB2	0.917127072	0.8556701	0.988095238
ST8SIA2 AND NOT-ERBB2	0.922155689	0.92771084	0.916666667
VANGL2 AND NOT-ERBB2	0.918238994	0.97333333	0.869047619
CHRNA3 AND CHRNA3	0.913580247	0.94871795	0.880952381
DIABLO AND NOT-ERBB2	0.90797546	0.93670886	0.880952381
SLC10A4 AND NOT-TNFRSF10A	0.917127072	0.8556701	0.988095238
ST8SIA2 AND NOT-TNFRSF10A	0.950617284	0.98717949	0.916666667
GAL AND NOT-TNFRSF10A	0.903614458	0.91463415	0.892857143
VANGL2 AND NOT-TNFRSF10A	0.918238994	0.97333333	0.869047619
CHRNA3 AND NOT-TNFRSF10A	0.927710843	0.93902439	0.916666667
DIABLO AND NOT-TNFRSF10A	0.90797546	0.93670886	0.880952381
MARCH11 AND MARCH11	0.924855491	0.8988764	0.952380952
SLC10A4 AND NOT-L1CAM	0.917127072	0.8556701	0.988095238
ST8SIA2 AND L1CAM	0.922155689	0.92771084	0.916666667
NOT-GDAP1L1 AND L1CAM	0.901734104	0.87640449	0.928571429
CHRNA3 AND CHRNA3	0.913580247	0.94871795	0.880952381
DIABLO AND L1CAM	0.925	0.97368421	0.880952381
MARCH11 AND MARCH11	0.924855491	0.8988764	0.952380952
SLC10A4 AND NOT-CD22	0.917127072	0.8556701	0.988095238
ST8SIA2 AND NOT-ST8SIA2	0.904458599	0.97260274	0.845238095
CHRNA3 AND CHRNA3	0.913580247	0.94871795	0.880952381
MARCH11 AND NOT-B4GALNT1	0.901234568	0.93589744	0.869047619
SLC10A4 AND NOT-B4GALNT1	0.911111111	0.85416667	0.976190476
ST8SIA2 AND NOT-ST8SIA2	0.904458599	0.97260274	0.845238095
CHRNA3 AND CHRNA3	0.913580247	0.94871795	0.880952381

Antigen Pair Logic	F1	Precision	Recall
NLGN3 AND NOT-YIPF6	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-P2RY8	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-DEXI	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-OSTM1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SLC6A16	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-MAGEH1	0.933333333	0.933333333	0.933333333
GRIK3 AND NOT-KCNS3	0.909090909	0.833333333	1
GRIK3 AND NOT-SCN4B	0.9375	0.882352941	1
GRIK3 AND NOT-ELOVL7	0.909090909	0.833333333	1
GRIK3 AND NOT-PKDCC	0.9375	0.882352941	1
NLGN3 AND NOT-ANPEP	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-GNL2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ST8SIA5	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-GUSB	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ALG6	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-CERS2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SLC2A8	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-GZMK	0.965517241	1	0.933333333
NLGN3 AND NOT-HEXB	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-HGF	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-HLA-E	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-MR1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-HMGCR	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-HMOX1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-VN1R2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-HSD3B2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-DNAJC4	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-HTR2B	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-HTR4	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ICAM3	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TMEM52	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TSPAN33	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SERINC2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-AMIGO2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-IGF2R	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SLCO4C1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-IL1R1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-IL4R	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-IL6R	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-IL10RA	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-IL10RB	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-IL12B	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-IL15RA	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-INSR	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ITGA2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ITGA5	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ITGA9	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ITGAE	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ITGAM	0.933333333	0.933333333	0.933333333
GDAP1L1 AND TMEM179B	0.909090909	0.833333333	1
NLGN3 AND NOT-KCNA7	0.933333333	0.933333333	0.933333333
KCNQ2 AND NOT-KCNB1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-PEAR1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-MIA3	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TMEM110	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-KCNJ5	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-KCNJ8	0.933333333	0.933333333	0.933333333
KCNJ10 AND TSPAN11	0.909090909	0.833333333	1
KCNJ10 AND NAT10	0.909090909	0.833333333	1
KCNJ10 AND FAM57A	0.909090909	0.833333333	1
KCNJ10 AND GPAA1	0.909090909	0.833333333	1
NLGN3 AND NOT-KCNK2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-KCNK3	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-KCNQ1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-KDR	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-KIR3DL2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TMEM189	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TMEM220	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-LDLR	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-LHCGR	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-FADS3	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-HACD4	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-IFITM10	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-PPAPDC2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-LRP5	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-LRPAP1	0.933333333	0.933333333	0.933333333

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
MARCH11 AND MARCH11	0.924855491	0.8988764	0.952380952
SLC10A4 AND NOT-MUC13	0.917127072	0.8556701	0.988095238
ST8SIA2 AND NOT-MUC13	0.922155689	0.92771084	0.916666667
CHRNA3 AND NOT-MUC13	0.950617284	0.98717949	0.916666667
SLC10A4 AND NOT-GPA33	0.917127072	0.8556701	0.988095238
ST8SIA2 AND NOT-GPA33	0.95	1	0.904761905
CHRNA3 AND NOT-GPA33	0.950617284	0.98717949	0.916666667
SLC10A4 AND NOT-AXL	0.917127072	0.8556701	0.988095238
ST8SIA2 AND NOT-AXL	0.944785276	0.97468354	0.916666667
CHRNA3 AND NOT-AXL	0.924050633	0.98648649	0.869047619
SLC10A4 AND NOT-ITGB3	0.917127072	0.8556701	0.988095238
ST8SIA2 AND NOT-ITGB3	0.95	1	0.904761905
CHRNA3 AND NOT-ITGB3	0.95	1	0.904761905
PCDHB10 AND NOT-ITGB3	0.909090909	1	0.833333333
SLC10A4 AND NOT-CSPG4	0.917127072	0.8556701	0.988095238
ST8SIA2 AND NOT-CSPG4	0.917197452	0.98630137	0.857142857
CHRNA3 AND CHRNA3	0.913580247	0.94871795	0.880952381
CHRNA3 AND NOT-SMIM6	0.956521739	1	0.916666667
FAM163A AND NOT-SMIM6	0.970760234	0.95402299	0.988095238
SLC10A4 AND NOT-SMIM6	0.976470588	0.96511628	0.988095238
ST8SIA2 AND NOT-SMIM6	0.956521739	1	0.916666667
SLC10A4 AND NOT-NEMP2	0.922222222	0.86458333	0.988095238
MARCH11 AND NOT-NEMP2	0.946745562	0.94117647	0.952380952
CHRNA3 AND NOT-HCN4	0.95	1	0.904761905
MARCH11 AND NOT-PET117	0.952380952	0.95238095	0.952380952
FAM163A AND NOT-APELA	0.958083832	0.96385542	0.952380952
ST8SIA2 AND NOT-APELA	0.950617284	0.98717949	0.916666667
SLC10A4 AND NOT-TMEM178B	0.922222222	0.86458333	0.988095238
MARCH11 AND NOT-TMEM178B	0.946745562	0.94117647	0.952380952
CHRNA3 AND NOT-MICA	0.950617284	0.98717949	0.916666667
SLC10A4 AND NOT-MICA	0.922222222	0.86458333	0.988095238
CHRNA3 AND NOT-100507547?	0.943396226	1	0.892857143
SLC10A4 AND NOT-100507547?	0.927374302	0.87368421	0.988095238
CHRNA3 AND UBA2	0.956521739	1	0.916666667
SLC10A4 AND UBA2	0.922222222	0.86458333	0.988095238
ST8SIA2 AND UBA2	0.956521739	1	0.916666667
CHRNA3 AND NOT-ABCC9	0.950617284	0.98717949	0.916666667
SLC10A4 AND NOT-MUC12	0.936416185	0.91011236	0.964285714
SLC10A4 AND NOT-UST	0.970414201	0.96470588	0.976190476
ST8SIA2 AND NOT-UST	0.943396226	1	0.892857143
ST8SIA2 AND NOT-PDZK1IP1	0.943396226	1	0.892857143
CHRNA3 AND NOT-DHRS9	0.950617284	0.98717949	0.916666667
GPR19 AND NOT-DHRS9	0.963414634	0.9875	0.94047619
GAL AND NOT-DHRS9	0.936708861	1	0.880952381
ST8SIA2 AND NOT-DHRS9	0.956521739	1	0.916666667
FAM163A AND CNIH1	0.93258427	0.88297872	0.988095238
SLC10A4 AND NOT-TENM1	0.927374302	0.87368421	0.988095238
CHRNA3 AND NOT-101805491?	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-GPA33	0.950617284	0.98717949	0.916666667
ST8SIA2 AND NOT-GPA33	0.95	1	0.904761905
SLC10A4 AND NOT-MSLN	0.927374302	0.87368421	0.988095238
CHRNA3 AND NOT-PRSS16	0.950617284	0.98717949	0.916666667
FAM163A AND NOT-PRSS16	0.937853107	0.89247312	0.988095238
SLC10A4 AND NOT-PRSS16	0.927374302	0.87368421	0.988095238
MARCH11 AND NOT-PRSS16	0.946745562	0.94117647	0.952380952
SLC10A4 AND NOT-RTN3	0.922222222	0.86458333	0.988095238
CHRNA3 AND NOT-NMUR1	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-B3GALT5	0.950617284	0.98717949	0.916666667
SLC10A4 AND NOT-B3GALT5	0.93258427	0.88297872	0.988095238
ST8SIA2 AND NOT-B3GALT5	0.950617284	0.98717949	0.916666667
FAM163A AND NOT-SIRPB1	0.942528736	0.91111111	0.976190476
SLC10A4 AND NOT-SIRPB1	0.937142857	0.9010989	0.976190476
CHRNA3 AND NOT-B3GNT3	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-TLR6	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-PKDREJ	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-CDS1	0.95	1	0.904761905
FAM163A AND NOT-CDS1	0.943181818	0.90217391	0.988095238
SLC10A4 AND NOT-CDS1	0.959537572	0.93258427	0.988095238
MARCH11 AND NOT-CDS1	0.946745562	0.94117647	0.952380952
ST8SIA2 AND NOT-CDS1	0.95	1	0.904761905
CHRNA3 AND NOT-CDSN	0.950617284	0.98717949	0.916666667
SLC10A4 AND NOT-TIMM17A	0.922222222	0.86458333	0.988095238
FAM163A AND CLGN	0.927374302	0.87368421	0.988095238
CHRNA3 AND NOT-CRTAP	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-SEMA6B	0.956521739	1	0.916666667
CHRNA3 AND NOT-SEMA4D	0.950617284	0.98717949	0.916666667

Antigen Pair Logic	F1	Precision	Recall
NLGN3 AND NOT-DUOXA2	0.965517241	1	0.933333333
NLGN3 AND NOT-LY75	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TACSTD2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-EPCAM	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SMAD2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-MAG	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-MAN1A1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-MAN2A1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-MARS	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ARSD	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-CD46	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-DNAJB9	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ART4	0.928571429	1	0.866666667
NLGN3 AND NOT-MGAT1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-MGST1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-C1orf186	0.965517241	1	0.933333333
NLGN3 AND NOT-SHISA8	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SMIM4	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TMEM150C	0.933333333	0.933333333	0.933333333
SLC1A2 AND TSPAN11	0.909090909	0.833333333	1
GDAP1L1 AND TSPAN11	0.909090909	0.833333333	1
NLGN3 AND NOT-ASPH	0.933333333	0.933333333	0.933333333
NLGN3 AND NDUFB1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-NDUFB8	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-NINJ1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-NPR3	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ROR2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-DDR2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-NUCB1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-GPX8	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ATP4A	0.933333333	0.933333333	0.933333333
OMG AND GPAA1	0.909090909	0.833333333	1
NLGN3 AND NOT-OXA1L	0.933333333	0.933333333	0.933333333
SLC1A2 AND P2RX4	0.909090909	0.833333333	1
GDAP1L1 AND P2RX4	0.909090909	0.833333333	1
LINGO1 AND P2RX4	0.909090909	0.833333333	1
NLGN3 AND NOT-CYHR1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TAS2R1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-F11R	0.933333333	0.933333333	0.933333333
NOT-F11R AND TMEM259	0.928571429	1	0.866666667
NLGN3 AND NOT-TMED5	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SLC35B3	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SLC35C2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TXNDC12	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ADIPOR1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SCCPDH	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TMEM69	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-NT5C3A	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-BET1L	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ERGIC2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-PCDH12	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SLC15A3	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ZDHHC3	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-MS4A4A	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TEX264	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TMEM138	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-MTFP1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-NBAS	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-LSR	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-PDGFRB	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SLC25A39	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-FKBP7	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ENPP1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ENPP3	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TMEM8B	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-PECAM1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ASIC5	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-PEX10	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-PF4	0.928571429	1	0.866666667
SLC4A4 AND NOT-ATP8B1	0.928571429	1	0.866666667
NLGN3 AND NOT-ABCB4	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-PIGA	0.933333333	0.933333333	0.933333333
SLC4A4 AND NOT-PIGR	0.928571429	1	0.866666667
NLGN3 AND NOT-ACP1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TPCN1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-PLIN1	0.933333333	0.933333333	0.933333333

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
CHRNA3 AND NOT-C6orf10	0.943396226	1	0.892857143
CHRNA3 AND NOT-CD226	0.950617284	0.98717949	0.916666667
SLC10A4 AND NOT-CSPG5	0.922222222	0.86458333	0.988095238
ST8SIA2 AND NOT-CLDN16	0.936708861	1	0.880952381
SLC10A4 AND NOT-FUT9	0.927374302	0.87368421	0.988095238
CHRNA3 AND NOT-RRH	0.950617284	0.98717949	0.916666667
FAM163A AND NOT-RRH	0.937853107	0.89247312	0.988095238
SLC10A4 AND NOT-RRH	0.948571429	0.91208791	0.988095238
ST8SIA2 AND NOT-RRH	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-CERS1	0.950617284	0.98717949	0.916666667
SLC10A4 AND NOT-CERS1	0.922222222	0.86458333	0.988095238
ST8SIA2 AND NOT-CERS1	0.950617284	0.98717949	0.916666667
SLC10A4 AND NOT-NRG3	0.922222222	0.86458333	0.988095238
SLC10A4 AND NOT-ZMYND11	0.922222222	0.86458333	0.988095238
SLC10A4 AND NOT-CYSLTR1	0.922222222	0.86458333	0.988095238
CHRNA3 AND NOT-CCR9	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-GJB6	0.95	1	0.904761905
NRSN1 AND NOT-GJB6	0.95	1	0.904761905
SMIM17 AND NOT-GJB6	0.95	1	0.904761905
FAM163A AND NOT-GJB6	0.970414201	0.96470588	0.976190476
SLC10A4 AND NOT-GJB6	0.982035928	0.98795181	0.976190476
GPR19 AND NOT-GJB6	0.958083832	0.96385542	0.952380952
MARCH11 AND NOT-GJB6	0.946745562	0.94117647	0.952380952
ST8SIA3 AND NOT-GJB6	0.952380952	0.95238095	0.952380952
HMP19 AND NOT-GJB6	0.956521739	1	0.916666667
ST8SIA2 AND NOT-GJB6	0.956521739	1	0.916666667
CHRNA3 AND NOT-SLC26A1	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-CD300C	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-EDDM3A	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-CEACAM7	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-LYVE1	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-OCLM	0.950617284	0.98717949	0.916666667
FAM163A AND NOT-OCLM	0.922222222	0.86458333	0.988095238
SLC10A4 AND NOT-OCLM	0.943181818	0.90217391	0.988095238
CHRNA3 AND NOT-CEACAM4	0.95	1	0.904761905
CHRNA3 AND NOT-ADCY2	0.950617284	0.98717949	0.916666667
SLC10A4 AND NOT-ADCY2	0.922222222	0.86458333	0.988095238
SLC10A4 AND NOT-MAN1A2	0.927374302	0.87368421	0.988095238
CHRNA3 AND NOT-EDAR	0.950617284	0.98717949	0.916666667
MARCH11 AND NOT-EDAR	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-GPR75	0.950617284	0.98717949	0.916666667
SLC10A4 AND NOT-GPR75	0.922222222	0.86458333	0.988095238
ST8SIA2 AND NOT-GPR75	0.950617284	0.98717949	0.916666667
MARCH11 AND NOT-AFG3L2	0.952380952	0.95238095	0.952380952
CHRNA3 AND NOT-TMED10	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-SLC38A3	0.956521739	1	0.916666667
SLC10A4 AND NOT-SLC27A4	0.922222222	0.86458333	0.988095238
SLC10A4 AND NOT-TMED1	0.922222222	0.86458333	0.988095238
CHRNA3 AND NOT-UPK1A	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-ADAM30	0.950617284	0.98717949	0.916666667
SLC10A4 AND NOT-PLA2G16	0.937142857	0.9010989	0.976190476
CHRNA3 AND NOT-HHLA2	0.950617284	0.98717949	0.916666667
ST8SIA2 AND NOT-HHLA2	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-ADCY5	0.956521739	1	0.916666667
CHRNA3 AND NOT-GPR45	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-PTGDR2	0.950617284	0.98717949	0.916666667
SLC10A4 AND NOT-PTGDR2	0.93258427	0.88297872	0.988095238
CHRNA3 AND NOT-SLC6A14	0.950617284	0.98717949	0.916666667
ST8SIA2 AND NOT-SLC6A14	0.950617284	0.98717949	0.916666667
SLC10A4 AND NOT-CMTM7	0.922222222	0.86458333	0.988095238
SCN3B AND NRM	0.947368421	0.93103448	0.964285714
CHRNA3 AND NOT-STX1B	0.943396226	1	0.892857143
FAM163A AND NOT-SCN11A	0.947368421	0.93103448	0.964285714
CHRNA3 AND NOT-HOGA1	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-CYP4F8	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-B4GALT7	0.950617284	0.98717949	0.916666667
SLC10A4 AND NOT-MGAT4A	0.927374302	0.87368421	0.988095238
CHRNA3 AND NOT-CHRM4	0.950617284	0.98717949	0.916666667
FAM163A AND NOT-CHRM5	0.942528736	0.91111111	0.976190476
SLC10A4 AND NOT-CHRM5	0.970414201	0.96470588	0.976190476
ST8SIA2 AND NOT-CHRM5	0.956521739	1	0.916666667
SLC10A4 AND NOT-RNF13	0.970414201	0.96470588	0.976190476
CHRNA3 AND NOT-CHRNA4	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-SLC2A13	0.936708861	1	0.880952381
CHRNA3 AND NOT-CHRND	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-PKD1L2	0.950617284	0.98717949	0.916666667

Antigen Pair Logic	F1	Precision	Recall
NLGN3 AND NOT-PLN	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-PLP2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-FXYD5	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-IL20RA	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ATP7A	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-DUOX1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-A4GALT	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-WNT4	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-YIPF1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-POR	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TMCO1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-VSIG10	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-P4HTM	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-LEPROT	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-LY6K	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-DNAJB12	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TRPM4	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-CNNM2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-PPIC	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-CDHR2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-WBP1L	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-PAQR5	0.965517241	1	0.933333333
NLGN3 AND NOT-DNAJC25	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TMEM104	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-RETSAT	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-RNF43	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-PQLC2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TMEM260	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TMEM161A	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-PARP16	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SLC35F6	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-MARC2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-C19orf24	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-STX17	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TMEM248	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TMEM45A	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TMEM51	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-AUP1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ZDHHC4	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-RMDN3	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-NAT10	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SLC38A7	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SLC47A1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TYW1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TMEM39A	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TMEM143	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TMEM140	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-MCOLN3	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-AVPR1A	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-DRAM1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SYNJ2BP	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-LRRC59	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-PARL	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ETNK1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SLC39A4	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-FLVCR2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-PIGV	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TMEM127	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SLC30A6	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-LMBR1L	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-HHAT	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-NGLY1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ERMARD	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ST6GALNAC1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-RNF130	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-GLT8D1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-EMC3	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TEX2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-GPRC5C	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ACSS2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SLC50A1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ALG1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-MOSPD1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-PRLR	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SUSD2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-MRAP	0.933333333	0.933333333	0.933333333

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
CHRNA3 AND NOT-CSMD3	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-ELFN2	0.956521739	1	0.916666667
CHRNA3 AND NOT-SLAMF6	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-C1QTNF1	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-SLC26A9	0.95	1	0.904761905
CHRNA3 AND NOT-OMA1	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-SLC5A11	0.956521739	1	0.916666667
CHRNA3 AND NOT-EVI5L	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-SLC22A12	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-ACSM1	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-PANX3	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-SLC26A8	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-GRIN3B	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-SLC18B1	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-CATSPER2	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-MRGPRX2	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-MRGPRX3	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-ADCYAP1R1	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-GPR62	0.956521739	1	0.916666667
CHRNA3 AND NOT-CLCNKB	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-OR5P2	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-OR5P3	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-TMEM52B	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-ANO4	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-SLC24A4	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-DEGS2	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-SLC51B	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-CCR3	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-ACKR2	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-CD300LB	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-B4GALNT2	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-SPNS2	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-GJD3	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-CNGA4	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-LRRC25	0.95	1	0.904761905
CHRNA3 AND NOT-CYP4F22	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-GJB4	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-DCST2	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-TMEM125	0.956521739	1	0.916666667
CHRNA3 AND NOT-FITM2	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-TMEM150A	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-CNTNAP5	0.943396226	1	0.892857143
CHRNA3 AND NOT-SGPP2	0.95	1	0.904761905
CHRNA3 AND NOT-ACVR1C	0.943396226	1	0.892857143
CHRNA3 AND NOT-LYPD6B	0.95	1	0.904761905
CHRNA3 AND NOT-CD200R1	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-TMEM207	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-TMPRSS11B	0.956521739	1	0.916666667
CHRNA3 AND NOT-EVC2	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-IL31RA	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-SLCO6A1	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-TMEM171	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-TMEM174	0.950617284	0.98717949	0.916666667
CHRNA3 AND COX7A2	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-TAAR9	0.956521739	1	0.916666667
CHRNA3 AND NOT-TAAR1	0.943396226	1	0.892857143
CHRNA3 AND NOT-ADORA1	0.943396226	1	0.892857143
CHRNA3 AND NOT-RAET1E	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-DPCR1	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-WBSCR28	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-TMEM139	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-OR6B1	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-OR2F2	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-SSMEM1	0.956521739	1	0.916666667
CHRNA3 AND NOT-RNF183	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-ADGRG4	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-CRHR1	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-CRHR2	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-AADAC	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-ASB11	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-ASB5	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-TRPM6	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-C20orf173	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-CST11	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-SIRPA	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-CRY2	0.950617284	0.98717949	0.916666667

Antigen Pair Logic	F1	Precision	Recall
NLGN3 AND NOT-GKN1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TRPV5	0.965517241	1	0.933333333
NLGN3 AND NOT-TMPRSS15	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-PSEN2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-GPR108	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-CHPT1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ENTPD7	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-PNPLA2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-CD248	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-RHBG	0.928571429	1	0.866666667
NLGN3 AND NOT-MAN1C1	0.933333333	0.933333333	0.933333333
NLGN3 AND RTN4	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TMEM159	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-DOLPP1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-MCOLN1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ATP10A	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ATP10D	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ADGRG6	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ERGIC1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SNX14	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-MRS2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-PTGFRN	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-RHBDD2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-GRAMD1B	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-RNF150	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-MTUS1	0.928571429	1	0.866666667
NLGN3 AND NOT-SEMA6A	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-PTPN2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-WDR19	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SLAMF7	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-PTPRH	0.965517241	1	0.933333333
NLGN3 AND NOT-PVR	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-PVRL2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-MS4A7	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-FAM3A	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-RNASE1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-RNASE4	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ELOVL5	0.933333333	0.933333333	0.933333333
NLGN3 AND RPL37A	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-BCS1L	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-RRBP1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SCN4B	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SCN7A	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SCNN1A	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-CEACAM1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SDC1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-THADA	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-BIK	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-MANBAL	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SELPLG	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-PERP	0.928571429	1	0.866666667
NLGN3 AND NOT-POPDC2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ELSPBP1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-XYLT2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-DNAJC1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ATL2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-MS4A6A	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SMIM5	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-LMBR1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-C1orf233	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-MOSPD3	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SYNDIG1L	0.965517241	1	0.933333333
NLGN3 AND NOT-C16orf58	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-FNDC3B	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ST3GAL1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-FNDC4	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-PORCN	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ST3GAL4	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TMEM135	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SLC1A5	0.965517241	1	0.933333333
NLGN3 AND NOT-SLC4A1	0.965517241	1	0.933333333
NLGN3 AND NOT-SLC6A2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SLC12A2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SLC16A2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SLCO2A1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SLC22A3	0.933333333	0.933333333	0.933333333

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
CHRNA3 AND NOT-CSF1	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-TMEM86A	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-OR10A5	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-ISM2	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-TMCO5A	0.956521739	1	0.916666667
CHRNA3 AND NOT-CMTM4	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-GSG1L	0.943396226	1	0.892857143
CHRNA3 AND NOT-MGAT5B	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-RBFOX3	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-SLC47A2	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-ADRA1D	0.956521739	1	0.916666667
CHRNA3 AND NOT-C19orf18	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-TMEM190	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-TMC4	0.956521739	1	0.916666667
CHRNA3 AND NOT-ZNRF4	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-MFSD4	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-DCST1	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-IL23R	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-CTLA4	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-C1orf210	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-ADIG	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-SLC9B1	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-PROM2	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-SLC23A3	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-RMDN2	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-PPM1L	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-ADRA2B	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-C4orf26	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-SLC25A48	0.956521739	1	0.916666667
CHRNA3 AND NOT-MARVELD2	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-SLC2A12	0.956521739	1	0.916666667
CHRNA3 AND NOT-NKAIN2	0.956521739	1	0.916666667
CHRNA3 AND NOT-CLEC2L	0.956521739	1	0.916666667
CHRNA3 AND NOT-CYP2A13	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-PEBP4	0.943396226	1	0.892857143
CHRNA3 AND NOT-CYP4B1	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-OR1Q1	0.956521739	1	0.916666667
CHRNA3 AND NOT-CYP19A1	0.95	1	0.904761905
CHRNA3 AND NOT-CCDC67	0.943396226	1	0.892857143
CHRNA3 AND NOT-CLECL1	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-SLC5A8	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-TMEM229B	0.95	1	0.904761905
CHRNA3 AND NOT-SYNE3	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-MDGA2	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-MFSD6L	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-TRPV3	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-SPPL2C	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-SYNE4	0.95	1	0.904761905
CHRNA3 AND NOT-DCT	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-TMPRSS6	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-ADGRF3	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-SPTSSB	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-RNF133	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-PKD1L1	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-TMEM252	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-CLEC4C	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-HTR3C	0.943396226	1	0.892857143
CHRNA3 AND NOT-CLYBL	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-DIO2	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-AGER	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-DRD1	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-DRD3	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-DRD4	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-DRD5	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-DSC1	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-DSCAM	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-EFNB3	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-EGF	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-MEGF8	0.95	1	0.904761905
CHRNA3 AND NOT-MEGF9	0.956521739	1	0.916666667
CHRNA3 AND NOT-EGFR	0.956521739	1	0.916666667
CHRNA3 AND NOT-GRAMD2	0.956521739	1	0.916666667
CHRNA3 AND NOT-TMEM61	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-CYP4Z1	0.943396226	1	0.892857143
CHRNA3 AND NOT-GABRR3	0.956521739	1	0.916666667
CHRNA3 AND NOT-TVP23C	0.950617284	0.98717949	0.916666667

Antigen Pair Logic	F1	Precision	Recall
NLGN3 AND NOT-CYP4F12	0.965517241	1	0.933333333
NLGN3 AND NOT-SMPD1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SMPD2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SIGLEC1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SOAT1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SPAG4	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SPG7	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SRD5A2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ST14	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-STX3	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-STX4	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SURF4	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-BST1	0.928571429	1	0.866666667
NLGN3 AND NOT-SYPL1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TCTA	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TGFB3	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TGFBR2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TGFBR3	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-THBD	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TLR2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TLR5	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TSPAN8	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TSPAN4	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TM7SF2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TNFRSF1B	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TPTE	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TLCD2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-GXYLT2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-C5AR1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-UGT2B4	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-VRK2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-WNT11	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-CACNA1D	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SLC30A1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-BSND	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-MOGS	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-IL1R2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-CXCR4	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-FZD5	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SEMA3B	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SLMAP	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SLC25A20	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TMEM106C	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ALG8	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ELOVL6	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TMEM109	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ALG12	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TMUB2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-MBOAT7	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-MUL1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-OCEL1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SRD5A3	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-STEAP4	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ACBD4	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ZDHHC11	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-RNF122	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-THSD4	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-CYBRD1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-KIAA0319L	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-UBXN8	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ERMP1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-CPED1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ELOVL7	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SLC8B1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-LRRC8E	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-CWH43	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-MYCT1	0.928571429	1	0.866666667
NLGN3 AND NOT-TMEM134	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ALPK1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-PAAF1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-HSD3B7	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SLC44A4	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-CYB5B	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SLC2A10	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-OR2C3	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-PTDSS2	0.933333333	0.933333333	0.933333333

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
CHRNA3 AND NOT-SPNS3	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-ADAM32	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-CT83	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-TMEM31	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-EPHA1	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-EPHA4	0.956521739	1	0.916666667
CHRNA3 AND NOT-EPHA8	0.936708861	1	0.880952381
CHRNA3 AND NOT-EPO	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-LVRN	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-ETFDH	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-EXTL3	0.956521739	1	0.916666667
CHRNA3 AND NOT-FAAH	0.956521739	1	0.916666667
CHRNA3 AND NOT-TMEM26	0.95	1	0.904761905
CHRNA3 AND NOT-GJD4	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-MS4A15	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-LRTOMT	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-FCAR	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-MS4A2	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-FCER2	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-SLC39A12	0.936708861	1	0.880952381
CHRNA3 AND NOT-OPN5	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-PI16	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-FCN2	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-GPRC6A	0.95	1	0.904761905
CHRNA3 AND NOT-ADGRF2	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-FGFR3	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-FGFR2	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-FGFR4	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-LMTK2	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-ENPP4	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-SORCS3	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-SV2C	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-NFASC	0.936708861	1	0.880952381
CHRNA3 AND NOT-FLT3	0.95	1	0.904761905
CHRNA3 AND NOT-FLT4	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-KIAA1024	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-FMO2	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-FMO4	0.943396226	1	0.892857143
CHRNA3 AND NOT-ACSL6	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-SLC9A8	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-FAM189A1	0.943396226	1	0.892857143
CHRNA3 AND NOT-CRB1	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-SLC7A8	0.936708861	1	0.880952381
CHRNA3 AND NOT-ATP1B4	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-SLC44A1	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-ABCB9	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-NPTXR	0.95	1	0.904761905
CHRNA3 AND NOT-FOLH1	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-SYNGR4	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-HYAL4	0.95	1	0.904761905
CHRNA3 AND NOT-FPR2	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-FLRT2	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-MS4A6E	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-IL27	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-FUT2	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-FUT3	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-FUT7	0.936708861	1	0.880952381
CHRNA3 AND NOT-TECRL	0.956521739	1	0.916666667
CHRNA3 AND NOT-NAALADL2	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-ZDHHC23	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-CALHM1	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-C19orf26	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-CCDC108	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-MCOLN2	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-MYADML2	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-RNF144B	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-GABRA1	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-GABRA2	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-GABRA4	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-GABRB2	0.943396226	1	0.892857143
CHRNA3 AND NOT-GABRD	0.95	1	0.904761905
CHRNA3 AND NOT-GABRG3	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-GABRR2	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-SLC24A2	0.936708861	1	0.880952381
CHRNA3 AND NOT-KLK5	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-TPSG1	0.950617284	0.98717949	0.916666667

Antigen Pair Logic	F1	Precision	Recall
NLGN3 AND NOT-SLCO5A1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-VANGL1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TMEM187	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SLC10A3	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-DYSF	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-FZD4	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-FZD6	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-APH1B	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-GLT8D2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-PLVAP	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TMUB1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-CRISPLD2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TMEM120A	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TSPAN10	0.965517241	1	0.933333333
NLGN3 AND NOT-SPNS1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ITFG3	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TMEM222	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SLC41A2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-MFSD7	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-LRRC8C	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-RHBDD1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TMEM107	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-C2orf40	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-JAGN1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TPST2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-MS4A14	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ABHD1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-PPAPDC3	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-IL17RC	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-AIFM2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ZDHHC12	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SPPL2A	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-FAM73B	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-CIRH1A	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TMEM209	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-FIBCD1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-RELT	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-DISP1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SLC43A1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ITGA8	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-APOL1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-LY6D	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-PPAP2A	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-UNC5C	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-AOC3	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-KCNK5	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-DGAT1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-B3GALT4	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ABCC3	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TNFRSF25	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TNFSF9	0.965517241	1	0.933333333
NLGN3 AND NOT-TNFRSF14	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TNFRSF10B	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-CREG1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-DPM2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-NRP2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-NRP1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-CD84	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ST3GAL5	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SLC5A6	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SGCE	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-CACNA1H	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-P4HA2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ABCC10	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-RHOT2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-PHLDB2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TSPAN18	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-PKD2L1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TMEM41A	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-GPRC5A	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SLC7A7	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-CLDN12	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-CLDN8	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-CLDN1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TMEM55B	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-LMF2	0.933333333	0.933333333	0.933333333

FIG. 1 (CONT.)

Antigen Pair Logic	F1	Precision	Recall
CHRNA3 AND NOT-METTL7A	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-SGMS1	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-WFDC11	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-WFDC9	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-TAS2R39	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-TAS2R40	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-TAS2R41	0.956521739	1	0.916666667
CHRNA3 AND NOT-FAM205A	0.956521739	1	0.916666667
CHRNA3 AND NOT-PNKD	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-LRIG1	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-CYP4X1	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-CFAP61	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-OR1C1	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-OR2F1	0.95	1	0.904761905
CHRNA3 AND NOT-OR1J4	0.95	1	0.904761905
CHRNA3 AND NOT-OR2M4	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-OR2L2	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-OR2K2	0.936708861	1	0.880952381
CHRNA3 AND NOT-SLC13A4	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-IL1RAPL2	0.956521739	1	0.916666667
CHRNA3 AND NOT-OR7A17	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-OR5K1	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-OR8G1	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-OR10A3	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-TSPAN16	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-OR12D2	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-OR10H3	0.956521739	1	0.916666667
CHRNA3 AND NOT-OR10H2	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-OR8B2	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-AMELX	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-GDF9	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-BEST4	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-HS6ST3	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-OR4D1	0.943396226	1	0.892857143
CHRNA3 AND NOT-GDNF	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-OR2W1	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-ADGRF1	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-OR2H1	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-OR1J2	0.936708861	1	0.880952381
CHRNA3 AND NOT-B4GALT1	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-GHRHR	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-GHSR	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-GJA3	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-KCNV1	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-PKD2L2	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-GJA8	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-GJB3	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-CACNG5	0.943396226	1	0.892857143
CHRNA3 AND NOT-GJB5	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-SIGLEC9	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-SIGLEC8	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-OXGR1	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-GPR78	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-C5AR2	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-BMP10	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-PCDH11X	0.943396226	1	0.892857143
CHRNA3 AND NOT-GLP1R	0.95	1	0.904761905
CHRNA3 AND NOT-GLRA1	0.950617284	0.98717949	0.916666667
CHRNA3 AND NOT-GNRHR	0.950617284	0.98717949	0.916666667

(Continued at top)

Antigen Pair Logic	F1	Precision	Recall
NLGN3 AND NOT-AIFM1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SYNGR2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-PKDCC	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-RFT1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SLC16A7	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SLC33A1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-CD4	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-MTDH	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TMEM88	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-DAPL1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-CCPG1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-DHRS3	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-CD8B	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TAAR2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ORAI3	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-GTF3C3	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-CD163	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-MMGT1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TAOK2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ECEL1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SIGLEC6	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-CD34	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-STX8	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-CD36	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SCARB1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ACVRL1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ADAMTS1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SPTLC2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-MPDU1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-C14orf2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ENTPD6	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ENTPD5	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-NFEZL3	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ABCG1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-KIAA0040	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-EDEM1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-CD69	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-CD72	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-MLEC	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-ADGRE5	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-PIEZO1	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-RNF144A	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-HEPH	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TOMM70A	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-FIG4	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-GOLGA5	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-SLC23A2	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-TNFSF15	0.933333333	0.933333333	0.933333333
NLGN3 AND NOT-CLEC2B	0.933333333	0.933333333	0.933333333
SLC1A2 AND NAT10	0.909090909	0.833333333	1
SLC1A2 AND DOLPP1	0.909090909	0.833333333	1
GDAP1L1 AND DOLPP1	0.909090909	0.833333333	1
SLC4A4 AND NOT-SCNN1A	0.928571429	1	0.866666667
GDAP1L1 AND MS4A6A	0.909090909	0.833333333	1
SLC1A2 AND MPPE1	0.909090909	0.833333333	1
SLC1A2 AND SOAT1	0.909090909	0.833333333	1
SLC1A2 AND JAGN1	0.909090909	0.833333333	1
SLC1A2 AND GPAA1	0.909090909	0.833333333	1
GDAP1L1 AND SLC22A4	0.909090909	0.833333333	1
SLC4A4 AND NOT-SPINT1	0.928571429	1	0.866666667
SLC4A4 AND NOT-TMPRSS2	0.928571429	1	0.866666667
SLC4A4 AND NOT-P4HA2	0.928571429	1	0.866666667

FIG. 2

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
100132596?	100132596	NR_003254.1	7098	XGY2 Xg pseudogene, Y-linked 2
100505984?	100505984	NR_103775.1	7103	uncharacterized LOC10050598
100507547?	100507547	NR_037169.1	7099	Homo sapiens uncharacterized LOC100507547 (LOC100507547), transcript variant 1, long non-coding RNA
101805491?	101805491	NR_103805.1	7110	uncharacterized LOC101805491
339166?	339166	NR_040000	7114	uncharacterized LOC339166
63914?	63914	NR_026784	7115	long intergenic non-protein coding RNA 1590
A4GALT	53947	NP_059132	4291	alpha 1,4-galactosyltransferase
A4GNT	51146	NP_057245	4179	alpha-1,4-N-acetylglucosaminyltransferase
AADAC	13	NP_001077	1067	arylacetamide deacetylase (esterase)
AASDH	132949	NP_861522	6638	aminoadipate-semialdehyde dehydrogenase
ABCA1	19	NP_005493	3298	ATP-binding cassette, sub-family A (ABC1), member 1
ABCA12	26154	NP_056472	4109	ATP-binding cassette, sub-family A (ABC1), member 12
ABCA12	26154	NP_775099	6332	ATP-binding cassette, sub-family A (ABC1), member 12
ABCA13	154664	NP_689914	6102	ATP-binding cassette, sub-family A (ABC1), member 13
ABCA2	20	NP_001597	2417	ATP-binding cassette, sub-family A (ABC1), member 2
ABCA2	20	NP_997698	7034	ATP-binding cassette, sub-family A (ABC1), member 2
ABCA3	21	NP_001080	1098	ATP-binding cassette, sub-family A (ABC1), member 3
ABCA4	24	NP_000341	104	ATP-binding cassette, sub-family A (ABC1), member 4
ABCA5	23461	NP_061142	4483	ATP-binding cassette, sub-family A (ABC1), member 5
ABCA5	23461	NP_758424	6299	ATP-binding cassette, sub-family A (ABC1), member 5
ABCA6	23460	NP_525023	5599	ATP-binding cassette, sub-family A (ABC1), member 6
ABCA7	10347	NP_061985	4567	ATP-binding cassette, sub-family A (ABC1), member 7
ABCA8	10351	NP_009099	3640	ATP-binding cassette, sub-family A (ABC1), member 8
ABCA9	10350	NP_525022	5598	ATP-binding cassette, sub-family A (ABC1), member 9
ABCB1	5243	NP_000918	331	ATP-binding cassette, sub-family B (MDR/TAP), member 1
ABCB10	23456	NP_036221	3701	ATP-binding cassette, sub-family B (MDR/TAP), member 10
ABCB11	8647	NP_003733	2884	ATP-binding cassette, sub-family B (MDR/TAP), member 11
ABCB4	5244	NP_000434	129	ATP-binding cassette, sub-family B (MDR/TAP), member 4
ABCB4	5244	NP_061337	4494	ATP-binding cassette, sub-family B (MDR/TAP), member 4
ABCB4	5244	NP_061338	4495	ATP-binding cassette, sub-family B (MDR/TAP), member 4
ABCB5	340273	NP_001157413.1	7412	ATP binding cassette subfamily B member 5
ABCB6	10058	NP_005680	3344	ATP-binding cassette, sub-family B (MDR/TAP), member 6
ABCB7	22	NP_004290	3010	ATP-binding cassette, sub-family B (MDR/TAP), member 7
ABCB8	11194	NP_009119	3645	ATP-binding cassette, sub-family B (MDR/TAP), member 8
ABCB9	23457	NP_062570	4578	ATP-binding cassette, sub-family B (MDR/TAP), member 9
ABCB9	23457	NP_062571	4579	ATP-binding cassette, sub-family B (MDR/TAP), member 9
ABCB9	23457	NP_982269	6936	ATP-binding cassette, sub-family B (MDR/TAP), member 9
ABCC1	4363	NP_004987	3193	ATP-binding cassette, sub-family C (CFTR/MRP), member 1
ABCC1	4363	NP_004987.2	4589	ATP-binding cassette, sub-family C (CFTR/MRP), member 1
ABCC1	4363	NP_004987.2	4594	ATP-binding cassette, sub-family C (CFTR/MRP), member 1
ABCC1	4363	NP_004987.2	4595	ATP-binding cassette, sub-family C (CFTR/MRP), member 1
ABCC1	4363	NP_004987.2	4596	ATP-binding cassette, sub-family C (CFTR/MRP), member 1
ABCC10	89845	NP_258261	5499	ATP-binding cassette, sub-family C (CFTR/MRP), member 10
ABCC11	85320	NP_115972	5374	ATP-binding cassette, sub-family C (CFTR/MRP), member 11
ABCC11	85320	NP_149163	5453	ATP-binding cassette, sub-family C (CFTR/MRP), member 11
ABCC11	85320	NP_660187	5934	ATP-binding cassette, sub-family C (CFTR/MRP), member 11
ABCC12	94160	NP_150229	5473	ATP-binding cassette, sub-family C (CFTR/MRP), member 12
ABCC2	1244	NP_000383	117	ATP-binding cassette, sub-family C (CFTR/MRP), member 2
ABCC3	8714	NP_001137542	1768	ATP-binding cassette, sub-family C (CFTR/MRP), member 3
ABCC3	8714	NP_003777	2895	ATP-binding cassette, sub-family C (CFTR/MRP), member 3
ABCC4	10257	NP_001098985	1223	ATP-binding cassette, sub-family C (CFTR/MRP), member 4
ABCC4	10257	NP_005836	3382	ATP-binding cassette, sub-family C (CFTR/MRP), member 4
ABCC5	10057	NP_001018881	675	ATP-binding cassette, sub-family C (CFTR/MRP), member 5
ABCC5	10057	NP_005679	3343	ATP-binding cassette, sub-family C (CFTR/MRP), member 5
ABCC6	368	NP_001072996	987	ATP-binding cassette, sub-family C, member 6 pseudogene 2
ABCC6	368	NP_001162	2233	ATP-binding cassette, sub-family C, member 6 pseudogene 2
ABCC8	6833	NP_000343	106	ATP-binding cassette, sub-family C (CFTR/MRP), member 8
ABCC9	10060	NP_005682	3345	ATP-binding cassette, sub-family C (CFTR/MRP), member 9
ABCC9	10060	NP_064693	4624	ATP-binding cassette, sub-family C (CFTR/MRP), member 9
ABCC9	10060	NP_005682.2	4625	ATP-binding cassette, sub-family C (CFTR/MRP), member 9
ABCD4	5826	NP_005041	3202	ATP-binding cassette, sub-family D (ALD), member 4
ABCG1	9619	NP_004906	3166	ATP-binding cassette, sub-family G (WHITE), member 1
ABCG1	9619	NP_058198	4272	ATP-binding cassette, sub-family G (WHITE), member 1
ABCG1	9619	NP_997057	7001	ATP-binding cassette, sub-family G (WHITE), member 1
ABCG1	9619	NP_997510	7029	ATP-binding cassette, sub-family G (WHITE), member 1
ABCG1	9619	NP_997511	7030	ATP-binding cassette, sub-family G (WHITE), member 1
ABCG1	9619	NP_997512	7031	ATP-binding cassette, sub-family G (WHITE), member 1
ABCG2	9429	NP_004818	3142	ATP-binding cassette, sub-family G (WHITE), member 2
ABCG4	64137	NP_001135977	1668	ATP-binding cassette, sub-family G (WHITE), member 4
ABCG4	64137	NP_071452	4893	ATP-binding cassette, sub-family G (WHITE), member 4
ABCG5	64240	NP_071881.1	7296	ATP binding cassette subfamily G member 5
ABCG8	64241	NP_071882	4913	ATP-binding cassette, sub-family G (WHITE), member 8
ABHD1	84696	NP_115993	5378	abhydrolase domain containing 1
ABHD12	26090	NP_001035937	892	abhydrolase domain containing 12
ABHD12	26090	NP_056415	4105	abhydrolase domain containing 12
ABHD13	84945	NP_116248	5418	abhydrolase domain containing 13
ABHD14A	25864	NP_056222	4077	abhydrolase domain containing 14A
ABHD15	116236	NP_937790	6744	abhydrolase domain containing 15

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
ABHD16A	7920	NP_066983	4783	HLA-B associated transcript 5
ABHD3	171586	NP_612213	5730	abhydrolase domain containing 3
ABHD6	57406	NP_065727	4696	abhydrolase domain containing 6
ABI3BP	25890	NP_056244	4081	ABI family, member 3 (NESH) binding protein
ABO	28	NP_065202	4666	ABO blood group (transferase A, alpha 1-3-N-acetylgalactosaminyltransferase transferase B, alpha 1-3-galactosyltransferase)
ACACB	32	NP_001084	1099	acetyl-Coenzyme A carboxylase beta
ACBD4	79777	NP_001129177.1	1556	acyl-Coenzyme A binding domain containing 4
ACBD4	79777	NP_001129177	1557	acyl-Coenzyme A binding domain containing 4
ACBD4	79777	NP_001129178	1558	acyl-Coenzyme A binding domain containing 4
ACBD4	79777	NP_001129179	1559	acyl-Coenzyme A binding domain containing 4
ACBD4	79777	NP_078998	5071	acyl-Coenzyme A binding domain containing 4
ACBD5	91452	NP_001035938	893	acyl-Coenzyme A binding domain containing 5
ACBD5	91452	NP_663736	5971	acyl-Coenzyme A binding domain containing 5
ACE	1636	NP_000780	249	angiotensin I converting enzyme (peptidyl-dipeptidase A) 1
ACE	1636	NP_690043	6116	angiotensin I converting enzyme (peptidyl-dipeptidase A) 1
ACE2	59272	NP_068576	4831	angiotensin I converting enzyme (peptidyl-dipeptidase A) 2
ACER1	125981	NP_597999	5696	alkaline ceramidase 1
ACER3	55331	NP_060837	4430	alkaline ceramidase 3
ACHE	43	NP_000656	189	acetylcholinesterase (Yt blood group)
ACHE	43	NP_056646	4121	acetylcholinesterase (Yt blood group)
ACKR1	2532	NP_001116423	1293	Duffy blood group, chemokine receptor
ACKR1	2532	NP_002027	2537	Duffy blood group, chemokine receptor
ACKR2	1238	NP_001287	2338	chemokine binding protein 2
ACKR3	57007	NP_064707	4628	chemokine (C-X-C motif) receptor 7
ACMSD	130013	NP_612199	5725	aminocarboxymuconate semialdehyde decarboxylase
ACP1	52	NP_001035739	876	acid phosphatase 1, soluble
ACP1	52	NP_004291	3011	acid phosphatase 1, soluble
ACP1	52	NP_009030	3629	acid phosphatase 1, soluble
ACP2	53	NP_001289418.1	1468	acid phosphatase 2, lysosomal
ACP2	53	NP_001601	2418	acid phosphatase 2, lysosomal
ACPP	55	NP_001090	1102	acid phosphatase, prostate
ACPP	55	NP_001127666	1471	acid phosphatase, prostate
ACPT	93650	NP_149059	5448	acid phosphatase, testicular
ACSBG2	81616	NP_112186	5194	acyl-CoA synthetase bubblegum family member 2
ACSL1	2180	NP_001986	2521	acyl-CoA synthetase long-chain family member 1
ACSL3	2181	NP_004448	3054	acyl-CoA synthetase long-chain family member 3
ACSL3	2181	NP_976251	6927	acyl-CoA synthetase long-chain family member 3
ACSL5	51703	NP_057318	4188	acyl-CoA synthetase long-chain family member 5
ACSL5	51703	NP_976313	6928	acyl-CoA synthetase long-chain family member 5
ACSL5	51703	NP_976314	6929	acyl-CoA synthetase long-chain family member 5
ACSL6	23305	NP_001009185	560	acyl-CoA synthetase long-chain family member 6
ACSL6	23305	NP_056071	4052	acyl-CoA synthetase long-chain family member 6
ACSM1	116285	NP_443188	5550	acyl-CoA synthetase medium-chain family member 1
ACSS1	84532	NP_115890	5362	acyl-CoA synthetase short-chain family member 1
ACSS2	55902	NP_001070020	947	acyl-CoA synthetase short-chain family member 2
ACSS2	55902	NP_061147	4484	acyl-CoA synthetase short-chain family member 2
ACSS3	79611	NP_078836	5048	acyl-CoA synthetase short-chain family member 3
ACVR1	90	NP_001096	1206	activin A receptor, type I
ACVR1	90	NP_001104537	1246	activin A receptor, type I
ACVR1B	91	NP_004293	3012	activin A receptor, type IB
ACVR1B	91	NP_064732	4629	activin A receptor, type IB
ACVR1B	91	NP_064733	4630	activin A receptor, type IB
ACVR1C	130399	NP_001104501	1243	activin A receptor, type IC
ACVR1C	130399	NP_001104502	1244	activin A receptor, type IC
ACVR1C	130399	NP_001104503	1245	activin A receptor, type IC
ACVR1C	130399	NP_660302	5944	activin A receptor, type IC
ACVR2A	92	NP_001607	2419	activin A receptor, type IIA
ACVR2B	93	NP_001097	1212	activin A receptor, type IIB
ACVRL1	94	NP_000011	2	activin A receptor type II-like 1
ACVRL1	94	NP_001070869	965	activin A receptor type II-like 1
ADAM10	102	NP_001101	1233	ADAM metallopeptidase domain 10
ADAM11	4185	NP_002381	2640	ADAM metallopeptidase domain 11
ADAM12	8038	NP_003465	2839	ADAM metallopeptidase domain 12
ADAM12	8038	NP_067673	4815	ADAM metallopeptidase domain 12
ADAM15	8751	NP_001248393.1	7434	ADAM metallopeptidase domain 15
ADAM17	6868	NP_003174	2796	ADAM metallopeptidase domain 17
ADAM18	8749	NP_055052	3893	ADAM metallopeptidase domain 18
ADAM19	8728	NP_150377.1	4970	ADAM metallopeptidase domain 19 (meltrin beta)
ADAM19	8728	NP_150377	5478	ADAM metallopeptidase domain 19 (meltrin beta)
ADAM2	2515	NP_001455	2381	ADAM metallopeptidase domain 2
ADAM20	8748	NP_003805	2903	ADAM metallopeptidase domain 20
ADAM21	8747	NP_003804	2902	ADAM metallopeptidase domain 21 pseudogene
ADAM22	53616	NP_004185	2983	ADAM metallopeptidase domain 22
ADAM22	53616	NP_057435	4207	ADAM metallopeptidase domain 22
ADAM22	53616	NP_068367	4819	ADAM metallopeptidase domain 22
ADAM22	53616	NP_068368	4820	ADAM metallopeptidase domain 22

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
ADAM22	53616	NP_068369	4821	ADAM metallopeptidase domain 22
ADAM23	8745	NP_003803	2901	ADAM metallopeptidase domain 23
ADAM28	10863	NP_055080	3901	ADAM metallopeptidase domain 28
ADAM28	10863	NP_068547	4827	ADAM metallopeptidase domain 28
ADAM29	11086	NP_001124175	1435	ADAM metallopeptidase domain 29
ADAM29	11086	NP_001124176	1436	ADAM metallopeptidase domain 29
ADAM29	11086	NP_001124177	1437	ADAM metallopeptidase domain 29
ADAM29	11086	NP_055084	3903	ADAM metallopeptidase domain 29
ADAM30	11085	NP_068566	4829	ADAM metallopeptidase domain 30
ADAM32	203102	NP_659441	5915	ADAM metallopeptidase domain 32
ADAM33	80332	NP_079496	5135	ADAM metallopeptidase domain 33
ADAM33	80332	NP_694882	6157	ADAM metallopeptidase domain 33
ADAM7	8756	NP_003808	2905	ADAM metallopeptidase domain 7
ADAM8	101	NP_001100	1232	ADAM metallopeptidase domain 8
ADAM8	101	NP_001157961	2084	ADAM metallopeptidase domain 8
ADAM8	101	NP_001157962	2085	ADAM metallopeptidase domain 8
ADAM9	8754	NP_001005845	472	ADAM metallopeptidase domain 9 (meltrin gamma)
ADAM9	8754	NP_003807	2904	ADAM metallopeptidase domain 9 (meltrin gamma)
ADAMTS1	9510	NP_008919	3607	ADAM metallopeptidase with thrombospondin type 1 motif, 1
ADAMTS13	11093	NP_620594	5829	ADAM metallopeptidase with thrombospondin type 1 motif, 13
ADAMTS13	11093	NP_620595	5830	ADAM metallopeptidase with thrombospondin type 1 motif, 13
ADAMTS13	11093	NP_620596	5831	ADAM metallopeptidase with thrombospondin type 1 motif, 13
ADCY1	107	NP_066939	4777	adenylate cyclase 1 (brain)
ADCY10	55811	NP_001161221	2196	adenylate cyclase 10 (soluble)
ADCY10	55811	NP_060887	4445	adenylate cyclase 10 (soluble)
ADCY2	108	NP_065433	4682	adenylate cyclase 2 (brain)
ADCY3	109	NP_004027	2949	adenylate cyclase 3
ADCY4	196883	NP_640340	5849	adenylate cyclase 4
ADCY5	111	NP_899200	6706	adenylate cyclase 5
ADCY6	112	NP_056085	4056	adenylate cyclase 6
ADCY6	112	NP_056085.1	4753	adenylate cyclase 6
ADCY7	113	NP_001105	1247	adenylate cyclase 7
ADCY8	114	NP_001106	1248	adenylate cyclase 8 (brain)
ADCY9	115	NP_001107	1260	adenylate cyclase 9
ADCYAP1R1	117	NP_001109	1279	adenylate cyclase activating polypeptide 1 (pituitary) receptor type I
ADGRA1	84435	NP_001077378	1081	G protein-coupled receptor 123
ADGRA2	25960	NP_116166	5398	G protein-coupled receptor 124
ADGRA3	166647	NP_660333	5954	G protein-coupled receptor 125
ADGRB1	575	NP_001693	2439	brain-specific angiogenesis inhibitor 1
ADGRB3	577	NP_001695	2440	brain-specific angiogenesis inhibitor 3
ADGRD1	283383	NP_942122	6838	G protein-coupled receptor 133
ADGRE1	2015	NP_001965	2514	egf-like module containing, mucin-like,Hormone receptor-like 1
ADGRE2	30817	NP_038475	3825	egf-like module containing, mucin-like,Hormone receptor-like 2
ADGRE2	30817	NP_001257981.1	6133	egf-like module containing, mucin-like,Hormone receptor-like 2
ADGRE2	30817	NP_001257981.1	6134	egf-like module containing, mucin-like,Hormone receptor-like 2
ADGRE2	30817	NP_001257981.1	6135	egf-like module containing, mucin-like,Hormone receptor-like 2
ADGRE2	30817	NP_001257981.1	6136	egf-like module containing, mucin-like,Hormone receptor-like 2
ADGRE2	30817	NP_001257981.1	6137	egf-like module containing, mucin-like,Hormone receptor-like 2
ADGRE2	30817	NP_001257981.1	6138	egf-like module containing, mucin-like,Hormone receptor-like 2
ADGRE3	84658	NP_115960	5372	egf-like module containing, mucin-like,Hormone receptor-like 3
ADGRE5	976	NP_001020331	698	CD97 molecule
ADGRE5	976	NP_001775	2471	CD97 molecule
ADGRE5	976	NP_510966	5592	CD97 molecule
ADGRF1	266977	NP_079324	5114	G protein-coupled receptor 110
ADGRF1	266977	NP_722582	6242	G protein-coupled receptor 110
ADGRF2	222611	NP_722581	6241	G protein-coupled receptor 111
ADGRF3	165082	NP_001138640	1824	G protein-coupled receptor 113
ADGRF3	165082	NP_001138641	1825	G protein-coupled receptor 113
ADGRF3	165082	NP_722577	6238	G protein-coupled receptor 113
ADGRF4	221393	NP_722580	6240	G protein-coupled receptor 115
ADGRF5	221395	NP_001091988	1131	G protein-coupled receptor 116
ADGRF5	221395	NP_056049	4049	G protein-coupled receptor 116
ADGRG2	10149	NP_001073327	1003	G protein-coupled receptor 64
ADGRG2	10149	NP_001073328	1004	G protein-coupled receptor 64
ADGRG2	10149	NP_001073329	1005	G protein-coupled receptor 64
ADGRG2	10149	NP_005747	3357	G protein-coupled receptor 64
ADGRG3	222487	NP_740746	6263	G protein-coupled receptor 97
ADGRG4	139378	NP_722576	6237	G protein-coupled receptor 112
ADGRG5	221188	NP_722579	6239	G protein-coupled receptor 114
ADGRG6	57211	NP_001027566	745	G protein-coupled receptor 126
ADGRG6	57211	NP_001027567	746	G protein-coupled receptor 126
ADGRG6	57211	NP_065188	4663	G protein-coupled receptor 126
ADGRG6	57211	NP_940971	6818	G protein-coupled receptor 126
ADGRG7	84873	NP_116176	5401	G protein-coupled receptor 128
ADGRL1	22859	NP_001008701	551	latrophilin 1
ADGRL1	22859	NP_055736	4013	latrophilin 1
ADGRL2	23266	NP_036434	3740	latrophilin 2

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
ADGRL3	23284	NP_056051	4050	latrophilin 3
ADGRL4	64123	NP_071442	4890	EGF, latrophilin and seven transmembrane domain containing 1
ADGRV1	84059	NP_115495	5299	G protein-coupled receptor 98
ADIG	149685	NP_001018092	668	adipogenin
ADIPOR1	51094	NP_001277482.1	1346	adiponectin receptor 1
ADIPOR1	51094	NP_057083	4149	adiponectin receptor 1
ADIPOR2	79602	NP_078827	5045	adiponectin receptor 2
ADORA1	134	NP_000665	190	adenosine A1 receptor
ADORA1	134	NP_001041695	931	adenosine A1 receptor
ADORA2B	136	NP_000667	191	hypothetical LOC100131909 adenosine A2b receptor
ADORA3	140	NP_000668	192	adenosine A3 receptor
ADORA3	140	NP_001075445	1060	adenosine A3 receptor
ADORA3	140	NP_065734	4698	adenosine A3 receptor
ADPRM	56985	NP_064618	4621	chromosome 17 open reading frame 48
ADRA1A	148	NP_000671	195	adrenergic, alpha-1A-, receptor
ADRA1A	148	NP_150645	5483	adrenergic, alpha-1A-, receptor
ADRA1A	148	NP_150646	5484	adrenergic, alpha-1A-, receptor
ADRA1A	148	NP_150647	5485	adrenergic, alpha-1A-, receptor
ADRA1B	147	NP_000670	194	adrenergic, alpha-1B-, receptor
ADRA1D	146	NP_000669	193	adrenergic, alpha-1D-, receptor
ADRA2A	150	NP_000672	196	adrenergic, alpha-2A-, receptor
ADRA2B	151	NP_000673	197	adrenergic, alpha-2B-, receptor
ADRA2C	152	NP_000674	198	adrenergic, alpha-2C-, receptor
ADRB1	153	NP_000675	199	adrenergic, beta-1-, receptor
ADRB2	154	NP_000015	5	adrenergic, beta-2-, receptor, surface
ADRB3	155	NP_000016	6	adrenergic, beta-3-, receptor
ADSSL1	122622	NP_689541	6034	adenylosuccinate synthase like 1
ADSSL1	122622	NP_954634	6860	adenylosuccinate synthase like 1
ADTRP	84830	NP_001137420	1738	chromosome 6 open reading frame 105
ADTRP	84830	NP_116133	5396	chromosome 6 open reading frame 105
AES	166	NP_001121	1333	amino-terminal enhancer of split
AES	166	NP_945320	6848	amino-terminal enhancer of split
AES	166	NP_945321	6849	amino-terminal enhancer of split
AFG3L2	10939	NP_006787	3565	AFG3 ATPase family gene 3-like 2 (yeast)
AFP	174	NP_001125.1	7252	alpha fetoprotein
AGER	177	NP_001127	1470	advanced glycosylation end product-specific receptor
AGER	177	NP_751947	6286	advanced glycosylation end product-specific receptor
AGPAT1	10554	NP_006402	3485	1-acylglycerol-3-phosphate O-acyltransferase 1 (lysophosphatidic acid acyltransferase, alpha)
AGPAT1	10554	NP_116130	5395	1-acylglycerol-3-phosphate O-acyltransferase 1 (lysophosphatidic acid acyltransferase, alpha)
AGPAT2	10555	NP_001012745	606	1-acylglycerol-3-phosphate O-acyltransferase 2 (lysophosphatidic acid acyltransferase, beta)
AGPAT2	10555	NP_006403	3486	1-acylglycerol-3-phosphate O-acyltransferase 2 (lysophosphatidic acid acyltransferase, beta)
AGPAT3	56894	NP_001032642	777	1-acylglycerol-3-phosphate O-acyltransferase 3
AGPAT3	56894	NP_064517	4601	1-acylglycerol-3-phosphate O-acyltransferase 3
AGPAT4	56895	NP_064518	4602	1-acylglycerol-3-phosphate O-acyltransferase 4 (lysophosphatidic acid acyltransferase, delta)
AGPAT5	55326	NP_060831	4429	1-acylglycerol-3-phosphate O-acyltransferase 5 (lysophosphatidic acid acyltransferase, epsilon)
AGPAT6	137964	NP_848934	6564	1-acylglycerol-3-phosphate O-acyltransferase 6 (lysophosphatidic acid acyltransferase, zeta)
AGPAT9	84803	NP_116106	5388	1-acylglycerol-3-phosphate O-acyltransferase 9
AGTR1	185	NP_000676	200	angiotensin II receptor, type 1
AGTR1	185	NP_004826	3145	angiotensin II receptor, type 1
AGTR1	185	NP_033611	3693	angiotensin II receptor, type 1
AGTR1	185	NP_114038	5240	angiotensin II receptor, type 1
AGTR1	185	NP_114438	5287	angiotensin II receptor, type 1
AGTR2	186	NP_000677	201	angiotensin II receptor, type 2
AGTRAP	57085	NP_001035284	857	angiotensin II receptor-associated protein
AGTRAP	57085	NP_001035285	858	angiotensin II receptor-associated protein
AGTRAP	57085	NP_001035286	859	angiotensin II receptor-associated protein
AGTRAP	57085	NP_001035287	860	angiotensin II receptor-associated protein
AGTRAP	57085	NP_065083	4633	angiotensin II receptor-associated protein
AIFM1	9131	NP_001124318	1439	apoptosis-inducing factor, mitochondrion-associated, 1
AIFM1	9131	NP_001124319	1440	apoptosis-inducing factor, mitochondrion-associated, 1
AIFM1	9131	NP_004199	2988	apoptosis-inducing factor, mitochondrion-associated, 1
AIFM1	9131	NP_056997	4133	apoptosis-inducing factor, mitochondrion-associated, 1
AIFM1	9131	NP_665811	5980	apoptosis-inducing factor, mitochondrion-associated, 1
AIFM1	9131	NP_665812	5982	apoptosis-inducing factor, mitochondrion-associated, 1
AIFM2	84883	NP_116186	5403	apoptosis-inducing factor, mitochondrion-associated, 2
AIG1	51390	NP_057192	4170	androgen-induced 1
AJAP1	55966	NP_001035943	894	adherens junctions associated protein 1
AJAP1	55966	NP_061324	4491	adherens junctions associated protein 1
AKAP1	8165	NP_003479	2840	A kinase (PRKA) anchor protein 1
AKAP6	9472	NP_004265	3005	A kinase (PRKA) anchor protein 6
ALCAM	214	NP_001618	2420	hypothetical protein LOC100133690 activated leukocyte cell adhesion molecule
ALDH1A1	216	NP_000680.2	7313	aldehyde dehydrogenase 1 family member A1
ALDH3A2	224	NP_000373	113	aldehyde dehydrogenase 3 family, member A2
ALDH3A2	224	NP_001026976	734	aldehyde dehydrogenase 3 family, member A2
ALDH6A1	4329	NP_005580	3320	aldehyde dehydrogenase 6 family, member A1
ALG1	56052	NP_061982	4565	asparagine-linked glycosylation 1, beta-1,4-mannosyltransferaseHomolog (S. cerevisiae)
ALG10	84920	NP_116223	5415	asparagine-linked glycosylation 10, alpha-1,2-glucosyltransferaseHomolog (S. pombe)

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
ALG10B	144245	NP_001013642	616	asparagine-linked glycosylation 10, alpha-1,2-glucosyltransferaseHomolog B (yeast)
ALG12	79087	NP_077010	5012	asparagine-linked glycosylation 12, alpha-1,6-mannosyltransferaseHomolog (S. cerevisiae)
ALG3	10195	NP_001006942	494	asparagine-linked glycosylation 3, alpha-1,3- mannosyltransferaseHomolog (S. cerevisiae)
ALG3	10195	NP_005778	3368	asparagine-linked glycosylation 3, alpha-1,3- mannosyltransferaseHomolog (S. cerevisiae)
ALG5	29880	NP_001135836	1650	asparagine-linked glycosylation 5, dolichyl-phosphate beta-glucosyltransferaseHomolog (S. cerevisiae)
ALG5	29880	NP_037470	3806	asparagine-linked glycosylation 5, dolichyl-phosphate beta-glucosyltransferaseHomolog (S. cerevisiae)
ALG6	29929	NP_037471	3807	asparagine-linked glycosylation 6, alpha-1,3-glucosyltransferaseHomolog (S. cerevisiae)
ALG8	79053	NP_001007028	500	asparagine-linked glycosylation 8, alpha-1,3-glucosyltransferaseHomolog (S. cerevisiae)
ALG8	79053	NP_076984	5003	asparagine-linked glycosylation 8, alpha-1,3-glucosyltransferaseHomolog (S. cerevisiae)
ALG9	79796	NP_001071158	976	asparagine-linked glycosylation 9, alpha-1,2-mannosyltransferaseHomolog (S. cerevisiae)
ALG9	79796	NP_001071159	977	asparagine-linked glycosylation 9, alpha-1,2-mannosyltransferaseHomolog (S. cerevisiae)
ALG9	79796	NP_001071160	978	asparagine-linked glycosylation 9, alpha-1,2-mannosyltransferaseHomolog (S. cerevisiae)
ALG9	79796	NP_079016	5074	asparagine-linked glycosylation 9, alpha-1,2-mannosyltransferaseHomolog (S. cerevisiae)
ALK	238	NP_004295	3013	anaplastic lymphoma receptor tyrosine kinase
ALOX5AP	241	NP_001620	2421	arachidonate 5-lipoxygenase-activating protein
ALPK1	80216	NP_001095876	1205	alpha-kinase 1
ALPK1	80216	NP_079420	5125	alpha-kinase 1
AMELX	265	NP_001133	1622	amelogenin (amelogenesis imperfecta 1, X-linked)
AMELX	265	NP_872621	6673	amelogenin (amelogenesis imperfecta 1, X-linked)
AMELX	265	NP_872622	6674	amelogenin (amelogenesis imperfecta 1, X-linked)
AMFR	267	NP_001135	1623	autocrine motility factor receptor
AMHR2	269	NP_001158162	2094	anti-MullerianHormone receptor, type II
AMHR2	269	NP_001158163	2095	anti-MullerianHormone receptor, type II
AMHR2	269	NP_065434	4683	anti-MullerianHormone receptor, type II
AMICA1	120425	NP_001091996	1136	adhesion molecule, interacts with CXADR antigen 1
AMICA1	120425	NP_694938	6158	adhesion molecule, interacts with CXADR antigen 1
AMIGO2	347902	NP_001137140	1710	adhesion molecule with Ig-like domain 2
AMIGO2	347902	NP_862830	6640	adhesion molecule with Ig-like domain 2
AMIGO3	386724	NP_942015	6836	adhesion molecule with Ig-like domain 3
AMN	81693	NP_112205	5199	amnionlessHomolog (mouse)
ANG	283	NP_001091046	1103	angiogenin, ribonuclease, RNase A family, 5
ANG	283	NP_001136	1669	angiogenin, ribonuclease, RNase A family, 5
ANGPTL1	9068	NP_004664	3100	angiopoietin-like 1
ANGPTL7	10218	NP_066969	4781	angiopoietin-like 7
ANK1	286	NP_000028	7	ankyrin 1, erythrocytic
ANK1	286	NP_001135917	1659	ankyrin 1, erythrocytic
ANK1	286	NP_001135918	1660	ankyrin 1, erythrocytic
ANK1	286	NP_065208	4671	ankyrin 1, erythrocytic
ANK1	286	NP_065209	4672	ankyrin 1, erythrocytic
ANK1	286	NP_065210	4673	ankyrin 1, erythrocytic
ANK1	286	NP_065211	4674	ankyrin 1, erythrocytic
ANK1	286	NP_065213	4675	ankyrin 1, erythrocytic
ANKH	56172	NP_473368	5568	ankylosis, progressiveHomolog (mouse)
ANKLE1	126549	NP_689576	6043	ankyrin repeat and LEM domain containing 1
ANKLE2	23141	NP_055929	4034	ankyrin repeat and LEM domain containing 2
ANKRD29	147463	NP_775776	6348	ankyrin repeat domain 29
ANKRD46	157567	NP_940683	6791	ankyrin repeat domain 46
ANO1	55107	NP_060513	4383	anoctamin 1, calcium activated chloride channel
ANO10	55129	NP_060545	4388	anoctamin 10
ANO2	57101	NP_005682.2	4637	anoctamin 2
ANO3	63982	NP_001300655.1	7457	anoctamin 3
ANO4	121601	NP_849148	6566	anoctamin 4
ANO5	203859	NP_001136121	1683	anoctamin 5
ANO5	203859	NP_998764	7040	anoctamin 5
ANO6	196527	NP_001020527	708	anoctamin 6
ANO6	196527	NP_001136150	1684	anoctamin 6
ANO6	196527	NP_001136151	1685	anoctamin 6
ANO6	196527	NP_001136152	1686	anoctamin 6
ANO7	50636	NP_001001666	372	anoctamin 7
ANO7	50636	NP_001001891	378	anoctamin 7
ANO8	57719	NP_066010.1	7338	anoctamin 8
ANO9	338440	NP_001012302	591	anoctamin 9
ANPEP	290	NP_001141	1936	alanyl (membrane) aminopeptidase
ANTXR1	84168	NP_060623	4399	anthrax toxin receptor 1
ANTXR1	84168	NP_115584	5307	anthrax toxin receptor 1
ANTXR1	84168	NP_444262	5557	anthrax toxin receptor 1
ANTXR2	118429	NP_001139266	1876	anthrax toxin receptor 2
ANTXR2	118429	NP_477520	5582	anthrax toxin receptor 2
ANXA1	301	NP_000691.1	7253	annexin A1
AOC2	314	NP_001149.2	7344	amine oxidase, copper containing 2
AOC3	8639	NP_003725	2877	amine oxidase, copper containing 3 (vascular adhesion protein 1)
APCDD1	147495	NP_694545	6146	adenomatosis polyposis coli down-regulated 1
APELA	100506013	NP_001284479	7104	apelin receptor early endogenous ligand
APH1A	51107	NP_001071096	974	anterior pharynx defective 1Homolog A (C. elegans)
APH1A	51107	NP_057106	4153	anterior pharynx defective 1Homolog A (C. elegans)
APH1B	83464	NP_001139118	1872	anterior pharynx defective 1Homolog B (C. elegans)
APH1B	83464	NP_112591	5218	anterior pharynx defective 1Homolog B (C. elegans)

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
APLF	200558	NP_775816	6356	aprataxin and PNKP like factor
APLN	8862	NP_059109	4283	apelin
APLNR	187	NP_005152	3221	apelin receptor
APLP1	333	NP_001019978	686	amyloid beta (A4) precursor-like protein 1
APLP1	333	NP_005157	3222	amyloid beta (A4) precursor-like protein 1
APLP2	334	NP_001135748	1631	amyloid beta (A4) precursor-like protein 2
APLP2	334	NP_001135749	1632	amyloid beta (A4) precursor-like protein 2
APLP2	334	NP_001135750	1633	amyloid beta (A4) precursor-like protein 2
APLP2	334	NP_001633	2422	amyloid beta (A4) precursor-like protein 2
APMAP	57136	NP_065392	4679	chromosome 20 open reading frame 3
APOB	338	NP_000375	114	apolipoprotein B (including Ag(x) antigen)
APOC1	341	NP_001636	2423	apolipoprotein C-I
APOL1	8542	NP_001130012	1610	apolipoprotein L, 1
APOL1	8542	NP_001130013	1611	apolipoprotein L, 1
APOL1	8542	NP_003652	2865	apolipoprotein L, 1
APOL1	8542	NP_663318	5962	apolipoprotein L, 1
APOL2	23780	NP_112092	5184	apolipoprotein L, 2
APOL2	23780	NP_663612	5963	apolipoprotein L, 2
APOL4	80832	NP_085146	5153	apolipoprotein L, 4
APOL4	80832	NP_663693	5970	apolipoprotein L, 4
APOLD1	81575	NP_001123887	1423	apolipoprotein L domain containing 1
APOLD1	81575	NP_110444	5181	apolipoprotein L domain containing 1
APOO	79135	NP_077027	5014	apolipoprotein O
APP	351	NP_000475.1	7254	amyloid beta precursor protein
AQP1	358	NP_932766	6740	aquaporin 1 (Colton blood group)
AQP10	89872	NP_536354.2	7316	aquaporin 10
AQP11	282679	NP_766627	6326	aquaporin 11
AQP2	359	NP_000477	137	aquaporin 2 (collecting duct)
AQP3	360	NP_004916	3168	aquaporin 3 (Gill blood group)
AQP4	361	NP_001641	2424	aquaporin 4
AQP4	361	NP_004019	2948	aquaporin 4
AQP5	362	NP_001642	2425	aquaporin 5
AQP6	363	NP_001643	2426	aquaporin 6, kidney specific
AQP7	364	NP_001161	2180	aquaporin 7
AQP8	343	NP_001160	2172	aquaporin 8
AQP9	366	NP_066190.2	7378	aquaporin 9
AREL1	9870	NP_001034568	794	KIAA0317
ARHGAP1	392	NP_004299	3014	Rho GTPase activating protein 1
ARHGAP36	158763	NP_659404	5908	hypothetical protein FLJ30058
ARL10	285598	NP_775935	6377	ADP-ribosylation factor-like 10
ARL6IP1	23204	NP_055976	4041	ADP-ribosylation factor-like 6 interacting protein 1
ARL6IP5	10550	NP_006398	3483	ADP-ribosylation-like factor 6 interacting protein 5
ARL6IP6	151188	NP_689735	6074	ADP-ribosylation-like factor 6 interacting protein 6
ARMC10	83787	NP_001154481	1996	armadillo repeat containing 10
ARMC10	83787	NP_001154482	1997	armadillo repeat containing 10
ARMC10	83787	NP_001154483	1998	armadillo repeat containing 10
ARMC10	83787	NP_001154484	1999	armadillo repeat containing 10
ARMC10	83787	NP_001154485	2000	armadillo repeat containing 10
ARMC10	83787	NP_114111	5266	armadillo repeat containing 10
ARMCX1	51309	NP_057692	4261	armadillo repeat containing, X-linked 1
ARMCX2	9823	NP_055597	3988	armadillo repeat containing, X-linked 2
ARMCX2	9823	NP_808818	6507	armadillo repeat containing, X-linked 2
ARMCX3	51566	NP_057691	4260	armadillo repeat containing, X-linked 3
ARMCX3	51566	NP_808816	6505	armadillo repeat containing, X-linked 3
ARMCX3	51566	NP_808817	6506	armadillo repeat containing, X-linked 3
ARMCX4	100131755	XP_001713936	7051	similar toHCG1792883
ARMCX4	100131755	XP_001717051	7056	similar toHCG1792883
ARMCX4	100131755	XP_001717934	7057	similar toHCG1792883
ARMCX4	100131755	XP_002346361	7082	similar toHCG1792883
ARSB	411	NP_000037	8	arylsulfatase B
ARSB	411	NP_942002	6826	arylsulfatase B
ARSD	414	NP_001660	2427	arylsulfatase D
ARSD	414	NP_033667	3695	arylsulfatase D
ARSE	415	NP_000038	9	arylsulfatase E (chondrodysplasia punctata 1)
ARSF	416	NP_004033	2950	arylsulfatase F
ARSK	153642	NP_937793	6747	arylsulfatase family, member K
ART3	419	NP_001123488	1409	ADP-ribosyltransferase 3
ART3	419	NP_001123489	1410	ADP-ribosyltransferase 3
ART3	419	NP_001170	2309	ADP-ribosyltransferase 3
ART4	420	NP_066549	4762	ADP-ribosyltransferase 4 (Dombrock blood group)
ASB11	140456	NP_001012428	595	ankyrin repeat and SOCS box-containing 11
ASB11	140456	NP_543149	5642	ankyrin repeat and SOCS box-containing 11
ASB5	140458	NP_543150	5643	ankyrin repeat and SOCS box-containing 5
ASGR1	432	NP_001662	2428	asialoglycoprotein receptor 1
ASIC5	51802	NP_059115	4286	amiloride-sensitive cation channel 5, intestinal
ASPH	444	NP_001158222	2105	aspartate beta-hydroxylase
ASPH	444	NP_001158223	2106	aspartate beta-hydroxylase

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
ASPH	444	NP_001158224	2107	aspartate beta-hydroxylase
ASPH	444	NP_001158225	2108	aspartate beta-hydroxylase
ASPH	444	NP_001158226	2109	aspartate beta-hydroxylase
ASPH	444	NP_001158227	2110	aspartate beta-hydroxylase
ASPH	444	NP_001158228	2111	aspartate beta-hydroxylase
ASPH	444	NP_004309	3015	aspartate beta-hydroxylase
ASPH	444	NP_064549	4609	aspartate beta-hydroxylase
ASPH	444	NP_115855	5356	aspartate beta-hydroxylase
ASPH	444	NP_115856	5357	aspartate beta-hydroxylase
ASPH	444	NP_115857	5358	aspartate beta-hydroxylase
ASTN1	460	NP_004310	3016	astrotactin 1
ASTN1	460	NP_996991	6999	astrotactin 1
ASTN2	23245	NP_054729	3853	astrotactin 2
ASTN2	23245	NP_937829	6753	astrotactin 2
ASTN2	23245	NP_937830	6754	astrotactin 2
ASTN2	23245	NP_937831	6755	astrotactin 2
ATCAY	85300	NP_149053	5447	ataxia, cerebellar, Cayman type
ATF7	11016	NP_001123532.1	1413	activating transcription factor 7
ATF7	11016	NP_001123532	1414	activating transcription factor 7
ATF7	11016	NP_006847	3583	activating transcription factor 7
ATG9A	79065	NP_001070666	955	ATG9 autophagy related 9Homolog A (S. cerevisiae)
ATG9A	79065	NP_076990	5007	ATG9 autophagy related 9Homolog A (S. cerevisiae)
ATG9B	285973	NP_001303985.1	6379	ATG9 autophagy related 9Homolog B (S. cerevisiae)
ATL1	51062	NP_001121185	1357	atlastin GTPase 1
ATL1	51062	NP_056999	4134	atlastin GTPase 1
ATL1	51062	NP_853629	6612	atlastin GTPase 1
ATL2	64225	NP_001129145	1548	atlastin GTPase 2
ATL2	64225	NP_071769	4909	atlastin GTPase 2
ATL3	25923	NP_056274	4087	atlastin GTPase 3
ATP10A	57194	NP_077816	5034	ATPase, class V, type 10A
ATP10B	23120	NP_079429	5126	ATPase, class V, type 10B
ATP10D	57205	NP_065186	4662	ATPase, class V, type 10D
ATP11A	23250	NP_056020	4045	ATPase, class VI, type 11A
ATP11A	23250	NP_115565	5306	ATPase, class VI, type 11A
ATP11B	23200	NP_055431	3964	ATPase, class VI, type 11B
ATP11C	286410	NP_001010986	584	ATPase, class VI, type 11C
ATP11C	286410	NP_775965	6382	ATPase, class VI, type 11C
ATP12A	479	NP_001667	2429	ATPase,H+/K+ transporting, nongastric, alpha polypeptide
ATP13A1	57130	NP_065143	4652	ATPase type 13A1
ATP13A2	23400	NP_001135445.1	7400	ATPase 13A2
ATP13A3	79572	NP_078800	5040	ATPase type 13A3
ATP13A4	84239	NP_115655	5318	ATPase type 13A4
ATP13A5	344905	NP_940907	6805	ATPase type 13A5
ATP1A1	476	NP_000692	202	ATPase, Na+/K+ transporting, alpha 1 polypeptide
ATP1A1	476	NP_001001586	371	ATPase, Na+/K+ transporting, alpha 1 polypeptide
ATP1A1	476	NP_001153705	1983	ATPase, Na+/K+ transporting, alpha 1 polypeptide
ATP1A1	476	NP_001153706	1984	ATPase, Na+/K+ transporting, alpha 1 polypeptide
ATP1A2	477	NP_000693	203	ATPase, Na+/K+ transporting, alpha 2 (+) polypeptide
ATP1A3	478	NP_689509	6027	ATPase, Na+/K+ transporting, alpha 3 polypeptide
ATP1A4	480	NP_001001734	376	ATPase, Na+/K+ transporting, alpha 4 polypeptide
ATP1A4	480	NP_653300	5893	ATPase, Na+/K+ transporting, alpha 4 polypeptide
ATP1B1	481	NP_001668.1	7255	ATPase Na+/K+ transporting subunit beta 1
ATP1B2	482	NP_001669	2430	ATPase, Na+/K+ transporting, beta 2 polypeptide
ATP1B3	483	NP_001670	2431	ATPase, Na+/K+ transporting, beta 3 polypeptide
ATP1B4	23439	NP_001135919	1661	ATPase, (Na+)/K+ transporting, beta 4 polypeptide
ATP1B4	23439	NP_036201	3697	ATPase, (Na+)/K+ transporting, beta 4 polypeptide
ATP2A1	487	NP_004311	3017	ATPase, Ca++ transporting, cardiac muscle, fast twitch 1
ATP2A1	487	NP_775293	6339	ATPase, Ca++ transporting, cardiac muscle, fast twitch 1
ATP2A2	488	NP_001672.1	1561	ATPase, Ca++ transporting, cardiac muscle, slow twitch 2
ATP2A2	488	NP_001672	2432	ATPase, Ca++ transporting, cardiac muscle, slow twitch 2
ATP2A2	488	NP_733765	6245	ATPase, Ca++ transporting, cardiac muscle, slow twitch 2
ATP2A3	489	NP_005164	3223	ATPase, Ca++ transporting, ubiquitous
ATP2A3	489	NP_777613	6417	ATPase, Ca++ transporting, ubiquitous
ATP2A3	489	NP_777614	6418	ATPase, Ca++ transporting, ubiquitous
ATP2A3	489	NP_777615	6419	ATPase, Ca++ transporting, ubiquitous
ATP2A3	489	NP_777616	6420	ATPase, Ca++ transporting, ubiquitous
ATP2A3	489	NP_777617	6421	ATPase, Ca++ transporting, ubiquitous
ATP2A3	489	NP_777618	6422	ATPase, Ca++ transporting, ubiquitous
ATP2B1	490	NP_001001323	351	ATPase, Ca++ transporting, plasma membrane 1
ATP2B1	490	NP_001673	2433	ATPase, Ca++ transporting, plasma membrane 1
ATP2B2	491	NP_001001331	353	ATPase, Ca++ transporting, plasma membrane 2
ATP2B2	491	NP_001674	2434	ATPase, Ca++ transporting, plasma membrane 2
ATP2B3	492	NP_001001344	354	ATPase, Ca++ transporting, plasma membrane 3
ATP2B3	492	NP_068768	4852	ATPase, Ca++ transporting, plasma membrane 3
ATP2C1	27032	NP_001001485	363	ATPase, Ca++ transporting, type 2C, member 1
ATP2C1	27032	NP_001001486	364	ATPase, Ca++ transporting, type 2C, member 1
ATP2C1	27032	NP_001001487	365	ATPase, Ca++ transporting, type 2C, member 1

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
ATP2C1	27032	NP_055197	3926	ATPase, Ca++ transporting, type 2C, member 1
ATP2C2	9914	NP_055676	4001	ATPase, Ca++ transporting, type 2C, member 2
ATP4A	495	NP_000695	204	ATPase,H+/K+ exchanging, alpha polypeptide
ATP4B	496	NP_000696	205	ATPase,H+/K+ exchanging, beta polypeptide
ATP5F1	515	NP_001679	2435	ATP synthase,H+ transporting, mitochondrial F0 complex, subunit B1
ATP5G3	518	NP_001002258	392	ATP synthase,H+ transporting, mitochondrial F0 complex, subunit C3 (subunit 9)
ATP5G3	518	NP_001680	2436	ATP synthase,H+ transporting, mitochondrial F0 complex, subunit C3 (subunit 9)
ATP5J2	9551	NP_001003713	420	ATP synthase,H+ transporting, mitochondrial F0 complex, subunit F2
ATP5J2	9551	NP_001003714	421	ATP synthase,H+ transporting, mitochondrial F0 complex, subunit F2
ATP5J2	9551	NP_001034267	788	ATP synthase,H+ transporting, mitochondrial F0 complex, subunit F2
ATP5J2	9551	NP_004880	3162	ATP synthase,H+ transporting, mitochondrial F0 complex, subunit F2
ATP6AP1	537	NP_001174	2310	ATPase,H+ transporting, lysosomal accessory protein 1
ATP6AP1L	92270	NP_001017971	655	ATPase,H+ transporting, lysosomal accessory protein 1-like
ATP6AP2	10159	NP_005756	3360	ATPase,H+ transporting, lysosomal accessory protein 2
ATP6V0A1	535	NP_001123492.1	7393	ATPase H+ transporting V0 subunit a1
ATP6V0A2	23545	NP_036595	3782	ATPase,H+ transporting, lysosomal V0 subunit a2
ATP6V0A4	50617	NP_065683	4686	ATPase,H+ transporting, lysosomal V0 subunit a4
ATP6V0A4	50617	NP_570855	5669	ATPase,H+ transporting, lysosomal V0 subunit a4
ATP6V0A4	50617	NP_570856	5670	ATPase,H+ transporting, lysosomal V0 subunit a4
ATP6V0B	533	NP_001034546	793	ATPase,H+ transporting, lysosomal 21kDa, V0 subunit b
ATP6V0B	533	NP_004038	2951	ATPase,H+ transporting, lysosomal 21kDa, V0 subunit b
ATP6V0C	527	NP_001685	2437	ATPase,H+ transporting, lysosomal 16kDa, V0 subunit c
ATP6V0E1	8992	NP_003936	2928	ATPase,H+ transporting, lysosomal 9kDa, V0 subunit e1
ATP6V0E2	155066	NP_001094062	1189	ATPase,H+ transporting V0 subunit e2
ATP6V0E2	155066	NP_660265	5936	ATPase,H+ transporting V0 subunit e2
ATP7A	538	NP_000043	10	ATPase, Cu++ transporting, alpha polypeptide
ATP7B	540	NP_000044	11	ATPase, Cu++ transporting, beta polypeptide
ATP7B	540	NP_001005918	477	ATPase, Cu++ transporting, beta polypeptide
ATP8A1	10396	NP_001098999	1224	ATPase, aminophospholipid transporter (APLT), class I, type 8A, member 1
ATP8A1	10396	NP_006086	3433	ATPase, aminophospholipid transporter (APLT), class I, type 8A, member 1
ATP8A2	51761	NP_057613	4237	ATPase, aminophospholipid transporter-like, class I, type 8A, member 2
ATP8B1	5205	NP_005594	3323	ATPase, class I, type 8B, member 1
ATP8B2	57198	NP_001005855.1	7339	ATPase phospholipid transporting 8B2
ATP8B3	148229	NP_620168	5801	ATPase, class I, type 8B, member 3
ATP8B4	79895	NP_079113	5090	ATPase, class I, type 8B, member 4
ATP9A	10079	NP_006036	3420	ATPase, class II, type 9A
ATP9B	374868	NP_940933	6809	ATPase, class II, type 9B
ATRAID	51374	NP_001164266	2256	chromosome 2 open reading frame 28
ATRAID	51374	NP_057169	4166	chromosome 2 open reading frame 28
ATRAID	51374	NP_542159	5608	chromosome 2 open reading frame 28
ATRN	8455	NP_647537	5855	attractin
ATRN	8455	NP_647538	5856	attractin
ATRNL1	26033	NP_997186	7004	attractin-like 1
AUP1	550	NP_853553	6611	ancient ubiquitous protein 1
AVPR1A	552	NP_000697	206	arginine vasopressin receptor 1A
AVPR1B	553	NP_000698	207	arginine vasopressin receptor 1B
AVPR2	554	NP_000045	12	arginine vasopressin receptor 2
AVPR2	554	NP_001139623	1907	arginine vasopressin receptor 2
AWAT1	158833	NP_001013597	615	acyl-CoA wax alcohol acyltransferase 1
AXL	558	NP_001690	2438	AXL receptor tyrosine kinase
AXL	558	NP_068713	4847	AXL receptor tyrosine kinase
B3GALNT1	8706	NP_001033717	784	beta-1,3-N-acetylgalactosaminyltransferase 1 (globoside blood group)
B3GALNT1	8706	NP_003772	2892	beta-1,3-N-acetylgalactosaminyltransferase 1 (globoside blood group)
B3GALNT1	8706	NP_149357	5455	beta-1,3-N-acetylgalactosaminyltransferase 1 (globoside blood group)
B3GALNT1	8706	NP_149358	5456	beta-1,3-N-acetylgalactosaminyltransferase 1 (globoside blood group)
B3GALNT1	8706	NP_149359	5457	beta-1,3-N-acetylgalactosaminyltransferase 1 (globoside blood group)
B3GALT1	8708	NP_066191	4751	UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase, polypeptide 1
B3GALT2	8707	NP_003774	2894	UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase, polypeptide 2
B3GALT4	8705	NP_003773	2893	UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase, polypeptide 4
B3GALT5	10317	NP_006048	3424	UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase, polypeptide 5
B3GALT5	10317	NP_149360	5458	UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase, polypeptide 5
B3GALT5	10317	NP_149361	5459	UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase, polypeptide 5
B3GALT5	10317	NP_149362	5460	UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase, polypeptide 5
B3GALT5	10317	NP_149363	5461	UDP-Gal:betaGlcNAc beta 1,3-galactosyltransferase, polypeptide 5
B3GALT6	126792	NP_542172	5610	UDP-Gal:betaGal beta 1,3-galactosyltransferase polypeptide 6
B3GAT1	27087	NP_061114	4475	beta-1,3-glucuronyltransferase 1 (glucuronosyltransferase P)
B3GAT1	27087	NP_473366	5567	beta-1,3-glucuronyltransferase 1 (glucuronosyltransferase P)
B3GAT2	135152	NP_542780	5625	beta-1,3-glucuronyltransferase 2 (glucuronosyltransferase S)
B3GAT3	26229	NP_036332	3720	beta-1,3-glucuronyltransferase 3 (glucuronosyltransferase I)
B3GLCT	145173	NP_919299	6718	beta 1,3-galactosyltransferase-like
B3GNT2	10678	NP_001306004.1	3518	UDP-GlcNAc:betaGal beta-1,3-N-acetylglucosaminyltransferase 1
B3GNT2	10678	NP_006867	3588	UDP-GlcNAc:betaGal beta-1,3-N-acetylglucosaminyltransferase 1
B3GNT3	10331	NP_055071	3898	UDP-GlcNAc:betaGal beta-1,3-N-acetylglucosaminyltransferase 3
B3GNT5	84002	NP_114436	5286	UDP-GlcNAc:betaGal beta-1,3-N-acetylglucosaminyltransferase 5
B3GNT6	192134	NP_619651	5773	UDP-GlcNAc:betaGal beta-1,3-N-acetylglucosaminyltransferase 6 (core 3 synthase)
B4GALNT1	2583	NP_001469	2385	beta-1,4-N-acetyl-galactosaminyl transferase 1
B4GALNT2	124872	NP_001152859	1939	beta-1,4-N-acetyl-galactosaminyl transferase 2

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
B4GALNT2	124872	NP_001152860	1940	beta-1,4-N-acetyl-galactosaminyl transferase 2
B4GALNT2	124872	NP_703147	6193	beta-1,4-N-acetyl-galactosaminyl transferase 2
B4GALNT3	283358	NP_775864	6361	beta-1,4-N-acetyl-galactosaminyl transferase 3
B4GALNT4	338707	NP_848632	6551	beta-1,4-N-acetyl-galactosaminyl transferase 4
B4GALT1	2683	NP_001488	2390	UDP-Gal:betaGlcNAc beta 1,4- galactosyltransferase, polypeptide 1
B4GALT3	8703	NP_003770	2891	UDP-Gal:betaGlcNAc beta 1,4- galactosyltransferase, polypeptide 3
B4GALT4	8702	NP_003769	2890	UDP-Gal:betaGlcNAc beta 1,4- galactosyltransferase, polypeptide 4
B4GALT4	8702	NP_997708	7035	UDP-Gal:betaGlcNAc beta 1,4- galactosyltransferase, polypeptide 4
B4GALT5	9334	NP_004767	3122	UDP-Gal:betaGlcNAc beta 1,4- galactosyltransferase, polypeptide 5
B4GALT6	9331	NP_004766	3121	UDP-Gal:betaGlcNAc beta 1,4- galactosyltransferase, polypeptide 6
B4GALT7	11285	NP_009186	3661	xylosylprotein beta 1,4-galactosyltransferase, polypeptide 7 (galactosyltransferase I)
B4GAT1	11041	NP_006867.1	3517	UDP-GlcNAc:betaGal beta-1,3-N-acetylglucosaminyltransferase 1
B4GAT1	11041	NP_006867.1	3587	UDP-GlcNAc:betaGal beta-1,3-N-acetylglucosaminyltransferase 1
BACE1	23621	NP_036236	3704	beta-site APP-cleaving enzyme 1
BACE1	23621	NP_620427	5811	beta-site APP-cleaving enzyme 1
BACE1	23621	NP_620428	5812	beta-site APP-cleaving enzyme 1
BACE1	23621	NP_620429	5813	beta-site APP-cleaving enzyme 1
BACE2	25825	NP_036237	3705	beta-site APP-cleaving enzyme 2
BACE2	25825	NP_620476	5814	beta-site APP-cleaving enzyme 2
BACE2	25825	NP_620477	5815	beta-site APP-cleaving enzyme 2
BAGE	574	NP_001178.1	7269	B melanoma antigen
BAMBI	25805	NP_036474	3751	hypothetical LOC729590 BMP and activin membrane-bound inhibitorHomolog (Xenopus laevis)
BANP	54971	NP_060339	4361	BTG3 associated nuclear protein
BANP	54971	NP_524576	5597	BTG3 associated nuclear protein
BAX	581	NP_004315	3018	BCL2-associated X protein
BAX	581	NP_620116	5786	BCL2-associated X protein
BAX	581	NP_620118	5787	BCL2-associated X protein
BAX	581	NP_620119	5788	BCL2-associated X protein
BAX	581	NP_620120	5789	BCL2-associated X protein
BBS4	585	NP_149017	5439	Bardet-Biedl syndrome 4
BCAM	4059	NP_001013275	611	basal cell adhesion molecule (Lutheran blood group)
BCAM	4059	NP_005572	3317	basal cell adhesion molecule (Lutheran blood group)
BCAN	63827	NP_068767.3	7332	brevican
BCAP29	55973	NP_001008405	534	B-cell receptor-associated protein 29
BCAP29	55973	NP_001008406	535	B-cell receptor-associated protein 29
BCAP29	55973	NP_001008407	536	B-cell receptor-associated protein 29
BCAP29	55973	NP_061332	4492	B-cell receptor-associated protein 29
BCAP31	10134	NP_001132913	1616	B-cell receptor-associated protein 31
BCAP31	10134	NP_001132929	1619	B-cell receptor-associated protein 31
BCAP31	10134	NP_005736	3355	B-cell receptor-associated protein 31
BCHE	590	NP_000046	13	butyrylcholinesterase
BCL2	596	NP_000624	182	B-cell CLL/lymphoma 2
BCL2	596	NP_000648	188	B-cell CLL/lymphoma 2
BCL2L1	598	NP_001182	2311	BCL2-like 1
BCL2L1	598	NP_612815	5763	BCL2-like 1
BCL2L10	10017	NP_065129	4645	BCL2-like 10 (apoptosis facilitator)
BCL2L13	23786	NP_056182	4069	BCL2-like 13 (apoptosis facilitator)
BCL2L2	599	NP_004041	2952	BCL2-like 2
BCS1L	617	NP_001073335	1008	BCS1-like (yeast)
BCS1L	617	NP_004319	3019	BCS1-like (yeast)
BDKRB1	623	NP_000701	208	bradykinin receptor B1
BDKRB2	624	NP_000614	177	bradykinin receptor B2
BDNF	627	NP_001137277	1717	brain-derived neurotrophic factor
BDNF	627	NP_001137278	1718	brain-derived neurotrophic factor
BDNF	627	NP_001137279	1719	brain-derived neurotrophic factor
BDNF	627	NP_001137280	1720	brain-derived neurotrophic factor
BDNF	627	NP_001137281	1721	brain-derived neurotrophic factor
BDNF	627	NP_001137282	1722	brain-derived neurotrophic factor
BDNF	627	NP_001137283	1723	brain-derived neurotrophic factor
BDNF	627	NP_001137284	1724	brain-derived neurotrophic factor
BDNF	627	NP_001137285	1725	brain-derived neurotrophic factor
BDNF	627	NP_001137286	1726	brain-derived neurotrophic factor
BDNF	627	NP_001137277.1	1727	brain-derived neurotrophic factor
BDNF	627	NP_001137288	1728	brain-derived neurotrophic factor
BDNF	627	NP_001700	2442	brain-derived neurotrophic factor
BDNF	627	NP_733927	6251	brain-derived neurotrophic factor
BDNF	627	NP_733928	6252	brain-derived neurotrophic factor
BDNF	627	NP_733929	6253	brain-derived neurotrophic factor
BDNF	627	NP_733930	6254	brain-derived neurotrophic factor
BDNF	627	NP_733931	6255	brain-derived neurotrophic factor
BEST1	7439	NP_001132915	1617	bestrophin 1
BEST1	7439	NP_004174	2982	bestrophin 1
BEST2	54831	NP_060152	4323	bestrophin 2
BEST3	144453	NP_116124	5392	bestrophin 3
BEST3	144453	NP_689652	6059	bestrophin 3
BEST4	266675	NP_695006	6171	bestrophin 4
BET1	10282	NP_001304668.1	2028	hypothetical protein LOC100128542 blocked early in transport 1Homolog (S. cerevisiae)

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
BET1	10282	NP_005859	3391	hypothetical protein LOC100128542 blocked early in transport 1Homolog (S. cerevisiae)
BET1L	51272	NP_001092257	1145	blocked early in transport 1Homolog (S. cerevisiae)-like
BET1L	51272	NP_057610	4236	blocked early in transport 1Homolog (S. cerevisiae)-like
BFAR	51283	NP_057645	4243	bifunctional apoptosis regulator
BIK	638	NP_001188	2313	BCL2-interacting killer (apoptosis-inducing)
BIRC5	332	NP_001012270.1	7342	baculoviral IAP repeat containing 5
BLCAP	10904	NP_001161292	2198	bladder cancer associated protein
BLCAP	10904	NP_001161293	2199	bladder cancer associated protein
BLCAP	10904	NP_001161294	2200	bladder cancer associated protein
BLCAP	10904	NP_001161295	2201	bladder cancer associated protein
BLCAP	10904	NP_006689	3549	bladder cancer associated protein
BMP10	27302	NP_055297	3945	bone morphogenetic protein 10
BMP2	650	NP_001191	2314	bone morphogenetic protein 2
BMPR1A	657	NP_004320	3020	bone morphogenetic protein receptor, type IA
BMPR1B	658	NP_001194	2315	bone morphogenetic protein receptor, type IB
BMPR2	659	NP_001195	2316	bone morphogenetic protein receptor, type II (serine/threonine kinase)
BMX	660	NP_001712	2444	BMX non-receptor tyrosine kinase
BMX	660	NP_975010	6920	BMX non-receptor tyrosine kinase
BNIP1	662	NP_001196	2317	BCL2/adenovirus E1B 19kDa interacting protein 1
BNIP1	662	NP_053581	3842	BCL2/adenovirus E1B 19kDa interacting protein 1
BNIP1	662	NP_053582	3843	BCL2/adenovirus E1B 19kDa interacting protein 1
BNIP1	662	NP_053583	3844	BCL2/adenovirus E1B 19kDa interacting protein 1
BNIP2	663	NP_004321	3021	BCL2/adenovirus E1B 19kDa interacting protein 2
BNIP3	664	NP_004043	2953	BCL2/adenovirus E1B 19kDa interacting protein 3
BNIP3L	665	NP_004322	3022	BCL2/adenovirus E1B 19kDa interacting protein 3-like
BOC	91653	NP_150279	5474	BocHomolog (mouse)
BPI	671	NP_001716	2446	bactericidal/permeability-increasing protein
BPIFB1	92747	NP_149974	5465	chromosome 20 open reading frame 114
BPIFB2	80341	NP_079503	5136	bactericidal/permeability-increasing protein-like 1
BRCA1	672	NP_009225	3672	breast cancer 1, early onset
BRCA1	672	NP_009226	3673	breast cancer 1, early onset
BRCA1	672	NP_009227	3674	breast cancer 1, early onset
BRCA1	672	NP_009228	3675	breast cancer 1, early onset
BRCA1	672	NP_009229	3676	breast cancer 1, early onset
BRCA1	672	NP_009230	3677	breast cancer 1, early onset
BRCA1	672	NP_009231	3678	breast cancer 1, early onset
BRCA1	672	NP_009233	3679	breast cancer 1, early onset
BRCA1	672	NP_009234	3680	breast cancer 1, early onset
BRCA1	672	NP_009235	3681	breast cancer 1, early onset
BRCA1	672	NP_009236	3682	breast cancer 1, early onset
BRI3	25798	NP_001152963	1944	brain protein I3 brain protein I3 pseudogene 1
BRI3	25798	NP_056194	4072	brain protein I3 brain protein I3 pseudogene 1
BRICD5	283870	NP_872369	6661	chromosome 16 open reading frame 79
BRINP2	57795	NP_066988	4785	family with sequence similarity 5, member B
BRS3	680	NP_001718	2447	bombesin-like receptor 3
BSG	682	NP_001719	2448	basigin (Ok blood group)
BSG	682	NP_940991	6821	basigin (Ok blood group)
BSG	682	NP_940993	6822	basigin (Ok blood group)
BSND	7809	NP_476517	5579	Bartter syndrome, infantile, with sensorineural deafness (Barttin)
BST1	683	NP_004325	3023	bone marrow stromal cell antigen 1
BST2	684	NP_004326	3024	NPC-A-7, bone marrow stromal cell antigen 2
BTC	685	NP_001720	2449	betacellulin
BTLA	151888	NP_001078826	1090	B and T lymphocyte associated
BTLA	151888	NP_861445	6632	B and T lymphocyte associated
BTN1A1	696	NP_001723	2450	butyrophilin, subfamily 1, member A1
BTN2A1	11120	NP_008980	3621	butyrophilin, subfamily 2, member A1
BTN2A1	11120	NP_510961	5591	butyrophilin, subfamily 2, member A1
BTN2A2	10385	NP_008926	3609	butyrophilin, subfamily 2, member A2
BTN2A2	10385	NP_853509	6607	butyrophilin, subfamily 2, member A2
BTN3A1	11119	NP_001138480	1804	butyrophilin, subfamily 3, member A1
BTN3A1	11119	NP_001138481	1805	butyrophilin, subfamily 3, member A1
BTN3A1	11119	NP_008979	3620	butyrophilin, subfamily 3, member A1
BTN3A1	11119	NP_919423	6725	butyrophilin, subfamily 3, member A1
BTN3A2	11118	NP_008978	3619	butyrophilin, subfamily 3, member A2
BTN3A3	10384	NP_008925	3608	butyrophilin, subfamily 3, member A3
BTN3A3	10384	NP_932078	6736	butyrophilin, subfamily 3, member A3
BTNL2	56244	NP_062548	4575	butyrophilin-like 2 (MHC class II associated)
BTNL3	10917	NP_932079	6737	butyrophilin-like 3
BTNL8	79908	NP_001035552	874	butyrophilin-like 8
BTNL8	79908	NP_001153179	1952	butyrophilin-like 8
BTNL8	79908	NP_001153180	1953	butyrophilin-like 8
BTNL8	79908	NP_001153181	1954	butyrophilin-like 8
BTNL8	79908	NP_001153182	1955	butyrophilin-like 8
BTNL8	79908	NP_079126	5093	butyrophilin-like 8
BTNL9	153579	NP_689760	6079	butyrophilin-like 9
BVES	11149	NP_009004	3627	blood vessel epicardial substance
BVES	11149	NP_671488	5990	blood vessel epicardial substance

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
C10orf35	219738	NP_660349	5959	chromosome 10 open reading frame 35
C10orf54	64115	NP_071436	4888	chromosome 10 open reading frame 54
C10orf99	387695	NP_997256	7018	chromosome 10 open reading frame 99
C11orf24	53838	NP_071733	4898	chromosome 11 open reading frame 24
C11orf87	399947	NP_997528	7032	chromosome 11 open reading frame 87
C12orf49	79794	NP_079014	5073	chromosome 12 open reading frame 49
C12orf73	728568	NP_001129042	1540	chromosome 12 open reading frame 73
C14orf1	11161	NP_009107	3644	chromosome 14 open reading frame 1
C14orf180	400258	NP_001008404	533	chromosome 14 open reading frame 180
C14orf2	9556	NP_001120865	1324	chromosome 14 open reading frame 2
C14orf2	9556	NP_004885	3163	chromosome 14 open reading frame 2
C14orf37	145407	NP_001001872	377	chromosome 14 open reading frame 37
C15orf27	123591	NP_689548	6035	chromosome 15 open reading frame 27
C15orf48	84419	NP_115789	5352	chromosome 15 open reading frame 48
C15orf48	84419	NP_922946	6733	chromosome 15 open reading frame 48
C16orf54	283897	NP_787096	6466	chromosome 16 open reading frame 54
C16orf58	64755	NP_073581	4941	chromosome 16 open reading frame 58
C16orf91	283951	NP_001258980.1	574	chromosome 16 open reading frame 91
C16orf92	146378	NP_001103129	1234	chromosome 16 open reading frame 92
C16orf92	146378	NP_001103130	1235	chromosome 16 open reading frame 92
C17orf62	79415	NP_001028218	752	chromosome 17 open reading frame 62
C17orf62	79415	NP_001093877	1186	chromosome 17 open reading frame 62
C17orf62	79415	NP_001093878	1187	chromosome 17 open reading frame 62
C17orf74	201243	NP_783861	6456	chromosome 17 open reading frame 74
C17orf78	284099	NP_775896	6365	chromosome 17 open reading frame 78
C19orf12	83636	NP_001026896	729	chromosome 19 open reading frame 12
C19orf12	83636	NP_113636	5226	chromosome 19 open reading frame 12
C19orf18	147685	NP_689687	6066	chromosome 19 open reading frame 18
C19orf24	55009	NP_060384	4369	chromosome 19 open reading frame 24
C19orf26	255057	NP_689982	6113	chromosome 19 open reading frame 26
C1GALT1	56913	NP_064541	4606	core 1 synthase, glycoprotein-N-acetylgalactosamine 3-beta-galactosyltransferase, 1
C1GALT1C1	29071	NP_001011551	588	C1GALT1-specific chaperone 1
C1GALT1C1	29071	NP_689905	6100	C1GALT1-specific chaperone 1
C1orf101	257044	NP_001124429	1450	chromosome 1 open reading frame 101
C1orf101	257044	NP_776168	6388	chromosome 1 open reading frame 101
C1orf115	79762	NP_078985	5068	chromosome 1 open reading frame 115
C1orf159	54991	NP_060361	4364	chromosome 1 open reading frame 159
C1orf162	128346	NP_777556	6405	chromosome 1 open reading frame 162
C1orf186	440712	NP_001007545	523	chromosome 1 open reading frame 186
C1orf210	149466	NP_001158301	2116	chromosome 1 open reading frame 210
C1orf210	149466	NP_872323	6648	chromosome 1 open reading frame 210
C1orf233	643988	NP_001229588	7101	C1orf233 chromosome 1 open reading frame 233, fibronectin type-III domain-containing transmembrane protein C1orf233 precurso
C1orf234	729059	NP_001229450	7116	chromosome 1 open reading frame 234
C1orf27	54953	NP_001157717	2070	chromosome 1 open reading frame 27
C1orf27	54953	NP_001157718	2071	chromosome 1 open reading frame 27
C1orf27	54953	NP_060317	4355	chromosome 1 open reading frame 27
C1orf43	25912	NP_001092086	1140	chromosome 1 open reading frame 43
C1orf43	25912	NP_056264	4085	chromosome 1 open reading frame 43
C1orf43	25912	NP_620077	5785	chromosome 1 open reading frame 43
C1orf95	375057	NP_001003665	410	chromosome 1 open reading frame 95
C1QB	713	NP_000482	138	complement component 1, q subcomponent, B chain
C1QTNF1	114897	NP_112230	5208	C1q and tumor necrosis factor related protein 1
C1QTNF1	114897	NP_940995	6823	C1q and tumor necrosis factor related protein 1
C1QTNF1	114897	NP_940996	6824	C1q and tumor necrosis factor related protein 1
C1QTNF6	114904	NP_114116	5267	C1q and tumor necrosis factor related protein 6
C1QTNF6	114904	NP_872292	6644	C1q and tumor necrosis factor related protein 6
C20orf173	140873	NP_001138822	1854	chromosome 20 open reading frame 173
C2CD2	25966	NP_056315	4092	C2 calcium-dependent domain containing 2
C2CD2	25966	NP_950251	6853	C2 calcium-dependent domain containing 2
C2CD2L	9854	NP_055622	3992	C2CD2-like
C2CD5	9847	NP_055617	3991	KIAA0528
C2orf40	84417	NP_115787	5351	chromosome 2 open reading frame 40
C2orf74	339804	NP_001137431	1741	chromosome 2 open reading frame 74
C2orf74	339804	NP_001137432	1742	chromosome 2 open reading frame 74
C2orf82	389084	NP_996778	6976	chromosome 2 open reading frame 82
C2orf83	56918	NP_001155955	2029	chromosome 2 open reading frame 83
C2orf83	56918	NP_064546	4608	chromosome 2 open reading frame 83
C3AR1	719	NP_004045	2954	complement component 3a receptor 1
C3orf18	51161	NP_001165211	2280	chromosome 3 open reading frame 18
C3orf18	51161	NP_001165212	2281	chromosome 3 open reading frame 18
C3orf18	51161	NP_001165214	2282	chromosome 3 open reading frame 18
C3orf18	51161	NP_057294	4184	chromosome 3 open reading frame 18
C3orf52	79669	NP_001165218	2283	chromosome 3 open reading frame 52
C3orf52	79669	NP_078892	5056	chromosome 3 open reading frame 52
C3orf80	401097	NP_001161686	2210	Similar to LOC166075
C4orf26	152816	NP_848592	6544	chromosome 4 open reading frame 26

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
C4orf3	401152	NP_001001701	374	chromosome 4 open reading frame 3
C4orf3	401152	NP_001163801	2239	chromosome 4 open reading frame 3
C4orf32	132720	NP_689613	6051	chromosome 4 open reading frame 32
C5AR1	728	NP_001727	2451	complement component 5a receptor 1
C5AR2	27202	NP_060955	4461	G protein-coupled receptor 77
C5orf15	56951	NP_064584	4616	chromosome 5 open reading frame 15
C5orf28	64417	NP_071928	4924	chromosome 5 open reading frame 28
C5orf46	389336	NP_996849	6994	chromosome 5 open reading frame 46
C6orf10	10665	NP_006772	3561	chromosome 6 open reading frame 10
C6orf136	221545	NP_001103408	1237	chromosome 6 open reading frame 136
C6orf136	221545	NP_001154848	2006	chromosome 6 open reading frame 136
C6orf136	221545	NP_659466	5920	chromosome 6 open reading frame 136
C6orf25	80739	NP_079536	5143	chromosome 6 open reading frame 25
C6orf25	80739	NP_612116	5713	chromosome 6 open reading frame 25
C6orf25	80739	NP_612117	5714	chromosome 6 open reading frame 25
C6orf25	80739	NP_612118	5715	chromosome 6 open reading frame 25
C6orf25	80739	NP_612119	5716	chromosome 6 open reading frame 25
C6orf25	80739	NP_612121	5717	chromosome 6 open reading frame 25
C6orf89	221477	NP_689947	6107	chromosome 6 open reading frame 89
C8B	732	NP_000057	14	complement component 8, beta polypeptide
C9orf57	138240	NP_001122090	1375	chromosome 9 open reading frame 57
C9orf69	90120	NP_690046	6117	chromosome 9 open reading frame 69
C9orf89	84270	NP_115686	5322	chromosome 9 open reading frame 89
C9orf91	203197	NP_694590	6153	chromosome 9 open reading frame 91
CA12	771	NP_001209	2319	carbonic anhydrase XII
CA12	771	NP_996808	6982	carbonic anhydrase XII
CA14	23632	NP_036245	3708	carbonic anhydrase XIV
CA4	762	NP_000708	210	carbonic anhydrase IV
CA9	768	NP_001207	2318	carbonic anhydrase IX
CABP7	164633	NP_872333	6651	calcium binding protein 7
CACFD1	11094	NP_001129247	1566	chromosome 9 open reading frame 7
CACFD1	11094	NP_060056	4310	chromosome 9 open reading frame 7
CACHD1	57685	NP_065976	4741	cache domain containing 1
CACNA1A	773	NP_001120693	1313	calcium channel, voltage-dependent, P/Q type, alpha 1A subunit
CACNA1A	773	NP_001120694	1314	calcium channel, voltage-dependent, P/Q type, alpha 1A subunit
CACNA1B	774	NP_000709	211	calcium channel, voltage-dependent, N type, alpha 1B subunit
CACNA1C	775	NP_000710	212	hypothetical protein LOC100131098 calcium channel, voltage-dependent, L type, alpha 1C subunit
CACNA1C	775	NP_001123299	1384	hypothetical protein LOC100131098 calcium channel, voltage-dependent, L type, alpha 1C subunit
CACNA1C	775	NP_001123301	1385	hypothetical protein LOC100131098 calcium channel, voltage-dependent, L type, alpha 1C subunit
CACNA1C	775	NP_001123302	1386	hypothetical protein LOC100131098 calcium channel, voltage-dependent, L type, alpha 1C subunit
CACNA1C	775	NP_001123303	1387	hypothetical protein LOC100131098 calcium channel, voltage-dependent, L type, alpha 1C subunit
CACNA1C	775	NP_001123304	1388	hypothetical protein LOC100131098 calcium channel, voltage-dependent, L type, alpha 1C subunit
CACNA1C	775	NP_001123305	1389	hypothetical protein LOC100131098 calcium channel, voltage-dependent, L type, alpha 1C subunit
CACNA1C	775	NP_001123306	1390	hypothetical protein LOC100131098 calcium channel, voltage-dependent, L type, alpha 1C subunit
CACNA1C	775	NP_001123307	1391	hypothetical protein LOC100131098 calcium channel, voltage-dependent, L type, alpha 1C subunit
CACNA1C	775	NP_001123308	1392	hypothetical protein LOC100131098 calcium channel, voltage-dependent, L type, alpha 1C subunit
CACNA1C	775	NP_001123309	1393	hypothetical protein LOC100131098 calcium channel, voltage-dependent, L type, alpha 1C subunit
CACNA1C	775	NP_001123310	1394	hypothetical protein LOC100131098 calcium channel, voltage-dependent, L type, alpha 1C subunit
CACNA1C	775	NP_001123311	1395	hypothetical protein LOC100131098 calcium channel, voltage-dependent, L type, alpha 1C subunit
CACNA1C	775	NP_001123312	1396	hypothetical protein LOC100131098 calcium channel, voltage-dependent, L type, alpha 1C subunit
CACNA1C	775	NP_001123313	1397	hypothetical protein LOC100131098 calcium channel, voltage-dependent, L type, alpha 1C subunit
CACNA1C	775	NP_001123314	1398	hypothetical protein LOC100131098 calcium channel, voltage-dependent, L type, alpha 1C subunit
CACNA1C	775	NP_001123315	1399	hypothetical protein LOC100131098 calcium channel, voltage-dependent, L type, alpha 1C subunit
CACNA1C	775	NP_001123316	1400	hypothetical protein LOC100131098 calcium channel, voltage-dependent, L type, alpha 1C subunit
CACNA1C	775	NP_001123318	1401	hypothetical protein LOC100131098 calcium channel, voltage-dependent, L type, alpha 1C subunit
CACNA1C	775	NP_001161095	2189	hypothetical protein LOC100131098 calcium channel, voltage-dependent, L type, alpha 1C subunit
CACNA1C	775	NP_001161096	2190	hypothetical protein LOC100131098 calcium channel, voltage-dependent, L type, alpha 1C subunit
CACNA1C	775	NP_001161097	2191	hypothetical protein LOC100131098 calcium channel, voltage-dependent, L type, alpha 1C subunit
CACNA1C	775	NP_955630	6886	hypothetical protein LOC100131098 calcium channel, voltage-dependent, L type, alpha 1C subunit
CACNA1D	776	NP_000711	213	calcium channel, voltage-dependent, L type, alpha 1D subunit
CACNA1D	776	NP_001122311	1377	calcium channel, voltage-dependent, L type, alpha 1D subunit
CACNA1D	776	NP_001122312	1378	calcium channel, voltage-dependent, L type, alpha 1D subunit
CACNA1E	777	NP_000712	214	calcium channel, voltage-dependent, R type, alpha 1E subunit
CACNA1F	778	NP_005174	3224	calcium channel, voltage-dependent, L type, alpha 1F subunit
CACNA1G	8913	NP_061496	4496	calcium channel, voltage-dependent, T type, alpha 1G subunit
CACNA1G	8913	NP_938190	6774	calcium channel, voltage-dependent, T type, alpha 1G subunit
CACNA1G	8913	NP_938191	6775	calcium channel, voltage-dependent, T type, alpha 1G subunit
CACNA1G	8913	NP_938192	6776	calcium channel, voltage-dependent, T type, alpha 1G subunit
CACNA1G	8913	NP_938193	6777	calcium channel, voltage-dependent, T type, alpha 1G subunit
CACNA1G	8913	NP_938194	6778	calcium channel, voltage-dependent, T type, alpha 1G subunit
CACNA1G	8913	NP_938196	6779	calcium channel, voltage-dependent, T type, alpha 1G subunit
CACNA1G	8913	NP_938197	6780	calcium channel, voltage-dependent, T type, alpha 1G subunit
CACNA1G	8913	NP_938198	6781	calcium channel, voltage-dependent, T type, alpha 1G subunit
CACNA1G	8913	NP_938199	6782	calcium channel, voltage-dependent, T type, alpha 1G subunit
CACNA1G	8913	NP_938200	6783	calcium channel, voltage-dependent, T type, alpha 1G subunit
CACNA1G	8913	NP_938201	6784	calcium channel, voltage-dependent, T type, alpha 1G subunit
CACNA1G	8913	NP_938202	6785	calcium channel, voltage-dependent, T type, alpha 1G subunit

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
CACNA1G	8913	NP_938406	6788	calcium channel, voltage-dependent, T type, alpha 1G subunit
CACNA1G	8913	NP_938407	6789	calcium channel, voltage-dependent, T type, alpha 1G subunit
CACNA1H	8912	NP_001005407	458	calcium channel, voltage-dependent, T type, alpha 1H subunit
CACNA1H	8912	NP_066921	4770	calcium channel, voltage-dependent, T type, alpha 1H subunit
CACNA1I	8911	NP_001003406	408	calcium channel, voltage-dependent, T type, alpha 1I subunit
CACNA1I	8911	NP_066919	4768	calcium channel, voltage-dependent, T type, alpha 1I subunit
CACNA1S	779	NP_000060	15	calcium channel, voltage-dependent, L type, alpha 1S subunit
CACNG1	786	NP_000718	215	calcium channel, voltage-dependent, gamma subunit 1
CACNG2	10369	NP_006069	3430	calcium channel, voltage-dependent, gamma subunit 2
CACNG3	10368	NP_006530	3510	calcium channel, voltage-dependent, gamma subunit 3
CACNG4	27092	NP_055220	3934	calcium channel, voltage-dependent, gamma subunit 4
CACNG5	27091	NP_665810.1	3933	calcium channel, voltage-dependent, gamma subunit 5
CACNG5	27091	NP_665810	5978	calcium channel, voltage-dependent, gamma subunit 5
CACNG6	59285	NP_114103	5265	calcium channel, voltage-dependent, gamma subunit 6
CACNG6	59285	NP_665813	5983	calcium channel, voltage-dependent, gamma subunit 6
CACNG6	59285	NP_665814	5984	calcium channel, voltage-dependent, gamma subunit 6
CACNG7	59284	NP_114102	5264	calcium channel, voltage-dependent, gamma subunit 7
CACNG8	59283	NP_114101	5263	calcium channel, voltage-dependent, gamma subunit 8
CADM1	23705	NP_001091987	1130	cell adhesion molecule 1
CADM1	23705	NP_055148	3916	cell adhesion molecule 1
CADM2	253559	NP_001161146	2194	cell adhesion molecule 2
CADM2	253559	NP_001161147	2195	cell adhesion molecule 2
CADM2	253559	NP_694854	6154	cell adhesion molecule 2
CADM3	57863	NP_001120645	1307	cell adhesion molecule 3
CADM3	57863	NP_067012	4790	cell adhesion molecule 3
CADM4	199731	NP_660339	5956	cell adhesion molecule 4
CALCR	799	NP_001158209	2103	calcitonin receptor
CALCR	799	NP_001158210	2104	calcitonin receptor
CALCR	799	NP_001733	2452	calcitonin receptor
CALCRL	10203	NP_005786	3370	calcitonin receptor-like
CALHM1	255022	NP_001001412	359	calciumHomeostasis modulator 1
CALHM2	51063	NP_057000.2	7314	calcium homeostasis modulator 2
CALHM3	119395	NP_001123214	1382	calciumHomeostasis modulator 3
CALN1	83698	NP_001017440	645	calneuron 1
CALN1	83698	NP_113656	5230	calneuron 1
CALY	50632	NP_056537	4119	calcyon neuron-specific vesicular protein
CAMKMT	79823	NP_079042	5079	chromosome 2 open reading frame 34
CAMP	820	NP_004336	3027	cathelicidin antimicrobial peptide
CANT1	124583	NP_001153244	1960	calcium activated nucleotidase 1
CANT1	124583	NP_001153245	1961	calcium activated nucleotidase 1
CANT1	124583	NP_620148	5797	calcium activated nucleotidase 1
CANX	821	NP_001019820	683	calnexin
CANX	821	NP_001737	2453	calnexin
CAPZA2	830	NP_006127	3441	capping protein (actin filament) muscle Z-line, alpha 2
CARKD	55739	NP_060680	4403	carbohydrate kinase domain containing
CASC4	113201	NP_612432	5747	cancer susceptibility candidate 4
CASC4	113201	NP_816929	6511	cancer susceptibility candidate 4
CASD1	64921	NP_075051	4958	CAS1 domain containing 1
CASR	846	NP_000379	116	calcium-sensing receptor
CATSPER1	117144	NP_444282	5560	cation channel, sperm associated 1
CATSPER2	117155	NP_473361	5564	cation channel, sperm associated 2
CATSPER2	117155	NP_742093	6273	cation channel, sperm associated 2
CATSPER2	117155	NP_742095	6274	cation channel, sperm associated 2
CATSPER3	347732	NP_821138	6514	cation channel, sperm associated 3
CATSPERB	79820	NP_079040	5078	cation channel, sperm-associated, beta
CATSPERD	257062	NP_689997.3	7356	cation channel sperm associated auxiliary subunit delta
CATSPERG	57828	NP_067008	4789	chromosome 19 open reading frame 15
CAV1	857	NP_001744	2454	caveolin 1, caveolae protein, 22kDa
CAV2	858	NP_001224	2320	caveolin 2
CAV2	858	NP_937855	6759	caveolin 2
CAV3	859	NP_001225	2321	caveolin 3
CAV3	859	NP_203123	5489	caveolin 3
CBX3	11335	NP_009207.2	7304	chromobox 3
CCDC107	203260	NP_777583	6411	coiled-coil domain containing 107
CCDC108	255101	NP_689602	6048	coiled-coil domain containing 108
CCDC108	255101	NP_919278	6716	coiled-coil domain containing 108
CCDC109B	55013	NP_060388	4370	coiled-coil domain containing 109B
CCDC112	153733	NP_001035530	868	coiled-coil domain containing 112
CCDC112	153733	NP_689762	6080	coiled-coil domain containing 112
CCDC126	90693	NP_620126	5792	coiled-coil domain containing 126
CCDC127	133957	NP_660308	5946	coiled-coil domain containing 127
CCDC134	79879	NP_079097	5088	coiled-coil domain containing 134
CCDC136	64753	NP_073579	4940	coiled-coil domain containing 136
CCDC141	285025	NP_775919	6371	coiled-coil domain containing 141
CCDC167	154467	NP_612502	5757	chromosome 6 open reading frame 129
CCDC168	643677	NP_001139669	1913	hypothetical protein LOC643677
CCDC51	79714	NP_078937	5065	coiled-coil domain containing 51

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
CCDC67	159989	NP_857596	6618	coiled-coil domain containing 67
CCDC80	151887	NP_955805	6889	coiled-coil domain containing 80
CCDC80	151887	NP_955806	6890	coiled-coil domain containing 80
CCKAR	886	NP_000721	216	cholecystokinin A receptor
CCKBR	887	NP_795344	6481	cholecystokinin B receptor
CCL11	6356	NP_002977	2744	chemokine (C-C motif) ligand 11
CCL22	6367	NP_002981	2745	chemokine (C-C motif) ligand 22
CCL4	6351	NP_001001435	360	chemokine (C-C motif) ligand 4
CCL4	6351	NP_002975	2743	chemokine (C-C motif) ligand 4
CCL4	6351	NP_996890	6996	chemokine (C-C motif) ligand 4
CCPG1	9236	NP_004739	3115	cell cycle progression 1
CCPG1	9236	NP_065790	4713	cell cycle progression 1
CCR1	1230	NP_001286	2337	chemokine (C-C motif) receptor 1
CCR10	2826	NP_057686	4258	chemokine (C-C motif) receptor 10
CCR2	729230	NP_001116513	1295	chemokine (C-C motif) receptor 2
CCR2	729230	NP_001116868	1297	chemokine (C-C motif) receptor 2
CCR3	1232	NP_001158152	2093	chemokine (C-C motif) receptor 3
CCR3	1232	NP_001828	2484	chemokine (C-C motif) receptor 3
CCR3	1232	NP_847898	6537	chemokine (C-C motif) receptor 3
CCR3	1232	NP_847899	6538	chemokine (C-C motif) receptor 3
CCR4	1233	NP_005499	3301	chemokine (C-C motif) receptor 4
CCR5	1234	NP_000570	167	chemokine (C-C motif) receptor 5
CCR5	1234	NP_001093638	1184	chemokine (C-C motif) receptor 5
CCR6	1235	NP_001034666	801	cydin L2 chemokine (C-C motif) receptor 6
CCR6	1235	NP_004358.2	1777	cydin L2 chemokine (C-C motif) receptor 6
CCR6	1235	NP_004358.2	1778	cydin L2 chemokine (C-C motif) receptor 6
CCR6	1235	NP_004358	3034	cydin L2 chemokine (C-C motif) receptor 6
CCR6	1235	NP_112199	5198	cydin L2 chemokine (C-C motif) receptor 6
CCR6	1235	NP_113597	5221	cydin L2 chemokine (C-C motif) receptor 6
CCR7	1236	NP_001829	2485	chemokine (C-C motif) receptor 7
CCR8	1237	NP_005192	3228	chemokine (C-C motif) receptor 8
CCR9	10803	NP_006632	3535	chemokine (C-C motif) receptor 9
CCR9	10803	NP_112477	5210	chemokine (C-C motif) receptor 9
CCRL2	9034	NP_001124382	1446	chemokine (C-C motif) receptor-like 2
CCRL2	9034	NP_003956	2933	chemokine (C-C motif) receptor-like 2
CD101	9398	NP_004249	3001	immunoglobulin superfamily, member 2
CD151	977	NP_001034579	796	CD151 molecule (Raph blood group)
CD151	977	NP_004348	3029	CD151 molecule (Raph blood group)
CD151	977	NP_620598	5832	CD151 molecule (Raph blood group)
CD151	977	NP_620599	5833	CD151 molecule (Raph blood group)
CD160	11126	NP_008984	3624	CD160 molecule
CD163	9332	NP_004235	2996	CD163 molecule
CD163	9332	NP_981961	6934	CD163 molecule
CD163L1	283316	NP_777601	6416	CD163 molecule-like 1
CD164	8763	NP_001135873	1655	CD164 molecule, sialomucin
CD164	8763	NP_001135874	1656	CD164 molecule, sialomucin
CD164	8763	NP_001135875	1657	CD164 molecule, sialomucin
CD164	8763	NP_001135876	1658	CD164 molecule, sialomucin
CD164	8763	NP_006007	3414	CD164 molecule, sialomucin
CD180	4064	NP_005573	3318	CD180 molecule
CD19	930	NP_001761	2460	CD19 molecule
CD1A	909	NP_001754	2455	CD1a molecule
CD1B	910	NP_001755	2456	CD1b molecule
CD1C	911	NP_001756	2457	CD1c molecule
CD1D	912	NP_001757	2458	CD1d molecule
CD1E	913	NP_001036048	906	CD1e molecule
CD1E	913	NP_001036049	907	CD1e molecule
CD1E	913	NP_001036050	908	CD1e molecule
CD1E	913	NP_001036051	909	CD1e molecule
CD1E	913	NP_001036052	910	CD1e molecule
CD1E	913	NP_112155	5187	CD1e molecule
CD2	914	NP_001758	2459	CD2 molecule
CD200	4345	NP_001004196	428	CD200 molecule
CD200	4345	NP_005935	3404	CD200 molecule
CD200R1	131450	NP_620161	5800	CD200 receptor 1
CD200R1	131450	NP_620385	5806	CD200 receptor 1
CD200R1	131450	NP_620386	5807	CD200 receptor 1
CD200R1	131450	NP_740750	6264	CD200 receptor 1
CD207	50489	NP_056532	4117	CD207 molecule, langerin
CD22	933	NP_001762	2461	CD22 molecule
CD226	10666	NP_006557	3513	CD226 molecule
CD24	100133941	NP_001278666.1	7446	CD24 molecule
CD244	51744	NP_001160135	2175	CD244 molecule, natural killer cell receptor 2B4
CD244	51744	NP_001160136	2176	CD244 molecule, natural killer cell receptor 2B4
CD244	51744	NP_057466	4215	CD244 molecule, natural killer cell receptor 2B4
CD247	919	NP_000725.1	7270	CD247 molecule
CD248	57124	NP_065137	4649	CD248 molecule, endosialin

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
CD27	939	NP_001233	2322	CD27 molecule
CD274	29126	NP_054862	3872	CD274 molecule
CD276	80381	NP_001019907	685	CD276 molecule
CD276	80381	NP_079516	5140	CD276 molecule
CD28	940	NP_006130	3444	CD28 molecule
CD300A	11314	NP_009192	3663	CD300a molecule
CD300C	10871	NP_006669	3543	CD300c molecule
CD300LB	124599	NP_777552	6404	CD300 molecule-like family member b
CD300LF	146722	NP_620587	5827	CD300 molecule-like family member f
CD300LG	146894	NP_001161794	2217	CD300 molecule-like family member g
CD300LG	146894	NP_001161795	2218	CD300 molecule-like family member g
CD300LG	146894	NP_001161796	2219	CD300 molecule-like family member g
CD300LG	146894	NP_660316	5948	CD300 molecule-like family member g
CD320	51293	NP_001159367	2127	CD320 molecule
CD320	51293	NP_057663	4249	CD320 molecule
CD33	945	NP_001076087	1062	CD33 molecule
CD33	945	NP_001763	2462	CD33 molecule
CD34	947	NP_001020280	697	CD34 molecule
CD34	947	NP_001764	2463	CD34 molecule
CD36	948	NP_000063	16	CD36 molecule (thrombospondin receptor)
CD36	948	NP_001001547	367	CD36 molecule (thrombospondin receptor)
CD36	948	NP_001001548	368	CD36 molecule (thrombospondin receptor)
CD36	948	NP_001120915	1327	CD36 molecule (thrombospondin receptor)
CD36	948	NP_001120916	1328	CD36 molecule (thrombospondin receptor)
CD37	951	NP_001035120	823	CD37 molecule
CD37	951	NP_001765	2464	CD37 molecule
CD38	952	NP_001766	2465	CD38 molecule
CD3D	915	NP_000723	217	CD3d molecule, delta (CD3-TCR complex)
CD3D	915	NP_001035741	877	CD3d molecule, delta (CD3-TCR complex)
CD3G	917	NP_000064	17	CD3g molecule, gamma (CD3-TCR complex)
CD4	920	NP_000607	174	CD4 molecule
CD40	958	NP_001241.1	7267	CD40 molecule
CD40LG	959	NP_000065	18	CD40 ligand
CD44	960	NP_000601	172	CD44 molecule (Indian blood group)
CD44	960	NP_001001389	355	CD44 molecule (Indian blood group)
CD44	960	NP_001001390	356	CD44 molecule (Indian blood group)
CD44	960	NP_001001391	357	CD44 molecule (Indian blood group)
CD44	960	NP_001001392	358	CD44 molecule (Indian blood group)
CD46	4179	NP_002380	2639	CD46 molecule, complement regulatory protein
CD46	4179	NP_722548	6234	CD46 molecule, complement regulatory protein
CD46	4179	NP_758860	6309	CD46 molecule, complement regulatory protein
CD46	4179	NP_758861	6310	CD46 molecule, complement regulatory protein
CD46	4179	NP_758862	6311	CD46 molecule, complement regulatory protein
CD46	4179	NP_758863	6312	CD46 molecule, complement regulatory protein
CD46	4179	NP_758864	6313	CD46 molecule, complement regulatory protein
CD46	4179	NP_758865	6314	CD46 molecule, complement regulatory protein
CD46	4179	NP_758866	6315	CD46 molecule, complement regulatory protein
CD46	4179	NP_758867	6316	CD46 molecule, complement regulatory protein
CD46	4179	NP_758868	6317	CD46 molecule, complement regulatory protein
CD46	4179	NP_758869	6318	CD46 molecule, complement regulatory protein
CD46	4179	NP_758870	6319	CD46 molecule, complement regulatory protein
CD46	4179	NP_758871	6320	CD46 molecule, complement regulatory protein
CD47	961	NP_001768.1	693	CD47 molecule
CD47	961	NP_001768	2467	CD47 molecule
CD47	961	NP_942088	6837	CD47 molecule
CD48	962	NP_001769.2	7315	CD48 molecule
CD5	921	NP_055022	3881	CD5 molecule
CD52	1043	NP_001794	2478	CD52 molecule
CD53	963	NP_000551	161	CD53 molecule
CD53	963	NP_001035122	824	CD53 molecule
CD58	965	NP_001138294.1	7405	CD58 molecule
CD63	967	NP_001244318.1	825	CD63 molecule
CD63	967	NP_001771	2468	CD63 molecule
CD68	968	NP_001035148	826	CD68 molecule
CD68	968	NP_001242	2330	CD68 molecule
CD69	969	NP_001772	2469	CD69 molecule
CD7	924	NP_006128	3442	CD7 molecule
CD70	970	NP_001243	2331	CD70 molecule
CD72	971	NP_001773	2470	CD72 molecule
CD74	972	NP_001020329.1	7351	CD74 molecule
CD79A	973	NP_001774.1	7257	CD79a molecule
CD79B	974	NP_000617	178	CD79b molecule, immunoglobulin-associated beta
CD79B	974	NP_001035022	817	CD79b molecule, immunoglobulin-associated beta
CD79B	974	NP_067613	4808	CD79b molecule, immunoglobulin-associated beta
CD80	941	NP_005182	3225	CD80 molecule
CD81	975	NP_004347	3028	CD81 molecule
CD82	3732	NP_001020015	687	CD82 molecule

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
CD82	3732	NP_002222	2587	CD82 molecule
CD83	9308	NP_001035370	865	CD83 molecule
CD83	9308	NP_004224	2995	CD83 molecule
CD84	8832	NP_003865	2924	CD84 molecule
CD86	942	NP_008820	3589	CD86 molecule
CD86	942	NP_787058	6463	CD86 molecule
CD8A	925	NP_001139345.1	7408	CD8a molecule
CD8B	926	NP_004922	3169	CD8b molecule
CD8B	926	NP_742097	6275	CD8b molecule
CD8B	926	NP_742099	6276	CD8b molecule
CD8B	926	NP_742100	6277	CD8b molecule
CD8B	926	NP_757362	6295	CD8b molecule
CD9	928	NP_001317241.1	7464	CD9 molecule
CD93	22918	NP_036204	3698	CD93 molecule
CD96	10225	NP_005807	3375	CD96 molecule
CD96	10225	NP_937839	6757	CD96 molecule
CD99	4267	NP_001116370	1292	CD99 molecule
CD99	4267	NP_002405	2647	CD99 molecule
CD99L2	83692	NP_113650	5228	CD99 molecule-like 2
CD99L2	83692	NP_604394	5710	CD99 molecule-like 2
CD99L2	83692	NP_604395	5711	CD99 molecule-like 2
CDC14B	8555	NP_001070649	952	CDC14 cell division cycle 14Homolog B (S. cerevisiae)
CDC14B	8555	NP_003662	2867	CDC14 cell division cycle 14Homolog B (S. cerevisiae)
CDC14B	8555	NP_201588	5488	CDC14 cell division cycle 14Homolog B (S. cerevisiae)
CDCA7L	55536	NP_001120842	1319	cell division cycle associated 7-like
CDCA7L	55536	NP_001120843	1320	cell division cycle associated 7-like
CDCA7L	55536	NP_061189	4488	cell division cycle associated 7-like
CDCP1	64866	NP_073753	4954	CUB domain containing protein 1
CDCP1	64866	NP_835488	6529	CUB domain containing protein 1
CDH1	999	NP_004351	3030	cadherin 1, type 1, E-cadherin (epithelial)
CDH10	1008	NP_006718	3552	cadherin 10, type 2 (T2-cadherin)
CDH11	1009	NP_001788	2477	cadherin 11, type 2, OB-cadherin (osteoblast)
CDH12	1010	NP_004052	2955	cadherin 12, type 2 (N-cadherin 2)
CDH15	1013	NP_004924	3171	cadherin 15, type 1, M-cadherin (myotubule)
CDH16	1014	NP_004053	2956	cadherin 16, KSP-cadherin
CDH17	1015	NP_001138135	1770	cadherin 17, LI cadherin (liver-intestine)
CDH17	1015	NP_004054	2957	cadherin 17, LI cadherin (liver-intestine)
CDH18	1016	NP_001161139	2192	cadherin 18, type 2
CDH18	1016	NP_004925	3172	cadherin 18, type 2
CDH19	28513	NP_066976	4782	cadherin 19, type 2
CDH2	1000	NP_001783	2472	cadherin 2, type 1, N-cadherin (neuronal)
CDH20	28316	NP_114097	5261	cadherin 20, type 2
CDH22	64405	NP_067071	4803	cadherin-like 22
CDH23	64072	NP_001165401	2291	cadherin-like 23
CDH23	64072	NP_001165402	2292	cadherin-like 23
CDH23	64072	NP_001165403	2293	cadherin-like 23
CDH23	64072	NP_001165404	2294	cadherin-like 23
CDH23	64072	NP_001165405	2295	cadherin-like 23
CDH23	64072	NP_001165406	2296	cadherin-like 23
CDH23	64072	NP_001165407	2297	cadherin-like 23
CDH23	64072	NP_071407	4877	cadherin-like 23
CDH23	64072	NP_443068	5515	cadherin-like 23
CDH24	64403	NP_071923	4922	cadherin-like 24
CDH24	64403	NP_659422	5913	cadherin-like 24
CDH26	60437	NP_068582	4835	cadherin-like 26
CDH26	60437	NP_817089	6512	cadherin-like 26
CDH3	1001	NP_001784	2473	cadherin 3, type 1, P-cadherin (placental)
CDH4	1002	NP_001785	2474	cadherin 4, type 1, R-cadherin (retinal)
CDH5	1003	NP_001786	2475	cadherin 5, type 2 (vascular endothelium)
CDH6	1004	NP_004923	3170	cadherin 6, type 2, K-cadherin (fetal kidney)
CDH7	1005	NP_004352	3031	cadherin 7, type 2
CDH7	1005	NP_387450	5510	cadherin 7, type 2
CDH8	1006	NP_001787	2476	cadherin 8, type 2
CDH9	1007	NP_057363	4198	cadherin 9, type 2 (T1-cadherin)
CDHR1	92211	NP_001165442	2305	protocadherin 21
CDHR1	92211	NP_149091	5449	protocadherin 21
CDHR2	54825	NP_001165447	2306	protocadherin 24
CDHR2	54825	NP_060145	4321	protocadherin 24
CDHR3	222256	NP_001288090.1	7449	cadherin related family member 3
CDHR5	53841	NP_001165439	2304	mucin-like protocadherin
CDHR5	53841	NP_068743	4849	mucin-like protocadherin
CDHR5	53841	NP_112554	5212	mucin-like protocadherin
CDIPT	10423	NP_006310	3465	CDP-diacylglycerol--inositol 3-phosphatidyltransferase (phosphatidylinositol synthase)
CDKAL1	54901	NP_060244	4341	CDK5 regulatory subunit associated protein 1-like 1
CDKN3	1033	NP_001124323	1441	cyclin-dependent kinase inhibitor 3
CDKN3	1033	NP_005183	3226	cyclin-dependent kinase inhibitor 3
CDON	50937	NP_058648	4281	CdonHomolog (mouse)

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
CDRT15L2	256223	XP_001725473	7073	CMT1A duplicated region transcript 15-like 1 CMT1A duplicated region transcript 15-like 2
CDRT15L2	256223	XP_001725838	7075	CMT1A duplicated region transcript 15-like 1 CMT1A duplicated region transcript 15-like 2
CDRT15L2	256223	XP_170840	7086	CMT1A duplicated region transcript 15-like 1 CMT1A duplicated region transcript 15-like 2
CDRT15L2	256223	XP_937402	7091	CMT1A duplicated region transcript 15-like 1 CMT1A duplicated region transcript 15-like 2
CDS1	1040	NP_001254	2332	CDP-diacylglycerol synthase (phosphatidate cytidylyltransferase) 1
CDS2	8760	NP_003809	2906	CDP-diacylglycerol synthase (phosphatidate cytidylyltransferase) 2
CDSN	1041	NP_001255	2333	corneodesmosin
CDYL	9425	NP_001137442	1747	chromodomain protein, Y-like
CDYL	9425	NP_001137443	1748	chromodomain protein, Y-like
CDYL	9425	NP_004815	3140	chromodomain protein, Y-like
CEACAM1	634	NP_001020083	690	carcinoembryonic antigen-related cell adhesion molecule 1 (biliary glycoprotein)
CEACAM1	634	NP_001703	2443	carcinoembryonic antigen-related cell adhesion molecule 1 (biliary glycoprotein)
CEACAM19	56971	NP_001121365	1358	carcinoembryonic antigen-related cell adhesion molecule 19
CEACAM19	56971	NP_064604	4619	carcinoembryonic antigen-related cell adhesion molecule 19
CEACAM21	90273	NP_001091976	1128	carcinoembryonic antigen-related cell adhesion molecule 21
CEACAM21	90273	NP_291021	5506	carcinoembryonic antigen-related cell adhesion molecule 21
CEACAM3	1084	NP_001806	2479	carcinoembryonic antigen-related cell adhesion molecule 3
CEACAM4	1089	NP_001808	2480	carcinoembryonic antigen-related cell adhesion molecule 4
CEACAM5	1048	NP_001278413.1	7444	carcinoembryonic antigen related cell adhesion molecule 5
CEACAM6	4680	NP_002474.4	7445	carcinoembryonic antigen related cell adhesion molecule 6
CEACAM7	1087	NP_008821	3590	carcinoembryonic antigen-related cell adhesion molecule 7
CECR1	51816	NP_001269154.1	4288	cat eye syndrome chromosome region, candidate 1
CECR1	51816	NP_803124	6490	cat eye syndrome chromosome region, candidate 1
CECR6	27439	NP_001156551	2041	cat eye syndrome chromosome region, candidate 6
CECR6	27439	NP_114096	5260	cat eye syndrome chromosome region, candidate 6
CELF4	56853	NP_001020258	694	bruno-like 4, RNA binding protein (Drosophila)
CELF4	56853	NP_001020259	695	bruno-like 4, RNA binding protein (Drosophila)
CELF4	56853	NP_001020260	696	bruno-like 4, RNA binding protein (Drosophila)
CELF4	56853	NP_064565	4612	bruno-like 4, RNA binding protein (Drosophila)
CELSR1	9620	NP_055061	3895	cadherin, EGF LAG seven-pass G-type receptor 1 (flamingoHomolog, Drosophila)
CELSR2	1952	NP_001399	2365	cadherin, EGF LAG seven-pass G-type receptor 2 (flamingoHomolog, Drosophila)
CELSR3	1951	NP_001398.2	870	solute carrier family 26, member 6 cadherin, EGF LAG seven-pass G-type receptor 3 (flamingoHomolog, Drosophila)
CELSR3	1951	NP_001398	2364	solute carrier family 26, member 6 cadherin, EGF LAG seven-pass G-type receptor 3 (flamingoHomolog, Drosophila)
CELSR3	1951	NP_075062	4963	solute carrier family 26, member 6 cadherin, EGF LAG seven-pass G-type receptor 3 (flamingoHomolog, Drosophila)
CELSR3	1951	NP_001398.2	5704	solute carrier family 26, member 6 cadherin, EGF LAG seven-pass G-type receptor 3 (flamingoHomolog, Drosophila)
CELSR3	1951	NP_001398.2	5709	solute carrier family 26, member 6 cadherin, EGF LAG seven-pass G-type receptor 3 (flamingoHomolog, Drosophila)
CEND1	51286	NP_057648	4245	cell cycle exit and neuronal differentiation 1
CEP95	90799	NP_612372	5734	coiled-coil domain containing 45
CEPT1	10390	NP_001007795	526	choline/ethanolamine phosphotransferase 1
CEPT1	10390	NP_006081	3432	choline/ethanolamine phosphotransferase 1
CERS1	10715	NP_001483	2389	growth differentiation factor 1 LAG1Homolog, ceramide synthase 1
CERS1	10715	NP_067090	4805	growth differentiation factor 1 LAG1Homolog, ceramide synthase 1
CERS1	10715	NP_937850	6758	growth differentiation factor 1 LAG1Homolog, ceramide synthase 1
CERS2	29956	NP_071358	4872	LAG1Homolog, ceramide synthase 2
CERS2	29956	NP_859530	6629	LAG1Homolog, ceramide synthase 2
CERS3	204219	NP_849164	6571	LAG1Homolog, ceramide synthase 3
CERS4	79603	NP_078828	5046	LAG1Homolog, ceramide synthase 4
CERS5	91012	NP_671723	5995	LAG1Homolog, ceramide synthase 5
CERS6	253782	NP_982288	6939	LAG1Homolog, ceramide synthase 6
CES5A	221223	NP_001137157	1711	carboxylesterase 7
CES5A	221223	NP_659461	5919	carboxylesterase 7
CFAP47	286464	NP_001291477.1	6383	chromosome X open reading frame 59
CFAP61	26074	NP_001161288	2197	chromosome 20 open reading frame 26
CFAP61	26074	NP_056400	4104	chromosome 20 open reading frame 26
CFTR	1080	NP_000483	139	cystic fibrosis transmembrane conductance regulator (ATP-binding cassette sub-family C, member 7)
CGA	1081	NP_000726	218	glycoproteinHormones, alpha polypeptide
CGRRF1	10668	NP_006559	3514	cell growth regulator with ring finger domain 1
CHI3L2	1117	NP_001020368	699	chitinase 3-like 2
CHI3L2	1117	NP_001020370	700	chitinase 3-like 2
CHI3L2	1117	NP_003991	2944	chitinase 3-like 2
CHIC1	53344	NP_001034929	813	cysteine-richHydrophobic domain 1
CHIC2	26511	NP_036242	3707	cysteine-richHydrophobic domain 2
CHIT1	1118	NP_003456	2835	chitinase 1 (chitotriosidase)
CHL1	10752	NP_006605	3528	cell adhesion molecule withHomology to L1CAM (closeHomolog of L1)
CHODL	140578	NP_079220	5106	chondrolectin
CHPT1	56994	NP_064629	4622	choline phosphotransferase 1
CHRM1	1128	NP_000729	219	cholinergic receptor, muscarinic 1
CHRM2	1129	NP_000730	220	cholinergic receptor, muscarinic 2
CHRM2	1129	NP_001006627	483	cholinergic receptor, muscarinic 2
CHRM2	1129	NP_001006628	484	cholinergic receptor, muscarinic 2
CHRM2	1129	NP_001006629	485	cholinergic receptor, muscarinic 2
CHRM2	1129	NP_001006630	486	cholinergic receptor, muscarinic 2

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
CHRM2	1129	NP_001006631	487	cholinergic receptor, muscarinic 2
CHRM2	1129	NP_001006632	488	cholinergic receptor, muscarinic 2
CHRM2	1129	NP_001006633	489	cholinergic receptor, muscarinic 2
CHRM3	1131	NP_000731	221	cholinergic receptor, muscarinic 3
CHRM4	1132	NP_000732	222	cholinergic receptor, muscarinic 4
CHRM5	1133	NP_036257	3709	cholinergic receptor, muscarinic 5
CHRNA1	1134	NP_000070	19	cholinergic receptor, nicotinic, alpha 1 (muscle)
CHRNA1	1134	NP_001034612	798	cholinergic receptor, nicotinic, alpha 1 (muscle)
CHRNA10	57053	NP_065135	4647	cholinergic receptor, nicotinic, alpha 10
CHRNA2	1135	NP_000733	223	cholinergic receptor, nicotinic, alpha 2 (neuronal)
CHRNA3	1136	NP_000734	224	cholinergic receptor, nicotinic, alpha 3
CHRNA3	1136	NP_001160166	2177	cholinergic receptor, nicotinic, alpha 3
CHRNA4	1137	NP_000735	225	cholinergic receptor, nicotinic, alpha 4
CHRNA5	1138	NP_000736	226	cholinergic receptor, nicotinic, alpha 5
CHRNA6	8973	NP_001186208.1	7425	cholinergic receptor nicotinic alpha 6 subunit
CHRNA9	55584	NP_060051	4308	cholinergic receptor, nicotinic, alpha 9
CHRNB1	1140	NP_000738	227	cholinergic receptor, nicotinic, beta 1 (muscle)
CHRNB2	1141	NP_000739	228	cholinergic receptor, nicotinic, beta 2 (neuronal)
CHRNB3	1142	NP_000740	229	cholinergic receptor, nicotinic, beta 3
CHRNB4	1143	NP_000741	230	cholinergic receptor, nicotinic, beta 4
CHRND	1144	NP_000742	231	cholinergic receptor, nicotinic, delta
CHRNE	1145	NP_000071	20	cholinergic receptor, nicotinic, epsilon
CHRNG	1146	NP_005190	3227	cholinergic receptor, nicotinic, gamma
CHST1	8534	NP_003645	2864	carbohydrate (keratan sulfate Gal-6) sulfotransferase 1
CHST10	9486	NP_004845	3148	carbohydrate sulfotransferase 10
CHST11	50515	NP_060883	4443	carbohydrate (chondroitin 4) sulfotransferase 11
CHST12	55501	NP_061111	4473	carbohydrate (chondroitin 4) sulfotransferase 12
CHST13	166012	NP_690849	6128	carbohydrate (chondroitin 4) sulfotransferase 13
CHST15	51363	NP_056976	4129	carbohydrate (N-acetylgalactosamine 4-sulfate 6-O) sulfotransferase 15
CHST3	9469	NP_004264	3004	carbohydrate (chondroitin 6) sulfotransferase 3
CHST4	10164	NP_001159867	2166	carbohydrate (N-acetylglucosamine 6-O) sulfotransferase 4
CHST4	10164	NP_005760	3363	carbohydrate (N-acetylglucosamine 6-O) sulfotransferase 4
CHST8	64377	NP_001121367	1359	carbohydrate (N-acetylgalactosamine 4-0) sulfotransferase 8
CHST8	64377	NP_001121368	1360	carbohydrate (N-acetylgalactosamine 4-0) sulfotransferase 8
CHST8	64377	NP_071912	4921	carbohydrate (N-acetylgalactosamine 4-0) sulfotransferase 8
CHST9	83539	NP_113610	5222	carbohydrate (N-acetylgalactosamine 4-0) sulfotransferase 9
CHSY1	22856	NP_055733	4012	chondroitin sulfate synthase 1
CHSY3	337876	NP_787052	6461	chondroitin sulfate synthase 3
CIRH1A	84916	NP_116219	5412	cirrhosis, autosomal recessive 1A (cirhin)
CISD2	493856	NP_001008389	530	CDGSH iron sulfur domain 2
CKAP4	10970	NP_006816	3573	cytoskeleton-associated protein 4
CKLF	51192	NP_001035228	841	chemokine-like factor
CKLF	51192	NP_057410	4204	chemokine-like factor
CKLF	51192	NP_058647	4280	chemokine-like factor
CKLF	51192	NP_857591	6614	chemokine-like factor
CKLF	51192	NP_857592	6615	chemokine-like factor
CLCA2	9635	NP_006527	3509	chloride channel accessory 2
CLCA4	22802	NP_036260	3710	chloride channel accessory 4
CLCC1	23155	NP_001041675	930	chloride channel CLIC-like 1
CLCC1	23155	NP_055942	4037	chloride channel CLIC-like 1
CLCN1	1180	NP_000074	21	chloride channel 1, skeletal muscle
CLCN2	1181	NP_001164558	2264	chloride channel 2
CLCN2	1181	NP_001164559	2265	chloride channel 2
CLCN2	1181	NP_001164560	2266	chloride channel 2
CLCN2	1181	NP_004357	3033	chloride channel 2
CLCN3	1182	NP_001820	2481	chloride channel 3
CLCN3	1182	NP_776297	6399	chloride channel 3
CLCN4	1183	NP_001821	2482	chloride channel 4
CLCN5	1184	NP_000075	22	chloride channel 5
CLCN5	1184	NP_001121370	1361	chloride channel 5
CLCN5	1184	NP_001121371	1362	chloride channel 5
CLCN6	1185	NP_001277	2334	chloride channel 6
CLCN6	1185	NP_001243888.1	4824	chloride channel 6
CLCN6	1185	NP_001243888.1	4825	chloride channel 6
CLCN6	1185	NP_001243888.1	4826	chloride channel 6
CLCN7	1186	NP_001107803	1272	chloride channel 7
CLCN7	1186	NP_001278	2335	chloride channel 7
CLCNKB	1188	NP_000076	23	chloride channel Kb
CLCNKB	1188	NP_001159417	2131	chloride channel Kb
CLDN1	9076	NP_066924	4771	claudin 1
CLDN10	9071	NP_001153572	1975	claudin 10
CLDN10	9071	NP_008915	3606	claudin 10
CLDN10	9071	NP_878268	6689	claudin 10
CLDN11	5010	NP_005593	3322	claudin 11
CLDN12	9069	NP_036261	3711	claudin 12
CLDN14	23562	NP_001139549	1902	claudin 14
CLDN14	23562	NP_001139550	1903	claudin 14

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
CLDN14	23562	NP_001139551	1904	claudin 14
CLDN14	23562	NP_036262	3712	claudin 14
CLDN14	23562	NP_652763	5857	claudin 14
CLDN15	24146	NP_055158	3920	claudin 15
CLDN16	10686	NP_006571	3520	claudin 16
CLDN17	26285	NP_036263	3713	claudin 17
CLDN18	51208	NP_001002026	387	claudin 18
CLDN18	51208	NP_057453	4213	claudin 18
CLDN19	149461	NP_001116867	1296	claudin 19
CLDN19	149461	NP_683763	6010	claudin 19
CLDN2	9075	NP_001164563	2267	claudin 2
CLDN2	9075	NP_001164566	2268	claudin 2
CLDN2	9075	NP_065117	4642	claudin 2
CLDN20	49861	NP_001001346.1	7334	claudin 20
CLDN22	53842	NP_001104789.1	7381	claudin 22
CLDN23	137075	NP_919260	6713	claudin 23
CLDN24	100132463	NP_001172078.1	7422	claudin 24
CLDN3	1365	NP_001297	2343	claudin 3
CLDN4	1364	NP_001296	2342	claudin 4
CLDN5	7122	NP_001124333	1443	claudin 5
CLDN5	7122	NP_003268	2815	claudin 5
CLDN6	9074	NP_067018	4792	claudin 6
CLDN7	1366	NP_001298	2344	claudin 7
CLDN8	9073	NP_955360	6874	claudin 8
CLDN9	9080	NP_066192	4752	claudin 9
CLDND1	56650	NP_001035271	854	claudin domain containing 1
CLDND1	56650	NP_001035272	855	claudin domain containing 1
CLDND1	56650	NP_001035273	856	claudin domain containing 1
CLDND1	56650	NP_001035289	861	claudin domain containing 1
CLDND1	56650	NP_001035290	862	claudin domain containing 1
CLDND1	56650	NP_063948	4593	claudin domain containing 1
CLDND2	125875	NP_689566	6040	claudin domain containing 2
CLEC10A	10462	NP_006335	3472	C-type lectin domain family 10, member A
CLEC10A	10462	NP_878910	6690	C-type lectin domain family 10, member A
CLEC12A	160364	NP_001123470	1406	C-type lectin domain family 12, member A C-type lectin domain family 12, member B
CLEC12A	160364	NP_612210	5728	C-type lectin domain family 12, member A C-type lectin domain family 12, member B
CLEC12A	160364	NP_963917	6914	C-type lectin domain family 12, member A C-type lectin domain family 12, member B
CLEC12A	160364	NP_001193939.1	6958	C-type lectin domain family 12, member A C-type lectin domain family 12, member B
CLEC12B	387837	NP_001123470	1407	C-type lectin domain family 12, member A
CLEC12B	387837	NP_001123470.1	5729	C-type lectin domain family 12, member A
CLEC12B	387837	NP_001123470.1	6915	C-type lectin domain family 12, member A
CLEC12B	387837	NP_995324	6959	C-type lectin domain family 12, member A
CLEC14A	161198	NP_778230	6440	C-type lectin domain family 14, member A
CLEC1A	51267	NP_057595	4232	C-type lectin domain family 1, member A
CLEC1B	51266	NP_001092901	1159	C-type lectin domain family 1, member B
CLEC1B	51266	NP_057593	4231	C-type lectin domain family 1, member B
CLEC2B	9976	NP_005118	3217	C-type lectin domain family 2, member B
CLEC2D	29121	NP_001004419	433	C-type lectin domain family 2, member D
CLEC2D	29121	NP_037401	3791	C-type lectin domain family 2, member D
CLEC2L	154790	NP_001073980	1035	C-type lectin domain family 2, member L
CLEC3A	10143	NP_005743	3356	C-type lectin domain family 3, member A
CLEC4A	50856	NP_057268	4182	C-type lectin domain family 4, member A
CLEC4A	50856	NP_919429	6727	C-type lectin domain family 4, member A
CLEC4A	50856	NP_919430	6728	C-type lectin domain family 4, member A
CLEC4A	50856	NP_919432	6729	C-type lectin domain family 4, member A
CLEC4C	170482	NP_569708	5657	C-type lectin domain family 4, member C
CLEC4C	170482	NP_987099	6948	C-type lectin domain family 4, member C
CLEC4D	338339	NP_525126	5600	C-type lectin domain family 4, member D
CLEC4E	26253	NP_055173	3922	C-type lectin domain family 4, member E
CLEC4F	165530	NP_775806	6353	C-type lectin domain family 4, member F
CLEC4G	339390	NP_940894	6801	C-type lectin domain family 4, member G
CLEC4M	10332	NP_001138376	1784	C-type lectin domain family 4, member M
CLEC4M	10332	NP_001138377	1785	C-type lectin domain family 4, member M
CLEC4M	10332	NP_001138378	1786	C-type lectin domain family 4, member M
CLEC4M	10332	NP_001138379	1787	C-type lectin domain family 4, member M
CLEC4M	10332	NP_001138380	1788	C-type lectin domain family 4, member M
CLEC4M	10332	NP_001138381	1789	C-type lectin domain family 4, member M
CLEC4M	10332	NP_001138382	1790	C-type lectin domain family 4, member M
CLEC4M	10332	NP_001138383	1791	C-type lectin domain family 4, member M
CLEC4M	10332	NP_055072	3899	C-type lectin domain family 4, member M
CLEC5A	23601	NP_037384	3790	C-type lectin domain family 5, member A
CLEC7A	64581	NP_072092.2	7301	C-type lectin domain containing 7A
CLECL1	160365	NP_742001	6267	C-type lectin-like 1
CLGN	1047	NP_001124147	1433	calmegin
CLGN	1047	NP_004353	3032	calmegin
CLMN	79789	NP_079010	5072	calmin (calponin-like, transmembrane)
CLMP	79827	NP_079045.1	7299	CXADR like membrane protein

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
CLN3	1201	NP_000077	24	ceroid-lipofuscinosis, neuronal 3
CLN3	1201	NP_001035897	886	ceroid-lipofuscinosis, neuronal 3
CLN5	1203	NP_006484	3497	ceroid-lipofuscinosis, neuronal 5
CLN8	2055	NP_061764	4532	ceroid-lipofuscinosis, neuronal 8 (epilepsy, progressive with mental retardation)
CLPS	1208	NP_001823	2483	colipase, pancreatic
CLPTM1	1209	NP_001285	2336	cleft lip and palate associated transmembrane protein 1
CLPTM1L	81037	NP_110409	5171	CLPTM1-like
CLRN1	7401	NP_443721	5554	clarin 1
CLRN1	7401	NP_777367	6402	clarin 1
CLRN3	119467	NP_689524	6029	clarin 3
CLSTN1	22883	NP_001009566	562	calsyntenin 1
CLSTN1	22883	NP_055759	4018	calsyntenin 1
CLSTN2	64084	NP_071414	4879	calsyntenin 2
CLSTN3	9746	NP_055533	3977	calsyntenin 3
CLYBL	171425	NP_996531	6965	citrate lyase beta like
CMKLR1	1240	NP_001135815	1643	chemokine-like receptor 1
CMKLR1	1240	NP_001135816	1644	chemokine-like receptor 1
CMKLR1	1240	NP_001135817	1645	chemokine-like receptor 1
CMKLR1	1240	NP_004063	2958	chemokine-like receptor 1
CMTM1	113540	NP_443725	5555	CKLF-like MARVEL transmembrane domain containing 1
CMTM1	113540	NP_851785	6582	CKLF-like MARVEL transmembrane domain containing 1
CMTM1	113540	NP_851786	6583	CKLF-like MARVEL transmembrane domain containing 1
CMTM1	113540	NP_851787	6584	CKLF-like MARVEL transmembrane domain containing 1
CMTM1	113540	NP_851788	6585	CKLF-like MARVEL transmembrane domain containing 1
CMTM1	113540	NP_851789	6586	CKLF-like MARVEL transmembrane domain containing 1
CMTM1	113540	NP_851800	6587	CKLF-like MARVEL transmembrane domain containing 1
CMTM1	113540	NP_851813	6588	CKLF-like MARVEL transmembrane domain containing 1
CMTM2	146225	NP_653274	5887	CKLF-like MARVEL transmembrane domain containing 2
CMTM3	123920	NP_653202.1	933	CKLF-like MARVEL transmembrane domain containing 3
CMTM3	123920	NP_653202	5873	CKLF-like MARVEL transmembrane domain containing 3
CMTM3	123920	NP_853531	6609	CKLF-like MARVEL transmembrane domain containing 3
CMTM3	123920	NP_853532	6610	CKLF-like MARVEL transmembrane domain containing 3
CMTM4	146223	NP_848933	6563	CKLF-like MARVEL transmembrane domain containing 4
CMTM4	146223	NP_852662	6606	CKLF-like MARVEL transmembrane domain containing 4
CMTM5	116173	NP_001032365	772	CKLF-like MARVEL transmembrane domain containing 5
CMTM5	116173	NP_612469	5755	CKLF-like MARVEL transmembrane domain containing 5
CMTM6	54918	NP_060271	4346	CKLF-like MARVEL transmembrane domain containing 6
CMTM7	112616	NP_612419	5745	CKLF-like MARVEL transmembrane domain containing 7
CMTM7	112616	NP_852137	6599	CKLF-like MARVEL transmembrane domain containing 7
CMTM8	152189	NP_849199	6576	CKLF-like MARVEL transmembrane domain containing 8
CNEP1R1	255919	NP_694993	6168	transmembrane protein 188
CNGA1	1259	NP_000078	25	cyclic nucleotide gated channel alpha 1
CNGA1	1259	NP_001136036	1674	cyclic nucleotide gated channel alpha 1
CNGA3	1261	NP_001073347	1010	cyclic nucleotide gated channel alpha 3
CNGA3	1261	NP_001289	2339	cyclic nucleotide gated channel alpha 3
CNGA4	1262	NP_001032406	775	cyclic nucleotide gated channel alpha 4
CNGB3	54714	NP_061971.3	7366	cyclic nucleotide gated channel beta 3
CNIH1	10175	NP_005767	3365	cornichonHomolog (Drosophila)
CNIH2	254263	NP_872359	6658	cornichonHomolog 2 (Drosophila)
CNIH3	149111	NP_689708	6071	cornichonHomolog 3 (Drosophila)
CNIH4	29097	NP_054903	3877	cornichonHomolog 4 (Drosophila)
CNNM2	54805	NP_060119	4317	cyclin M2
CNNM2	54805	NP_951058	6854	cyclin M2
CNNM2	54805	NP_951059	6855	cyclin M2
CNNM3	26505	NP_060093	4313	cyclin M3
CNNM3	26505	NP_951060	6856	cyclin M3
CNNM4	26504	NP_064569	4614	cyclin M4
CNOT1	23019	NP_057368	4199	CCR4-NOT transcription complex, subunit 1
CNOT1	23019	NP_996882	6995	CCR4-NOT transcription complex, subunit 1
CNPPD1	27013	NP_056495	4113	chromosome 2 open reading frame 24
CNR1	1268	NP_001153698	1982	cannabinoid receptor 1 (brain)
CNR1	1268	NP_001153730	1985	cannabinoid receptor 1 (brain)
CNR1	1268	NP_001153731	1986	cannabinoid receptor 1 (brain)
CNR1	1268	NP_001153698.1	1987	cannabinoid receptor 1 (brain)
CNR1	1268	NP_057167	4165	cannabinoid receptor 1 (brain)
CNR1	1268	NP_149421	5464	cannabinoid receptor 1 (brain)
CNR2	1269	NP_001832	2486	cannabinoid receptor 2 (macrophage)
CNST	163882	NP_001132931	1620	chromosome 1 open reading frame 71
CNST	163882	NP_689822	6088	chromosome 1 open reading frame 71
CNTN1	1272	NP_001834	2487	contactin 1
CNTN1	1272	NP_778203	6435	contactin 1
CNTNAP1	8506	NP_003623	2857	contactin associated protein 1
CNTNAP2	26047	NP_054860	3871	contactin associated protein-like 2
CNTNAP4	85445	NP_207837	5492	contactin associated protein-like 4
CNTNAP4	85445	NP_620481	5816	contactin associated protein-like 4
CNTNAP5	129684	NP_570129	5663	contactin associated protein-like 5
COA1	55744	NP_060694	4404	chromosome 7 open reading frame 44

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
COA3	28958	NP_001035521	867	coiled-coil domain containing 56
COL11A2	1302	NP_001157243	2057	collagen, type XI, alpha 2
COL11A2	1302	NP_542410	5617	collagen, type XI, alpha 2
COL11A2	1302	NP_542411	5618	collagen, type XI, alpha 2
COL11A2	1302	NP_542412	5619	collagen, type XI, alpha 2
COL14A1	7373	NP_066933	4773	collagen, type XIV, alpha 1
COL17A1	1308	NP_000485	140	collagen, type XVII, alpha 1
COL24A1	255631	NP_690850	6129	collagen, type XXIV, alpha 1
COL25A1	84570	NP_115907	5365	collagen, type XXV, alpha 1
COL25A1	84570	NP_942014	6835	collagen, type XXV, alpha 1
COL26A1	136227	NP_597714	5694	EMI domain containing 2
COL4A2	1284	NP_001837	2488	collagen, type IV, alpha 2
COLEC11	78989	NP_076932	4996	collectin sub-family member 11
COLEC11	78989	NP_954705	6868	collectin sub-family member 11
COLEC12	81035	NP_569057	5650	collectin sub-family member 12
COLQ	8292	NP_005668.2	7306	collagen like tail subunit of asymmetric acetylcholinesterase
COMT	1312	NP_000745	232	catechol-O-methyltransferase
COMT	1312	NP_001128633	1531	catechol-O-methyltransferase
COMT	1312	NP_001128634	1532	catechol-O-methyltransferase
COMT	1312	NP_009294	3683	catechol-O-methyltransferase
COMTD1	118881	NP_653190	5871	catechol-O-methyltransferase domain containing 1
COQ2	27235	NP_056512	4115	coenzyme Q2Homolog, prenyltransferase (yeast)
COQ7	10229	NP_057222	4175	coenzyme Q7Homolog, ubiquinone (yeast)
CORIN	10699	NP_006578	3523	corin, serine peptidase
COX10	1352	NP_001294	2340	COX10Homolog, cytochrome c oxidase assembly protein,Heme A: farnesyltransferase (yeast)
COX11	1353	NP_001156333	2033	COX11Homolog, cytochrome c oxidase assembly protein (yeast)
COX11	1353	NP_001156334	2034	COX11Homolog, cytochrome c oxidase assembly protein (yeast)
COX11	1353	NP_004366	3035	COX11Homolog, cytochrome c oxidase assembly protein (yeast)
COX14	84987	NP_116290	5422	chromosome 12 open reading frame 62
COX15	1355	NP_004367	3036	COX15Homolog, cytochrome c oxidase assembly protein (yeast)
COX15	1355	NP_510870	5589	COX15Homolog, cytochrome c oxidase assembly protein (yeast)
COX18	285521	NP_776188	6390	COX18 cytochrome c oxidase assemblyHomolog (S. cerevisiae)
COX4I1	1327	NP_001852	2491	cytochrome c oxidase subunit IV isoform 1
COX4I2	84701	NP_115998	5379	cytochrome c oxidase subunit IV isoform 2 (lung)
COX7A1	1346	NP_001855	2492	cytochrome c oxidase subunit VIIa polypeptide 1 (muscle)
COX7A2	1347	NP_001856	2493	cytochrome c oxidase subunit VIIa polypeptide 2 (liver)
COX7A2L	9167	NP_004709	3109	cytochrome c oxidase subunit VIIa polypeptide 2 like
COX8A	1351	NP_004065	2959	cytochrome c oxidase subunit 8A (ubiquitous)
COX8C	341947	NP_892016	6697	cytochrome c oxidase subunit 8C
CPD	1362	NP_001295	2341	carboxypeptidase D
CPED1	79974	NP_001099003	1225	chromosome 7 open reading frame 58
CPED1	79974	NP_079189	5102	chromosome 7 open reading frame 58
CPT1A	1374	NP_001027017	738	carnitine palmitoyltransferase 1A (liver)
CPT1A	1374	NP_001867	2494	carnitine palmitoyltransferase 1A (liver)
CPT1C	126129	NP_001129524	1596	carnitine palmitoyltransferase 1C
CPT1C	126129	NP_689572	6042	carnitine palmitoyltransferase 1C
CPXM2	119587	NP_937791	6745	carboxypeptidase X (M14 family), member 2
CR1	1378	NP_000564	165	complement component (3b/4b) receptor 1 (Knops blood group)
CR1	1378	NP_000642	186	complement component (3b/4b) receptor 1 (Knops blood group)
CR2	1380	NP_001006659	493	complement component (3d/Epstein Barr virus) receptor 2
CR2	1380	NP_001868	2495	complement component (3d/Epstein Barr virus) receptor 2
CRB1	23418	NP_957705	6891	crumbsHomolog 1 (Drosophila)
CRB2	286204	NP_775960	6381	crumbsHomolog 2 (Drosophila)
CRB3	92359	NP_631900	5838	crumbsHomolog 3 (Drosophila)
CRB3	92359	NP_777377	6403	crumbsHomolog 3 (Drosophila)
CREG1	8804	NP_003842	2914	cellular repressor of E1A-stimulated genes 1
CRELD1	78987	NP_001026887	727	cysteine-rich with EGF-like domains 1
CRELD1	78987	NP_001070883	966	cysteine-rich with EGF-like domains 1
CRELD1	78987	NP_056328	4094	cysteine-rich with EGF-like domains 1
CRHR1	1394	NP_001138618	1818	corticotropin releasingHormone receptor 1
CRHR1	1394	NP_001138619	1819	corticotropin releasingHormone receptor 1
CRHR1	1394	NP_001138620	1820	corticotropin releasingHormone receptor 1
CRHR1	1394	NP_004373	3037	corticotropin releasingHormone receptor 1
CRHR2	1395	NP_001874	2496	corticotropin releasingHormone receptor 2
CRIM1	51232	NP_057525	4218	cysteine rich transmembrane BMP regulator 1 (chordin-like)
CRISP3	10321	NP_006052	3425	cysteine-rich secretory protein 3
CRISPLD2	83716	NP_113664	5231	cysteine-rich secretory protein LCCL domain containing 2
CRLS1	54675	NP_001120930	1330	cardiolipin synthase 1
CRLS1	54675	NP_061968	4564	cardiolipin synthase 1
CRTAM	56253	NP_001291711.1	7453	cytotoxic and regulatory T-cell molecule
CRTAP	10491	NP_006362	3477	cartilage associated protein
CRY2	1408	NP_001120929	1329	cryptochrome 2 (photolyase-like)
CRY2	1408	NP_066940	4778	cryptochrome 2 (photolyase-like)
CRYGC	1420	NP_066269	4754	crystallin, gamma C
CSF1	1435	NP_000748	233	colony stimulating factor 1 (macrophage)
CSF1	1435	NP_757349	6292	colony stimulating factor 1 (macrophage)
CSF1	1435	NP_757350	6293	colony stimulating factor 1 (macrophage)

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
CSF1	1435	NP_757351	6294	colony stimulating factor 1 (macrophage)
CSF1R	1436	NP_005202	3229	colony stimulating factor 1 receptor
CSF2RA	1438	NP_001155001	2017	colony stimulating factor 2 receptor, alpha, low-affinity (granulocyte-macrophage)
CSF2RA	1438	NP_001155002	2018	colony stimulating factor 2 receptor, alpha, low-affinity (granulocyte-macrophage)
CSF2RA	1438	NP_001155003	2019	colony stimulating factor 2 receptor, alpha, low-affinity (granulocyte-macrophage)
CSF2RA	1438	NP_001155004	2020	colony stimulating factor 2 receptor, alpha, low-affinity (granulocyte-macrophage)
CSF2RA	1438	NP_006131	3445	colony stimulating factor 2 receptor, alpha, low-affinity (granulocyte-macrophage)
CSF2RA	1438	NP_758448	6302	colony stimulating factor 2 receptor, alpha, low-affinity (granulocyte-macrophage)
CSF2RA	1438	NP_758449	6303	colony stimulating factor 2 receptor, alpha, low-affinity (granulocyte-macrophage)
CSF2RA	1438	NP_758450	6304	colony stimulating factor 2 receptor, alpha, low-affinity (granulocyte-macrophage)
CSF2RA	1438	NP_758452	6305	colony stimulating factor 2 receptor, alpha, low-affinity (granulocyte-macrophage)
CSF2RB	1439	NP_000386.1	7275	colony stimulating factor 2 receptor beta common subunit
CSF3	1440	NP_000750	234	colony stimulating factor 3 (granulocyte)
CSF3	1440	NP_757373	6296	colony stimulating factor 3 (granulocyte)
CSF3	1440	NP_757374	6297	colony stimulating factor 3 (granulocyte)
CSF3R	1441	NP_000751	235	colony stimulating factor 3 receptor (granulocyte)
CSF3R	1441	NP_724780	6243	colony stimulating factor 3 receptor (granulocyte)
CSF3R	1441	NP_724781	6244	colony stimulating factor 3 receptor (granulocyte)
CSF3R	1441	NP_758519	6306	colony stimulating factor 3 receptor (granulocyte)
CSGALNACT1	55790	NP_001123990	1431	chondroitin sulfate N-acetylgalactosaminyltransferase 1
CSGALNACT1	55790	NP_060841	4432	chondroitin sulfate N-acetylgalactosaminyltransferase 1
CSGALNACT2	55454	NP_061060	4469	chondroitin sulfate N-acetylgalactosaminyltransferase 2 novel protein similar to chondroitin sulfate GalNAcT-2 (GALNACT-2)
CSH1	1442	NP_001308	2345	chorionic somatomammotropinHormone 1 (placental lactogen)
CSH1	1442	NP_066271	4755	chorionic somatomammotropinHormone 1 (placental lactogen)
CSH1	1442	NP_001308.1	4934	chorionic somatomammotropinHormone 1 (placental lactogen)
CSH1	1442	NP_001308.1	4935	chorionic somatomammotropinHormone 1 (placental lactogen)
CSMD1	64478	NP_150094	5472	CUB and Sushi multiple domains 1
CSMD2	114784	NP_443128	5530	CUB and Sushi multiple domains 2
CSMD3	114788	NP_443132	5532	CUB and Sushi multiple domains 3
CSMD3	114788	NP_937756	6741	CUB and Sushi multiple domains 3
CSMD3	114788	NP_937757	6742	CUB and Sushi multiple domains 3
CSPG4	1464	NP_001888	2497	chondroitin sulfate proteoglycan 4
CSPG5	10675	NP_006565	3516	chondroitin sulfate proteoglycan 5 (neuroglycan C)
CST11	140880	NP_543020	5636	cystatin 11
CST11	140880	NP_570612	5665	cystatin 11
CST3	1471	NP_000090	26	cystatin C
CST9	128822	NP_001008693	549	cystatin 9 (testatin)
CST9L	128821	NP_542177	5611	cystatin 9-like
CT83	203413	NP_001017978	658	chromosome X open reading frame 61
CTAG1A	246100	NP_640343.1	7312	cancer/testis antigen 1
CTAG1B	1485	NP_001318.1	7258	cancer/testis antigen 1B
CTAG2	30848	NP_066274.2	7442	cancer/testis antigen 2
CTAGE1	64693	NP_758441	6300	cutaneous T-cell lymphoma-associated antigen 1
CTAGE5	4253	NP_001139131	1873	CTAGE family, member 5 pseudogene, CTAGE family, member 4
CTAGE5	4253	NP_005921	3402	CTAGE family, member 5 pseudogene, CTAGE family, member 4
CTAGE5	4253	NP_940897	6802	CTAGE family, member 5 pseudogene, CTAGE family, member 4
CTAGE5	4253	NP_976229	6924	CTAGE family, member 5 pseudogene, CTAGE family, member 4
CTAGE5	4253	NP_976230	6925	CTAGE family, member 5 pseudogene, CTAGE family, member 4
CTAGE5	4253	NP_976231	6926	CTAGE family, member 5 pseudogene, CTAGE family, member 4
CTDNEP1	23399	NP_001137247	1713	dullardHomolog (Xenopus laevis)
CTDNEP1	23399	NP_056158	4064	dullardHomolog (Xenopus laevis)
CTLA4	1493	NP_001032720	779	cytotoxic T-lymphocyte-associated protein 4
CTLA4	1493	NP_005205	3230	cytotoxic T-lymphocyte-associated protein 4
CTNS	1497	NP_001026851	722	cystinosis, nephropathic
CTNS	1497	NP_004928	3173	cystinosis, nephropathic
CTRB2	440387	NP_001020371	701	chymotrypsinogen B2
CTSA	5476	NP_000299	87	cathepsin A
CTSA	5476	NP_001121167	1348	cathepsin A
CTSA	5476	NP_001161066	2186	cathepsin A
CTSZ	1522	NP_001327	2346	cathepsin Z
CTXN1	404217	NP_996664	6973	cortexin 1
CTXN3	613212	NP_001041717	934	cortexin 3
CTXN3	613212	NP_001120857	1322	cortexin 3
CUTA	51596	NP_001014433	621	cutA divalent cation toleranceHomolog (E. coli)
CUTA	51596	NP_001014837	625	cutA divalent cation toleranceHomolog (E. coli)
CUTA	51596	NP_001014838	626	cutA divalent cation toleranceHomolog (E. coli)
CUTA	51596	NP_001014840	627	cutA divalent cation toleranceHomolog (E. coli)
CUTA	51596	NP_057005	4135	cutA divalent cation toleranceHomolog (E. coli)
CUX1	1523	NP_001904	2498	cut-likeHomeobox 1
CUX1	1523	NP_852477	6603	cut-likeHomeobox 1
CUX1	1523	NP_853530	6608	cut-likeHomeobox 1
CWH43	80157	NP_079363	5119	hypothetical protein FLJ21511
CX3CL1	6376	NP_002987	2746	chemokine (C-X3-C motif) ligand 1
CX3CR1	1524	NP_001164642	2272	chemokine (C-X3-C motif) receptor 1
CX3CR1	1524	NP_001164643	2273	chemokine (C-X3-C motif) receptor 1
CX3CR1	1524	NP_001164645	2274	chemokine (C-X3-C motif) receptor 1

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
CX3CR1	1524	NP_001328	2347	chemokine (C-X3-C motif) receptor 1
CXADR	1525	NP_001329	2348	coxsackie virus and adenovirus receptor pseudogene 2 coxsackie virus and adenovirus receptor
CXCL10	3627	NP_001556.2	7375	C-X-C motif chemokine ligand 10
CXCL14	9547	NP_004878	3161	chemokine (C-X-C motif) ligand 14
CXCL16	58191	NP_001094282	1193	chemokine (C-X-C motif) ligand 16
CXCL16	58191	NP_071342	4870	chemokine (C-X-C motif) ligand 16
CXCL8	3576	NP_000575	168	interleukin 8
CXCL9	4283	NP_002407	2648	chemokine (C-X-C motif) ligand 9
CXCR1	3577	NP_000625	183	interleukin 8 receptor, alpha
CXCR2	3579	NP_001161770	2215	interleukin 8 receptor, beta
CXCR2	3579	NP_001548	2412	interleukin 8 receptor, beta
CXCR3	2833	NP_001136269	1703	chemokine (C-X-C motif) receptor 3
CXCR3	2833	NP_001495	2393	chemokine (C-X-C motif) receptor 3
CXCR4	7852	NP_001008540	547	chemokine (C-X-C motif) receptor 4
CXCR4	7852	NP_003458	2836	chemokine (C-X-C motif) receptor 4
CXCR5	643	NP_001707.1	7256	C-X-C motif chemokine receptor 5
CXCR6	10663	NP_006555	3512	chemokine (C-X-C motif) receptor 6
CYB561	1534	NP_001017916	646	cytochrome b-561
CYB561	1534	NP_001017917	647	cytochrome b-561
CYB561	1534	NP_001906	2500	cytochrome b-561
CYB561A3	220002	NP_001154924	2012	cytochrome b, ascorbate dependent 3
CYB561A3	220002	NP_001154926	2013	cytochrome b, ascorbate dependent 3
CYB561A3	220002	NP_705839	6201	cytochrome b, ascorbate dependent 3
CYB561D1	284613	NP_001127872	1482	cytochrome b-561 domain containing 1
CYB561D1	284613	NP_001127874	1483	cytochrome b-561 domain containing 1
CYB561D1	284613	NP_001127875	1484	cytochrome b-561 domain containing 1
CYB561D1	284613	NP_001127876	1485	cytochrome b-561 domain containing 1
CYB561D1	284613	NP_872386	6664	cytochrome b-561 domain containing 1
CYB561D2	11068	NP_008953	3616	cytochrome b-561 domain containing 2
CYB5A	1528	NP_001905	2499	cytochrome b5 type A (microsomal)
CYB5A	1528	NP_683725	6008	cytochrome b5 type A (microsomal)
CYB5B	80777	NP_085056	5150	cytochrome b5 type B (outer mitochondrial membrane)
CYB5D2	124936	NP_653212	5874	cytochrome b5 domain containing 2
CYB5R1	51706	NP_057327	4192	cytochrome b5 reductase 1
CYB5R2	51700	NP_057313	4186	cytochrome b5 reductase 2
CYBA	1535	NP_000092	27	cytochrome b-245, alpha polypeptide
CYBB	1536	NP_000388	118	cytochrome b-245, beta polypeptide
CYBRD1	79901	NP_001120855	1321	cytochrome b reductase 1
CYBRD1	79901	NP_079119	5091	cytochrome b reductase 1
CYHR1	50626	NP_001123360	1403	cysteine/histidine-rich 1
CYHR1	50626	NP_116076	5387	cysteine/histidine-rich 1
CYHR1	50626	NP_612505	5758	cysteine/histidine-rich 1
CYP19A1	1588	NP_000094	28	cytochrome P450, family 19, subfamily A, polypeptide 1
CYP19A1	1588	NP_112503	5211	cytochrome P450, family 19, subfamily A, polypeptide 1
CYP1A1	1543	NP_000490	141	cytochrome P450, family 1, subfamily A, polypeptide 1
CYP1A2	1544	NP_000752	236	cytochrome P450, family 1, subfamily A, polypeptide 2
CYP1B1	1545	NP_000095	29	cytochrome P450, family 1, subfamily B, polypeptide 1
CYP20A1	57404	NP_803882	6496	cytochrome P450, family 20, subfamily A, polypeptide 1
CYP26B1	56603	NP_063938	4590	cytochrome P450, family 26, subfamily B, polypeptide 1
CYP2A13	1553	NP_000757	240	cytochrome P450, family 2, subfamily A, polypeptide 13
CYP2A6	1548	NP_000753	237	cytochrome P450, family 2, subfamily A, polypeptide 6
CYP2A7	1549	NP_000755	238	cytochrome P450, family 2, subfamily A, polypeptide 7
CYP2A7	1549	NP_085079	5151	cytochrome P450, family 2, subfamily A, polypeptide 7
CYP2C8	1558	NP_000761	241	cytochrome P450, family 2, subfamily C, polypeptide 8
CYP2E1	1571	NP_000764	242	cytochrome P450, family 2, subfamily E, polypeptide 1
CYP2J2	1573	NP_000766	243	cytochrome P450, family 2, subfamily J, polypeptide 2
CYP2R1	120227	NP_078790	5038	cytochrome P450, family 2, subfamily R, polypeptide 1
CYP2S1	29785	NP_085125	5152	cytochrome P450, family 2, subfamily S, polypeptide 1
CYP2U1	113612	NP_898898	6702	cytochrome P450, family 2, subfamily U, polypeptide 1
CYP39A1	51302	NP_057677	4254	cytochrome P450, family 39, subfamily A, polypeptide 1
CYP3A4	1576	NP_059488	4296	cytochrome P450, family 3, subfamily A, polypeptide 4
CYP3A43	64816	NP_073731	4948	cytochrome P450, family 3, subfamily A, polypeptide 43
CYP3A43	64816	NP_476436	5575	cytochrome P450, family 3, subfamily A, polypeptide 43
CYP3A43	64816	NP_476437	5576	cytochrome P450, family 3, subfamily A, polypeptide 43
CYP3A5	1577	NP_000768	244	cytochrome P450, family 3, subfamily A, polypeptide 5
CYP3A7	1551	NP_000756	239	cytochrome P450, family 3, subfamily A, polypeptide 7
CYP4A11	1579	NP_000769	245	cytochrome P450, family 4, subfamily A, polypeptide 11
CYP4B1	1580	NP_000770	246	cytochrome P450, family 4, subfamily B, polypeptide 1
CYP4B1	1580	NP_001093242	1172	cytochrome P450, family 4, subfamily B, polypeptide 1
CYP4F12	66002	NP_076433	4987	similar to cytochrome P450, family 4, subfamily F, polypeptide 12
CYP4F2	8529	NP_001073	988	cytochrome P450, family 4, subfamily F, polypeptide 2
CYP4F22	126410	NP_775754	6344	cytochrome P450, family 4, subfamily F, polypeptide 22
CYP4F8	11283	NP_009184	3660	cytochrome P450, family 4, subfamily F, polypeptide 8
CYP4V2	285440	NP_997235	7015	cytochrome P450, family 4, subfamily V, polypeptide 2
CYP4X1	260293	NP_828847	6516	cytochrome P450, family 4, subfamily X, polypeptide 1
CYP4Z1	199974	NP_835235	6519	cytochrome P450, family 4, subfamily Z, polypeptide 1
CYP51A1	1595	NP_000777	247	cytochrome P450, family 51, subfamily A, polypeptide 1

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
CYP51A1	1595	NP_001139624	1908	cytochrome P450, family 51, subfamily A, polypeptide 1
CYP7B1	9420	NP_004811	3137	cytochrome P450, family 7, subfamily B, polypeptide 1
CYP8B1	1582	NP_004382	3038	cytochrome P450, family 8, subfamily B, polypeptide 1
CYSLTR1	10800	NP_006630	3534	cysteinyl leukotriene receptor 1
CYSLTR2	57105	NP_065110	4639	cysteinyl leukotriene receptor 2
CYYR1	116159	NP_443186	5549	cysteine/tyrosine-rich 1
DAD1	1603	NP_001335	2349	defender against cell death 1
DAG1	1605	NP_001159400	2128	dystroglycan 1 (dystrophin-associated glycoprotein 1)
DAG1	1605	NP_004384	3039	dystroglycan 1 (dystrophin-associated glycoprotein 1)
DAGLA	747	NP_006124	3439	diacylglycerol lipase, alpha
DAGLB	221955	NP_001136408	1708	diacylglycerol lipase, beta
DAGLB	221955	NP_631918	5846	diacylglycerol lipase, beta
DAPL1	92196	NP_001017920	648	death associated protein-like 1
DBH	1621	NP_000778	248	dopamine beta-hydroxylase (dopamine beta-monooxygenase)
DBI	1622	NP_001073331	1006	diazepam binding inhibitor (GABA receptor modulator, acyl-Coenzyme A binding protein)
DBI	1622	NP_001073332	1007	diazepam binding inhibitor (GABA receptor modulator, acyl-Coenzyme A binding protein)
DBI	1622	NP_065438	4684	diazepam binding inhibitor (GABA receptor modulator, acyl-Coenzyme A binding protein)
DCBLD1	285761	NP_775945	6378	discoidin, CUB and LCCL domain containing 1
DCBLD2	131566	NP_563615	5648	discoidin, CUB and LCCL domain containing 2
DCC	1630	NP_005206	3231	deleted in colorectal carcinoma
DCHS1	8642	NP_003728	2881	dachsous 1 (Drosophila)
DCHS2	54798	NP_001136024	1672	dachsous 2 (Drosophila)
DCHS2	54798	NP_001136025	1673	dachsous 2 (Drosophila)
DCHS2	54798	NP_060109	4316	dachsous 2 (Drosophila)
DCST1	149095	NP_001137159	1712	DC-STAMP domain containing 1
DCST1	149095	NP_689707	6070	DC-STAMP domain containing 1
DCST2	127579	NP_653223	5876	DC-STAMP domain containing 2
DCSTAMP	81501	NP_001244246.1	7433	dendrocyte expressed seven transmembrane protein
DCT	1638	NP_001123361	1404	dopachrome tautomerase (dopachrome delta-isomerase, tyrosine-related protein 2)
DCT	1638	NP_001913	2501	dopachrome tautomerase (dopachrome delta-isomerase, tyrosine-related protein 2)
DCUN1D5	84259	NP_115675	5321	DCN1, defective in cullin neddylation 1, domain containing 5 (S. cerevisiae)
DDB1	1642	NP_001914	2502	damage-specific DNA binding protein 1, 127kDa
DDB2	1643	NP_000098	30	damage-specific DNA binding protein 2, 48kDa
DDC	1644	NP_000781	250	dopa decarboxylase (aromatic L-amino acid decarboxylase)
DDC	1644	NP_001076440	1065	dopa decarboxylase (aromatic L-amino acid decarboxylase)
DDOST	1650	NP_005207	3232	dolichyl-diphosphooligosaccharide-protein glycosyltransferase
DDR1	780	NP_001945	2509	discoidin domain receptor tyrosine kinase 1
DDR1	780	NP_054699	3849	discoidin domain receptor tyrosine kinase 1
DDR1	780	NP_054700	3850	discoidin domain receptor tyrosine kinase 1
DDR2	4921	NP_001014796	623	discoidin domain receptor tyrosine kinase 2
DDR2	4921	NP_006173	3452	discoidin domain receptor tyrosine kinase 2
DDX3X	1654	NP_001180345.1	7424	DEAD-box helicase 3, X-linked
DDX59	83479	NP_001026895	728	DEAD (Asp-Glu-Ala-Asp) box polypeptide 59
DEAF1	10522	NP_066288	4758	deformed epidermal autoregulatory factor 1 (Drosophila)
DEFB126	81623	NP_112193	5197	defensin, beta 126
DEGS1	8560	NP_003667	2868	degenerative spermatocyteHomolog 1, lipid desaturase (Drosophila)
DEGS1	8560	NP_001308470.1	5905	degenerative spermatocyteHomolog 1, lipid desaturase (Drosophila)
DEGS2	123099	NP_996801	6980	degenerative spermatocyteHomolog 2, lipid desaturase (Drosophila)
DENND1B	163486	NP_001182144.1	1702	DENN/MADD domain containing 1B
DENND1B	163486	NP_659414	5911	DENN/MADD domain containing 1B
DENND5A	23258	NP_056028	4047	DENN/MADD domain containing 5A
DENND5B	160518	NP_659410	5910	DENN/MADD domain containing 5B
DERL1	79139	NP_001128143	1502	Der1-like domain family, member 1
DERL1	79139	NP_077271	5016	Der1-like domain family, member 1
DERL2	51009	NP_057125	4158	Der1-like domain family, member 2
DEXI	28955	NP_054734	3854	dexamethasone-induced transcript
DGAT1	8694	NP_036211	3699	diacylglycerol O-acyltransferaseHomolog 1 (mouse)
DGAT2	84649	NP_115953	5371	diacylglycerol O-acyltransferaseHomolog 2 (mouse)
DGCR2	9993	NP_005128	3219	DiGeorge syndrome critical region gene 2
DGKE	8526	NP_003638	2863	diacylglycerol kinase, epsilon 64kDa
DHCR24	1718	NP_055577	3985	24-dehydrocholesterol reductase
DHCR7	1717	NP_001157289	2058	7-dehydrocholesterol reductase
DHCR7	1717	NP_001351	2350	7-dehydrocholesterol reductase
DHODH	1723	NP_001352	2351	dihydroorotate dehydrogenase
DHRS11	79154	NP_077284	5021	dehydrogenase/reductase (SDR family) member 11
DHRS13	147015	NP_653284	5891	dehydrogenase/reductase (SDR family) member 13
DHRS2	10202	NP_005785	3369	dehydrogenase/reductase (SDR family) member 2
DHRS2	10202	NP_878912	6691	dehydrogenase/reductase (SDR family) member 2
DHRS3	9249	NP_004744	3117	dehydrogenase/reductase (SDR family) member 3
DHRS7	51635	NP_057113	4155	dehydrogenase/reductase (SDR family) member 7
DHRS7B	25979	NP_056325	4093	dehydrogenase/reductase (SDR family) member 7B
DHRS7C	201140	NP_001099041	1229	dehydrogenase/reductase (SDR family) member 7C
DHRS9	10170	NP_001135742	1628	dehydrogenase/reductase (SDR family) member 9
DHRS9	10170	NP_001135743	1629	dehydrogenase/reductase (SDR family) member 9
DHRS9	10170	NP_001135742.1	3364	dehydrogenase/reductase (SDR family) member 9
DHRS9	10170	NP_954674	6865	dehydrogenase/reductase (SDR family) member 9
DIABLO	56616	NP_063940	4591	diabloHomolog (Drosophila)

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
DIABLO	56616	NP_001265231.1	5805	diabloHomolog (Drosophila)
DIO1	1733	NP_000783	251	deiodinase, iodothyronine, type I
DIO1	1733	NP_001034804	810	deiodinase, iodothyronine, type I
DIO1	1733	NP_001034805	811	deiodinase, iodothyronine, type I
DIO1	1733	NP_998758	7039	deiodinase, iodothyronine, type I
DIO2	1734	NP_000784	252	deiodinase, iodothyronine, type II
DIO2	1734	NP_001007024	497	deiodinase, iodothyronine, type II
DIO2	1734	NP_054644	3848	deiodinase, iodothyronine, type II
DIO3	1735	NP_001353	2352	deiodinase, iodothyronine, type III
DIP2C	22982	NP_055789	4020	DIP2 disco-interacting protein 2Homolog C (Drosophila)
DISP1	84976	NP_116279	5420	dispatchedHomolog 1 (Drosophila)
DISP2	85455	NP_277045.1	7320	dispatched RND transporter family member 2
DKK1	22943	NP_036374.1	7282	dickkopf WNT signaling pathway inhibitor 1
DKKL1	27120	NP_055234	3935	dickkopf-like 1 (soggy)
DLK2	65989	NP_076421	4986	delta-like 2Homolog (Drosophila)
DLK2	65989	NP_996262	6964	delta-like 2Homolog (Drosophila)
DLL1	28514	NP_005609	3326	delta-like 1 (Drosophila)
DLL3	10683	NP_058637	4275	delta-like 3 (Drosophila)
DLL3	10683	NP_982353	6946	delta-like 3 (Drosophila)
DLL4	54567	NP_061947	4556	delta-like 4 (Drosophila)
DMRT2	10655	NP_001124337	1445	doublesex and mab-3 related transcription factor 2
DMRT2	10655	NP_006548	3511	doublesex and mab-3 related transcription factor 2
DMRT2	10655	NP_870987	6641	doublesex and mab-3 related transcription factor 2
DNAH10	196385	NP_997320	7021	dynein, axonemal,Heavy chain 10
DNAJB11	51726	NP_057390	4200	DnaJ (Hsp40)Homolog, subfamily B, member 11
DNAJB12	54788	NP_001002762	403	DnaJ (Hsp40)Homolog, subfamily B, member 12
DNAJB12	54788	NP_060096	4314	DnaJ (Hsp40)Homolog, subfamily B, member 12
DNAJB8	165721	NP_699161.1	7318	DnaJ heat shock protein family (Hsp40) member B8
DNAJB9	4189	NP_036460	3747	DnaJ (Hsp40)Homolog, subfamily B, member 9
DNAJC1	64215	NP_071760	4905	DnaJ (Hsp40)Homolog, subfamily C, member 1
DNAJC10	54431	NP_061854	4545	DnaJ (Hsp40)Homolog, subfamily C, member 10
DNAJC14	85406	NP_115740	5334	DnaJ (Hsp40)Homolog, subfamily C, member 14
DNAJC15	29103	NP_037370	3788	DnaJ (Hsp40)Homolog, subfamily C, member 15
DNAJC16	23341	NP_056106	4058	DnaJ (Hsp40)Homolog, subfamily C, member 16
DNAJC18	202052	NP_689899	6098	DnaJ (Hsp40)Homolog, subfamily C, member 18
DNAJC19	131118	NP_660304	5945	similar to translocase of the inner mitochondrial membrane 14, DnaJ (Hsp40)Homolog, subfamily C, member 19
DNAJC22	79962	NP_079178	5100	DnaJ (Hsp40)Homolog, subfamily C, member 22
DNAJC25	548645	NP_001015882	636	DnaJ (Hsp40)Homolog, subfamily C , member 25, guanine nucleotide binding protein (G protein), gamma 10
DNAJC25	548645	NP_001017998	660	DnaJ (Hsp40)Homolog, subfamily C , member 25, guanine nucleotide binding protein (G protein), gamma 10
DNAJC25	548645	NP_004116	2971	DnaJ (Hsp40)Homolog, subfamily C , member 25, guanine nucleotide binding protein (G protein), gamma 10
DNAJC30	84277	NP_115693	5325	DnaJ (Hsp40)Homolog, subfamily C, member 30
DNAJC4	3338	NP_005519	3308	DnaJ (Hsp40)Homolog, subfamily C, member 4
DNAJC5	80331	NP_079495	5134	DnaJ (Hsp40)Homolog, subfamily C, member 5
DNAJC5G	285126	NP_775921	6372	DnaJ (Hsp40)Homolog, subfamily C, member 5 gamma
DNASE1	1773	NP_005214	3233	deoxyribonuclease I
DNASE2B	58511	NP_067056	4799	deoxyribonuclease II beta
DNASE2B	58511	NP_490649	5588	deoxyribonuclease II beta
DNER	92737	NP_620711.3	7365	delta/notch like EGF repeat containing
DOLK	22845	NP_055723	4010	dolichol kinase
DOLPP1	57171	NP_001129389	1575	dolichyl pyrophosphate phosphatase 1
DOLPP1	57171	NP_065171	4659	dolichyl pyrophosphate phosphatase 1
DPAGT1	1798	NP_001373	2353	dolichyl-phosphate (UDP-N-acetylglucosamine) N-acetylglucosaminephosphotransferase 1 (GlcNAc-1-P transferase)
DPCR1	135656	NP_543146	5641	diffuse panbronchiolitis critical region 1
DPEP1	1800	NP_001121613.1	7391	dipeptidase 1 (renal)
DPEP2	64174	NP_071750	4901	dipeptidase 2
DPM2	8818	NP_003854	2920	dolichyl-phosphate mannosyltransferase polypeptide 2, regulatory subunit
DPP10	57628	NP_001004360	432	dipeptidyl-peptidase 10
DPP10	57628	NP_065919	4736	dipeptidyl-peptidase 10
DPP4	1803	NP_001926	2503	dipeptidyl-peptidase 4
DPP6	1804	NP_001034439	789	dipeptidyl-peptidase 6
DPP6	1804	NP_001927	2504	dipeptidyl-peptidase 6
DPP6	1804	NP_570629	5666	dipeptidyl-peptidase 6
DPP7	29952	NP_037511	3814	dipeptidyl-peptidase 7
DPP8	54878	NP_060213	4333	dipeptidyl-peptidase 8
DPP8	54878	NP_569118	5654	dipeptidyl-peptidase 8
DPP8	54878	NP_932064	6734	dipeptidyl-peptidase 8
DPP8	54878	NP_932065	6735	dipeptidyl-peptidase 8
DPY19L1	23333	NP_056098	4057	dpy-19-like 1 (C. elegans)
DPY19L2	283417	NP_776173	6389	dpy-19-like 2 (C. elegans)
DPY19L3	147991	NP_997208	7008	dpy-19-like 3 (C. elegans)
DPY19L4	286148	NP_861452	6635	dpy-19-like 4 (C. elegans)
DRAM1	55332	NP_060840	4431	DNA-damage regulated autophagy modulator 1
DRAM2	128338	NP_848549	6542	DNA-damage regulated autophagy modulator 2
DRAXIN	374946	NP_940947	6812	chromosome 1 open reading frame 187
DRD1	1812	NP_000785	253	dopamine receptor D1
DRD2	1813	NP_000786	254	dopamine receptor D2

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
DRD2	1813	NP_057658	4248	dopamine receptor D2
DRD3	1814	NP_000787	255	dopamine receptor D3
DRD3	1814	NP_387512	5511	dopamine receptor D3
DRD4	1815	NP_000788	256	dopamine receptor D4
DRD5	1816	NP_000789	257	dopamine receptor D5
DSC1	1823	NP_004939	3174	desmocollin 1
DSC1	1823	NP_077739	5028	desmocollin 1
DSC2	1824	NP_004940	3175	desmocollin 2
DSC2	1824	NP_077740	5029	desmocollin 2
DSC3	1825	NP_001932	2505	desmocollin 3
DSC3	1825	NP_077741	5030	desmocollin 3
DSCAM	1826	NP_001380	2354	Down syndrome cell adhesion molecule
DSCAML1	57453	NP_065744	4701	Down syndrome cell adhesion molecule like 1
DSE	29940	NP_001074445	1043	dermatan sulfate epimerase
DSE	29940	NP_037484	3810	dermatan sulfate epimerase
DSEL	92126	NP_115536	5303	dermatan sulfate epimerase-like
DSG2	1829	NP_001934	2506	desmoglein 2
DST	667	NP_001138241	1772	dystonin
DST	667	NP_001138242	1773	dystonin
DST	667	NP_001138243	1774	dystonin
DST	667	NP_001714	2445	dystonin
DST	667	NP_056363	4099	dystonin
DST	667	NP_001138241.1	4643	dystonin
DST	667	NP_899236	6707	dystonin
DTD1	92675	NP_036340	3722	histidyl-tRNA synthetase 2, mitochondrial (putative) D-tyrosyl-tRNA deacylase 1Homolog (S. cerevisiae)
DTD1	92675	NP_543010	5634	histidyl-tRNA synthetase 2, mitochondrial (putative) D-tyrosyl-tRNA deacylase 1Homolog (S. cerevisiae)
DTNA	1837	NP_001121647	1365	dystrobrevin, alpha
DTNA	1837	NP_001381	2355	dystrobrevin, alpha
DTNA	1837	NP_001382	2356	dystrobrevin, alpha
DTNA	1837	NP_001383	2357	dystrobrevin, alpha
DTNA	1837	NP_116757	5432	dystrobrevin, alpha
DTNA	1837	NP_116760	5433	dystrobrevin, alpha
DTNA	1837	NP_116761	5434	dystrobrevin, alpha
DTNA	1837	NP_116762	5435	dystrobrevin, alpha
DTNA	1837	NP_116763	5436	dystrobrevin, alpha
DUOX1	53905	NP_059130	4289	dual oxidase 1
DUOX1	53905	NP_787954	6472	dual oxidase 1
DUOX2	50506	NP_054799	3868	dual oxidase 2
DUOXA1	90527	NP_653166	5862	dual oxidase maturation factor 1
DUOXA2	405753	NP_997464	7025	dual oxidase maturation factor 2
DUSP13	51207	NP_001007272	513	dual specificity phosphatase 13
DUSP13	51207	NP_001007273	514	dual specificity phosphatase 13
DUSP13	51207	NP_001007274	515	dual specificity phosphatase 13
DUSP13	51207	NP_057448	4212	dual specificity phosphatase 13
DYSF	8291	NP_001123927	1425	dysferlin, limb girdle muscular dystrophy 2B (autosomal recessive)
DYSF	8291	NP_001124448	1453	dysferlin, limb girdle muscular dystrophy 2B (autosomal recessive)
DYSF	8291	NP_001124449	1454	dysferlin, limb girdle muscular dystrophy 2B (autosomal recessive)
DYSF	8291	NP_001124450	1455	dysferlin, limb girdle muscular dystrophy 2B (autosomal recessive)
DYSF	8291	NP_001124451	1456	dysferlin, limb girdle muscular dystrophy 2B (autosomal recessive)
DYSF	8291	NP_001124452	1457	dysferlin, limb girdle muscular dystrophy 2B (autosomal recessive)
DYSF	8291	NP_001124453	1458	dysferlin, limb girdle muscular dystrophy 2B (autosomal recessive)
DYSF	8291	NP_001124454	1459	dysferlin, limb girdle muscular dystrophy 2B (autosomal recessive)
DYSF	8291	NP_001124455	1460	dysferlin, limb girdle muscular dystrophy 2B (autosomal recessive)
DYSF	8291	NP_001124456	1461	dysferlin, limb girdle muscular dystrophy 2B (autosomal recessive)
DYSF	8291	NP_001124457	1462	dysferlin, limb girdle muscular dystrophy 2B (autosomal recessive)
DYSF	8291	NP_001124458	1463	dysferlin, limb girdle muscular dystrophy 2B (autosomal recessive)
DYSF	8291	NP_001124459	1464	dysferlin, limb girdle muscular dystrophy 2B (autosomal recessive)
DYSF	8291	NP_003485	2842	dysferlin, limb girdle muscular dystrophy 2B (autosomal recessive)
E2F5	1875	NP_001077057	1068	E2F transcription factor 5, p130-binding
E2F5	1875	NP_001077058	1069	E2F transcription factor 5, p130-binding
E2F5	1875	NP_001942	2508	E2F transcription factor 5, p130-binding
EBP	10682	NP_006570	3519	emopamil binding protein (sterol isomerase)
ECE1	1889	NP_001106818	1254	endothelin converting enzyme 1
ECE1	1889	NP_001106819	1255	endothelin converting enzyme 1
ECE1	1889	NP_001106820	1256	endothelin converting enzyme 1
ECE1	1889	NP_001388	2358	endothelin converting enzyme 1
ECE2	9718	NP_001032401	774	endothelin converting enzyme 2
ECE2	9718	NP_001093590	1180	endothelin converting enzyme 2
ECE2	9718	NP_001093591	1181	endothelin converting enzyme 2
ECE2	9718	NP_055508	3974	endothelin converting enzyme 2
ECE2	9718	NP_115707	5329	endothelin converting enzyme 2
ECEL1	9427	NP_004817	3141	endothelin converting enzyme-like 1
ECHDC1	55862	NP_001002030	388	enoyl Coenzyme AHydratase domain containing 1
ECHDC1	55862	NP_001099014	1227	enoyl Coenzyme AHydratase domain containing 1
ECHDC1	55862	NP_001099015	1228	enoyl Coenzyme AHydratase domain containing 1
ECHDC1	55862	NP_001132982	1621	enoyl Coenzyme AHydratase domain containing 1
ECHDC1	55862	NP_060949	4458	enoyl Coenzyme AHydratase domain containing 1

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
ECM1	1893	NP_004416	3040	extracellular matrix protein 1
ECM1	1893	NP_073155	4936	extracellular matrix protein 1
ECSCR	641700	NP_001071161	979	endothelial cell-specific chemotaxis regulator
EDA	1896	NP_001005609	464	ectodysplasin A
EDA	1896	NP_001005610	465	ectodysplasin A
EDA	1896	NP_001005611	466	ectodysplasin A
EDA	1896	NP_001005612	467	ectodysplasin A
EDA	1896	NP_001005615	468	ectodysplasin A
EDA	1896	NP_001390	2359	ectodysplasin A
EDA2R	60401	NP_068555	4828	ectodysplasin A2 receptor
EDAR	10913	NP_071731	4897	ectodysplasin A receptor
EDDM3A	10876	NP_006674	3545	family with sequence similarity 12, member A
EDDM3B	64184	NP_071755	4903	family with sequence similarity 12, member B (epididymal)
EDEM1	9695	NP_055489	3972	ER degradation enhancer, mannosidase alpha-like 1
EDNRA	1909	NP_001159527	2141	endothelin receptor type A
EDNRA	1909	NP_001948	2510	endothelin receptor type A
EDNRB	1910	NP_000106	33	endothelin receptor type B
EDNRB	1910	NP_001116131	1285	endothelin receptor type B
EDNRB	1910	NP_003982	2941	endothelin receptor type B
EEF1E1	9521	NP_001129122	1544	eukaryotic translation elongation factor 1 epsilon 1
EEF1E1	9521	NP_004271	3007	eukaryotic translation elongation factor 1 epsilon 1
EFCAB14	9813	NP_055589	3987	KIAA0494
EFNA5	1946	NP_001953	2511	ephrin-A5
EFNB1	1947	NP_004420	3041	ephrin-B1
EFNB2	1948	NP_004084	2961	ephrin-B2
EFNB3	1949	NP_001397	2362	ephrin-B3
EGF	1950	NP_001954	2512	epidermal growth factor (beta-urogastrone)
EGFLAM	133584	NP_689616	6053	EGF-like, fibronectin type III and laminin G domains
EGFLAM	133584	NP_877950	6683	EGF-like, fibronectin type III and laminin G domains
EGFLAM	133584	NP_877951	6684	EGF-like, fibronectin type III and laminin G domains
EGFLAM	133584	NP_877953	6685	EGF-like, fibronectin type III and laminin G domains
EGFR	1956	NP_005219	3235	epidermal growth factor receptor (erythroblastic leukemia viral (v-erb-b) oncogeneHomolog, avian)
EGFR	1956	NP_958439	6896	epidermal growth factor receptor (erythroblastic leukemia viral (v-erb-b) oncogeneHomolog, avian)
EGFR	1956	NP_958440	6897	epidermal growth factor receptor (erythroblastic leukemia viral (v-erb-b) oncogeneHomolog, avian)
EGFR	1956	NP_958441	6898	epidermal growth factor receptor (erythroblastic leukemia viral (v-erb-b) oncogeneHomolog, avian)
EI24	9538	NP_001007278	516	etoposide induced 2.4 mRNA
EI24	9538	NP_004870	3157	etoposide induced 2.4 mRNA
EIF2AK3	9451	NP_004827	3146	eukaryotic translation initiation factor 2-alpha kinase 3
ELANE	1991	NP_001963	2513	elastase, neutrophil expressed
ELFN2	114794	NP_443138	5534	extracellular leucine-rich repeat and fibronectin type III domain containing 2
ELN	2006	NP_000492	142	elastin
ELN	2006	NP_001075221	1056	elastin
ELN	2006	NP_001075222	1057	elastin
ELN	2006	NP_001075223	1058	elastin
ELN	2006	NP_001075224	1059	elastin
ELOVL1	64834	NP_073732	4949	elongation of very long chain fatty acids (FEN1/Elo2, SUR4/Elo3, yeast)-like 1
ELOVL2	54898	NP_060240	4340	elongation of very long chain fatty acids (FEN1/Elo2, SUR4/Elo3, yeast)-like 2
ELOVL3	83401	NP_689523	6028	elongation of very long chain fatty acids (FEN1/Elo2, SUR4/Elo3, yeast)-like 3
ELOVL4	6785	NP_073563	4937	elongation of very long chain fatty acids (FEN1/Elo2, SUR4/Elo3, yeast)-like 4
ELOVL5	60481	NP_068586	4836	ELOVL family member 5, elongation of long chain fatty acids (FEN1/Elo2, SUR4/Elo3-like, yeast)
ELOVL6	79071	NP_001124193	1438	ELOVL family member 6, elongation of long chain fatty acids (FEN1/Elo2, SUR4/Elo3-like, yeast)
ELOVL6	79071	NP_076995	5008	ELOVL family member 6, elongation of long chain fatty acids (FEN1/Elo2, SUR4/Elo3-like, yeast)
ELOVL7	79993	NP_001098028	1216	ELOVL family member 7, elongation of long chain fatty acids (yeast)
ELOVL7	79993	NP_079206	5104	ELOVL family member 7, elongation of long chain fatty acids (yeast)
ELSPBP1	64100	NP_071425	4883	epididymal sperm binding protein 1
EMB	133418	NP_940851	6796	embiginHomolog (mouse)
EMC3	55831	NP_060917	4453	transmembrane protein 111
EMC4	51234	NP_057538	4220	transmembrane protein 85
EMC6	83460	NP_001014764	622	transmembrane protein 93
EMC6	83460	NP_112588	5217	transmembrane protein 93
EMC7	56851	NP_064539	4604	chromosome 15 open reading frame 24
EMCN	51705	NP_001153166	1951	endomucin
EMCN	51705	NP_057326	4191	endomucin
EMID1	129080	NP_597712	5693	EMI domain containing 1
EML2	24139	NP_036287	3715	echinoderm microtubule associated protein like 2
EMP1	2012	NP_001414	2368	epithelial membrane protein 1
EMP2	2013	NP_001415	2369	epithelial membrane protein 2
EMP3	2014	NP_001416	2370	epithelial membrane protein 3
ENDOD1	23052	NP_055851	4027	endonuclease domain containing 1
ENG	2022	NP_000109	35	endoglin
ENG	2022	NP_001108225	1278	endoglin
ENHO	375704	NP_940975	6819	energyHomeostasis associated
ENPEP	2028	NP_001968	2515	glutamyl aminopeptidase (aminopeptidase A)
ENPP1	5167	NP_006199	3456	ectonucleotide pyrophosphatase/phosphodiesterase 1
ENPP2	5168	NP_001035181	827	ectonucleotide pyrophosphatase/phosphodiesterase 2
ENPP2	5168	NP_001124335	1444	ectonucleotide pyrophosphatase/phosphodiesterase 2
ENPP2	5168	NP_006200	3457	ectonucleotide pyrophosphatase/phosphodiesterase 2

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
ENPP3	5169	NP_005012	3199	ectonucleotide pyrophosphatase/phosphodiesterase 3
ENPP4	22875	NP_055751	4017	ectonucleotide pyrophosphatase/phosphodiesterase 4 (putative function)
ENPP5	59084	NP_067547	4807	ectonucleotide pyrophosphatase/phosphodiesterase 5 (putative function)
ENTPD1	953	NP_001091645	1115	ectonucleoside triphosphate diphosphohydrolase 1
ENTPD1	953	NP_001157650	2062	ectonucleoside triphosphate diphosphohydrolase 1
ENTPD1	953	NP_001157651	2063	ectonucleoside triphosphate diphosphohydrolase 1
ENTPD1	953	NP_001157653	2064	ectonucleoside triphosphate diphosphohydrolase 1
ENTPD1	953	NP_001157654	2065	ectonucleoside triphosphate diphosphohydrolase 1
ENTPD1	953	NP_001157655	2066	ectonucleoside triphosphate diphosphohydrolase 1
ENTPD1	953	NP_001767	2466	ectonucleoside triphosphate diphosphohydrolase 1
ENTPD2	954	NP_001237	2326	ectonucleoside triphosphate diphosphohydrolase 2
ENTPD2	954	NP_982293	6940	ectonucleoside triphosphate diphosphohydrolase 2
ENTPD3	956	NP_001239	2328	ectonucleoside triphosphate diphosphohydrolase 3
ENTPD4	9583	NP_001122402	1381	ectonucleoside triphosphate diphosphohydrolase 4
ENTPD4	9583	NP_004892	3164	ectonucleoside triphosphate diphosphohydrolase 4
ENTPD5	957	NP_001240	2329	ectonucleoside triphosphate diphosphohydrolase 5
ENTPD6	955	NP_001107561	1265	ectonucleoside triphosphate diphosphohydrolase 6 (putative function)
ENTPD6	955	NP_001238	2327	ectonucleoside triphosphate diphosphohydrolase 6 (putative function)
ENTPD7	57089	NP_065087	4634	ectonucleoside triphosphate diphosphohydrolase 7
EOGT	285203	NP_775925	6374	chromosome 3 open reading frame 64
EPCAM	4072	NP_002345	2632	epithelial cell adhesion molecule
EPDR1	54749	NP_060019	4304	ependymin related protein 1 (zebrafish)
EPHA1	2041	NP_005223	3236	EPH receptor A1
EPHA10	284656	NP_001092909	1160	EPH receptor A10
EPHA10	284656	NP_775912	6369	EPH receptor A10
EPHA2	1969	NP_004422	3042	EPH receptor A2
EPHA3	2042	NP_005224	3237	EPH receptor A3
EPHA3	2042	NP_872585	6669	EPH receptor A3
EPHA4	2043	NP_004429	3043	EPH receptor A4
EPHA5	2044	NP_004430	3044	EPH receptor A5
EPHA5	2044	NP_872272	6643	EPH receptor A5
EPHA6	285220	NP_001073917	1023	EPH receptor A6
EPHA6	285220	NP_775926	6375	EPH receptor A6
EPHA7	2045	NP_004431	3045	EPH receptor A7
EPHA8	2046	NP_001006944	495	EPH receptor A8
EPHA8	2046	NP_065387	4678	EPH receptor A8
EPHB1	2047	NP_004432	3046	EPH receptor B1
EPHB2	2048	NP_004433	3047	EPH receptor B2
EPHB2	2048	NP_059145	4294	EPH receptor B2
EPHB3	2049	NP_004434	3048	EPH receptor B3
EPHB4	2050	NP_004435	3049	EPH receptor B4
EPHB6	2051	NP_004436	3050	EPH receptor B6
EPHX3	79852	NP_001136358	1707	epoxideHydrolase 3
EPHX3	79852	NP_079070	5084	epoxideHydrolase 3
EPHX4	253152	NP_775838	6359	epoxideHydrolase 4
EPO	2056	NP_000790	258	erythropoietin
EPOR	2057	NP_000112	36	erythropoietin receptor
EPSTI1	94240	NP_001002264	395	epithelial stromal interaction 1 (breast)
EPSTI1	94240	NP_150280	5475	epithelial stromal interaction 1 (breast)
EPT1	85465	NP_277040	5503	selenoprotein I
EQTN	54586	NP_001155057	2021	chromosome 9 open reading frame 11
EQTN	54586	NP_065692	4687	chromosome 9 open reading frame 11
ERBB2	2064	NP_001005862	474	v-erb-b2 erythroblastic leukemia viral oncogeneHomolog 2, neuro/glioblastoma derived oncogeneHomolog (avian)
ERBB2	2064	NP_004439	3051	v-erb-b2 erythroblastic leukemia viral oncogeneHomolog 2, neuro/glioblastoma derived oncogeneHomolog (avian)
ERBB3	2065	NP_001005915	476	v-erb-b2 erythroblastic leukemia viral oncogeneHomolog 3 (avian)
ERBB3	2065	NP_001973	2517	v-erb-b2 erythroblastic leukemia viral oncogeneHomolog 3 (avian)
ERBB4	2066	NP_001036064	914	v-erb-a erythroblastic leukemia viral oncogeneHomolog 4 (avian)
ERBB4	2066	NP_005226	3238	v-erb-a erythroblastic leukemia viral oncogeneHomolog 4 (avian)
EREG	2069	NP_001423	2371	epiregulin
ERF	2077	NP_006485	3498	Ets2 repressor factor
ERGIC1	57222	NP_001026881	725	endoplasmic reticulum-golgi intermediate compartment (ERGIC) 1
ERGIC2	51290	NP_057654	4247	ERGIC and golgi 2
ERGIC3	51614	NP_057050	4144	ERGIC and golgi 3
ERGIC3	51614	NP_938408	6790	ERGIC and golgi 3
ERLIN1	10613	NP_001094096	1192	ER lipid raft associated 1
ERLIN1	10613	NP_006450	3494	ER lipid raft associated 1
ERMAP	114625	NP_001017922	649	erythroblast membrane-associated protein (Scianna blood group)
ERMAP	114625	NP_061008	4465	erythroblast membrane-associated protein (Scianna blood group)
ERMARD	55780	NP_060811	4424	chromosome 6 open reading frame 70
ERMP1	79956	NP_079172	5098	endoplasmic reticulum metallopeptidase 1
ERN1	2081	NP_001424	2372	endoplasmic reticulum to nucleus signaling 1
ERP29	10961	NP_001029197	758	endoplasmic reticulum protein 29
ERP29	10961	NP_006808	3572	endoplasmic reticulum protein 29
ERVFRD1	405754	NP_997465.1	7333	syncytin-2 preproprotein
ERVFRD-1	405754	NP_997465	7026	HERV-FRD provirus ancestral Env polyprotein

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
ERVMER34-1	100288413	XP_002342492	7076	similar to Uncharacterized protein LP9056
ERVMER34-1	100288413	XP_002345794	7081	similar to Uncharacterized protein LP9056
ERVMER34-1	100288413	XP_002346661	7083	similar to Uncharacterized protein LP9056
ERVW-1	30816	NP_001124397	1449	endogenous retroviral family W, env(C7), member 1
ERVW-1	30816	NP_055405	3963	endogenous retroviral family W, env(C7), member 1
ESAM	90952	NP_620411.2	7372	endothelial cell adhesion molecule
ESYT1	23344	NP_056107	4059	family with sequence similarity 62 (C2 domain containing), member A
ESYT2	57488	NP_065779	4709	family with sequence similarity 62 (C2 domain containing), member B
ESYT3	83850	NP_114119	5269	family with sequence similarity 62 (C2 domain containing), member C
ETFDH	2110	NP_004444	3052	electron-transferring-flavoprotein dehydrogenase
ETNK1	55500	NP_001034570	795	ethanolamine kinase 1
ETNK1	55500	NP_061108	4472	ethanolamine kinase 1
ETV6	2120	NP_001978	2518	ets variant 6
EVA1A	84141	NP_001128504	1512	family with sequence similarity 176, member A
EVA1A	84141	NP_115557	5305	family with sequence similarity 176, member A
EVA1B	55194	NP_060636	4400	family with sequence similarity 176, member B
EVA1C	59271	NP_478067	5585	chromosome 21 open reading frame 63
EVC	2121	NP_714928	6220	Ellis van Creveld syndrome
EVC2	132884	NP_001159608	2153	Ellis van Creveld syndrome 2
EVC2	132884	NP_667338	5988	Ellis van Creveld syndrome 2
EVI2A	2123	NP_001003927	426	ecotropic viral integration site 2A
EVI2A	2123	NP_055025	3882	ecotropic viral integration site 2A
EVI2B	2124	NP_006486	3499	ecotropic viral integration site 2B
EVI5L	115704	NP_001153416	1963	ecotropic viral integration site 5-like
EVI5L	115704	NP_660288	5939	ecotropic viral integration site 5-like
EXOG	9941	NP_001138936	1865	endo/exonuclease (5'-3'), endonuclease G-like
EXOG	9941	NP_005098	3212	endo/exonuclease (5'-3'), endonuclease G-like
EXT1	2131	NP_000118	37	exostoses (multiple) 1
EXTL1	2134	NP_004446	3053	exostoses (multiple)-like 1
EXTL2	2135	NP_001028197	749	exostoses (multiple)-like 2
EXTL2	2135	NP_001430	2373	exostoses (multiple)-like 2
EXTL3	2137	NP_001431	2374	exostoses (multiple)-like 3
F10	2159	NP_000495	143	coagulation factor X
F11R	50848	NP_058642	4278	F11 receptor
F2R	2149	NP_001983	2519	coagulation factor II (thrombin) receptor
F2RL1	2150	NP_005233	3239	coagulation factor II (thrombin) receptor-like 1
F2RL2	2151	NP_004092	2963	coagulation factor II (thrombin) receptor-like 2
F3	2152	NP_001984	2520	coagulation factor III (thromboplastin, tissue factor)
FA2H	79152	NP_077282	5019	fatty acid 2-hydroxylase
FAAH	2166	NP_001432	2375	fatty acid amideHydrolase
FAAH2	158584	NP_777572	6408	fatty acid amideHydrolase 2
FADS2	9415	NP_001268430.1	7439	fatty acid desaturase 2
FADS3	3995	NP_068373	4822	fatty acid desaturase 3
FAIM2	23017	NP_036438	3741	Fas apoptotic inhibitory molecule 2
FAM105A	54491	NP_061891	4548	family with sequence similarity 105, member A
FAM132A	388581	NP_001014980	629	family with sequence similarity 132, member A
FAM132B	151176	XP_001130886	7050	family with sequence similarity 132, member B
FAM132B	151176	XP_001714437	7052	family with sequence similarity 132, member B
FAM134A	79137	NP_077269	5015	family with sequence similarity 134, member A
FAM134B	54463	NP_001030022	762	family with sequence similarity 134, member B
FAM134B	54463	NP_061873	4547	family with sequence similarity 134, member B
FAM134C	162427	NP_835227	6517	family with sequence similarity 134, member C
FAM151A	338094	NP_788954	6475	family with sequence similarity 151, member A
FAM159A	348378	NP_001036158	916	family with sequence similarity 159, member A
FAM162A	26355	NP_055182	3923	family with sequence similarity 162, member A
FAM162B	221303	NP_001078949	1095	family with sequence similarity 162, member B
FAM163A	148753	NP_775780	6349	family with sequence similarity 163, member A
FAM171A1	221061	NP_001010924	578	family with sequence similarity 171, member A1
FAM171A2	284069	NP_940877	6798	family with sequence similarity 171, member A2
FAM171B	165215	NP_803237	6494	family with sequence similarity 171, member B
FAM173B	134145	NP_954584	6858	family with sequence similarity 173, member B
FAM174A	345757	NP_940909	6806	family with sequence similarity 174, member A
FAM174B	400451	NP_997329	7022	family with sequence similarity 174, member B
FAM187B	148109	NP_689694	6067	family with sequence similarity 187, member B
FAM189A1	23359	NP_056122	4060	KIAA0574 protein
FAM189A2	9413	NP_001121080	1334	chromosome 9 open reading frame 61
FAM189A2	9413	NP_004807	3136	chromosome 9 open reading frame 61
FAM189B	10712	NP_006580	3524	chromosome 1 open reading frame 2
FAM189B	10712	NP_937995	6763	chromosome 1 open reading frame 2
FAM198A	729085	NP_001123380	1405	chromosome 3 open reading frame 41Hypothetical protein LOC729085
FAM198B	51313	NP_001026870	724	chromosome 4 open reading frame 18
FAM198B	51313	NP_001121896	1372	chromosome 4 open reading frame 18
FAM198B	51313	NP_057697	4263	chromosome 4 open reading frame 18
FAM19A1	407738	NP_998774	7041	family with sequence similarity 19 (chemokine (C-C motif)-like), member A1
FAM19A2	338811	NP_848634	6552	family with sequence similarity 19 (chemokine (C-C motif)-like), member A2
FAM19A4	151647	NP_001005527	462	family with sequence similarity 19 (chemokine (C-C motif)-like), member A4
FAM19A4	151647	NP_872328	6649	family with sequence similarity 19 (chemokine (C-C motif)-like), member A4

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
FAM19A5	25817	NP_001076436	1064	family with sequence similarity 19 (chemokine (C-C motif)-like), member A5
FAM19A5	25817	NP_056196	4073	family with sequence similarity 19 (chemokine (C-C motif)-like), member A5
FAM200A	221786	NP_659802	5927	chromosome 7 open reading frame 38
FAM205A	259308	NP_001135389	1624	hypothetical protein LOC259308 chromosome 9 open reading frame 144
FAM209A	200232	NP_001012989	608	chromosome 20 open reading frame 106
FAM209B	388799	NP_001013668	617	chromosome 20 open reading frame 107
FAM20A	54757	NP_060035	4306	family with sequence similarity 20, member A
FAM20B	9917	NP_055679	4002	family with sequence similarity 20, member B
FAM20C	56975	NP_064608	4620	family with sequence similarity 20, member C
FAM210A	125228	NP_001092271	1146	chromosome 18 open reading frame 19
FAM210A	125228	NP_689565	6039	chromosome 18 open reading frame 19
FAM210B	116151	NP_543011	5635	chromosome 20 open reading frame 108
FAM213A	84293	NP_079401	5123	chromosome 10 open reading frame 58
FAM213A	84293	NP_115709	5331	chromosome 10 open reading frame 58
FAM24A	118670	NP_001025059	711	family with sequence similarity 24, member A
FAM24B	196792	NP_689857	6092	family with sequence similarity 24, member B
FAM26D	221301	NP_694581	6152	family with sequence similarity 26, member D
FAM26E	254228	NP_714922	6219	family with sequence similarity 26, member E
FAM26F	441168	NP_001010919	577	family with sequence similarity 26, member F
FAM3A	60343	NP_001164603	2269	family with sequence similarity 3, member A
FAM3A	60343	NP_001164604	2270	family with sequence similarity 3, member A
FAM3A	60343	NP_001164605	2271	family with sequence similarity 3, member A
FAM3A	60343	NP_068578	4833	family with sequence similarity 3, member A
FAM3B	54097	NP_478066	5584	family with sequence similarity 3, member B
FAM3B	54097	NP_996847	6993	family with sequence similarity 3, member B
FAM3D	131177	NP_620160	5799	family with sequence similarity 3, member D
FAM57A	79850	NP_079068	5083	family with sequence similarity 57, member A
FAM57B	83723	NP_113666	5232	family with sequence similarity 57, member B
FAM69A	388650	NP_001006606	478	family with sequence similarity 69, member A
FAM69B	138311	NP_689634	6058	family with sequence similarity 69, member B
FAM69C	125704	NP_001037834	920	chromosome 18 open reading frame 51
FAM73A	374986	NP_940951	6813	family with sequence similarity 73, member A
FAM73B	84895	NP_116198	5407	family with sequence similarity 73, member B
FAM76B	143684	NP_653265	5886	family with sequence similarity 76, member B
FAM8A1	51439	NP_057339	4195	family with sequence similarity 8, member A1
FAM9C	171484	NP_777561	6406	family with sequence similarity 9, member C
FANCM	57697	NP_065988	4743	Fanconi anemia, complementation group M
FAP	2191	NP_004451	3055	fibroblast activation protein, alpha
FAR1	84188	NP_115604	5309	fatty acyl CoA reductase 1
FAR2	55711	NP_060569	4391	fatty acyl CoA reductase 2
FASLG	356	NP_000630	184	Fas ligand (TNF superfamily, member 6)
FAT1	2195	NP_005236	3240	FAT tumor suppressorHomolog 1 (Drosophila)
FAT2	2196	NP_001438	2376	FAT tumor suppressorHomolog 2 (Drosophila)
FAT3	120114	NP_001008781	553	FAT tumor suppressorHomolog 3 (Drosophila)
FAT4	79633	NP_078858	5052	FAT tumor suppressorHomolog 4 (Drosophila)
FAXDC2	10826	NP_115761.2	4206	chromosome 5 open reading frame 4
FAXDC2	10826	NP_115761	5337	chromosome 5 open reading frame 4
FBN2	2201	NP_001990	2522	fibrillin 2
FBXL17	64839	NP_001156787	2051	F-box and leucine-rich repeat protein 17
FBXL17	64839	NP_001156787.2	4951	F-box and leucine-rich repeat protein 17
FBXW7	55294	NP_001013433	612	F-box and WD repeat domain containing 7
FBXW7	55294	NP_060785	4419	F-box and WD repeat domain containing 7
FBXW7	55294	NP_361014	5509	F-box and WD repeat domain containing 7
FCAMR	83953	NP_001116451.1	7388	Fc fragment of IgA and IgM receptor
FCAR	2204	NP_001991	2523	Fc fragment of IgA, receptor for
FCAR	2204	NP_579803	5679	Fc fragment of IgA, receptor for
FCAR	2204	NP_579805	5680	Fc fragment of IgA, receptor for
FCAR	2204	NP_579806	5681	Fc fragment of IgA, receptor for
FCAR	2204	NP_579807	5682	Fc fragment of IgA, receptor for
FCAR	2204	NP_579808	5683	Fc fragment of IgA, receptor for
FCAR	2204	NP_579811	5684	Fc fragment of IgA, receptor for
FCAR	2204	NP_579812	5685	Fc fragment of IgA, receptor for
FCAR	2204	NP_579813	5686	Fc fragment of IgA, receptor for
FCER1A	2205	NP_001992	2524	Fc fragment of IgE,High affinity I, receptor for alpha polypeptide
FCER1G	2207	NP_004097	2964	Fc fragment of IgE,High affinity I, receptor for gamma polypeptide
FCER2	2208	NP_001993	2525	Fc fragment of IgE, low affinity II, receptor for (CD23)
FCGR2B	2213	NP_001002273	398	Fc fragment of IgG, low affinity IIb, receptor (CD32) Fc fragment of IgG, low affinity IIc, receptor for (CD32)
FCGR2B	2213	NP_001002274	399	Fc fragment of IgG, low affinity IIb, receptor (CD32) Fc fragment of IgG, low affinity IIc, receptor for (CD32)
FCGR2B	2213	NP_001002275	400	Fc fragment of IgG, low affinity IIb, receptor (CD32) Fc fragment of IgG, low affinity IIc, receptor for (CD32)
FCGR2B	2213	NP_003992	2945	Fc fragment of IgG, low affinity IIb, receptor (CD32) Fc fragment of IgG, low affinity IIc, receptor for (CD32)
FCGR2B	2213	NP_963857	6909	Fc fragment of IgG, low affinity IIb, receptor (CD32) Fc fragment of IgG, low affinity IIc, receptor for (CD32)
FCGR3B	2215	NP_000561	164	Fc fragment of IgG, low affinity IIIb, receptor (CD16b)
FCGRT	2217	NP_001129491	1589	Fc fragment of IgG, receptor, transporter, alpha
FCGRT	2217	NP_004098	2965	Fc fragment of IgG, receptor, transporter, alpha
FCMR	9214	NP_001135945	1666	Fas apoptotic inhibitory molecule 3
FCMR	9214	NP_001135945	1667	Fas apoptotic inhibitory molecule 3
FCMR	9214	NP_005440	3287	Fas apoptotic inhibitory molecule 3

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
FCN2	2220	NP_004099	2966	ficolin (collagen/fibrinogen domain containing lectin) 2 (hucolin)
FCN2	2220	NP_056652	4122	ficolin (collagen/fibrinogen domain containing lectin) 2 (hucolin)
FCRL1	115350	NP_001152869	1941	Fc receptor-like 1
FCRL1	115350	NP_001152870	1942	Fc receptor-like 1
FCRL1	115350	NP_443170	5544	Fc receptor-like 1
FCRL2	79368	NP_110391	5166	Fc receptor-like 2
FCRL3	115352	NP_443171	5545	Fc receptor-like 3
FCRL4	83417	NP_112572	5214	Fc receptor-like 4
FCRL5	83416	NP_112571	5213	Fc receptor-like 5
FCRLA	84824	NP_116127	5393	Fc receptor-like A
FDCSP	260436	NP_694542	6144	chromosome 4 open reading frame 7
FDPS	2224	NP_001129293	1570	farnesyl diphosphate synthase (farnesyl pyrophosphate synthetase, dimethylallyltranstransferase, geranyltranstransferase)
FDPS	2224	NP_001129294	1571	farnesyl diphosphate synthase (farnesyl pyrophosphate synthetase, dimethylallyltranstransferase, geranyltranstransferase)
FDPS	2224	NP_001995	2526	farnesyl diphosphate synthase (farnesyl pyrophosphate synthetase, dimethylallyltranstransferase, geranyltranstransferase)
FER1L5	90342	NP_001280012.1	964	fer-1-like 5 (C. elegans)
FER1L5	90342	NP_001280012.1	1257	fer-1-like 5 (C. elegans)
FFAR1	2864	NP_005294	3269	free fatty acid receptor 1
FFAR2	2867	NP_005297	3270	free fatty acid receptor 2
FFAR4	338557	NP_859529	6628	G protein-coupled receptor 120
FGF10	2255	NP_004456	3056	fibroblast growth factor 10
FGF19	9965	NP_005108	3215	fibroblast growth factor 19
FGF4	2249	NP_001998	2527	fibroblast growth factor 4
FGF6	2251	NP_066276	4756	fibroblast growth factor 6
FGF7	2252	NP_002000	2528	hypothetical LOC100132771 fibroblast growth factor 7 (keratinocyte growth factor)
FGFR1	2260	NP_056934	4124	fibroblast growth factor receptor 1
FGFR1	2260	NP_075593	4973	fibroblast growth factor receptor 1
FGFR1	2260	NP_075594	4974	fibroblast growth factor receptor 1
FGFR1	2260	NP_001167534.1	4975	fibroblast growth factor receptor 1
FGFR1	2260	NP_001167534.1	4976	fibroblast growth factor receptor 1
FGFR1	2260	NP_075598	4977	fibroblast growth factor receptor 1
FGFR1	2260	NP_075599	4978	fibroblast growth factor receptor 1
FGFR2	2263	NP_000132	39	fibroblast growth factor receptor 2
FGFR2	2263	NP_001138385	1792	fibroblast growth factor receptor 2
FGFR2	2263	NP_001138386	1793	fibroblast growth factor receptor 2
FGFR2	2263	NP_001138387	1794	fibroblast growth factor receptor 2
FGFR2	2263	NP_001138388	1795	fibroblast growth factor receptor 2
FGFR2	2263	NP_001138389	1796	fibroblast growth factor receptor 2
FGFR2	2263	NP_001138390	1797	fibroblast growth factor receptor 2
FGFR2	2263	NP_001138391	1798	fibroblast growth factor receptor 2
FGFR2	2263	NP_075259	4967	fibroblast growth factor receptor 2
FGFR3	2261	NP_000133	40	fibroblast growth factor receptor 3
FGFR3	2261	NP_001156685	2048	fibroblast growth factor receptor 3
FGFR3	2261	NP_075254	4966	fibroblast growth factor receptor 3
FGFR4	2264	NP_002002	2529	fibroblast growth factor receptor 4
FGFR4	2264	NP_075252	4965	fibroblast growth factor receptor 4
FGFR4	2264	NP_998812	7045	fibroblast growth factor receptor 4
FGFRL1	53834	NP_001004356	430	fibroblast growth factor receptor-like 1
FGFRL1	53834	NP_001004358	431	fibroblast growth factor receptor-like 1
FGFRL1	53834	NP_068742	4848	fibroblast growth factor receptor-like 1
FGG	2266	NP_000500	144	fibrinogen gamma chain
FGG	2266	NP_068656	4838	fibrinogen gamma chain
FIBCD1	84929	NP_001138578	1810	fibrinogen C domain containing 1
FIBCD1	84929	NP_116232	5417	fibrinogen C domain containing 1
FICD	11153	NP_009007	3628	FIC domain containing
FIG4	9896	NP_055660	3997	FIG4Homolog (S. cerevisiae)
FIS1	51024	NP_057152	4160	fission 1 (mitochondrial outer membrane)Homolog (S. cerevisiae)
FITM1	161247	NP_981947	6932	fat storage-inducing transmembrane protein 1
FITM2	128486	NP_001073941	1028	fat storage-inducing transmembrane protein 2
FJX1	24147	NP_055159	3921	four jointed box 1 (Drosophila)
FKBP11	51303	NP_001137253	1715	FK506 binding protein 11, 19 kDa
FKBP11	51303	NP_001137254	1716	FK506 binding protein 11, 19 kDa
FKBP11	51303	NP_057678	4255	FK506 binding protein 11, 19 kDa
FKBP1A	2280	NP_000792	259	FK506 binding protein 1A, 12kDa
FKBP1A	2280	NP_463460	5563	FK506 binding protein 1A, 12kDa
FKBP7	51661	NP_001128684	1538	FK506 binding protein 7
FKBP7	51661	NP_851939	6594	FK506 binding protein 7
FKBP8	23770	NP_036313	3716	FK506 binding protein 8, 38kDa
FKRP	79147	NP_001034974	815	fukutin related protein
FKRP	79147	NP_077277	5018	fukutin related protein
FKTN	2218	NP_001073270	993	fukutin
FKTN	2218	NP_006722	3553	fukutin
FLOT2	2319	NP_004466.2	7359	flotillin 2
FLRT1	23769	NP_037412	3795	fibronectin leucine rich transmembrane protein 1
FLRT2	23768	NP_037363	3787	fibronectin leucine rich transmembrane protein 2

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
FLRT3	23767	NP_037413	3796	fibronectin leucine rich transmembrane protein 3
FLRT3	23767	NP_938205	6787	fibronectin leucine rich transmembrane protein 3
FLT3	2322	NP_004110	2967	fms-related tyrosine kinase 3
FLT3LG	2323	NP_001450	2377	fms-related tyrosine kinase 3 ligand
FLT4	2324	NP_002011	2530	fms-related tyrosine kinase 4
FLT4	2324	NP_891555	6692	fms-related tyrosine kinase 4
FLVCR1	28982	NP_054772	3864	feline leukemia virus subgroup C cellular receptor 1
FLVCR2	55640	NP_060261	4344	feline leukemia virus subgroup C cellular receptor family, member 2
FMNL1	752	NP_005883	3395	formin-like 1
FMO2	2327	NP_001451	2378	flavin containing monooxygenase 2 (non-functional)
FMO3	2328	NP_001002294	402	flavin containing monooxygenase 3
FMO3	2328	NP_008825	3591	flavin containing monooxygenase 3
FMO4	2329	NP_002013	2531	flavin containing monooxygenase 4
FMO5	2330	NP_001138301	1775	flavin containing monooxygenase 5
FMO5	2330	NP_001138302	1776	flavin containing monooxygenase 5
FMO5	2330	NP_001452	2379	flavin containing monooxygenase 5
FMR1NB	158521	NP_689791	6083	fragile X mental retardation 1 neighbor
FNDC3A	22862	NP_001073141	991	fibronectin type III domain containing 3A
FNDC3A	22862	NP_055738	4014	fibronectin type III domain containing 3A
FNDC3B	64778	NP_001128567	1520	fibronectin type III domain containing 3B
FNDC3B	64778	NP_073600	4944	fibronectin type III domain containing 3B
FNDC4	64838	NP_073734	4950	fibronectin type III domain containing 4
FNDC5	252995	NP_001165411	2298	fibronectin type III domain containing 5
FNDC5	252995	NP_001165412	2299	fibronectin type III domain containing 5
FNDC5	252995	NP_715637	6224	fibronectin type III domain containing 5
FOLH1	2346	NP_001014986	631	folateHydrolase (prostate-specific membrane antigen) 1
FOLH1	2346	NP_004467	3057	folateHydrolase (prostate-specific membrane antigen) 1
FOLR1	2348	NP_000793	260	folate receptor 1 (adult)
FOLR1	2348	NP_057936	4267	folate receptor 1 (adult)
FOLR1	2348	NP_057937	4268	folate receptor 1 (adult)
FOLR1	2348	NP_057941	4269	folate receptor 1 (adult)
FOLR1	2348	NP_057941	4270	folate receptor 1 (adult)
FOLR1	2348	NP_057941	4271	folate receptor 1 (adult)
FOLR2	2350	NP_000794	261	folate receptor 2 (fetal)
FOLR2	2350	NP_001107006	1261	folate receptor 2 (fetal)
FOLR2	2350	NP_001107007	1262	folate receptor 2 (fetal)
FOLR2	2350	NP_001107008	1263	folate receptor 2 (fetal)
FPR1	2357	NP_002020	2532	formyl peptide receptor 1
FPR2	2358	NP_001005738	469	formyl peptide receptor 2
FPR2	2358	NP_001453	2380	formyl peptide receptor 2
FPR3	2359	NP_002021	2533	formyl peptide receptor 3
FRAS1	80144	NP_001159605	2152	Fraser syndrome 1
FRAS1	80144	NP_079350	5116	Fraser syndrome 1
FREM1	158326	NP_659403	5907	FRAS1 related extracellular matrix 1
FREM2	341640	NP_997244	7016	FRAS1 related extracellular matrix protein 2
FRMD3	257019	NP_777598	6414	FERM domain containing 3
FRMD5	84978	NP_116281	5421	FERM domain containing 5
FRRS1L	23732	NP_055149	3917	chromosome 9 open reading frame 4
FSHR	2492	NP_000136	41	follicle stimulatingHormone receptor
FSHR	2492	NP_852111	6598	follicle stimulatingHormone receptor
FTHL17	53940	NP_114100	5262	ferritin,Heavy polypeptide-like 17
FURIN	5045	NP_002560	2685	furin (paired basic amino acid cleaving enzyme)
FUT1	2523	NP_000139	42	fucosyltransferase 1 (galactoside 2-alpha-L-fucosyltransferase,H blood group)
FUT10	84750	NP_116053	5386	fucosyltransferase 10 (alpha (1,3) fucosyltransferase)
FUT2	2524	NP_000502	145	fucosyltransferase 2 (secretor status included)
FUT2	2524	NP_001091107	1110	fucosyltransferase 2 (secretor status included)
FUT3	2525	NP_000140	43	fucosyltransferase 3 (galactoside 3(4)-L-fucosyltransferase, Lewis blood group)
FUT3	2525	NP_001091108	1111	fucosyltransferase 3 (galactoside 3(4)-L-fucosyltransferase, Lewis blood group)
FUT3	2525	NP_001091109	1112	fucosyltransferase 3 (galactoside 3(4)-L-fucosyltransferase, Lewis blood group)
FUT3	2525	NP_001091110	1113	fucosyltransferase 3 (galactoside 3(4)-L-fucosyltransferase, Lewis blood group)
FUT4	2526	NP_002024	2534	fucosyltransferase 4 (alpha (1,3) fucosyltransferase, myeloid-specific)
FUT5	2527	NP_002025	2535	fucosyltransferase 5 (alpha (1,3) fucosyltransferase)
FUT6	2528	NP_000141	44	fucosyltransferase 6 (alpha (1,3) fucosyltransferase)
FUT6	2528	NP_001035791	881	fucosyltransferase 6 (alpha (1,3) fucosyltransferase)
FUT7	2529	NP_004470	3058	fucosyltransferase 7 (alpha (1,3) fucosyltransferase)
FUT8	2530	NP_004471	3059	fucosyltransferase 8 (alpha (1,6) fucosyltransferase)
FUT8	2530	NP_835367	6521	fucosyltransferase 8 (alpha (1,6) fucosyltransferase)
FUT8	2530	NP_835368	6522	fucosyltransferase 8 (alpha (1,6) fucosyltransferase)
FUT8	2530	NP_835369	6523	fucosyltransferase 8 (alpha (1,6) fucosyltransferase)
FUT8	2530	NP_835370	6524	fucosyltransferase 8 (alpha (1,6) fucosyltransferase)
FUT9	10690	NP_006572	3521	fucosyltransferase 9 (alpha (1,3) fucosyltransferase)
FXR1	8087	NP_001013456	613	fragile X mental retardation, autosomalHomolog 1
FXR1	8087	NP_001013457	614	fragile X mental retardation, autosomalHomolog 1
FXR1	8087	NP_005078	3210	fragile X mental retardation, autosomalHomolog 1
FXYD1	5348	NP_005022	3200	FXYD domain containing ion transport regulator 1
FXYD1	5348	NP_068702	4839	FXYD domain containing ion transport regulator 1
FXYD3	5349	NP_001129479	1583	FXYD domain containing ion transport regulator 3

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
FXYD3	5349	NP_001129480	1584	FXYD domain containing ion transport regulator 3
FXYD3	5349	NP_001129481	1585	FXYD domain containing ion transport regulator 3
FXYD3	5349	NP_001129482	1586	FXYD domain containing ion transport regulator 3
FXYD3	5349	NP_001129483	1587	FXYD domain containing ion transport regulator 3
FXYD3	5349	NP_001129484	1588	FXYD domain containing ion transport regulator 3
FXYD3	5349	NP_005962	3408	FXYD domain containing ion transport regulator 3
FXYD3	5349	NP_068710	4844	FXYD domain containing ion transport regulator 3
FXYD4	53828	NP_775183	6336	FXYD domain containing ion transport regulator 4
FXYD5	53827	NP_001158077	2086	FXYD domain containing ion transport regulator 5
FXYD5	53827	NP_054883	3874	FXYD domain containing ion transport regulator 5
FXYD5	53827	NP_659003	5904	FXYD domain containing ion transport regulator 5
FXYD6	53826	NP_001158303	2117	FXYD domain containing ion transport regulator 6
FXYD6	53826	NP_001158304	2118	FXYD domain containing ion transport regulator 6
FXYD6	53826	NP_001158308	2119	FXYD domain containing ion transport regulator 6
FXYD6	53826	NP_001158309	2120	FXYD domain containing ion transport regulator 6
FXYD6	53826	NP_071286	4863	FXYD domain containing ion transport regulator 6
FXYD7	53822	NP_071289	4864	FXYD domain containing ion transport regulator 7
FZD1	8321	NP_003496.1	7259	frizzled class receptor 1
FZD10	11211	NP_009128	3647	frizzledHomolog 10 (Drosophila)
FZD2	2535	NP_001457	2382	frizzledHomolog 2 (Drosophila)
FZD3	7976	NP_059108	4282	frizzledHomolog 3 (Drosophila)
FZD4	8322	NP_036325	3717	frizzledHomolog 4 (Drosophila)
FZD5	7855	NP_003459	2837	frizzledHomolog 5 (Drosophila)
FZD6	8323	NP_001158087	2087	frizzledHomolog 6 (Drosophila)
FZD6	8323	NP_001158088	2088	frizzledHomolog 6 (Drosophila)
FZD6	8323	NP_003497	2844	frizzledHomolog 6 (Drosophila)
FZD7	8324	NP_003498	2845	frizzledHomolog 7 (Drosophila)
FZD8	8325	NP_114072	5252	frizzledHomolog 8 (Drosophila)
FZD9	8326	NP_003499	2846	frizzledHomolog 9 (Drosophila)
G6PC	2538	NP_000142	45	glucose-6-phosphatase, catalytic subunit
G6PC2	57818	NP_001075155	1055	glucose-6-phosphatase, catalytic, 2
G6PC2	57818	NP_066999	4787	glucose-6-phosphatase, catalytic, 2
G6PC3	92579	NP_612396	5739	glucose 6 phosphatase, catalytic, 3
GAA	2548	NP_000143	46	glucosidase, alpha acid
GAA	2548	NP_001073271	994	glucosidase, alpha acid
GAA	2548	NP_001073272	995	glucosidase, alpha acid
GABBR1	2550	NP_001461	2384	gamma-aminobutyric acid (GABA) B receptor, 1
GABBR1	2550	NP_068703	4840	gamma-aminobutyric acid (GABA) B receptor, 1
GABBR1	2550	NP_068704	4841	gamma-aminobutyric acid (GABA) B receptor, 1
GABBR1	2550	NP_001305982.1	4842	gamma-aminobutyric acid (GABA) B receptor, 1
GABBR2	9568	NP_005449	3288	gamma-aminobutyric acid (GABA) B receptor, 2
GABRA1	2554	NP_000797	262	gamma-aminobutyric acid (GABA) A receptor, alpha 1
GABRA1	2554	NP_001121115	1335	gamma-aminobutyric acid (GABA) A receptor, alpha 1
GABRA1	2554	NP_001121116	1336	gamma-aminobutyric acid (GABA) A receptor, alpha 1
GABRA1	2554	NP_001121117	1337	gamma-aminobutyric acid (GABA) A receptor, alpha 1
GABRA1	2554	NP_000797.2	1338	gamma-aminobutyric acid (GABA) A receptor, alpha 1
GABRA1	2554	NP_000797.2	1339	gamma-aminobutyric acid (GABA) A receptor, alpha 1
GABRA1	2554	NP_001121120	1340	gamma-aminobutyric acid (GABA) A receptor, alpha 1
GABRA2	2555	NP_000798	263	gamma-aminobutyric acid (GABA) A receptor, alpha 2
GABRA2	2555	NP_001107647	1269	gamma-aminobutyric acid (GABA) A receptor, alpha 2
GABRA3	2556	NP_000799	264	gamma-aminobutyric acid (GABA) A receptor, alpha 3
GABRA4	2557	NP_000800	265	gamma-aminobutyric acid (GABA) A receptor, alpha 4
GABRA5	2558	NP_000801	266	gamma-aminobutyric acid (GABA) A receptor, alpha 5
GABRA5	2558	NP_001158509	2125	gamma-aminobutyric acid (GABA) A receptor, alpha 5
GABRA6	2559	NP_000802	267	gamma-aminobutyric acid (GABA) A receptor, alpha 6
GABRB1	2560	NP_000803	268	gamma-aminobutyric acid (GABA) A receptor, beta 1
GABRB2	2561	NP_000804	269	gamma-aminobutyric acid (GABA) A receptor, beta 2
GABRB2	2561	NP_068711	4845	gamma-aminobutyric acid (GABA) A receptor, beta 2
GABRB3	2562	NP_000805	270	gamma-aminobutyric acid (GABA) A receptor, beta 3
GABRB3	2562	NP_068712	4846	gamma-aminobutyric acid (GABA) A receptor, beta 3
GABRD	2563	NP_000806	271	gamma-aminobutyric acid (GABA) A receptor, delta
GABRE	2564	NP_004952	3178	gamma-aminobutyric acid (GABA) A receptor, epsilon
GABRG1	2565	NP_775807	6354	gamma-aminobutyric acid (GABA) A receptor, gamma 1
GABRG2	2566	NP_000807	272	gamma-aminobutyric acid (GABA) A receptor, gamma 2
GABRG2	2566	NP_944493	6842	gamma-aminobutyric acid (GABA) A receptor, gamma 2
GABRG2	2566	NP_944494	6843	gamma-aminobutyric acid (GABA) A receptor, gamma 2
GABRG3	2567	NP_150092	5471	gamma-aminobutyric acid (GABA) A receptor, gamma 3
GABRP	2568	NP_055026	3883	gamma-aminobutyric acid (GABA) A receptor, pi
GABRQ	55879	NP_061028	4468	gamma-aminobutyric acid (GABA) receptor, theta
GABRR1	2569	NP_002033	2539	gamma-aminobutyric acid (GABA) receptor, rho 1
GABRR2	2570	NP_002034	2540	gamma-aminobutyric acid (GABA) receptor, rho 2
GABRR3	200959	NP_001099050	1231	gamma-aminobutyric acid (GABA) receptor, rho 3
GAGE1	2543	NP_001035753.1	7360	G antigen 1
GAL	51083	NP_057057	4145	galanin prepropeptide
GAL3ST1	9514	NP_001305032.1	7460	galactose-3-O-sulfotransferase 1
GAL3ST2	64090	NP_071417	4880	galactose-3-O-sulfotransferase 2
GAL3ST4	79690	NP_078913	5061	galactose-3-O-sulfotransferase 4

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
GALC	2581	NP_000144	47	galactosylceramidase
GALC	2581	NP_000144.2	776	galactosylceramidase
GALNS	2588	NP_000503	146	galactosamine (N-acetyl)-6-sulfate sulfatase
GALNT1	2589	NP_065207	4670	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 13 (GalNAc-T13)
GALNT1	2589	NP_443149	5539	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 13 (GalNAc-T13)
GALNT10	55568	NP_938080.1	4303	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 10 (GalNAc-T10)
GALNT10	55568	NP_938080	6767	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 10 (GalNAc-T10)
GALNT11	63917	NP_071370	4875	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 11 (GalNAc-T11)
GALNT12	79695	NP_078918	5063	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 12 (GalNAc-T12)
GALNT13	114805	NP_001288556.1	4669	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 13 (GalNAc-T13)
GALNT13	114805	NP_065207.2	5538	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 13 (GalNAc-T13)
GALNT14	79623	NP_078848	5050	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 14 (GalNAc-T14)
GALNT15	117248	NP_473451	5574	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase-like 2
GALNT16	57452	NP_001161840	2226	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase-like 1
GALNT16	57452	NP_065743	4700	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase-like 1
GALNT18	374378	NP_940918	6807	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase-like 4
GALNT2	2590	NP_004472	3060	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 2 (GalNAc-T2)
GALNT3	2591	NP_004473	3061	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 3 (GalNAc-T3)
GALNT5	11227	NP_055383	3957	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 5 (GalNAc-T5)
GALNT6	11226	NP_009141	3648	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 6 (GalNAc-T6)
GALNT7	51809	NP_059119	4287	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 7 (GalNAc-T7)
GALNT8	26290	NP_059113	4285	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 8 (GalNAc-T8)
GALNT9	50614	NP_001116108	1284	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 9 (GalNAc-T9)
GALNT9	50614	NP_068580	4834	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 9 (GalNAc-T9)
GALNTL5	168391	NP_660335	5955	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase-like 5
GALR1	2587	NP_001471	2386	galanin receptor 1
GALR2	8811	NP_003848	2919	galanin receptor 2
GALR3	8484	NP_003605	2854	galanin receptor 3
GANAB	23193	NP_938148	6770	glucosidase, alpha neutral AB
GANAB	23193	NP_938149	6771	glucosidase, alpha neutral AB
GAPT	202309	NP_689900	6099	GRB2-binding adaptor protein, transmembrane
GBGT1	26301	NP_068836	4861	globoside alpha-1,3-N-acetylgalactosaminyltransferase 1
GCG	2641	NP_002045	2541	glucagon
GCGR	2642	NP_000151	48	glucagon receptor
GCLC	2729	NP_001489	2391	glutamate-cysteine ligase, catalytic subunit
GCNT1	2650	NP_001091102	1106	glucosaminyl (N-acetyl) transferase 1, core 2 (beta-1,6-N-acetylglucosaminyltransferase)
GCNT1	2650	NP_001091103	1107	glucosaminyl (N-acetyl) transferase 1, core 2 (beta-1,6-N-acetylglucosaminyltransferase)
GCNT1	2650	NP_001091104	1108	glucosaminyl (N-acetyl) transferase 1, core 2 (beta-1,6-N-acetylglucosaminyltransferase)
GCNT1	2650	NP_001091105	1109	glucosaminyl (N-acetyl) transferase 1, core 2 (beta-1,6-N-acetylglucosaminyltransferase)
GCNT1	2650	NP_001481	2387	glucosaminyl (N-acetyl) transferase 1, core 2 (beta-1,6-N-acetylglucosaminyltransferase)
GCNT2	2651	NP_001482	2388	glucosaminyl (N-acetyl) transferase 2, I-branching enzyme (I blood group)
GCNT2	2651	NP_663624	5965	glucosaminyl (N-acetyl) transferase 2, I-branching enzyme (I blood group)
GCNT2	2651	NP_663630	5968	glucosaminyl (N-acetyl) transferase 2, I-branching enzyme (I blood group)
GCNT3	9245	NP_004742	3116	glucosaminyl (N-acetyl) transferase 3, mucin type
GCNT4	51301	NP_057675	4253	glucosaminyl (N-acetyl) transferase 4, core 2 (beta-1,6-N-acetylglucosaminyltransferase)
GDAP1	54332	NP_001035808	884	ganglioside-induced differentiation-associated protein 1
GDAP1	54332	NP_061845	4541	ganglioside-induced differentiation-associated protein 1
GDAP1L1	78997	NP_076939	4998	ganglioside-induced differentiation-associated protein 1-like 1
GDE1	51573	NP_057725	4265	glycerophosphodiester phosphodiesterase 1
GDF9	2661	NP_005251	3241	growth differentiation factor 9
GDNF	2668	NP_000505	147	glial cell derived neurotrophic factor
GDNF	2668	NP_954701	6866	glial cell derived neurotrophic factor
GDNF	2668	NP_954704	6867	glial cell derived neurotrophic factor
GDPD1	284161	NP_001159465	2139	glycerophosphodiester phosphodiesterase domain containing 1
GDPD1	284161	NP_001159466	2140	glycerophosphodiester phosphodiesterase domain containing 1
GDPD1	284161	NP_872375	6662	glycerophosphodiester phosphodiesterase domain containing 1
GDPD2	54857	NP_001164662	2275	glycerophosphodiester phosphodiesterase domain containing 2
GDPD2	54857	NP_001164663	2276	glycerophosphodiester phosphodiesterase domain containing 2
GDPD2	54857	NP_001164664	2277	glycerophosphodiester phosphodiesterase domain containing 2
GDPD2	54857	NP_060181	4327	glycerophosphodiester phosphodiesterase domain containing 2
GDPD3	79153	NP_077283	5020	glycerophosphodiester phosphodiesterase domain containing 3
GDPD5	81544	NP_110419	5177	glycerophosphodiester phosphodiesterase domain containing 5
GGCX	2677	NP_000812	273	gamma-glutamyl carboxylase
GGCX	2677	NP_001135741	1627	gamma-glutamyl carboxylase
GGT5	2687	NP_001093251	1173	gamma-glutamyltransferase 5
GGT5	2687	NP_001093252	1174	gamma-glutamyltransferase 5
GGT5	2687	NP_004112	2968	gamma-glutamyltransferase 5
GGT7	2686	NP_821158	6515	gamma-glutamyltransferase 7
GGTLC1	92086	NP_842563	6534	gamma-glutamyltransferase light chain 1
GGTLC1	92086	NP_842564	6535	gamma-glutamyltransferase light chain 1
GHITM	27069	NP_055209	3930	growthHormone inducible transmembrane protein
GHR	2690	NP_000154	49	growthHormone receptor
GHRHR	2692	NP_000814	274	growthHormone releasingHormone receptor
GHRHR	2692	NP_000814.2	564	growthHormone releasingHormone receptor
GHRL	51738	NP_001128413	1508	ghrelin/obestatin prepropeptide
GHRL	51738	NP_001128416	1509	ghrelin/obestatin prepropeptide
GHRL	51738	NP_001128417	1510	ghrelin/obestatin prepropeptide

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
GHRL	51738	NP_001128418	1511	ghrelin/obestatin prepropeptide
GHRL	51738	NP_057446	4210	ghrelin/obestatin prepropeptide
GHSR	2693	NP_004113	2969	growthHormone secretagogue receptor
GHSR	2693	NP_940799	6794	growthHormone secretagogue receptor
GIMAP1	170575	NP_570115	5659	GTPase, IMAP family member 1
GIMAP2	26157	NP_056475	4110	GTPase, IMAP family member 2
GINM1	116254	NP_620140	5793	chromosome 6 open reading frame 72
GIPR	2696	NP_000155	50	gastric inhibitory polypeptide receptor
GJA1	2697	NP_000156	51	gap junction protein, alpha 1, 43kDa
GJA10	84694	NP_115991	5377	gap junction protein, alpha 10, 62kDa
GJA3	2700	NP_068773	4854	gap junction protein, alpha 3, 46kDa
GJA4	2701	NP_002051	2542	gap junction protein, alpha 4, 37kDa
GJA5	2702	NP_005257	3242	gap junction protein, alpha 5, 40kDa
GJA5	2702	NP_859054	6622	gap junction protein, alpha 5, 40kDa
GJA8	2703	NP_005258	3243	gap junction protein, alpha 8, 50kDa
GJB1	2705	NP_000157	52	gap junction protein, beta 1, 32kDa
GJB1	2705	NP_001091111	1114	gap junction protein, beta 1, 32kDa
GJB2	2706	NP_003995	2946	gap junction protein, beta 2, 26kDa
GJB3	2707	NP_001005752	470	gap junction protein, beta 3, 31kDa
GJB3	2707	NP_076872	4992	gap junction protein, beta 3, 31kDa
GJB4	127534	NP_694944	6159	gap junction protein, beta 4, 30.3kDa
GJB5	2709	NP_005259	3244	gap junction protein, beta 5, 31.1kDa
GJB6	10804	NP_001103689	1238	gap junction protein, beta 6, 30kDa
GJB6	10804	NP_001103690	1239	gap junction protein, beta 6, 30kDa
GJB6	10804	NP_001103691	1240	gap junction protein, beta 6, 30kDa
GJB6	10804	NP_006774	3563	gap junction protein, beta 6, 30kDa
GJC1	10052	NP_001073852	1017	gap junction protein, gamma 1, 45kDa
GJC1	10052	NP_005488	3295	gap junction protein, gamma 1, 45kDa
GJC2	57165	NP_065168	4658	gap junction protein, gamma 2, 47kDa
GJD2	57369	NP_065711	4693	gap junction protein, delta 2, 36kDa
GJD3	125111	NP_689343	6018	gap junction protein, delta 3, 31.9kDa
GJD4	219770	NP_699199	6185	gap junction protein, delta 4, 40.1kDa
GKN1	56287	NP_062563	4576	gastrokine 1
GLB1L2	89944	NP_612351	5732	galactosidase, beta 1-like 2
GLCCI1	113263	NP_612435	5748	glucocorticoid induced transcript 1
GLCE	26035	NP_056369	4101	glucuronic acid epimerase
GLDN	342035	NP_861454	6636	gliomedin
GLG1	2734	NP_001139138	1874	golgi apparatus protein 1
GLG1	2734	NP_001139139	1875	golgi apparatus protein 1
GLG1	2734	NP_036333	3721	golgi apparatus protein 1
GLIPR1	11010	NP_006842	3580	GLI pathogenesis-related 1
GLMP	112770	NP_653181	5866	chromosome 1 open reading frame 85
GLP1R	2740	NP_002053	2543	glucagon-like peptide 1 receptor
GLP2R	9340	NP_004237	2997	glucagon-like peptide 2 receptor
GLRA1	2741	NP_000162	53	glycine receptor, alpha 1
GLRA1	2741	NP_001139512	1899	glycine receptor, alpha 1
GLRA2	2742	NP_001112357	1281	glycine receptor, alpha 2
GLRA2	2742	NP_001112358	1282	glycine receptor, alpha 2
GLRA2	2742	NP_001165413	2300	glycine receptor, alpha 2
GLRA2	2742	NP_002054	2544	glycine receptor, alpha 2
GLRA3	8001	NP_001036008	904	glycine receptor, alpha 3
GLRA3	8001	NP_006520	3508	glycine receptor, alpha 3
GLRB	2743	NP_000815	275	glycine receptor, beta
GLRB	2743	NP_001159532	2143	glycine receptor, beta
GLRB	2743	NP_001159533	2144	glycine receptor, beta
GLT8D1	55830	NP_001010983	583	glycosyltransferase 8 domain containing 1
GLT8D1	55830	NP_060916	4452	glycosyltransferase 8 domain containing 1
GLT8D1	55830	NP_690909	6139	glycosyltransferase 8 domain containing 1
GLT8D2	83468	NP_112592	5219	glycosyltransferase 8 domain containing 2
GLTPD2	388323	NP_001014985	630	glycolipid transfer protein domain containing 2
GMFB	2764	NP_004115	2970	glia maturation factor, beta
GNB2	2783	NP_005264	3245	guanine nucleotide binding protein (G protein), beta polypeptide 2
GNL2	29889	NP_037417	3797	guanine nucleotide binding protein-like 2 (nucleolar)
GNPTAB	79158	NP_077288	5023	N-acetylglucosamine-1-phosphate transferase, alpha and beta subunits
GNRHR	2798	NP_000397	119	gonadotropin-releasingHormone receptor
GNRHR	2798	NP_001012781	607	gonadotropin-releasingHormone receptor
GNS	2799	NP_002067	2545	glucosamine (N-acetyl)-6-sulfatase
GOLGA5	9950	NP_005104	3213	golgi autoantigen, golgin subfamily a, 5
GOLGB1	2804	NP_004478	3063	golgin B1, golgi integral membrane protein
GOLIM4	27333	NP_055313	3947	golgi integral membrane protein 4
GOLM1	51280	NP_057632	4241	golgi membrane protein 1
GOLM1	51280	NP_808800	6502	golgi membrane protein 1
GOLT1A	127845	NP_940849	6795	golgi transport 1Homolog A (S. cerevisiae)
GOLT1B	51026	NP_057156	4161	golgi transport 1Homolog B (S. cerevisiae)
GOSR1	9527	NP_001007025	498	golgi SNAP receptor complex member 1
GOSR1	9527	NP_001007026	499	golgi SNAP receptor complex member 1
GOSR1	9527	NP_004862	3155	golgi SNAP receptor complex member 1

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
GOSR2	9570	NP_001012529	598	golgi SNAP receptor complex member 2
GOSR2	9570	NP_004278	3008	golgi SNAP receptor complex member 2
GOSR2	9570	NP_473363	5566	golgi SNAP receptor complex member 2
GP1BA	2811	NP_000164	54	glycoprotein Ib (platelet), alpha polypeptide
GP2	2813	NP_001007241	509	glycoprotein 2 (zymogen granule membrane)
GP2	2813	NP_001007242	510	glycoprotein 2 (zymogen granule membrane)
GP2	2813	NP_001007243	511	glycoprotein 2 (zymogen granule membrane)
GP2	2813	NP_001493	2392	glycoprotein 2 (zymogen granule membrane)
GP5	2814	NP_004479	3064	glycoprotein V (platelet)
GP6	51206	NP_001077368	1080	glycoprotein VI (platelet)
GP6	51206	NP_057447	4211	glycoprotein VI (platelet)
GP9	2815	NP_000165	55	glycoprotein IX (platelet)
GPA33	10223	NP_005805	3374	glycoprotein A33 (transmembrane)
GPAA1	8733	NP_003792	2897	glycosylphosphatidylinositol anchor attachment protein 1Homolog (yeast)
GPATCH2L	55668	NP_060396	4372	chromosome 14 open reading frame 118
GPATCH2L	55668	NP_060442	4376	chromosome 14 open reading frame 118
GPBAR1	151306	NP_001070659	953	G protein-coupled bile acid receptor 1
GPBAR1	151306	NP_001070662	954	G protein-coupled bile acid receptor 1
GPBAR1	151306	NP_733800	6248	G protein-coupled bile acid receptor 1
GPC3	2719	NP_001158089	2089	glypican 3
GPC3	2719	NP_001158090	2090	glypican 3
GPC3	2719	NP_001158091	2091	glypican 3
GPC3	2719	NP_004475	3062	glypican 3
GPD2	2820	NP_000399	120	glycerol-3-phosphate dehydrogenase 2 (mitochondrial)
GPD2	2820	NP_001076581	1066	glycerol-3-phosphate dehydrogenase 2 (mitochondrial)
GPER1	2852	NP_001035055	819	G protein-coupled estrogen receptor 1
GPER1	2852	NP_001091671	1118	G protein-coupled estrogen receptor 1
GPER1	2852	NP_001496	2394	G protein-coupled estrogen receptor 1
GPHB5	122876	NP_660154	5932	glycoproteinHormone beta 5
GPM6A	2823	NP_005268	3246	glycoprotein M6A
GPM6A	2823	NP_963885	6912	glycoprotein M6A
GPM6A	2823	NP_963886	6913	glycoprotein M6A
GPM6B	2824	NP_001001994	382	glycoprotein M6B
GPM6B	2824	NP_001001995	383	glycoprotein M6B
GPM6B	2824	NP_001001996	384	glycoprotein M6B
GPM6B	2824	NP_005269	3247	glycoprotein M6B
GPNMB	10457	NP_001005340	454	glycoprotein (transmembrane) nmb
GPNMB	10457	NP_002501	2664	glycoprotein (transmembrane) nmb
GPR1	2825	NP_001091669	1116	G protein-coupled receptor 1
GPR1	2825	NP_005270	3248	G protein-coupled receptor 1
GPR101	83550	NP_473362	5565	G protein-coupled receptor 101
GPR107	57720	NP_001130029	1612	G protein-coupled receptor 107
GPR107	57720	NP_001130030	1613	G protein-coupled receptor 107
GPR107	57720	NP_066011	4747	G protein-coupled receptor 107
GPR108	56927	NP_001073921	1024	G protein-coupled receptor 108
GPR108	56927	NP_001073921.1	4611	G protein-coupled receptor 108
GPR12	2835	NP_005279	3255	G protein-coupled receptor 12
GPR135	64582	NP_072093	4932	G protein-coupled receptor 135
GPR137	56834	NP_001164197	2252	G protein-coupled receptor 137
GPR137	56834	NP_001164351	2258	G protein-coupled receptor 137
GPR137	56834	NP_001164352	2259	G protein-coupled receptor 137
GPR137	56834	NP_064540	4605	G protein-coupled receptor 137
GPR137B	7107	NP_003263	2812	G protein-coupled receptor 137B
GPR137C	283554	NP_001093122	1164	G protein-coupled receptor 137C
GPR143	4935	NP_000264	79	G protein-coupled receptor 143
GPR146	115330	NP_612454	5754	G protein-coupled receptor 146
GPR15	2838	NP_005281	3256	G protein-coupled receptor 15
GPR153	387509	NP_997253	7017	G protein-coupled receptor 153
GPR155	151556	NP_001028217	751	G protein-coupled receptor 155
GPR155	151556	NP_689742	6076	G protein-coupled receptor 155
GPR156	165829	NP_001161743	2213	G protein-coupled receptor 156
GPR156	165829	NP_694547	6147	G protein-coupled receptor 156
GPR158	57512	NP_065803	4716	G protein-coupled receptor 158
GPR160	26996	NP_055188	3924	G protein-coupled receptor 160
GPR161	23432	NP_722561	6236	G protein-coupled receptor 161
GPR162	27239	NP_055264	3942	G protein-coupled receptor 162
GPR162	27239	NP_062832	4586	G protein-coupled receptor 162
GPR17	2840	NP_001154887	2008	G protein-coupled receptor 17
GPR17	2840	NP_001154888	2009	G protein-coupled receptor 17
GPR17	2840	NP_001154889	2010	G protein-coupled receptor 17
GPR17	2840	NP_005282	3257	G protein-coupled receptor 17
GPR171	29909	NP_037440	3800	G protein-coupled receptor 171
GPR173	54328	NP_061842	4538	G protein-coupled receptor 173
GPR174	84636	NP_115942.1	7302	G protein-coupled receptor 174
GPR176	11245	NP_009154	3654	G protein-coupled receptor 176
GPR179	440435	NP_001004334.3	7447	G protein-coupled receptor 179
GPR18	2841	NP_001091670	1117	G protein-coupled receptor 18

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
GPR18	2841	NP_005283	3258	G protein-coupled receptor 18
GPR180	160897	NP_851320	6577	G protein-coupled receptor 180
GPR182	11318	NP_009195	3664	G protein-coupled receptor 182
GPR183	1880	NP_004942	3176	G protein-coupled receptor 183
GPR19	2842	NP_006134	3446	G protein-coupled receptor 19
GPR20	2843	NP_005284	3259	G protein-coupled receptor 20
GPR21	2844	NP_005285	3260	G protein-coupled receptor 21
GPR22	2845	NP_005286	3261	G protein-coupled receptor 22
GPR25	2848	NP_005289	3264	G protein-coupled receptor 25
GPR26	2849	NP_703143	6190	G protein-coupled receptor 26
GPR27	2850	NP_061844	4540	G protein-coupled receptor 27
GPR3	2827	NP_005272	3249	G protein-coupled receptor 3
GPR31	2853	NP_005290	3265	G protein-coupled receptor 31
GPR32	2854	NP_001497	2395	G protein-coupled receptor 32
GPR34	2857	NP_001091048	1104	G protein-coupled receptor 34
GPR34	2857	NP_005291	3266	G protein-coupled receptor 34
GPR35	2859	NP_005292	3267	G protein-coupled receptor 35
GPR37	2861	NP_005293	3268	G protein-coupled receptor 37 (endothelin receptor type B-like)
GPR37L1	9283	NP_004758	3118	G protein-coupled receptor 37 like 1
GPR39	2863	NP_001499	2397	G protein-coupled receptor 39
GPR4	2828	NP_005273	3250	G protein-coupled receptor 4
GPR45	11250	NP_009158	3655	G protein-coupled receptor 45
GPR50	9248	NP_004215	2993	G protein-coupled receptor 50
GPR52	9293	NP_005675	3342	G protein-coupled receptor 52
GPR55	9290	NP_005674	3341	G protein-coupled receptor 55
GPR6	2830	NP_005275	3252	G protein-coupled receptor 6
GPR61	83873	NP_114142	5271	G protein-coupled receptor 61
GPR62	118442	NP_543141	5639	G protein-coupled receptor 62
GPR63	81491	NP_001137429	1739	G protein-coupled receptor 63
GPR63	81491	NP_110411	5173	G protein-coupled receptor 63
GPR65	8477	NP_003599.2	7328	G protein-coupled receptor 65
GPR68	8111	NP_001171147.1	7420	G protein-coupled receptor 68
GPR75	10936	NP_006785	3564	G protein-coupled receptor 75
GPR78	27201	NP_543009	5633	G protein-coupled receptor 78
GPR82	27197	NP_543007	5631	G protein-coupled receptor 82
GPR83	10888	NP_057624	4239	G protein-coupled receptor 83
GPR84	53831	NP_065103	4635	G protein-coupled receptor 84
GPR85	54329	NP_001139737	1920	G protein-coupled receptor 85
GPR85	54329	NP_001139738	1921	G protein-coupled receptor 85
GPR85	54329	NP_001139739	1922	G protein-coupled receptor 85
GPR85	54329	NP_061843	4539	G protein-coupled receptor 85
GPR87	53836	NP_076404	4979	G protein-coupled receptor 87
GPR88	54112	NP_071332	4868	G protein-coupled receptor 88
GPRC5A	9052	NP_003970	2937	G protein-coupled receptor, family C, group 5, member A
GPRC5B	51704	NP_057319	4189	G protein-coupled receptor, family C, group 5, member B
GPRC5C	55890	NP_061123	4478	G protein-coupled receptor, family C, group 5, member C
GPRC5C	55890	NP_071319	4865	G protein-coupled receptor, family C, group 5, member C
GPRC5D	55507	NP_061124	4479	G protein-coupled receptor, family C, group 5, member D
GPRC6A	222545	NP_683766	6012	G protein-coupled receptor, family C, group 6, member A
GPX7	2882	NP_056511	4114	glutathione peroxidase 7
GPX8	493869	NP_001008398	532	glutathione peroxidase 8 (putative)
GRAMD1B	57476	NP_065767	4707	GRAM domain containing 1B
GRAMD1C	54762	NP_001165576	2307	GRAM domain containing 1C
GRAMD1C	54762	NP_060047	4307	GRAM domain containing 1C
GRAMD2	196996	NP_001012660	599	GRAM domain containing 2
GRAMD3	65983	NP_001139791	1927	GRAM domain containing 3
GRAMD3	65983	NP_001139792	1928	GRAM domain containing 3
GRAMD3	65983	NP_001139793	1929	GRAM domain containing 3
GRAMD3	65983	NP_001139794	1930	GRAM domain containing 3
GRAMD3	65983	NP_076416	4985	GRAM domain containing 3
GRAMD4	23151	NP_001015881	635	TSC22 domain family, member 3 GRAM domain containing 4
GRAMD4	23151	NP_004080	2960	TSC22 domain family, member 3 GRAM domain containing 4
GRAMD4	23151	NP_055939	4036	TSC22 domain family, member 3 GRAM domain containing 4
GRAMD4	23151	NP_932174	6738	TSC22 domain family, member 3 GRAM domain containing 4
GREM1	26585	NP_037504	3812	gremlin 1, cysteine knot superfamily,Homolog (Xenopus laevis)
GRHL1	29841	NP_937825.2	3954	grainyhead-like 1 (Drosophila)
GRHL1	29841	NP_937825	6752	grainyhead-like 1 (Drosophila)
GRIA1	2890	NP_000818	277	glutamate receptor, ionotropic, AMPA 1
GRIA1	2890	NP_001107655	1270	glutamate receptor, ionotropic, AMPA 1
GRIA2	2891	NP_000817	276	glutamate receptor, ionotropic, AMPA 2
GRIA2	2891	NP_001077088	1078	glutamate receptor, ionotropic, AMPA 2
GRIA2	2891	NP_001077089	1079	glutamate receptor, ionotropic, AMPA 2
GRIA3	2892	NP_000819	278	glutamate receptor, ionotrophic, AMPA 3
GRIA3	2892	NP_015564	3686	glutamate receptor, ionotrophic, AMPA 3
GRIA4	2893	NP_000820	279	glutamate receptor, ionotrophic, AMPA 4
GRIA4	2893	NP_001070711	958	glutamate receptor, ionotrophic, AMPA 4
GRIA4	2893	NP_001070712	959	glutamate receptor, ionotrophic, AMPA 4

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
GRIA4	2893	NP_001106283	1253	glutamate receptor, ionotrophic, AMPA 4
GRID1	2894	NP_060021	4305	glutamate receptor, ionotropic, delta 1
GRID2	2895	NP_001501	2398	glutamate receptor, ionotropic, delta 2
GRIK1	2897	NP_000821	280	glutamate receptor, ionotropic, kainate 1
GRIK1	2897	NP_783300	6445	glutamate receptor, ionotropic, kainate 1
GRIK2	2898	NP_001159719	2155	glutamate receptor, ionotropic, kainate 2
GRIK2	2898	NP_068775	4855	glutamate receptor, ionotropic, kainate 2
GRIK2	2898	NP_786944	6460	glutamate receptor, ionotropic, kainate 2
GRIK3	2899	NP_000822	281	glutamate receptor, ionotropic, kainate 3
GRIK4	2900	NP_055434	3965	glutamate receptor, ionotropic, kainate 4
GRIK5	2901	NP_002079	2546	glutamate receptor, ionotropic, kainate 5
GRIN1	2902	NP_000823	282	glutamate receptor, ionotropic, N-methyl D-aspartate 1
GRIN1	2902	NP_015566	3687	glutamate receptor, ionotropic, N-methyl D-aspartate 1
GRIN1	2902	NP_067544	4806	glutamate receptor, ionotropic, N-methyl D-aspartate 1
GRIN2A	2903	NP_000824	283	glutamate receptor, ionotropic, N-methyl D-aspartate 2A
GRIN2A	2903	NP_001127879	1486	glutamate receptor, ionotropic, N-methyl D-aspartate 2A
GRIN2A	2903	NP_001127880	1487	glutamate receptor, ionotropic, N-methyl D-aspartate 2A
GRIN2B	2904	NP_000825	284	glutamate receptor, ionotropic, N-methyl D-aspartate 2B
GRIN2C	2905	NP_000826	285	glutamate receptor, ionotropic, N-methyl D-aspartate 2C
GRIN2D	2906	NP_000827	286	glutamate receptor, ionotropic, N-methyl D-aspartate 2D
GRIN3A	116443	NP_597702	5691	glutamate receptor, ionotropic, N-methyl-D-aspartate 3A
GRIN3B	116444	NP_619635	5769	glutamate receptor, ionotropic, N-methyl-D-aspartate 3B
GRINA	2907	NP_000828	287	glutamate receptor, ionotropic, N-methyl D-aspartate-associated protein 1 (glutamate binding)
GRINA	2907	NP_001009184	559	glutamate receptor, ionotropic, N-methyl D-aspartate-associated protein 1 (glutamate binding)
GRM1	2911	NP_000829	288	glutamate receptor, metabotropic 1
GRM1	2911	NP_001264993.1	1271	glutamate receptor, metabotropic 1
GRM2	2912	NP_000830	289	glutamate receptor, metabotropic 2
GRM2	2912	NP_001123535	1415	glutamate receptor, metabotropic 2
GRM3	2913	NP_000831	290	glutamate receptor, metabotropic 3
GRM4	2914	NP_000832	291	glutamate receptor, metabotropic 4
GRM5	2915	NP_000833	292	glutamate receptor, metabotropic 5
GRM5	2915	NP_001137303	1731	glutamate receptor, metabotropic 5
GRM6	2916	NP_000834	293	glutamate receptor, metabotropic 6
GRM7	2917	NP_000835	294	glutamate receptor, metabotropic 7
GRM7	2917	NP_870989	6642	glutamate receptor, metabotropic 7
GRM8	2918	NP_000836.2	7390	glutamate metabotropic receptor 8
GRPR	2925	NP_005305	3271	gastrin-releasing peptide receptor
GSG1	83445	NP_001074023	1038	germ cell associated 1
GSG1	83445	NP_001074024	1039	germ cell associated 1
GSG1	83445	NP_112579	5215	germ cell associated 1
GSG1	83445	NP_722545	6233	germ cell associated 1
GSG1L	146395	NP_001103233	1236	GSG1-like
GSG1L	146395	NP_653276	5888	GSG1-like
GTF3C3	9330	NP_036218	3700	general transcription factor IIIC, polypeptide 3, 102kDa
GUCA2A	2980	NP_291031	5507	guanylate cyclase activator 2A (guanylin)
GUCA2B	2981	NP_009033	3630	guanylate cyclase activator 2B (uroguanylin)
GUCY2C	2984	NP_004954	3179	guanylate cyclase 2C (heat stable enterotoxin receptor)
GUCY2D	3000	NP_000171	56	guanylate cyclase 2D, membrane (retina-specific)
GUCY2F	2986	NP_001513	2399	guanylate cyclase 2F, retinal
GULP1	51454	NP_057399	4201	GULP, engulfment adaptor PTB domain containing 1
GUSB	2990	NP_000172	57	glucuronidase, beta
GXYLT1	283464	NP_001093120	1163	glycosyltransferase 8 domain containing 3
GXYLT1	283464	NP_775872	6363	glycosyltransferase 8 domain containing 3
GXYLT2	727936	NP_001073862	1018	glycosyltransferase 8 domain containing 4
GYLTL1B	120071	NP_689525	6030	glycosyltransferase-like 1B
GYPA	2993	NP_002090	2547	glycophorin A (MNS blood group)
GYPB	2994	NP_001291311.1	7452	glycophorin B (MNS blood group)
GYPC	2995	NP_001243513.1	7432	glycophorin C (Gerbich blood group)
GYPE	2996	NP_002091	2548	glycophorin E glycophorin B (MNS blood group)
GYPE	2996	NP_002093	2549	glycophorin E glycophorin B (MNS blood group)
GYPE	2996	NP_941391	6825	glycophorin E glycophorin B (MNS blood group)
GZMK	3003	NP_002095	2550	granzyme K (granzyme 3 tryptase II)
HACD1	9200	NP_055056	3894	protein tyrosine phosphatase-like (proline instead of catalytic arginine), member A
HACD4	401494	NP_001010915	576	protein tyrosine phosphatase-like A domain containing 2
HADH	3033	NP_005318	3272	hydroxyacyl-Coenzyme A dehydrogenase
HAMP	57817	NP_066998	4786	hepcidin antimicrobial peptide
HAPLN4	404037	NP_075378	4968	hyaluronan and proteoglycan link protein 4
HAS1	3036	NP_001514	2400	hyaluronan synthase 1
HAS2	3037	NP_005319	3273	hyaluronan synthase 2
HAS3	3038	NP_005320	3274	hyaluronan synthase 3
HAS3	3038	NP_619515	5764	hyaluronan synthase 3
HAVCR1	26762	NP_001166864.1	7417	hepatitis A virus cellular receptor 1
HAVCR2	84868	NP_116171.3	7336	hepatitis A virus cellular receptor 2
HBD	3045	NP_000510	148	hemoglobin, delta
HBEGF	1839	NP_001936	2507	heparin-binding EGF-like growth factor
HBS1L	10767	NP_001138630	1823	HBS1-like (S. cerevisiae)
HBS1L	10767	NP_001138679	1827	HBS1-like (S. cerevisiae)

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
HBS1L	10767	NP_006611	3529	HBS1-like (S. cerevisiae)
HCAR1	27198	NP_115943	5369	G protein-coupled receptor 81
HCN1	348980	NP_066550	4763	hyperpolarization activated cyclic nucleotide-gated potassium channel 1
HCN2	610	NP_001185.3	7377	hyperpolarization activated cyclic nucleotide gated potassium channel 2
HCN3	57657	NP_065948	4738	hyperpolarization activated cyclic nucleotide-gated potassium channel 3
HCN4	10021	NP_005468	3292	hyperpolarization activated cyclic nucleotide-gated potassium channel 4
HCRT	3060	NP_001515	2401	hypocretin (orexin) neuropeptide precursor
HCRTR1	3061	NP_001516	2402	hypocretin (orexin) receptor 1
HCRTR2	3062	NP_001517	2403	hypocretin (orexin) receptor 2
HCST	10870	NP_001007470	518	hematopoietic cell signal transducer
HCST	10870	NP_055081	3902	hematopoietic cell signal transducer
HDAC2	3066	NP_001518	2404	histone deacetylase 2
HEG1	57493	NP_065784	4710	HEGHomolog 1 (zebrafish)
HELZ	9931	NP_055692	4005	helicase with zinc finger
HEPACAM	220296	NP_001032647	778	hepatocyte cell adhesion moleculeHEPACAM opposite strand 1
HEPACAM	220296	NP_689935	6105	hepatocyte cell adhesion moleculeHEPACAM opposite strand 1
HEPACAM2	253012	NP_001034461	790	HEPACAM family member 2
HEPACAM2	253012	NP_937794	6748	HEPACAM family member 2
HEPH	9843	NP_001124332	1442	hephaestin
HEPH	9843	NP_055614	3989	hephaestin
HEPH	9843	NP_620074	5784	hephaestin
HERC4	26091	NP_056416	4106	hect domain and RLD 4
HERC4	26091	NP_071362	4874	hect domain and RLD 4
HERPUD1	9709	NP_001010989	585	homocysteine-inducible, endoplasmic reticulum stress-inducible, ubiquitin-like domain member 1
HERPUD1	9709	NP_001010989.1	586	homocysteine-inducible, endoplasmic reticulum stress-inducible, ubiquitin-like domain member 1
HERPUD1	9709	NP_055500	3973	homocysteine-inducible, endoplasmic reticulum stress-inducible, ubiquitin-like domain member 1
HERPUD2	64224	NP_071768	4908	HERPUD family member 2
HEXB	3074	NP_000512	149	hexosaminidase B (beta polypeptide)
HFE	3077	NP_000401	121	hemochromatosis
HFE	3077	NP_620572	5817	hemochromatosis
HFE	3077	NP_620573	5818	hemochromatosis
HFE	3077	NP_620575	5819	hemochromatosis
HFE	3077	NP_620576	5820	hemochromatosis
HFE	3077	NP_620577	5821	hemochromatosis
HFE	3077	NP_620578	5822	hemochromatosis
HFE	3077	NP_620579	5823	hemochromatosis
HFE	3077	NP_620580	5824	hemochromatosis
HFE2	148738	NP_660320	5950	hemochromatosis type 2 (juvenile)
HFE2	148738	NP_973733	6919	hemochromatosis type 2 (juvenile)
HFE2	148738	NP_998817	7047	hemochromatosis type 2 (juvenile)
HFE2	148738	NP_998818	7048	hemochromatosis type 2 (juvenile)
HGF	3082	NP_000592	171	hepatocyte growth factor (hepapoietin A scatter factor)
HGF	3082	NP_001010931	579	hepatocyte growth factor (hepapoietin A scatter factor)
HGF	3082	NP_001010932	580	hepatocyte growth factor (hepapoietin A scatter factor)
HGF	3082	NP_001010933	581	hepatocyte growth factor (hepapoietin A scatter factor)
HGF	3082	NP_001010934	582	hepatocyte growth factor (hepapoietin A scatter factor)
HGFAC	3083	NP_001519	2405	HGF activator
HGSNAT	138050	NP_689632	6057	heparan-alpha-glucosaminide N-acetyltransferase
HHAT	55733	NP_001116306	1286	hedgehog acyltransferase
HHAT	55733	NP_001164035	2245	hedgehog acyltransferase
HHAT	55733	NP_001164051	2246	hedgehog acyltransferase
HHAT	55733	NP_001164058	2248	hedgehog acyltransferase
HHAT	55733	NP_001164059	2249	hedgehog acyltransferase
HHAT	55733	NP_060664	4401	hedgehog acyltransferase
HHATL	57467	NP_065758	4705	hedgehog acyltransferase-like
HHIP	64399	NP_071920.1	7311	hedgehog interacting protein
HHIPL2	79802	NP_079022	5076	HHIP-like 2
HHLA2	11148	NP_009003	3626	HERV-H LTR-associating 2
HIATL1	84641	NP_115947	5370	hippocampus abundant transcript-like 1
HIGD1A	25994	NP_001093138	1168	similar toHIG1 domain family, member 1A
HIGD1A	25994	NP_001093139	1169	similar toHIG1 domain family, member 1A
HIGD1A	25994	NP_054775	3865	similar toHIG1 domain family, member 1A
HIGD1B	51751	NP_057522	4217	HIG1Hypoxia inducible domain family, member 1B
HIGD2A	192286	NP_620175	5802	HIG1Hypoxia inducible domain family, member 2A
HILPDA	29923	NP_001092256	1144	chromosome 7 open reading frame 68
HILPDA	29923	NP_037464	3803	chromosome 7 open reading frame 68
HLA-A	3105	NP_002107.3	7319	major histocompatibility complex, class I, A
HLA-B	3106	NP_005505.2	2552	majorHistocompatibility complex, class I, C
HLA-B	3106	NP_005505	3305	majorHistocompatibility complex, class I, C
HLA-C	3107	NP_002108	2551	majorHistocompatibility complex, class I, C
HLA-C	3107	NP_001229971.1	3304	majorHistocompatibility complex, class I, C
HLA-DMA	3108	NP_006111	3437	majorHistocompatibility complex, class II, DM alpha
HLA-DMB	3109	NP_002109	2553	majorHistocompatibility complex, class II, DM beta
HLA-DOA	3111	NP_002110	2554	majorHistocompatibility complex, class II, DO alpha
HLA-DOB	3112	NP_002111	2555	majorHistocompatibility complex, class II, DO beta
HLA-DQB2	3120	NP_001185787	7118	major histocompatibility complex, class II, DQ beta 2
HLA-DRA	3122	NP_061984	4566	majorHistocompatibility complex, class II, DR alpha

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
HLA-DRB1	3123	NP_002115	2556	majorHistocompatibility complex, class II, DR beta 4
HLA-DRB1	3123	NP_068818	4859	majorHistocompatibility complex, class II, DR beta 4
HLA-E	3133	NP_005507	3306	majorHistocompatibility complex, class I, E
HLA-F	3134	NP_001091948	1123	majorHistocompatibility complex, class I, F
HLA-F	3134	NP_001091949	1124	majorHistocompatibility complex, class I, F
HLA-F	3134	NP_061823	4535	majorHistocompatibility complex, class I, F
HLA-G	3135	NP_002118	2557	majorHistocompatibility complex, class I, G
HM13	81502	NP_110416	5174	histocompatibility (minor) 13
HM13	81502	NP_848695	6558	histocompatibility (minor) 13
HM13	81502	NP_848696	6559	histocompatibility (minor) 13
HM13	81502	NP_848697	6560	histocompatibility (minor) 13
HMGCR	3156	NP_000850	295	3-hydroxy-3-methylglutaryl-Coenzyme A reductase
HMGCR	3156	NP_001124468	1466	3-hydroxy-3-methylglutaryl-Coenzyme A reductase
HMOX1	3162	NP_002124	2558	heme oxygenase (decycling) 1
HMOX2	3163	NP_001120676	1310	heme oxygenase (decycling) 2
HMOX2	3163	NP_001120677	1311	heme oxygenase (decycling) 2
HMOX2	3163	NP_001120678	1312	heme oxygenase (decycling) 2
HMOX2	3163	NP_002125	2559	heme oxygenase (decycling) 2
HMP19	51617	NP_057064	4146	HMP19 protein
HOGA1	112817	NP_001128142	1501	N-acetylneuraminate pyruvate lyase 2 (putative)
HOGA1	112817	NP_612422	5746	N-acetylneuraminate pyruvate lyase 2 (putative)
HPN	3249	NP_002142	2560	hepsin
HPN	3249	NP_892028	6699	hepsin
HRASLS2	54979	NP_060348	4363	HRAS-like suppressor 2
HRCT1	646962	NP_001034881	812	histidine rich carboxyl terminus 1
HRH1	3269	NP_000852	296	histamine receptorH1
HRH1	3269	NP_001091681	1119	histamine receptorH1
HRH1	3269	NP_001091682	1120	histamine receptorH1
HRH1	3269	NP_001091683	1121	histamine receptorH1
HRH2	3274	NP_001124527	1467	histamine receptorH2
HRH2	3274	NP_071640	4895	histamine receptorH2
HRH3	11255	NP_009163	3657	histamine receptorH3
HRH4	59340	NP_001137300	1730	histamine receptorH4
HRH4	59340	NP_001153638	1977	histamine receptorH4
HRH4	59340	NP_067637	4810	histamine receptorH4
HS3ST2	9956	NP_006034	3419	heparan sulfate (glucosamine) 3-O-sulfotransferase 2
HS3ST3B1	9953	NP_006032	3418	heparan sulfate (glucosamine) 3-O-sulfotransferase 3B1
HS3ST5	222537	NP_705840	6202	heparan sulfate (glucosamine) 3-O-sulfotransferase 5
HS6ST3	266722	NP_703157	6194	heparan sulfate 6-O-sulfotransferase 3
HSD11B1	3290	NP_005516	3307	hydroxysteroid (11-beta) dehydrogenase 1
HSD11B1	3290	NP_861420	6630	hydroxysteroid (11-beta) dehydrogenase 1
HSD17B12	51144	NP_057226	4176	hydroxysteroid (17-beta) dehydrogenase 12
HSD17B2	3294	NP_002144	2561	hydroxysteroid (17-beta) dehydrogenase 2
HSD17B3	3293	NP_000188	59	hydroxysteroid (17-beta) dehydrogenase 3
HSD3B2	3284	NP_000189	60	hydroxy-delta-5-steroid dehydrogenase, 3 beta- and steroid delta-isomerase 2
HSD3B2	3284	NP_001159592	2151	hydroxy-delta-5-steroid dehydrogenase, 3 beta- and steroid delta-isomerase 2
HSD3B7	80270	NP_001129193	1560	hydroxy-delta-5-steroid dehydrogenase, 3 beta- and steroid delta-isomerase 7
HSD3B7	80270	NP_001136249	1699	hydroxy-delta-5-steroid dehydrogenase, 3 beta- and steroid delta-isomerase 7
HSD3B7	80270	NP_001136250	1700	hydroxy-delta-5-steroid dehydrogenase, 3 beta- and steroid delta-isomerase 7
HSD3B7	80270	NP_060538	4387	hydroxy-delta-5-steroid dehydrogenase, 3 beta- and steroid delta-isomerase 7
HSD3B7	80270	NP_079469	5132	hydroxy-delta-5-steroid dehydrogenase, 3 beta- and steroid delta-isomerase 7
HSPA13	6782	NP_008879	3601	heat shock protein 70kDa family, member 13
HSPA5	3309	NP_005338.1	7305	heat shock protein family A (Hsp70) member 5
HTATIP2	10553	NP_001091990	1132	HIV-1 Tat interactive protein 2, 30kDa
HTATIP2	10553	NP_001091991	1133	HIV-1 Tat interactive protein 2, 30kDa
HTATIP2	10553	NP_001091992	1134	HIV-1 Tat interactive protein 2, 30kDa
HTATIP2	10553	NP_001091993	1135	HIV-1 Tat interactive protein 2, 30kDa
HTATIP2	10553	NP_006401	3484	HIV-1 Tat interactive protein 2, 30kDa
HTR1A	3350	NP_000515	150	5-hydroxytryptamine (serotonin) receptor 1A
HTR1B	3351	NP_000854	297	5-hydroxytryptamine (serotonin) receptor 1B
HTR1D	3352	NP_000855	298	5-hydroxytryptamine (serotonin) receptor 1D
HTR1E	3354	NP_000856.1	7260	5-hydroxytryptamine receptor 1E
HTR1F	3355	NP_000857	299	5-hydroxytryptamine (serotonin) receptor 1F
HTR2A	3356	NP_000612	176	5-hydroxytryptamine (serotonin) receptor 2A
HTR2A	3356	NP_001159419	2132	5-hydroxytryptamine (serotonin) receptor 2A
HTR2B	3357	NP_000858	300	5-hydroxytryptamine (serotonin) receptor 2B
HTR2C	3358	NP_000859	301	5-hydroxytryptamine (serotonin) receptor 2C
HTR3A	3359	NP_000860	302	5-hydroxytryptamine (serotonin) receptor 3A
HTR3A	3359	NP_001155244	2027	5-hydroxytryptamine (serotonin) receptor 3A
HTR3A	3359	NP_998786	7044	5-hydroxytryptamine (serotonin) receptor 3A
HTR3B	9177	NP_006019	3417	5-hydroxytryptamine (serotonin) receptor 3B
HTR3C	170572	NP_570126	5662	5-hydroxytryptamine (serotonin) receptor 3, family member C
HTR4	3360	NP_000861	303	5-hydroxytryptamine (serotonin) receptor 4
HTR4	3360	NP_001035259	848	5-hydroxytryptamine (serotonin) receptor 4
HTR4	3360	NP_000861.1	849	5-hydroxytryptamine (serotonin) receptor 4
HTR4	3360	NP_001035262	850	5-hydroxytryptamine (serotonin) receptor 4
HTR4	3360	NP_001035263	851	5-hydroxytryptamine (serotonin) receptor 4

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
HTR4	3360	NP_001035264	852	5-hydroxytryptamine (serotonin) receptor 4
HTR4	3360	NP_955525	6885	5-hydroxytryptamine (serotonin) receptor 4
HTR5A	3361	NP_076917	4993	5-hydroxytryptamine (serotonin) receptor 5A
HTR6	3362	NP_000862	304	5-hydroxytryptamine (serotonin) receptor 6
HTR7	3363	NP_000863	305	5-hydroxytryptamine (serotonin) receptor 7 (adenylate cyclase-coupled)
HTR7	3363	NP_062873	4587	5-hydroxytryptamine (serotonin) receptor 7 (adenylate cyclase-coupled)
HTR7	3363	NP_062874	4588	5-hydroxytryptamine (serotonin) receptor 7 (adenylate cyclase-coupled)
HTRA2	27429	NP_037379	3789	HtrA serine peptidase 2
HTRA2	27429	NP_659540	5926	HtrA serine peptidase 2
HTRA4	203100	NP_710159	6215	HtrA serine peptidase 4
HVCN1	84329	NP_001035196	836	hydrogen voltage-gated channel 1
HVCN1	84329	NP_115745	5335	hydrogen voltage-gated channel 1
HYAL4	23553	NP_036401	3733	hyaluronoglucosaminidase 4
HYOU1	10525	NP_001124463	1465	hypoxia up-regulated 1
HYOU1	10525	NP_006380	3479	hypoxia up-regulated 1
ICA1	3382	NP_001129492	1590	islet cell autoantigen 1, 69kDa
ICA1	3382	NP_004959	3180	islet cell autoantigen 1, 69kDa
ICA1	3382	NP_071682	4896	islet cell autoantigen 1, 69kDa
ICAM1	3383	NP_000192.2	7382	intercellular adhesion molecule 1
ICAM2	3384	NP_000864	306	intercellular adhesion molecule 2
ICAM2	3384	NP_001093256	1175	intercellular adhesion molecule 2
ICAM2	3384	NP_001093257	1176	intercellular adhesion molecule 2
ICAM2	3384	NP_001093258	1177	intercellular adhesion molecule 2
ICAM2	3384	NP_001093259	1178	intercellular adhesion molecule 2
ICAM3	3385	NP_002153	2563	intercellular adhesion molecule 3
ICAM4	3386	NP_001034221	787	intercellular adhesion molecule 4 (Landsteiner-Wiener blood group)
ICAM4	3386	NP_001535	2410	intercellular adhesion molecule 4 (Landsteiner-Wiener blood group)
ICAM4	3386	NP_071772	4911	intercellular adhesion molecule 4 (Landsteiner-Wiener blood group)
ICAM5	7087	NP_003250	2804	intercellular adhesion molecule 5, telencephalin
ICMT	23463	NP_036537	3769	isoprenylcysteine carboxyl methyltransferase
ICOS	29851	NP_036224	3703	inducible T-cell co-stimulator
ICOSLG	23308	NP_056074	4054	inducible T-cell co-stimulator ligand
IER3IP1	51124	NP_057181	4168	immediate early response 3 interacting protein 1
IFI27	3429	NP_001123552	1417	interferon, alpha-inducible protein 27
IFI27	3429	NP_005523	3309	interferon, alpha-inducible protein 27
IFI27L1	122509	NP_660292	5940	interferon, alpha-inducible protein 27-like 1
IFI27L1	122509	NP_996832	6991	interferon, alpha-inducible protein 27-like 1
IFI6	2537	NP_002029	2538	interferon, alpha-inducible protein 6
IFI6	2537	NP_075010	4956	interferon, alpha-inducible protein 6
IFI6	2537	NP_075011	4957	interferon, alpha-inducible protein 6
IFITM1	8519	NP_003632	2861	interferon induced transmembrane protein 1 (9-27)
IFITM10	402778	NP_001164291	2257	similar to RIKEN cDNA 6330512M04 gene (mouse)
IFITM2	10581	NP_006426	3491	interferon induced transmembrane protein 2 (1-8D)
IFNAR1	3454	NP_000620	180	interferon (alpha, beta and omega) receptor 1
IFNAR2	3455	NP_000865	307	interferon (alpha, beta and omega) receptor 2
IFNAR2	3455	NP_997467	7027	interferon (alpha, beta and omega) receptor 2
IFNAR2	3455	NP_997468	7028	interferon (alpha, beta and omega) receptor 2
IFNB1	3456	NP_002167	2564	interferon, beta 1, fibroblast
IFNE	338376	NP_795372	6487	interferon, epsilon
IFNGR1	3459	NP_000407.1	7273	interferon gamma receptor 1
IFNGR2	3460	NP_005525	3310	interferon gamma receptor 2 (interferon gamma transducer 1)
IFNL1	282618	NP_742152	6284	interleukin 29 (interferon, lambda 1)
IFNL2	282616	NP_742150	6283	interleukin 28A (interferon, lambda 2)
IFNLR1	163702	NP_734464	6260	interleukin 28 receptor, alpha (interferon, lambda receptor)
IFNLR1	163702	NP_775087	6328	interleukin 28 receptor, alpha (interferon, lambda receptor)
IFNLR1	163702	NP_775088	6329	interleukin 28 receptor, alpha (interferon, lambda receptor)
IGB1	3688	NP_002202.2	7309	integrin beta-1 isoform 1A precursor
IGDCC3	9543	NP_004875	3159	immunoglobulin superfamily, DCC subclass, member 3
IGDCC4	57722	NP_066013	4748	immunoglobulin superfamily, DCC subclass, member 4
IGF1R	3480	NP_000866	308	insulin-like growth factor 1 receptor
IGF2R	3482	NP_000867	309	insulin-like growth factor 2 receptor
IGFL1	374918	NP_940943	6811	IGF-like family member 1
IGFLR1	79713	NP_001332933.1	7465	IGF like family receptor 1
IGK	50802	N/A	N/A	immunoglobulin kappa locus
IGSF1	3547	NP_001164432	2260	immunoglobulin superfamily, member 1
IGSF1	3547	NP_001164433	2261	immunoglobulin superfamily, member 1
IGSF1	3547	NP_001164434	2262	immunoglobulin superfamily, member 1
IGSF1	3547	NP_001546	2411	immunoglobulin superfamily, member 1
IGSF1	3547	NP_991402	6949	immunoglobulin superfamily, member 1
IGSF10	285313	NP_849144	6565	immunoglobulin superfamily, member 10
IGSF11	152404	NP_001015887	637	immunoglobulin superfamily, member 11
IGSF11	152404	NP_689751	6078	immunoglobulin superfamily, member 11
IGSF3	3321	NP_001007238	508	immunoglobulin superfamily, member 3
IGSF3	3321	NP_001533	2408	immunoglobulin superfamily, member 3
IGSF6	10261	NP_005840	3384	immunoglobulin superfamily, member 6
IGSF8	93185	NP_443100	5521	immunoglobulin superfamily, member 8
IGSF9	57549	NP_001128522.1	7396	immunoglobulin superfamily member 9

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
IHH	3549	NP_002172	2565	IndianHedgehogHomolog (Drosophila)
IL10RA	3587	NP_001549	2413	interleukin 10 receptor, alpha
IL10RB	3588	NP_000619	179	interleukin 10 receptor, beta
IL11RA	3590	NP_001136256	1701	interleukin 11 receptor, alpha
IL11RA	3590	NP_001136256.1	3066	interleukin 11 receptor, alpha
IL11RA	3590	NP_001136256.1	5993	interleukin 11 receptor, alpha
IL12B	3593	NP_002178	2570	interleukin 12B (natural killer cell stimulatory factor 2, cytotoxic lymphocyte maturation factor 2, p40)
IL12RB1	3594	NP_005526	3311	interleukin 12 receptor, beta 1
IL12RB1	3594	NP_714912	6216	interleukin 12 receptor, beta 1
IL12RB2	3595	NP_001550	2414	interleukin 12 receptor, beta 2
IL13	3596	NP_002179	2571	interleukin 13
IL13RA1	3597	NP_001551	2415	interleukin 13 receptor, alpha 1
IL13RA2	3598	NP_000631	185	interleukin 13 receptor, alpha 2
IL15RA	3601	NP_002180	2572	interleukin 15 receptor, alpha
IL15RA	3601	NP_751950	6288	interleukin 15 receptor, alpha
IL17D	53342	NP_612141	5718	interleukin 17D
IL17RA	23765	NP_055154	3918	interleukin 17 receptor A
IL17RB	55540	NP_061195	4489	interleukin 17 receptor B
IL17RC	84818	NP_116121	5391	interleukin 17 receptor C
IL17RC	84818	NP_703190	6195	interleukin 17 receptor C
IL17RC	84818	NP_703191	6196	interleukin 17 receptor C
IL17RE	132014	NP_705613	6197	interleukin 17 receptor E
IL17RE	132014	NP_705614	6198	interleukin 17 receptor E
IL17RE	132014	NP_705616	6199	interleukin 17 receptor E
IL18R1	8809	NP_003846	2917	interleukin 18 receptor 1
IL18RAP	8807	NP_003844	2915	interleukin 18 receptor accessory protein
IL1R1	3554	NP_000868	310	interleukin 1 receptor, type I
IL1R2	7850	NP_004624	3093	interleukin 1 receptor, type II
IL1R2	7850	NP_001248348.1	6340	interleukin 1 receptor, type II
IL1RAP	3556	NP_001161400	2204	interleukin 1 receptor accessory protein
IL1RAP	3556	NP_001161401	2205	interleukin 1 receptor accessory protein
IL1RAP	3556	NP_001161402	2206	interleukin 1 receptor accessory protein
IL1RAP	3556	NP_001161403	2207	interleukin 1 receptor accessory protein
IL1RAP	3556	NP_002173	2566	interleukin 1 receptor accessory protein
IL1RAP	3556	NP_608273	5712	interleukin 1 receptor accessory protein
IL1RAPL1	11141	NP_055086	3905	interleukin 1 receptor accessory protein-like 1
IL1RAPL2	26280	NP_059112	4284	interleukin 1 receptor accessory protein-like 2
IL1RL1	9173	NP_003847	2918	interleukin 1 receptor-like 1
IL1RL1	9173	NP_057316	4187	interleukin 1 receptor-like 1
IL1RL2	8808	NP_003845	2916	interleukin 1 receptor-like 2
IL20RA	53832	NP_055247	3936	interleukin 20 receptor, alpha
IL20RB	53833	NP_653318	5898	interleukin 20 receptor beta
IL21R	50615	NP_068570	4830	interleukin 21 receptor
IL21R	50615	NP_851564	6580	interleukin 21 receptor
IL21R	50615	NP_851565	6581	interleukin 21 receptor
IL22	50616	NP_065386	4677	interleukin 22
IL22RA1	58985	NP_067081	4804	interleukin 22 receptor, alpha 1
IL23R	149233	NP_653302	5894	interleukin 23 receptor
IL25	64806	NP_073626	4947	interleukin 25
IL25	64806	NP_758525	6307	interleukin 25
IL27	246778	NP_663634	5969	interleukin 27
IL27RA	9466	NP_004834.1	7276	interleukin 27 receptor subunit alpha
IL2RA	3559	NP_000408.1	7271	interleukin 2 receptor subunit alpha
IL2RB	3560	NP_000869.1	7261	interleukin 2 receptor subunit beta
IL2RG	3561	NP_000197.1	7274	interleukin 2 receptor subunit gamma
IL31RA	133396	NP_620586	5826	interleukin 31 receptor A
IL3RA	3563	NP_002174	2567	interleukin 3 receptor, alpha (low affinity)
IL4I1	259307	NP_690863	6130	interleukin 4 induced 1
IL4I1	259307	NP_758962	6323	interleukin 4 induced 1
IL4R	3566	NP_000409	122	interleukin 4 receptor
IL4R	3566	NP_001008699	550	interleukin 4 receptor
IL5RA	3568	NP_000555	162	interleukin 5 receptor, alpha
IL5RA	3568	NP_783851	6450	interleukin 5 receptor, alpha
IL5RA	3568	NP_783852	6451	interleukin 5 receptor, alpha
IL5RA	3568	NP_783853	6452	interleukin 5 receptor, alpha
IL5RA	3568	NP_783854	6453	interleukin 5 receptor, alpha
IL5RA	3568	NP_783855	6454	interleukin 5 receptor, alpha
IL6R	3570	NP_000556	163	interleukin 6 receptor
IL6R	3570	NP_852004	6596	interleukin 6 receptor
IL6ST	3572	NP_002175	2568	interleukin 6 signal transducer (gp130, oncostatin M receptor)
IL6ST	3572	NP_786943	6459	interleukin 6 signal transducer (gp130, oncostatin M receptor)
IL7R	3575	NP_002176	2569	interleukin 7 receptor
ILDR1	286676	NP_787120	6471	immunoglobulin-like domain containing receptor 1
ILDR2	387597	NP_955383	6880	immunoglobulin-like domain containing receptor 2
ILVBL	10994	NP_006835	3578	ilvB (bacterial acetolactate synthase)-like
IMPAD1	54928	NP_060283	4347	inositol monophosphatase domain containing 1
IMPG2	50939	NP_057331	4193	interphotoreceptor matrix proteoglycan 2

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
INAFM1	255783	NP_848606	6547	hypothetical protein LOC255783
INPP4A	3631	NP_001127696	1472	inositol polyphosphate-4-phosphatase, type I, 107kDa
INPP4A	3631	NP_001127697	1473	inositol polyphosphate-4-phosphatase, type I, 107kDa
INPP4A	3631	NP_001557	2416	inositol polyphosphate-4-phosphatase, type I, 107kDa
INPP4A	3631	NP_004018	2947	inositol polyphosphate-4-phosphatase, type I, 107kDa
INPP4B	8821	NP_001095139	1204	inositol polyphosphate-4-phosphatase, type II, 105kDa
INPP4B	8821	NP_003857	2921	inositol polyphosphate-4-phosphatase, type II, 105kDa
INSIG1	3638	NP_005533	3312	insulin induced gene 1
INSIG1	3638	NP_938150	6772	insulin induced gene 1
INSIG1	3638	NP_938151	6773	insulin induced gene 1
INSIG2	51141	NP_057217	4174	insulin induced gene 2
INSL3	3640	NP_005534	3313	insulin-like 3 (Leydig cell)
INSR	3643	NP_000199	61	insulin receptor
INSR	3643	NP_001073285	997	insulin receptor
INSRR	3645	NP_055030	3884	insulin receptor-related receptor
ISLR2	57611	NP_001123608	1418	immunoglobulin superfamily containing leucine-rich repeat 2
ISLR2	57611	NP_001123609	1419	immunoglobulin superfamily containing leucine-rich repeat 2
ISLR2	57611	NP_001123610	1420	immunoglobulin superfamily containing leucine-rich repeat 2
ISLR2	57611	NP_065902	4731	immunoglobulin superfamily containing leucine-rich repeat 2
ISM2	145501	NP_872315	6647	isthmin 2Homolog (zebrafish)
ISM2	145501	NP_954993	6873	isthmin 2Homolog (zebrafish)
ITFG1	81533	NP_110417	5175	integrin alpha FG-GAP repeat containing 1
ITFG3	83986	NP_114428	5284	integrin alpha FG-GAP repeat containing 3
ITGA10	8515	NP_003628	2859	integrin, alpha 10
ITGA11	22801	NP_001004439	435	integrin, alpha 11
ITGA2	3673	NP_002194	2573	integrin, alpha 2 (CD49B, alpha 2 subunit of VLA-2 receptor)
ITGA2B	3674	NP_000410	123	integrin, alpha 2b (platelet glycoprotein IIb of IIb/IIIa complex, antigen CD41)
ITGA3	3675	NP_002195	2574	integrin, alpha 3 (antigen CD49C, alpha 3 subunit of VLA-3 receptor)
ITGA3	3675	NP_002195.1	3297	integrin, alpha 3 (antigen CD49C, alpha 3 subunit of VLA-3 receptor)
ITGA4	3676	NP_000876	311	integrin, alpha 4 (antigen CD49D, alpha 4 subunit of VLA-4 receptor)
ITGA5	3678	NP_002196	2575	integrin, alpha 5 (fibronectin receptor, alpha polypeptide)
ITGA6	3655	NP_000201	62	integrin, alpha 6
ITGA6	3655	NP_001073286	998	integrin, alpha 6
ITGA7	3679	NP_001138468	1800	integrin, alpha 7
ITGA7	3679	NP_001138469	1801	integrin, alpha 7
ITGA7	3679	NP_002197	2576	integrin, alpha 7
ITGA8	8516	NP_003629	2860	integrin, alpha 8
ITGA9	3680	NP_002198	2577	integrin, alpha 9
ITGAD	3681	NP_005344	3275	integrin, alpha D
ITGAE	3682	NP_002199	2578	integrin, alpha E (antigen CD103,Human mucosal lymphocyte antigen 1 alpha polypeptide)
ITGAL	3683	NP_001107852.1	7383	integrin subunit alpha L
ITGAM	3684	NP_000623	181	integrin, alpha M (complement component 3 receptor 3 subunit)
ITGAM	3684	NP_001139280	1877	integrin, alpha M (complement component 3 receptor 3 subunit)
ITGAV	3685	NP_001138471	1802	integrin, alpha V (vitronectin receptor, alpha polypeptide, antigen CD51)
ITGAV	3685	NP_001138472	1803	integrin, alpha V (vitronectin receptor, alpha polypeptide, antigen CD51)
ITGAV	3685	NP_002201	2579	integrin, alpha V (vitronectin receptor, alpha polypeptide, antigen CD51)
ITGAX	3687	NP_000878	312	integrin, alpha X (complement component 3 receptor 4 subunit)
ITGB1	3688	NP_596867.1	7310	integrin subunit beta 1
ITGB2	3689	NP_000202	63	integrin, beta 2 (complement component 3 receptor 3 and 4 subunit)
ITGB2	3689	NP_001120963	1331	integrin, beta 2 (complement component 3 receptor 3 and 4 subunit)
ITGB3	3690	NP_000203	64	integrin, beta 3 (platelet glycoprotein IIIa, antigen CD61)
ITGB5	3693	NP_002204	2580	integrin, beta 5
ITGB6	3694	NP_000879	313	integrin, beta 6
ITGB7	3695	NP_000880	314	integrin, beta 7
ITGB8	3696	NP_002205	2581	integrin, beta 8
ITIH1	3697	NP_001159906	2168	inter-alpha (globulin) inhibitorH1
ITIH1	3697	NP_001159907	2169	inter-alpha (globulin) inhibitorH1
ITIH1	3697	NP_001159908	2170	inter-alpha (globulin) inhibitorH1
ITIH1	3697	NP_002206	2582	inter-alpha (globulin) inhibitorH1
ITM2A	9452	NP_001165052	2278	integral membrane protein 2A
ITM2A	9452	NP_004858	3153	integral membrane protein 2A
ITM2B	9445	NP_068839	4862	integral membrane protein 2B
ITM2C	81618	NP_001012532.1	7343	integral membrane protein 2C
ITPR1	3708	NP_001093422	1179	inositol 1,4,5-triphosphate receptor, type 1
ITPR1	3708	NP_001161744	2214	inositol 1,4,5-triphosphate receptor, type 1
ITPR1	3708	NP_002213	2583	inositol 1,4,5-triphosphate receptor, type 1
ITPR2	3709	NP_002214	2584	inositol 1,4,5-triphosphate receptor, type 2
ITPR3	3710	NP_002215	2585	inositol 1,4,5-triphosphate receptor, type 3
ITPRIPL1	150771	NP_001008949	558	inositol 1,4,5-triphosphate receptor interacting protein-like 1
ITPRIPL1	150771	NP_001156995	2054	inositol 1,4,5-triphosphate receptor interacting protein-like 1
ITPRIPL1	150771	NP_001156996	2055	inositol 1,4,5-triphosphate receptor interacting protein-like 1
ITPRIPL1	150771	NP_848590	6543	inositol 1,4,5-triphosphate receptor interacting protein-like 1
ITPRIPL2	162073	NP_001030013	761	inositol 1,4,5-triphosphate receptor interacting protein-like 2
ITSN1	6453	NP_001001132	350	intersectin 1 (SH3 domain protein)
ITSN1	6453	NP_003015	2756	intersectin 1 (SH3 domain protein)
IYD	389434	NP_001158166	2098	iodotyrosine deiodinase
IYD	389434	NP_001158167	2099	iodotyrosine deiodinase

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
IYD	389434	NP_981932	6930	iodotyrosine deiodinase
IZUMO1	284359	NP_872381	6663	izumo sperm-egg fusion 1
IZUMO2	126123	NP_689571	6041	chromosome 19 open reading frame 41
JAG1	182	NP_000205	65	jagged 1 (Alagille syndrome)
JAG2	3714	NP_002217	2586	jagged 2
JAG2	3714	NP_660142	5928	jagged 2
JAGN1	84522	NP_115881	5360	jagunalHomolog 1 (Drosophila)
JAM2	58494	NP_067042	4796	junctional adhesion molecule 2
JAM3	83700	NP_001192258.1	7430	junctional adhesion molecule 3
JPH1	56704	NP_065698	4689	junctophilin 1
JPH2	57158	NP_065166	4657	junctophilin 2
JPH2	57158	NP_787109	6469	junctophilin 2
JPH3	57338	NP_065706	4690	junctophilin 3
JTB	10899	NP_006685	3548	jumping translocation breakpoint
KANSL1L	151050	NP_689732	6073	chromosome 2 open reading frame 67
KCNA1	3736	NP_000208	66	potassium voltage-gated channel, shaker-related subfamily, member 1 (episodic ataxia with myokymia)
KCNA10	3744	NP_005540	3314	potassium voltage-gated channel, shaker-related subfamily, member 10
KCNA2	3737	NP_004965	3181	potassium voltage-gated channel, shaker-related subfamily, member 2
KCNA3	3738	NP_002223	2588	potassium voltage-gated channel, shaker-related subfamily, member 3
KCNA4	3739	NP_002224	2589	potassium voltage-gated channel, shaker-related subfamily, member 4
KCNA5	3741	NP_002225	2590	potassium voltage-gated channel, shaker-related subfamily, member 5
KCNA6	3742	NP_002226	2591	potassium voltage-gated channel, shaker-related subfamily, member 6
KCNA7	3743	NP_114092	5259	potassium voltage-gated channel, shaker-related subfamily, member 7
KCNB1	3745	NP_004966	3182	potassium voltage-gated channel, Shab-related subfamily, member 1
KCNB2	9312	NP_004761	3119	potassium voltage-gated channel, Shab-related subfamily, member 2
KCNC1	3746	NP_001106212	1249	potassium voltage-gated channel, Shaw-related subfamily, member 1
KCNC1	3746	NP_004967	3183	potassium voltage-gated channel, Shaw-related subfamily, member 1
KCNC2	3747	NP_631874	5836	potassium voltage-gated channel, Shaw-related subfamily, member 2
KCNC2	3747	NP_631875	5837	potassium voltage-gated channel, Shaw-related subfamily, member 2
KCNC2	3747	NP_715624	6223	potassium voltage-gated channel, Shaw-related subfamily, member 2
KCNC3	3748	NP_004968	3184	potassium voltage-gated channel, Shaw-related subfamily, member 3
KCNC4	3749	NP_001034663	800	potassium voltage-gated channel, Shaw-related subfamily, member 4
KCNC4	3749	NP_004969	3185	potassium voltage-gated channel, Shaw-related subfamily, member 4
KCNC4	3749	NP_001034663.1	6225	potassium voltage-gated channel, Shaw-related subfamily, member 4
KCND1	3750	NP_004970	3186	potassium voltage-gated channel, Shal-related subfamily, member 1
KCND2	3751	NP_036413.1	7288	potassium voltage-gated channel subfamily D member 2
KCND3	3752	NP_004971	3187	potassium voltage-gated channel, Shal-related subfamily, member 3
KCND3	3752	NP_751948	6287	potassium voltage-gated channel, Shal-related subfamily, member 3
KCNE1	3753	NP_000210	68	potassium voltage-gated channel, Isk-related family, member 1
KCNE1	3753	NP_001121140	1342	potassium voltage-gated channel, Isk-related family, member 1
KCNE1	3753	NP_001121141	1343	potassium voltage-gated channel, Isk-related family, member 1
KCNE1	3753	NP_001121142	1344	potassium voltage-gated channel, Isk-related family, member 1
KCNE2	9992	NP_751951	6289	potassium voltage-gated channel, Isk-related family, member 2
KCNE3	10008	NP_005463	3291	potassium voltage-gated channel, Isk-related family, member 3
KCNE4	23704	NP_542402	5616	potassium voltage-gated channel, Isk-related family, member 4
KCNE5	23630	NP_036414	3734	KCNE1-like
KCNF1	3754	NP_002227	2592	potassium voltage-gated channel, subfamily F, member 1
KCNG1	3755	NP_002228	2593	potassium voltage-gated channel, subfamily G, member 1
KCNG2	26251	NP_036415	3735	potassium voltage-gated channel, subfamily G, member 2
KCNG3	170850	NP_579875	5687	potassium voltage-gated channel, subfamily G, member 3
KCNG3	170850	NP_758847	6308	potassium voltage-gated channel, subfamily G, member 3
KCNH1	3756	NP_002229	2594	potassium voltage-gated channel, subfamilyH (eag-related), member 1
KCNH1	3756	NP_758872	6321	potassium voltage-gated channel, subfamilyH (eag-related), member 1
KCNH2	3757	NP_000229	75	potassium voltage-gated channel, subfamilyH (eag-related), member 2
KCNH2	3757	NP_742053	6271	potassium voltage-gated channel, subfamilyH (eag-related), member 2
KCNH2	3757	NP_742054	6272	potassium voltage-gated channel, subfamilyH (eag-related), member 2
KCNH3	23416	NP_036416	3736	potassium voltage-gated channel, subfamilyH (eag-related), member 3
KCNH4	23415	NP_036417	3737	potassium voltage-gated channel, subfamilyH (eag-related), member 4
KCNH5	27133	NP_647479	5853	potassium voltage-gated channel, subfamilyH (eag-related), member 5
KCNH5	27133	NP_758963	6324	potassium voltage-gated channel, subfamilyH (eag-related), member 5
KCNH5	27133	NP_758964	6325	potassium voltage-gated channel, subfamilyH (eag-related), member 5
KCNH6	81033	NP_110406	5170	potassium voltage-gated channel, subfamilyH (eag-related), member 6
KCNH6	81033	NP_775115	6335	potassium voltage-gated channel, subfamilyH (eag-related), member 6
KCNH7	90134	NP_150375	5477	potassium voltage-gated channel, subfamilyH (eag-related), member 7
KCNH7	90134	NP_775185	6337	potassium voltage-gated channel, subfamilyH (eag-related), member 7
KCNH8	131096	NP_653234	5880	potassium voltage-gated channel, subfamilyH (eag-related), member 8
KCNIP3	30818	NP_001030086	763	Kv channel interacting protein 3, calsenilin
KCNIP3	30818	NP_038462	3821	Kv channel interacting protein 3, calsenilin
KCNJ1	3758	NP_000211	69	potassium inwardly-rectifying channel, subfamily J, member 1
KCNJ1	3758	NP_722448	6226	potassium inwardly-rectifying channel, subfamily J, member 1
KCNJ1	3758	NP_722449	6227	potassium inwardly-rectifying channel, subfamily J, member 1
KCNJ1	3758	NP_722450	6228	potassium inwardly-rectifying channel, subfamily J, member 1
KCNJ1	3758	NP_722451	6229	potassium inwardly-rectifying channel, subfamily J, member 1
KCNJ10	3766	NP_002232	2597	potassium inwardly-rectifying channel, subfamily J, member 10
KCNJ11	3767	NP_000516	151	potassium inwardly-rectifying channel, subfamily J, member 11
KCNJ11	3767	NP_001159762	2157	potassium inwardly-rectifying channel, subfamily J, member 11
KCNJ12	3768	NP_066292	4759	similar toHkir2.2x

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
KCNJ13	3769	NP_002233	2598	potassium inwardly-rectifying channel, subfamily J, member 13
KCNJ14	3770	NP_037480	3809	potassium inwardly-rectifying channel, subfamily J, member 14
KCNJ14	3770	NP_037480.1	6249	potassium inwardly-rectifying channel, subfamily J, member 14
KCNJ15	3772	NP_002234	2599	potassium inwardly-rectifying channel, subfamily J, member 15
KCNJ15	3772	NP_733932	6256	potassium inwardly-rectifying channel, subfamily J, member 15
KCNJ15	3772	NP_733933	6257	potassium inwardly-rectifying channel, subfamily J, member 15
KCNJ16	3773	NP_061128	4481	potassium inwardly-rectifying channel, subfamily J, member 16
KCNJ16	3773	NP_733937	6258	potassium inwardly-rectifying channel, subfamily J, member 16
KCNJ16	3773	NP_733938	6259	potassium inwardly-rectifying channel, subfamily J, member 16
KCNJ2	3759	NP_000882	316	potassium inwardly-rectifying channel, subfamily J, member 2
KCNJ3	3760	NP_002230	2595	potassium inwardly-rectifying channel, subfamily J, member 3
KCNJ4	3761	NP_004972	3188	potassium inwardly-rectifying channel, subfamily J, member 4
KCNJ4	3761	NP_690607	6124	potassium inwardly-rectifying channel, subfamily J, member 4
KCNJ5	3762	NP_000881	315	potassium inwardly-rectifying channel, subfamily J, member 5
KCNJ6	3763	NP_002231	2596	potassium inwardly-rectifying channel, subfamily J, member 6
KCNJ8	3764	NP_004973	3189	potassium inwardly-rectifying channel, subfamily J, member 8
KCNJ9	3765	NP_004974	3190	potassium inwardly-rectifying channel, subfamily J, member 9
KCNK1	3775	NP_002236	2600	potassium channel, subfamily K, member 1
KCNK10	54207	NP_066984	4784	potassium channel, subfamily K, member 10
KCNK10	54207	NP_612190	5723	potassium channel, subfamily K, member 10
KCNK10	54207	NP_612191	5724	potassium channel, subfamily K, member 10
KCNK12	56660	NP_071338.1	7295	potassium two pore domain channel subfamily K member 12
KCNK13	56659	NP_071337	4869	potassium channel, subfamily K, member 13
KCNK15	60598	NP_071753	4902	potassium channel, subfamily K, member 15
KCNK16	83795	NP_001128577	1522	potassium channel, subfamily K, member 16
KCNK16	83795	NP_001128578	1523	potassium channel, subfamily K, member 16
KCNK16	83795	NP_001128579	1524	potassium channel, subfamily K, member 16
KCNK16	83795	NP_115491	5298	potassium channel, subfamily K, member 16
KCNK17	89822	NP_001128583.1	7397	potassium two pore domain channel subfamily K member 17
KCNK2	3776	NP_001017424	643	potassium channel, subfamily K, member 2
KCNK2	3776	NP_001017425	644	potassium channel, subfamily K, member 2
KCNK2	3776	NP_055032	3885	potassium channel, subfamily K, member 2
KCNK3	3777	NP_002237	2601	potassium channel, subfamily K, member 3
KCNK4	50801	NP_201567	5486	potassium channel, subfamily K, member 4
KCNK5	8645	NP_003731	2883	potassium channel, subfamily K, member 5
KCNK6	9424	NP_004814	3139	potassium channel, subfamily K, member 6
KCNK7	10089	NP_005705	3349	potassium channel, subfamily K, member 7
KCNK7	10089	NP_203133	5490	potassium channel, subfamily K, member 7
KCNK7	10089	NP_203134	5491	potassium channel, subfamily K, member 7
KCNK7	10089	NP_258416	5500	potassium channel, subfamily K, member 7
KCNK9	51305	NP_001269463.1	4257	potassium channel, subfamily K, member 9
KCNMA1	3778	NP_001014797	624	potassium large conductance calcium-activated channel, subfamily M, alpha member 1
KCNMA1	3778	NP_001154824	2002	potassium large conductance calcium-activated channel, subfamily M, alpha member 1
KCNMA1	3778	NP_001154825	2003	potassium large conductance calcium-activated channel, subfamily M, alpha member 1
KCNMA1	3778	NP_002238	2602	potassium large conductance calcium-activated channel, subfamily M, alpha member 1
KCNMB1	3779	NP_004128	2972	potassium large conductance calcium-activated channel, subfamily M, beta member 1
KCNMB2	10242	NP_005823	3379	potassium large conductance calcium-activated channel, subfamily M, beta member 2
KCNMB2	10242	NP_852006	6597	potassium large conductance calcium-activated channel, subfamily M, beta member 2
KCNMB4	27345	NP_055320	3949	potassium large conductance calcium-activated channel, subfamily M, beta member 4
KCNN2	3781	NP_067627	4809	potassium intermediate/small conductance calcium-activated channel, subfamily N, member 2
KCNN2	3781	NP_740721	6262	potassium intermediate/small conductance calcium-activated channel, subfamily N, member 2
KCNN3	3782	NP_002240	2603	potassium intermediate/small conductance calcium-activated channel, subfamily N, member 3
KCNN3	3782	NP_740752	6265	potassium intermediate/small conductance calcium-activated channel, subfamily N, member 3
KCNN4	3783	NP_002241	2604	potassium intermediate/small conductance calcium-activated channel, subfamily N, member 4
KCNQ1	3784	NP_000209	67	potassium voltage-gated channel, KQT-like subfamily, member 1
KCNQ1	3784	NP_861463	6637	potassium voltage-gated channel, KQT-like subfamily, member 1
KCNQ2	3785	NP_004509	3067	potassium voltage-gated channel, KQT-like subfamily, member 2
KCNQ2	3785	NP_742104	6278	potassium voltage-gated channel, KQT-like subfamily, member 2
KCNQ2	3785	NP_742105	6279	potassium voltage-gated channel, KQT-like subfamily, member 2
KCNQ2	3785	NP_742106	6280	potassium voltage-gated channel, KQT-like subfamily, member 2
KCNQ2	3785	NP_742107	6281	potassium voltage-gated channel, KQT-like subfamily, member 2
KCNQ3	3786	NP_004510	3068	potassium voltage-gated channel, KQT-like subfamily, member 3
KCNQ4	9132	NP_004691	3104	potassium voltage-gated channel, KQT-like subfamily, member 4
KCNQ4	9132	NP_751895	6285	potassium voltage-gated channel, KQT-like subfamily, member 4
KCNQ5	56479	NP_001153602.1	7409	potassium voltage-gated channel subfamily Q member 5
KCNS1	3787	NP_002242	2605	potassium voltage-gated channel, delayed-rectifier, subfamily S, member 1
KCNS2	3788	NP_065748	4702	potassium voltage-gated channel, delayed-rectifier, subfamily S, member 2
KCNS3	3790	NP_002243	2606	potassium voltage-gated channel, delayed-rectifier, subfamily S, member 3
KCNT1	57582	NP_001258932.1	7435	potassium sodium-activated channel subfamily T member 1
KCNT2	343450	NP_940905	6803	potassium channel, subfamily T, member 2
KCNU1	157855	NP_001027006	737	potassium channel, subfamily U, member 1
KCNV1	27012	NP_055194	3925	potassium channel, subfamily V, member 1
KCNV2	169522	NP_598004	5698	potassium channel, subfamily V, member 2
KCTD8	386617	NP_938167.1	7331	potassium channel tetramerization domain containing 8
KDELR1	10945	NP_006792	3566	KDEL (Lys-Asp-Glu-Leu) endoplasmic reticulum protein retention receptor 1
KDELR2	11014	NP_001094073	1191	KDEL (Lys-Asp-Glu-Leu) endoplasmic reticulum protein retention receptor 2
KDELR2	11014	NP_006845	3581	KDEL (Lys-Asp-Glu-Leu) endoplasmic reticulum protein retention receptor 2

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
KDELR3	11015	NP_006846	3582	KDEL (Lys-Asp-Glu-Leu) endoplasmic reticulum protein retention receptor 3
KDELR3	11015	NP_057839	4266	KDEL (Lys-Asp-Glu-Leu) endoplasmic reticulum protein retention receptor 3
KDR	3791	NP_002244	2607	kinase insert domain receptor (a type III receptor tyrosine kinase)
KDSR	2531	NP_002026	2536	3-ketodihydrosphingosine reductase
KEL	3792	NP_000411	124	Kell blood group, metallo-endopeptidase
KIAA0040	9674	NP_001156365	2035	KIAA0040
KIAA0040	9674	NP_001156366	2036	KIAA0040
KIAA0040	9674	NP_001156367	2037	KIAA0040
KIAA0040	9674	NP_055471	3971	KIAA0040
KIAA0319	9856	NP_001161846	2227	KIAA0319
KIAA0319	9856	NP_001161847	2228	KIAA0319
KIAA0319	9856	NP_001161848	2229	KIAA0319
KIAA0319	9856	NP_001161849	2230	KIAA0319
KIAA0319	9856	NP_055624	3993	KIAA0319
KIAA0319L	79932	NP_079150	5094	KIAA0319-like
KIAA0319L	79932	NP_079150.3	6676	KIAA0319-like
KIAA0754	643314	NP_055853.1	7374	KIAA0754
KIAA1024	23251	NP_056021	4046	KIAA1024
KIAA1109	84162	NP_056127	4061	KIAA1109
KIAA1161	57462	NP_065753	4704	KIAA1161
KIAA1324	57535	NP_065826	4719	KIAA1324
KIAA1324L	222223	NP_001136221	1687	KIAA1324-like
KIAA1324L	222223	NP_689961	6110	KIAA1324-like
KIAA1467	57613	NP_065904	4732	KIAA1467
KIAA1549	57670	NP_001158137	2092	KIAA1549
KIAA1549	57670	NP_065961	4739	KIAA1549
KIAA1549L	25758	NP_036326	3718	chromosome 11 open reading frame 41
KIAA1644	85352	NP_001092764	1155	KIAA1644
KIAA1715	80856	NP_085153	5154	KIAA1715
KIAA1919	91749	NP_699200	6186	KIAA1919
KIDINS220	57498	NP_065789	4712	kinase D-interacting substrate, 220kDa
KIR2DL1	3802	NP_055033.2	634	killer cell immunoglobulin-like receptor, two domains, long cytoplasmic tail, 2
KIR2DL1	3802	NP_055033	3886	killer cell immunoglobulin-like receptor, two domains, long cytoplasmic tail, 2
KIR2DL1	3802	NP_055034	3887	killer cell immunoglobulin-like receptor, two domains, long cytoplasmic tail, 2
KIR2DL3	3804	NP_056952.2	3952	killer cell immunoglobulin-like receptor, two domains, long cytoplasmic tail, 3
KIR2DL3	3804	NP_056952	4126	killer cell immunoglobulin-like receptor, two domains, long cytoplasmic tail, 3
KIR2DL4	3805	NP_001074239	1040	killer cell immunoglobulin-like receptor, two domains, long cytoplasmic tail, 4
KIR2DL4	3805	NP_001074241	1041	killer cell immunoglobulin-like receptor, two domains, long cytoplasmic tail, 4
KIR2DL4	3805	NP_002246	2608	killer cell immunoglobulin-like receptor, two domains, long cytoplasmic tail, 4
KIR2DS4	3809	NP_036444	3742	killer cell immunoglobulin-like receptor, two domains, short cytoplasmic tail, 2
KIR2DS4	3809	NP_036446	3743	killer cell immunoglobulin-like receptor, two domains, short cytoplasmic tail, 2
KIR3DL2	3812	NP_006728	3554	killer cell immunoglobulin-like receptor, three domains, long cytoplasmic tail, 2
KIRREL	55243	NP_060710	4406	kin of IRRE like (Drosophila)
KIRREL2	84063	NP_001316459.1	7463	kin of IRRE like 2 (Drosophila)
KIRREL3	84623	NP_001155179	2022	kin of IRRE like 3 (Drosophila)
KIRREL3	84623	NP_115920	5367	kin of IRRE like 3 (Drosophila)
KISS1R	84634	NP_115940.2	7349	KISS1 receptor
KIT	3815	NP_000213	70	similar to Mast/stem cell growth factor receptor precursor (SCFR) (Proto-oncogene tyrosine-protein kinase Kit) (c-kit) (CD117 antigen)
KIT	3815	NP_001087241	1101	similar to Mast/stem cell growth factor receptor precursor (SCFR) (Proto-oncogene tyrosine-protein kinase Kit) (c-kit) (CD117 antigen)
KITLG	4254	NP_000890	318	KIT ligand
KITLG	4254	NP_003985	2942	KIT ligand
KL	9365	NP_004786	3128	klotho
KLB	152831	NP_783864	6457	klotho beta
KLHDC10	23008	NP_055812	4024	kelch domain containing 10
KLHDC7A	127707	NP_689588	6045	kelch domain containing 7A
KLHL2	11275	NP_001154993	2015	kelch-like 2, Mayven (Drosophila)
KLHL2	11275	NP_001154994	2016	kelch-like 2, Mayven (Drosophila)
KLHL2	11275	NP_009177	3659	kelch-like 2, Mayven (Drosophila)
KLHL31	401265	NP_001003760	423	kelch-like 31 (Drosophila)
KLHL5	51088	NP_001007076	505	kelch-like 5 (Drosophila)
KLHL5	51088	NP_001165125	2279	kelch-like 5 (Drosophila)
KLHL5	51088	NP_057074	4147	kelch-like 5 (Drosophila)
KLHL5	51088	NP_950240	6852	kelch-like 5 (Drosophila)
KLK4	9622	NP_004908	3167	kallikrein-related peptidase 4
KLK5	25818	NP_001070959	971	kallikrein-related peptidase 5
KLK5	25818	NP_001070960	972	kallikrein-related peptidase 5
KLK5	25818	NP_036559	3773	kallikrein-related peptidase 5
KLK8	11202	NP_009127	3646	kallikrein-related peptidase 8
KLK8	11202	NP_653088	5858	kallikrein-related peptidase 8
KLK8	11202	NP_653089	5859	kallikrein-related peptidase 8
KLK8	11202	NP_653090	5860	kallikrein-related peptidase 8
KLRB1	3820	NP_002249	2609	killer cell lectin-like receptor subfamily B, member 1
KLRC3	3823	NP_002252	2610	killer cell lectin-like receptor subfamily C, member 3
KLRC3	3823	NP_031359	3689	killer cell lectin-like receptor subfamily C, member 3
KLRC4	8302	NP_038459	3820	killer cell lectin-like receptor subfamily C, member 4

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
KLRD1	3824	NP_001107868	1273	killer cell lectin-like receptor subfamily D, member 1
KLRD1	3824	NP_002253	2611	killer cell lectin-like receptor subfamily D, member 1
KLRD1	3824	NP_031360	3690	killer cell lectin-like receptor subfamily D, member 1
KLRF1	51348	NP_057607	4235	killer cell lectin-like receptor subfamily F, member 1
KLRG1	10219	NP_005801	3373	killer cell lectin-like receptor subfamily G, member 1
KMO	8564	NP_003670	2869	kynurenine 3-monooxygenase (kynurenine 3-hydroxylase)
KMT2E	55904	NP_061152	4485	myeloid/lymphoid or mixed-lineage leukemia 5 (trithoraxHomolog, Drosophila)
KMT2E	55904	NP_891847	6693	myeloid/lymphoid or mixed-lineage leukemia 5 (trithoraxHomolog, Drosophila)
KNCN	148930	NP_001091080	1105	kinocilin
KREMEN1	83999	NP_001034659	799	kringle containing transmembrane protein 1
KREMEN1	83999	NP_114434	5285	kringle containing transmembrane protein 1
KREMEN2	79412	NP_078783	5036	kringle containing transmembrane protein 2
KREMEN2	79412	NP_757384	6298	kringle containing transmembrane protein 2
KRT5	3852	NP_000415	125	keratin 5
KRTAP19-3	337970	NP_853640	6613	keratin associated protein 19-3
KRTCAP2	200185	NP_776251	6395	keratinocyte associated protein 2
KRTCAP3	200634	NP_001161836	2225	keratinocyte associated protein 3
KRTCAP3	200634	NP_776252	6396	keratinocyte associated protein 3
L1CAM	3897	NP_000416	126	L1 cell adhesion molecule
L1CAM	3897	NP_001137435	1743	L1 cell adhesion molecule
L1CAM	3897	NP_076493	4990	L1 cell adhesion molecule
L3MBTL2	83746	NP_113676	5235	l(3)mbt-like 2 (Drosophila)
LAG3	3902	NP_002277.4	7384	lymphocyte activating 3
LAIR1	3903	NP_002278	2612	leukocyte-associated immunoglobulin-like receptor 1
LAIR1	3903	NP_068352	4818	leukocyte-associated immunoglobulin-like receptor 1
LAMP1	3916	NP_005552	3315	lysosomal-associated membrane protein 1
LAMP2	3920	NP_001116078	1283	lysosomal-associated membrane protein 2
LAMP2	3920	NP_002285	2613	lysosomal-associated membrane protein 2
LAMP2	3920	NP_054701	3851	lysosomal-associated membrane protein 2
LAMP3	27074	NP_055213	3931	lysosomal-associated membrane protein 3
LAMP5	24141	NP_036393	3729	chromosome 20 open reading frame 103
LAPTM4A	9741	NP_055528	3976	lysosomal protein transmembrane 4 alpha
LAPTM4B	55353	NP_060877	4441	lysosomal protein transmembrane 4 beta
LAPTM5	7805	NP_006753	3558	lysosomal multispanning membrane protein 5
LARGE	9215	NP_004728	3113	like-glycosyltransferase
LARGE	9215	NP_598397	5701	like-glycosyltransferase
LAT	27040	NP_001014987.1	7347	linker for activation of T-cells
LAX1	54900	NP_001129662.1	7398	lymphocyte transmembrane adaptor 1
LAYN	143903	NP_849156	6568	layilin
LBR	3930	NP_002287	2614	lamin B receptor
LBR	3930	NP_919424	6726	lamin B receptor
LCAT	3931	NP_000220	71	lecithin-cholesterol acyltransferase
LCLAT1	253558	NP_001002257	391	lysocardiolipin acyltransferase 1
LCLAT1	253558	NP_872357	6657	lysocardiolipin acyltransferase 1
LCN10	414332	NP_001001712	375	lipocalin 10
LCT	3938	NP_002290	2615	lactase
LCTL	197021	NP_997221	7009	lactase-like
LDLR	3949	NP_000518	152	low density lipoprotein receptor
LDLRAD3	143458	NP_001291192.1	7451	low density lipoprotein receptor class A domain containing 3
LDLRAD4	753	NP_001003674	411	chromosome 18 open reading frame 1
LDLRAD4	753	NP_001003675	412	chromosome 18 open reading frame 1
LDLRAD4	753	NP_001003674.1	3025	chromosome 18 open reading frame 1
LDLRAD4	753	NP_852146	6600	chromosome 18 open reading frame 1
LDLRAD4	753	NP_852147	6601	chromosome 18 open reading frame 1
LDLRAD4	753	NP_852148	6602	chromosome 18 open reading frame 1
LECT1	11061	NP_001011705	590	leukocyte cell derived chemotaxin 1
LECT1	11061	NP_008946	3615	leukocyte cell derived chemotaxin 1
LEMD1	93273	NP_001001552	369	LEM domain containing 1
LEMD2	221496	NP_001137416	1737	LEM domain containing 2
LEMD2	221496	NP_851853	6593	LEM domain containing 2
LEMD3	23592	NP_001161086	2188	LEM domain containing 3
LEMD3	23592	NP_055134	3913	LEM domain containing 3
LEPR	3953	NP_001003679	413	leptin receptor
LEPR	3953	NP_001003680	414	leptin receptor
LEPR	3953	NP_002294	2616	leptin receptor
LEPROT	54741	NP_059996	4301	leptin receptor overlapping transcript
LEPROTL1	23484	NP_001121680	1367	leptin receptor overlapping transcript-like 1
LEPROTL1	23484	NP_056159	4065	leptin receptor overlapping transcript-like 1
LETM1	3954	NP_036450	3744	leucine zipper-EF-hand containing transmembrane protein 1
LETM2	137994	NP_653253	5884	leucine zipper-EF-hand containing transmembrane protein 2
LETMD1	25875	NP_001230618.1	684	LETM1 domain containing 1
LETMD1	25875	NP_056231	4079	LETM1 domain containing 1
LFNG	3955	NP_001035257	846	LFNG O-fucosylpeptide 3-beta-N-acetylglucosaminyltransferase
LFNG	3955	NP_001035258	847	LFNG O-fucosylpeptide 3-beta-N-acetylglucosaminyltransferase
LFNG	3955	NP_001159827	2163	LFNG O-fucosylpeptide 3-beta-N-acetylglucosaminyltransferase
LFNG	3955	NP_002295	2617	LFNG O-fucosylpeptide 3-beta-N-acetylglucosaminyltransferase
LGR4	55366	NP_060960	4462	leucine-rich repeat-containing G protein-coupled receptor 4

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
LGR5	8549	NP_003658	2866	leucine-rich repeat-containing G protein-coupled receptor 5
LGR6	59352	NP_001017403	641	leucine-rich repeat-containing G protein-coupled receptor 6
LGR6	59352	NP_001017404	642	leucine-rich repeat-containing G protein-coupled receptor 6
LGR6	59352	NP_067649	4813	leucine-rich repeat-containing G protein-coupled receptor 6
LHCGR	3973	NP_000224	74	luteinizingHormone/choriogonadotropin receptor
LHFP	10186	NP_005771	3367	lipomaHMGIC fusion partner
LHFPL1	340596	NP_835469	6527	lipomaHMGIC fusion partner-like 1
LHFPL2	10184	NP_005770	3366	lipomaHMGIC fusion partner-like 2
LHFPL3	375612	NP_945351	6850	lipomaHMGIC fusion partner-like 3
LHFPL4	375323	NP_940962	6815	lipomaHMGIC fusion partner-like 4
LHFPL5	222662	NP_872354	6656	lipomaHMGIC fusion partner-like 5
LIFR	3977	NP_001121143	1345	leukemia inhibitory factor receptor alpha
LIFR	3977	NP_002301	2618	leukemia inhibitory factor receptor alpha
LILRB1	10859	NP_001075106	1050	leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 1
LILRB1	10859	NP_001075107	1051	leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 1
LILRB1	10859	NP_001075108	1052	leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 1
LILRB1	10859	NP_006660	3537	leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 1
LILRB2	10288	NP_001074447	1044	leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 2
LILRB2	10288	NP_005865	3392	leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 2
LILRB3	11025	NP_001074919	1048	leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 3
LILRB3	11025	NP_006855	3586	leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 3
LILRB4	11006	NP_001265355.2	1045	leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 4
LILRB4	11006	NP_001265355.2	3579	leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 4
LILRB5	10990	NP_001074911	1046	leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 5
LILRB5	10990	NP_001074912	1047	leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 5
LILRB5	10990	NP_006831	3576	leukocyte immunoglobulin-like receptor, subfamily B (with TM and ITIM domains), member 5
LIM2	3982	NP_001155220	2026	lens intrinsic membrane protein 2, 19kDa
LIM2	3982	NP_085915	5156	lens intrinsic membrane protein 2, 19kDa
LINGO1	84894	NP_116197	5406	leucine rich repeat and Ig domain containing 1
LINGO2	158038	NP_689783	6082	leucine rich repeat and Ig domain containing 2
LINGO3	645191	NP_001094861	1202	leucine rich repeat and Ig domain containing 3
LITAF	9516	NP_001129944.1	7399	lipopolysaccharide induced TNF factor
LMAN2	10960	NP_006807	3571	lectin, mannose-binding 2
LMAN2L	81562	NP_001135764	1635	lectin, mannose-binding 2-like
LMAN2L	81562	NP_110432	5180	lectin, mannose-binding 2-like
LMBR1	64327	NP_071903	4920	limb region 1Homolog (mouse)
LMBR1L	55716	NP_060583	4395	limb region 1Homolog (mouse)-like
LMBRD1	55788	NP_060838.3	7414	LMBR1 domain containing 1
LMBRD2	92255	NP_001007528	521	LMBR1 domain containing 2
LMF1	64788	NP_073610	4945	lipase maturation factor 1
LMF2	91289	NP_149977	5468	lipase maturation factor 2
LMLN	89782	NP_001129521	1594	leishmanolysin-like (metallopeptidase M8 family)
LMLN	89782	NP_149018	5440	leishmanolysin-like (metallopeptidase M8 family)
LMO7	4008	NP_005349	3276	LIM domain 7
LMO7	4008	NP_056667	4123	LIM domain 7
LMTK2	22853	NP_055731	4011	lemur tyrosine kinase 2
LMTK3	114783	NP_001073903	1022	lemur tyrosine kinase 3
LNPEP	4012	NP_005566	3316	leucyl/cystinyl aminopeptidase
LNPEP	4012	NP_787116	6470	leucyl/cystinyl aminopeptidase
LPAR1	1902	NP_001392	2361	lysophosphatidic acid receptor 1
LPAR1	1902	NP_476500	5577	lysophosphatidic acid receptor 1
LPAR2	9170	NP_004711.2	7283	lysophosphatidic acid receptor 2
LPAR3	23566	NP_036284	3714	lysophosphatidic acid receptor 3
LPAR4	2846	NP_005287	3262	lysophosphatidic acid receptor 4
LPAR5	57121	NP_001136433	1709	lysophosphatidic acid receptor 5
LPAR5	57121	NP_065133	4646	lysophosphatidic acid receptor 5
LPAR6	10161	NP_001155969	2030	purinergic receptor P2Y, G-protein coupled, 5
LPAR6	10161	NP_001155970	2031	purinergic receptor P2Y, G-protein coupled, 5
LPAR6	10161	NP_005758	3361	purinergic receptor P2Y, G-protein coupled, 5
LPCAT1	79888	NP_079106	5089	lysophosphatidylcholine acyltransferase 1
LPCAT2	54947	NP_060309	4354	lysophosphatidylcholine acyltransferase 2
LPCAT3	10162	NP_005759	3362	lysophosphatidylcholine acyltransferase 3
LPCAT4	254531	NP_705841	6203	lysophosphatidylcholine acyltransferase 4
LPGAT1	9926	NP_055688	4003	lysophosphatidylglycerol acyltransferase 1
LPPR1	54886	NP_060223	4336	plasticity related gene 3
LPPR1	54886	NP_997182	7003	plasticity related gene 3
LPPR2	64748	NP_001164106	2250	lipid phosphate phosphatase-related protein type 2
LPPR2	64748	NP_073574	4939	lipid phosphate phosphatase-related protein type 2
LPPR3	79948	NP_079164	5096	plasticity-related gene 2
LPPR4	9890	NP_001159724	2156	plasticity related gene 1
LPPR4	9890	NP_055654	3996	plasticity related gene 1
LPPR5	163404	NP_001010861	570	phosphatidic acid phosphatase type 2
LPPR5	163404	NP_001032394	773	phosphatidic acid phosphatase type 2
LRCH1	23143	NP_001157683	2067	leucine-rich repeats and calponinHomology (CH) domain containing 1
LRCH1	23143	NP_001157685	2068	leucine-rich repeats and calponinHomology (CH) domain containing 1
LRCH1	23143	NP_055931	4035	leucine-rich repeats and calponinHomology (CH) domain containing 1
LRCH3	84859	NP_116162	5397	leucine-rich repeats and calponinHomology (CH) domain containing 3

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
LRFN1	57622	NP_065913	4735	leucine rich repeat and fibronectin type III domain containing 1
LRFN2	57497	NP_065788	4711	leucine rich repeat and fibronectin type III domain containing 2
LRFN3	79414	NP_078785	5037	leucine rich repeat and fibronectin type III domain containing 3
LRFN4	78999	NP_076941	4999	leucine rich repeat and fibronectin type III domain containing 4
LRFN5	145581	NP_689660	6060	leucine rich repeat and fibronectin type III domain containing 5
LRIG1	26018	NP_056356	4097	leucine-rich repeats and immunoglobulin-like domains 1
LRIG2	9860	NP_001299615.1	7456	leucine rich repeats and immunoglobulin like domains 2
LRIG3	121227	NP_001129523	1595	leucine-rich repeats and immunoglobulin-like domains 3
LRIG3	121227	NP_700356	6188	leucine-rich repeats and immunoglobulin-like domains 3
LRIT1	26103	NP_056428	4107	leucine-rich repeat, immunoglobulin-like and transmembrane domains 1
LRMP	4033	NP_006143	3447	lymphoid-restricted membrane protein
LRP1	4035	NP_002323	2619	low density lipoprotein-related protein 1 (alpha-2-macroglobulin receptor)
LRP10	26020	NP_054764	3862	low density lipoprotein receptor-related protein 10
LRP11	84918	NP_116221	5414	low density lipoprotein receptor-related protein 11
LRP12	29967	NP_001129175	1555	low density lipoprotein-related protein 12
LRP12	29967	NP_038465	3822	low density lipoprotein-related protein 12
LRP1B	53353	NP_061027	4467	low density lipoprotein-related protein 1B (deleted in tumors)
LRP2	4036	NP_004516	3069	low density lipoprotein-related protein 2
LRP3	4037	NP_002324	2620	low density lipoprotein receptor-related protein 3
LRP4	4038	NP_002325	2621	low density lipoprotein receptor-related protein 4
LRP5	4041	NP_002326	2622	low density lipoprotein receptor-related protein 5
LRP8	7804	NP_001018064	664	low density lipoprotein receptor-related protein 8, apolipoprotein e receptor
LRP8	7804	NP_004622	3092	low density lipoprotein receptor-related protein 8, apolipoprotein e receptor
LRP8	7804	NP_059992	4300	low density lipoprotein receptor-related protein 8, apolipoprotein e receptor
LRP8	7804	NP_150643	5482	low density lipoprotein receptor-related protein 8, apolipoprotein e receptor
LRPAP1	4043	NP_002328	2623	low density lipoprotein receptor-related protein associated protein 1
LRRC15	131578	NP_001128529	1517	leucine rich repeat containing 15
LRRC15	131578	NP_570843	5668	leucine rich repeat containing 15
LRRC19	64922	NP_075052	4959	leucine rich repeat containing 19
LRRC25	126364	NP_660299	5943	leucine rich repeat containing 25
LRRC3	81543	NP_112153	5186	leucine rich repeat containing 3
LRRC32	2615	NP_001122394	1379	leucine rich repeat containing 32
LRRC32	2615	NP_005503	3303	leucine rich repeat containing 32
LRRC37A3	374819	NP_955372	6878	leucine rich repeat containing 37, member A3
LRRC37B	114659	NP_443120	5528	similar toHCG1991475 leucine rich repeat containing 37B
LRRC3B	116135	NP_443185	5548	leucine rich repeat containing 3B
LRRC4	64101	NP_071426	4884	leucine rich repeat containing 4
LRRC4B	94030	NP_001073926	1026	leucine rich repeat containing 4B
LRRC4C	57689	NP_065980	4742	leucine rich repeat containing 4C
LRRC52	440699	NP_001005214	449	leucine rich repeat containing 52
LRRC55	219527	NP_001005210	448	leucine rich repeat containing 55
LRRC59	55379	NP_060979	4463	leucine rich repeat containing 59
LRRC66	339977	NP_001019782	679	leucine rich repeat containing 66
LRRC70	100130733	NP_852607	6604	leucine rich repeat containing 70
LRRC8A	56262	NP_001120716	1315	leucine rich repeat containing 8 family, member A
LRRC8A	56262	NP_001120717	1316	leucine rich repeat containing 8 family, member A
LRRC8A	56262	NP_062540	4572	leucine rich repeat containing 8 family, member A
LRRC8B	23507	NP_001127948	1497	leucine rich repeat containing 8 family, member B
LRRC8B	23507	NP_056165	4067	leucine rich repeat containing 8 family, member B
LRRC8C	84230	NP_115646	5315	leucine rich repeat containing 8 family, member C
LRRC8D	55144	NP_001127951	1499	leucine rich repeat containing 8 family, member D
LRRC8D	55144	NP_060573	4392	leucine rich repeat containing 8 family, member D
LRRC8E	80131	NP_079337	5115	leucine rich repeat containing 8 family, member E
LRRN1	57633	NP_065924	4737	leucine rich repeat neuronal 1
LRRN2	10446	NP_006329	3470	leucine rich repeat neuronal 2
LRRN2	10446	NP_963924	6916	leucine rich repeat neuronal 2
LRRN3	54674	NP_001093128	1165	leucine rich repeat neuronal 3
LRRN3	54674	NP_001093130	1166	leucine rich repeat neuronal 3
LRRN3	54674	NP_060804	4423	leucine rich repeat neuronal 3
LRRN4	164312	NP_689824	6089	leucine rich repeat neuronal 4
LRRN4CL	221091	NP_981967	6935	LRRN4 C-terminal like
LRRTM1	347730	NP_849161	6570	leucine rich repeat transmembrane neuronal 1
LRRTM2	26045	NP_056379	4102	leucine rich repeat transmembrane neuronal 2
LRRTM3	347731	NP_821079.3	7362	leucine rich repeat transmembrane neuronal 3
LRRTM4	80059	NP_001128217	1505	leucine rich repeat transmembrane neuronal 4
LRRTM4	80059	NP_079269	5113	leucine rich repeat transmembrane neuronal 4
LRTM1	57408	NP_065729	4697	leucine-rich repeats and transmembrane domains 1
LRTM2	654429	NP_001034118	786	leucine-rich repeats and transmembrane domains 2
LRTM2	654429	NP_001157397	2060	leucine-rich repeats and transmembrane domains 2
LRTM2	654429	NP_001157398	2061	leucine-rich repeats and transmembrane domains 2
LRTOMT	220074	NP_001138779	1849	leucine rich transmembrane and 0-methyltransferase domain containing
LRTOMT	220074	NP_001138780	1850	leucine rich transmembrane and 0-methyltransferase domain containing
LRTOMT	220074	NP_001138781	1851	leucine rich transmembrane and 0-methyltransferase domain containing
LRTOMT	220074	NP_001138782	1852	leucine rich transmembrane and 0-methyltransferase domain containing
LRTOMT	220074	NP_660352	5960	leucine rich transmembrane and 0-methyltransferase domain containing
LSAMP	4045	NP_002329	2624	limbic system-associated membrane protein
LSMEM1	286006	NP_001127940	1496	chromosome 7 open reading frame 53

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
LSMEM1	286006	NP_872403	6665	chromosome 7 open reading frame 53
LSMEM2	132228	NP_694947	6160	chromosome 3 open reading frame 45
LSR	51599	NP_057009	4137	lipolysis stimulated lipoprotein receptor
LSR	51599	NP_991403	6950	lipolysis stimulated lipoprotein receptor
LSR	51599	NP_991404	6951	lipolysis stimulated lipoprotein receptor
LST1	7940	NP_001160010	2173	leukocyte specific transcript 1
LST1	7940	NP_009092	3637	leukocyte specific transcript 1
LST1	7940	NP_995309	6952	leukocyte specific transcript 1
LST1	7940	NP_995310	6953	leukocyte specific transcript 1
LST1	7940	NP_995311	6954	leukocyte specific transcript 1
LST1	7940	NP_995312	6955	leukocyte specific transcript 1
LTA	4049	NP_000586	170	lymphotoxin alpha (TNF superfamily, member 1)
LTA	4049	NP_001153212	1958	lymphotoxin alpha (TNF superfamily, member 1)
LTB	4050	NP_002332	2625	lymphotoxin beta (TNF superfamily, member 3)
LTB	4050	NP_033666	3694	lymphotoxin beta (TNF superfamily, member 3)
LTB4R	1241	NP_001137391	1736	leukotriene B4 receptor
LTB4R	1241	NP_858043	6619	leukotriene B4 receptor
LTB4R2	56413	NP_001158164	2096	leukotriene B4 receptor 2
LTB4R2	56413	NP_001158164.1	2097	leukotriene B4 receptor 2
LTB4R2	56413	NP_062813	4580	leukotriene B4 receptor 2
LTBP3	4054	NP_001123616	1421	latent transforming growth factor beta binding protein 3
LTBP3	4054	NP_001157738	2072	latent transforming growth factor beta binding protein 3
LTBP3	4054	NP_066548	4761	latent transforming growth factor beta binding protein 3
LTBR	4055	NP_002333	2626	lymphotoxin beta receptor (TNFR superfamily, member 3)
LTC4S	4056	NP_665874	5985	leukotriene C4 synthase
LTF	4057	NP_002334	2627	lactotransferrin
LTK	4058	NP_001129157	1552	leukocyte receptor tyrosine kinase
LTK	4058	NP_002335	2628	leukocyte receptor tyrosine kinase
LTK	4058	NP_996844	6992	leukocyte receptor tyrosine kinase
LVRN	206338	NP_776161	6386	laeverin
LY6D	8581	NP_003686	2870	lymphocyte antigen 6 complex, locus D
LY6G5C	80741	NP_079538	5145	lymphocyte antigen 6 complex, locus G5C
LY6G6C	80740	NP_079537	5144	lymphocyte antigen 6 complex, locus G6C
LY6G6D	58530	NP_067069.2	417	lymphocyte antigen 6 complex, locus G6F
LY6G6D	58530	NP_067069	4802	lymphocyte antigen 6 complex, locus G6F
LY6G6F	259215	NP_001003693	416	lymphocyte antigen 6 complex, locus G6F
LY6G6F	259215	NP_001003693.1	4801	lymphocyte antigen 6 complex, locus G6F
LY6K	54742	NP_001153826	1992	lymphocyte antigen 6 complex, locus K
LY6K	54742	NP_001153827	1993	lymphocyte antigen 6 complex, locus K
LY6K	54742	NP_059997	4302	lymphocyte antigen 6 complex, locus K
LY75	4065	NP_002340	2630	CD302 molecule lymphocyte antigen 75
LY75	4065	NP_055695	4007	CD302 molecule lymphocyte antigen 75
LY9	4063	NP_001028839	755	lymphocyte antigen 9
LY9	4063	NP_002339	2629	lymphocyte antigen 9
LYPD1	116372	NP_001070895	970	LY6/PLAUR domain containing 1
LYPD1	116372	NP_653187	5869	LY6/PLAUR domain containing 1
LYPD6B	130576	NP_808879	6509	LY6/PLAUR domain containing 6B
LYPLA1	10434	NP_006321	3468	lysophospholipase I
LYSMD3	116068	NP_938014	6764	LysM, putative peptidoglycan-binding, domain containing 3
LYSMD4	145748	NP_689662	6061	LysM, putative peptidoglycan-binding, domain containing 4
LYVE1	10894	NP_006682	3547	lymphatic vessel endothelialHyaluronan receptor 1
M6PR	4074	NP_002346	2633	mannose-6-phosphate receptor (cation dependent)
MACF1	23499	NP_036222	3702	microtubule-actin crosslinking factor 1
MACF1	23499	NP_036222.3	5441	microtubule-actin crosslinking factor 1
MADCAM1	8174	NP_570116	5660	mucosal vascular addressin cell adhesion molecule 1
MADCAM1	8174	NP_570118	5661	mucosal vascular addressin cell adhesion molecule 1
MAG	4099	NP_002352	2634	myelin associated glycoprotein
MAG	4099	NP_542167	5609	myelin associated glycoprotein
MAGEA1	4100	NP_004979	3191	melanoma antigen family A, 1 (directs expression of antigen MZ2-E)
MAGEA11	4110	NP_001011544.1	7346	MAGE family member A11
MAGEA3	4102	NP_005353.1	7279	MAGE family member A3
MAGEA4	4103	NP_001011548.1	7341	MAGE family member A4
MAGEH1	28986	NP_054780	3867	melanoma antigen familyH, 1
MAL	4118	NP_002362	2635	mal, T-cell differentiation protein
MAL	4118	NP_071883	4914	mal, T-cell differentiation protein
MAL	4118	NP_071884	4915	mal, T-cell differentiation protein
MAL	4118	NP_071885	4916	mal, T-cell differentiation protein
MAL2	114569	NP_443118	5527	mal, T-cell differentiation protein 2
MALRD1	340895	XP_001716895	7055	chromosome 10 open reading frame 112
MALRD1	340895	XP_295865	7087	chromosome 10 open reading frame 112
MALRD1	340895	XP_944452	7093	chromosome 10 open reading frame 112
MAMDC4	158056	NP_996803	6981	MAM domain containing 4
MAN1A1	4121	NP_005898	3397	mannosidase, alpha, class 1A, member 1
MAN1A2	10905	NP_006690	3550	mannosidase, alpha, class 1A, member 2
MAN1B1	11253	NP_057303	4185	mannosidase, alpha, class 1B, member 1
MAN1C1	57134	NP_065112	4641	mannosidase, alpha, class 1C, member 1
MAN2A1	4124	NP_002363	2636	mannosidase, alpha, class 2A, member 1

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
MAN2A2	4122	NP_006113	3438	mannosidase, alpha, class 2A, member 2
MANBAL	63905	NP_001003897	425	mannosidase, beta A, lysosomal-like similar to mannosidase, beta A, lysosomal-like
MANBAL	63905	NP_071360	4873	mannosidase, beta A, lysosomal-like similar to mannosidase, beta A, lysosomal-like
MANEA	79694	NP_078917	5062	mannosidase, endo-alpha
MANEAL	149175	NP_001026910	732	mannosidase, endo-alpha-like
MANEAL	149175	NP_001106954	1258	mannosidase, endo-alpha-like
MANEAL	149175	NP_689709	6072	mannosidase, endo-alpha-like
MANSC1	54682	NP_060520	4384	MANSC domain containing 1
MAOB	4129	NP_000889	317	monoamine oxidase B
MARC1	64757	NP_073583	4942	MOCO sulphurase C-terminal domain containing 1
MARCH11	441061	NP_001096032	1209	membrane-associated ring finger (C3HC4) 11
MARCH2	54996	NP_060368	4366	MOCO sulphurase C-terminal domain containing 2
MARCH4	57574	NP_065865	4726	membrane-associated ring finger (C3HC4) 4
MARCH5	54708	NP_060294	4350	membrane-associated ring finger (C3HC4) 5
MARCH6	10299	NP_005876	3393	membrane-associated ring finger (C3HC4) 6
MARCH6	115123	NP_848545	6541	membrane-associated ring finger (C3HC4) 3
MARCH8	220972	NP_001002265	396	membrane-associated ring finger (C3HC4) 8
MARCH8	220972	NP_001002266	397	membrane-associated ring finger (C3HC4) 8
MARCH8	220972	NP_659458	5918	membrane-associated ring finger (C3HC4) 8
MARCH9	92979	NP_612405	5743	membrane-associated ring finger (C3HC4) 9
MARCO	8685	NP_006761	3560	macrophage receptor with collagenous structure
MARS	4141	NP_004981	3192	methionyl-tRNA synthetase
MARVELD1	83742	NP_113672	5234	hypothetical LOC100270710 MARVEL domain containing 1
MARVELD2	153562	NP_001033692	783	MARVEL domain containing 2
MARVELD2	153562	NP_001033692.2	5900	MARVEL domain containing 2
MARVELD3	91862	NP_001017967	653	MARVEL domain containing 3
MARVELD3	91862	NP_443090	5518	MARVEL domain containing 3
MAS1	4142	NP_002368	2637	MAS1 oncogene
MAS1L	116511	NP_443199	5553	MAS1 oncogene-like
MAVS	57506	NP_065797	4714	mitochondrial antiviral signaling protein
MBOAT1	154141	NP_001073949	1030	membrane bound O-acyltransferase domain containing 1
MBOAT2	129642	NP_620154	5798	membrane bound O-acyltransferase domain containing 2
MBOAT7	79143	NP_001139528	1900	membrane bound O-acyltransferase domain containing 7
MBOAT7	79143	NP_001139554	1905	membrane bound O-acyltransferase domain containing 7
MBOAT7	79143	NP_001139555	1906	membrane bound O-acyltransferase domain containing 7
MBOAT7	79143	NP_077274	5017	membrane bound O-acyltransferase domain containing 7
MBTPS2	51360	NP_056968	4128	membrane-bound transcription factor peptidase, site 2
MC1R	4157	NP_002377	2638	tubulin, beta 3 melanocortin 1 receptor (alpha melanocyte stimulatingHormone receptor)
MC1R	4157	NP_006077	3431	tubulin, beta 3 melanocortin 1 receptor (alpha melanocyte stimulatingHormone receptor)
MC2R	4158	NP_000520	153	melanocortin 2 receptor (adrenocorticotropicHormone)
MC3R	4159	NP_063941.3	7386	melanocortin 3 receptor
MC4R	4160	NP_005903	3398	melanocortin 4 receptor
MC5R	4161	NP_005904	3399	melanocortin 5 receptor
MCAM	4162	NP_006491	3500	melanoma cell adhesion molecule
MCEMP1	199675	NP_777578	6409	chromosome 19 open reading frame 59
MCHR1	2847	NP_005288	3263	melanin-concentratingHormone receptor 1
MCHR2	84539	NP_001035269	853	melanin-concentratingHormone receptor 2
MCHR2	84539	NP_115892	5363	melanin-concentratingHormone receptor 2
MCL1	4170	NP_068779	4856	myeloid cell leukemia sequence 1 (BCL2-related)
MCL1	4170	NP_877495	6681	myeloid cell leukemia sequence 1 (BCL2-related)
MCOLN1	57192	NP_065394	4681	mucolipin 1
MCOLN2	255231	NP_694991	6167	mucolipin 2
MCOLN3	55283	NP_060768	4417	mucolipin 3
MCTP1	79772	NP_001002796	404	multiple C2 domains, transmembrane 1
MCTP1	79772	NP_078993	5070	multiple C2 domains, transmembrane 1
MCTP2	55784	NP_001153115	1949	multiple C2 domains, transmembrane 2
MCTP2	55784	NP_001153116	1950	multiple C2 domains, transmembrane 2
MCTP2	55784	NP_060819	4427	multiple C2 domains, transmembrane 2
MCU	90550	NP_612366	5733	coiled-coil domain containing 109A
MCUR1	63933	NP_001026883	726	coiled-coil domain containing 90A
MDGA2	161357	NP_001106970	1259	MAM domain containing glycosylphosphatidylinositol anchor 2
MDGA2	161357	NP_878250	6687	MAM domain containing glycosylphosphatidylinositol anchor 2
MDM1	56890	NP_059136	4292	Mdm1 nuclear proteinHomolog (mouse)
MDM1	56890	NP_064513	4600	Mdm1 nuclear proteinHomolog (mouse)
MEGF10	84466	NP_115822	5355	multiple EGF-like-domains 10
MEGF11	84465	NP_115821	5354	multiple EGF-like-domains 11
MEGF6	1953	NP_001400	2366	multiple EGF-like-domains 6
MEGF8	1954	NP_001401	2367	multiple EGF-like-domains 8
MEGF9	1955	NP_001073966	1031	multiple EGF-like-domains 9
MEP1A	4224	NP_005579	3319	meprin A, alpha (PABA peptideHydrolase)
MEP1B	4225	NP_005916	3400	meprin A, beta
MERTK	10461	NP_006334	3471	c-mer proto-oncogene tyrosine kinase
MET	4233	NP_000236	76	met proto-oncogene (hepatocyte growth factor receptor)
MET	4233	NP_001120972	1332	met proto-oncogene (hepatocyte growth factor receptor)
METTL21A	151194	NP_001120867	1325	family with sequence similarity 119, member A
METTL21A	151194	NP_660323	5951	family with sequence similarity 119, member A
METTL7A	25840	NP_054752	3858	methyltransferase like 7A

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
METTL7B	196410	NP_689850	6091	methyltransferase like 7B
MFAP3	4238	NP_001128509	1514	microfibrillar-associated protein 3
MFAP3	4238	NP_005918	3401	microfibrillar-associated protein 3
MFAP3L	9848	NP_001009554	561	microfibrillar-associated protein 3-like
MFAP3L	9848	NP_067679	4816	microfibrillar-associated protein 3-like
MFF	56947	NP_064579	4615	mitochondrial fission factor
MFN2	9927	NP_001121132	1341	mitofusin 2
MFN2	9927	NP_055689	4004	mitofusin 2
MFSD1	64747	NP_001161375	2203	major facilitator superfamily domain containing 1
MFSD1	64747	NP_073573	4938	major facilitator superfamily domain containing 1
MFSD10	10227	NP_001111	1280	major facilitator superfamily domain containing 10
MFSD10	10227	NP_001139541	1901	major facilitator superfamily domain containing 10
MFSD11	79157	NP_077287	5022	major facilitator superfamily domain containing 11
MFSD12	126321	NP_001274458.1	915	chromosome 19 open reading frame 28
MFSD12	126321	NP_001274458.1	4823	chromosome 19 open reading frame 28
MFSD12	126321	NP_778148	6434	chromosome 19 open reading frame 28
MFSD2A	84879	NP_001129965	1606	major facilitator superfamily domain containing 2
MFSD2A	84879	NP_116182	5402	major facilitator superfamily domain containing 2
MFSD3	113655	NP_612440	5750	major facilitator superfamily domain containing 3
MFSD4	148808	NP_857595	6617	major facilitator superfamily domain containing 4
MFSD5	84975	NP_001164261.1	7415	major facilitator superfamily domain containing 5
MFSD6	54842	NP_060164	4324	major facilitator superfamily domain containing 6
MFSD6L	162387	NP_689812	6087	major facilitator superfamily domain containing 6-like
MFSD7	84179	NP_115595	5308	major facilitator superfamily domain containing 7
MFSD8	256471	NP_689991	6115	major facilitator superfamily domain containing 8
MFSD9	84804	NP_116107	5389	major facilitator superfamily domain containing 9
MGAM	8972	NP_004659	3099	maltase-glucoamylase (alpha-glucosidase)
MGAM2	93432	NP_001280555	7117	maltase-glucoamylase 2 (putative)
MGARP	84709	NP_116012	5380	chromosome 4 open reading frame 49
MGAT1	4245	NP_001108089	1274	mannosyl (alpha-1,3-)-glycoprotein beta-1,2-N-acetylglucosaminyltransferase
MGAT1	4245	NP_001108090	1275	mannosyl (alpha-1,3-)-glycoprotein beta-1,2-N-acetylglucosaminyltransferase
MGAT1	4245	NP_001108091	1276	mannosyl (alpha-1,3-)-glycoprotein beta-1,2-N-acetylglucosaminyltransferase
MGAT1	4245	NP_001108092	1277	mannosyl (alpha-1,3-)-glycoprotein beta-1,2-N-acetylglucosaminyltransferase
MGAT1	4245	NP_002397	2642	mannosyl (alpha-1,3-)-glycoprotein beta-1,2-N-acetylglucosaminyltransferase
MGAT2	4247	NP_002399	2643	mannosyl (alpha-1,6-)-glycoprotein beta-1,2-N-acetylglucosaminyltransferase
MGAT3	4248	NP_001091740	1122	mannosyl (beta-1,4-)-glycoprotein beta-1,4-N-acetylglucosaminyltransferase
MGAT3	4248	NP_002400	2644	mannosyl (beta-1,4-)-glycoprotein beta-1,4-N-acetylglucosaminyltransferase
MGAT4A	11320	NP_001153626	1976	mannosyl (alpha-1,3-)-glycoprotein beta-1,4-N-acetylglucosaminyltransferase, isozyme A
MGAT4A	11320	NP_036346	3723	mannosyl (alpha-1,3-)-glycoprotein beta-1,4-N-acetylglucosaminyltransferase, isozyme A
MGAT4B	11282	NP_055090	3906	mannosyl (alpha-1,3-)-glycoprotein beta-1,4-N-acetylglucosaminyltransferase, isozyme B
MGAT4B	11282	NP_463459	5562	mannosyl (alpha-1,3-)-glycoprotein beta-1,4-N-acetylglucosaminyltransferase, isozyme B
MGAT4D	152586	XP_001720251	7068	similar to RIKEN cDNA 4933434I20
MGAT4D	152586	XP_001720300	7069	similar to RIKEN cDNA 4933434I20
MGAT4D	152586	XP_001725525	7074	similar to RIKEN cDNA 4933434I20
MGAT5	4249	NP_002401	2645	mannosyl (alpha-1,6-)-glycoprotein beta-1,6-N-acetyl-glucosaminyltransferase
MGAT5B	146664	NP_653278	5890	mannosyl (alpha-1,6-)-glycoprotein beta-1,6-N-acetyl-glucosaminyltransferase, isozyme B
MGAT5B	146664	NP_945193	6847	mannosyl (alpha-1,6-)-glycoprotein beta-1,6-N-acetyl-glucosaminyltransferase, isozyme B
MGP	4256	NP_000891	319	matrix Gla protein
MGST1	4257	NP_064696	4626	microsomal glutathione S-transferase 1
MGST1	4257	NP_665707	5975	microsomal glutathione S-transferase 1
MGST1	4257	NP_665734	5976	microsomal glutathione S-transferase 1
MGST1	4257	NP_665735	5977	microsomal glutathione S-transferase 1
MGST2	4258	NP_002404	2646	microsomal glutathione S-transferase 2
MIA3	375056	NP_940953	6814	melanoma inhibitory activity family, member 3
MICA	100507436	NP_000238	7109	MHC class I polypeptide-related sequence A isoform 1
MICU3	286097	NP_859074	6625	EF-hand domain family, member A2
MID1	4281	NP_000372	112	midline 1 (Opitz/BBB syndrome)
MID1	4281	NP_001092094	1142	midline 1 (Opitz/BBB syndrome)
MID1	4281	NP_150632	5481	midline 1 (Opitz/BBB syndrome)
MIEF2	125170	NP_001138372	1782	Smith-Magenis syndrome chromosome region, candidate 7
MIEF2	125170	NP_631901	5839	Smith-Magenis syndrome chromosome region, candidate 7
MIEF2	125170	NP_683684	6003	Smith-Magenis syndrome chromosome region, candidate 7
MILR1	284021	NP_001078892	1093	chromosome 17 open reading frame 60
MINOS1	440574	NP_001027535	743	chromosome 1 open reading frame 151
MIP	4284	NP_036196	3696	major intrinsic protein of lens fiber
MLANA	2315	NP_005502	3302	melan-A
MLC1	23209	NP_055981	4042	megalencephalic leukoencephalopathy with subcortical cysts 1
MLC1	23209	NP_631941	5847	megalencephalic leukoencephalopathy with subcortical cysts 1
MLEC	9761	NP_055545	3978	malectin
MLF2	8079	NP_005430	3285	myeloid leukemia factor 2
MLLT10	8028	NP_001182555	563	myeloid/lymphoid or mixed-lineage leukemia (trithoraxHomolog, Drosophila)
MLLT10	8028	NP_004632	3095	myeloid/lymphoid or mixed-lineage leukemia (trithoraxHomolog, Drosophila)
MLN	4295	NP_001035198	837	motilin
MLN	4295	NP_002409	2649	motilin
MLNR	2862	NP_001498	2396	motilin receptor
MMD	23531	NP_036461	3748	monocyte to macrophage differentiation-associated
MMD2	221938	NP_001094070	1190	monocyte to macrophage differentiation-associated 2

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
MMD2	221938	NP_940685	6792	monocyte to macrophage differentiation-associated 2
MME	4311	NP_000893	320	membrane metallo-endopeptidase
MME	4311	NP_009218	3669	membrane metallo-endopeptidase
MME	4311	NP_009219	3670	membrane metallo-endopeptidase
MME	4311	NP_009220	3671	membrane metallo-endopeptidase
MMEL1	79258	NP_258428	5501	membrane metallo-endopeptidase-like 1
MMGT1	93380	NP_775741	6342	membrane magnesium transporter 1
MMP14	4323	NP_004986.1	7277	matrix metallopeptidase 14
MMP15	4324	NP_002419	2651	matrix metallopeptidase 15 (membrane-inserted)
MMP16	4325	NP_005932	3403	matrix metallopeptidase 16 (membrane-inserted)
MMP16	4325	NP_005932.2	4929	matrix metallopeptidase 16 (membrane-inserted)
MMP20	9313	NP_004762	3120	matrix metallopeptidase 20
MMP24	10893	NP_006681	3546	matrix metallopeptidase 24 (membrane-inserted)
MOG	4340	NP_001008229	528	myelin oligodendrocyte glycoprotein
MOG	4340	NP_001008230	529	myelin oligodendrocyte glycoprotein
MOG	4340	NP_001163888	2240	myelin oligodendrocyte glycoprotein
MOG	4340	NP_001008229.1	2241	myelin oligodendrocyte glycoprotein
MOG	4340	NP_002424	2652	myelin oligodendrocyte glycoprotein
MOG	4340	NP_996532	6966	myelin oligodendrocyte glycoprotein
MOG	4340	NP_996533	6967	myelin oligodendrocyte glycoprotein
MOG	4340	NP_996534	6968	myelin oligodendrocyte glycoprotein
MOG	4340	NP_996535	6969	myelin oligodendrocyte glycoprotein
MOG	4340	NP_996536	6970	myelin oligodendrocyte glycoprotein
MOG	4340	NP_996537	6971	myelin oligodendrocyte glycoprotein
MOGAT1	116255	NP_477513	5580	monoacylglycerol O-acyltransferase 1
MOGAT3	346606	NP_835470	6528	monoacylglycerol O-acyltransferase 3
MOGS	7841	NP_001139630	1909	mannosyl-oligosaccharide glucosidase
MOGS	7841	NP_006293	3463	mannosyl-oligosaccharide glucosidase
MOK	5891	NP_001258940.1	7436	MOK protein kinase
MOS	4342	NP_005363	3277	v-mos Moloney murine sarcoma viral oncogeneHomolog
MOSPD1	56180	NP_062456	4571	motile sperm domain containing 1
MOSPD2	158747	NP_689794	6084	motile sperm domain containing 2
MOSPD3	64598	NP_001035186	828	motile sperm domain containing 3
MOSPD3	64598	NP_001035187	829	motile sperm domain containing 3
MOSPD3	64598	NP_001035188	830	motile sperm domain containing 3
MOSPD3	64598	NP_076438	4989	motile sperm domain containing 3
MPC1	51660	NP_057182	4169	brain protein 44-like
MPDU1	9526	NP_004861	3154	mannose-P-dolichol utilization defect 1
MPEG1	219972	NP_001034485	792	macrophage expressed 1
MPG	4350	NP_001015052	632	N-methylpurine-DNA glycosylase
MPG	4350	NP_001015054	633	N-methylpurine-DNA glycosylase
MPG	4350	NP_002425	2653	N-methylpurine-DNA glycosylase
MPL	4352	NP_005364	3278	myeloproliferative leukemia virus oncogene
MPPE1	65258	NP_075563	4972	metallophosphoesterase 1
MPZ	4359	NP_000521	154	myelin protein zero
MPZL1	9019	NP_001139663	1912	myelin protein zero-like 1
MPZL1	9019	NP_003944	2929	myelin protein zero-like 1
MPZL1	9019	NP_078845	5049	myelin protein zero-like 1
MPZL2	10205	NP_005788	3371	myelin protein zero-like 2
MPZL2	10205	NP_658911	5902	myelin protein zero-like 2
MPZL3	196264	NP_938016	6765	myelin protein zero-like 3
MR1	3140	NP_001522	2406	majorHistocompatibility complex, class I-related
MRAP	56246	NP_848932	6562	melanocortin 2 receptor accessory protein
MRAP	56246	NP_996781	6977	melanocortin 2 receptor accessory protein
MRAP2	112609	NP_612418	5744	melanocortin 2 receptor accessory protein 2
MRGPRF	116535	NP_001091985	1129	MAS-related GPR, member F
MRGPRF	116535	NP_659452	5917	MAS-related GPR, member F
MRGPRX1	259249	NP_671732	5999	MAS-related GPR, member X1
MRGPRX2	117194	NP_473371	5570	MAS-related GPR, member X2
MRGPRX3	117195	NP_473372	5571	MAS-related GPR, member X3
MRGPRX4	117196	NP_473373	5572	MAS-related GPR, member X4
MRLN	100507027	NP_001291660	7105	myoregulin
MRS2	57380	NP_065713	4694	MRS2 magnesiumHomeostasis factorHomolog (S. cerevisiae)
MRVI1	10335	NP_001092049	1139	murine retrovirus integration site 1Homolog
MRVI1	10335	NP_001093633	1182	murine retrovirus integration site 1Homolog
MRVI1	10335	NP_001093637	1183	murine retrovirus integration site 1Homolog
MRVI1	10335	NP_569056	5649	murine retrovirus integration site 1Homolog
MS4A1	931	NP_068769	4853	membrane-spanning 4-domains, subfamily A, member 1
MS4A1	931	NP_690605	6123	membrane-spanning 4-domains, subfamily A, member 1
MS4A12	54860	NP_001157942	2082	membrane-spanning 4-domains, subfamily A, member 12
MS4A12	54860	NP_060186	4328	membrane-spanning 4-domains, subfamily A, member 12
MS4A14	84689	NP_001073160	992	membrane-spanning 4-domains, subfamily A, member 14
MS4A14	84689	NP_115986	5376	membrane-spanning 4-domains, subfamily A, member 14
MS4A15	219995	NP_001092305	1153	membrane-spanning 4-domains, subfamily A, member 15
MS4A15	219995	NP_689930	6104	membrane-spanning 4-domains, subfamily A, member 15
MS4A2	2206	NP_000130	38	membrane-spanning 4-domains, subfamily A, member 2 (Fc fragment of IgE,High affinity I, receptor for beta polypeptide)

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
MS4A2	2206	NP_000130.1	1637	membrane-spanning 4-domains, subfamily A, member 2 (Fc fragment of IgE,High affinity I, receptor for beta polypeptide)
MS4A3	932	NP_001026836	721	membrane-spanning 4-domains, subfamily A, member 3 (hematopoietic cell-specific)
MS4A3	932	NP_001026979	735	membrane-spanning 4-domains, subfamily A, member 3 (hematopoietic cell-specific)
MS4A3	932	NP_006129	3443	membrane-spanning 4-domains, subfamily A, member 3 (hematopoietic cell-specific)
MS4A4A	51338	NP_076926	4994	membrane-spanning 4-domains, subfamily A, member 4
MS4A4A	51338	NP_683876	6017	membrane-spanning 4-domains, subfamily A, member 4
MS4A5	64232	NP_076434	4988	membrane-spanning 4-domains, subfamily A, member 5
MS4A6A	64231	NP_071744	4900	membrane-spanning 4-domains, subfamily A, member 6A
MS4A6A	64231	NP_690590	6120	membrane-spanning 4-domains, subfamily A, member 6A
MS4A6A	64231	NP_690591	6121	membrane-spanning 4-domains, subfamily A, member 6A
MS4A6E	245802	NP_640342	5850	membrane-spanning 4-domains, subfamily A, member 6E
MS4A7	58475	NP_067024	4794	membrane-spanning 4-domains, subfamily A, member 7
MS4A7	58475	NP_996821	6983	membrane-spanning 4-domains, subfamily A, member 7
MS4A7	58475	NP_996822	6984	membrane-spanning 4-domains, subfamily A, member 7
MS4A7	58475	NP_996823	6985	membrane-spanning 4-domains, subfamily A, member 7
MS4A8	83661	NP_113645	5227	membrane-spanning 4-domains, subfamily A, member 8B
MSLN	10232	NP_005814	3377	mesothelin
MSLN	10232	NP_037536	3819	mesothelin
MSMB	4477	NP_002434	2654	microseminoprotein, beta-
MSMB	4477	NP_619540	5768	microseminoprotein, beta-
MSMO1	6307	NP_001017369.1	7348	methylsterol monooxygenase 1
MSR1	4481	NP_002436	2655	macrophage scavenger receptor 1
MSR1	4481	NP_619729	5774	macrophage scavenger receptor 1
MSR1	4481	NP_619730	5775	macrophage scavenger receptor 1
MST1R	4486	NP_002438	2656	macrophage stimulating 1 receptor (c-met-related tyrosine kinase)
MTCH1	23787	NP_055156	3919	mitochondrial carrierHomolog 1 (C. elegans)
MTDH	92140	NP_848927	6561	metadherin
MTFP1	51537	NP_001003704	418	mitochondrial protein 18 kDa
MTFP1	51537	NP_057582	4229	mitochondrial protein 18 kDa
MTHFD1	4522	NP_005947	3405	methylenetetrahydrofolate dehydrogenase (NADP+ dependent) 1, methenyltetrahydrofolate cyclohydrolase, formyltetrahydrofolate synthetase
MTNR1A	4543	NP_005949	3406	melatonin receptor 1A
MTNR1B	4544	NP_005950	3407	melatonin receptor 1B
MTUS1	57509	NP_001001924	379	mitochondrial tumor suppressor 1
MTUS1	57509	NP_001001925	380	mitochondrial tumor suppressor 1
MTUS1	57509	NP_001001931	381	mitochondrial tumor suppressor 1
MTUS1	57509	NP_001159865	2165	mitochondrial tumor suppressor 1
MTUS1	57509	NP_065800	4715	mitochondrial tumor suppressor 1
MUC1	4582	NP_001018016	661	mucin 1, cell surface associated
MUC1	4582	NP_001018017	662	mucin 1, cell surface associated
MUC1	4582	NP_001037855	922	mucin 1, cell surface associated
MUC1	4582	NP_001037856	923	mucin 1, cell surface associated
MUC1	4582	NP_001037857	924	mucin 1, cell surface associated
MUC1	4582	NP_001037858	925	mucin 1, cell surface associated
MUC1	4582	NP_002447	2657	mucin 1, cell surface associated
MUC12	10071	NP_001157934	2080	mucin 12, cell surface associated similar to mucin 11
MUC13	56667	NP_149038	5442	mucin 13, cell surface associated
MUC15	143662	NP_001128563	1518	mucin 15, cell surface associated
MUC15	143662	NP_001128564	1519	mucin 15, cell surface associated
MUC15	143662	NP_663625	5966	mucin 15, cell surface associated
MUC16	94025	NP_078966	5067	mucin 16, cell surface associated
MUC17	140453	NP_001035194	835	mucin 17, cell surface associated
MUC3A	4584	XP_001125753	7049	mucin 3B, cell surface associated
MUC3A	4584	XP_001718007	7058	mucin 3B, cell surface associated
MUC3A	4584	XP_001719378	7065	mucin 3B, cell surface associated
MUC3A	4584	XP_001719551	7066	mucin 3B, cell surface associated
MUC3A	4584	XP_001721642	7070	mucin 3B, cell surface associated
MUC3A	4584	XP_001725406	7072	mucin 3B, cell surface associated
MUC4	4585	NP_004523	3070	mucin 4, cell surface associated
MUC4	4585	NP_060876	4440	mucin 4, cell surface associated
MUC4	4585	NP_612154	5722	mucin 4, cell surface associated
MUL1	79594	NP_078820	5044	mitochondrial E3 ubiquitin ligase 1
MUSK	4593	NP_001159752.1	7413	muscle associated receptor tyrosine kinase
MXRA7	439921	NP_001008528	544	matrix-remodelling associated 7
MXRA7	439921	NP_001008529	545	matrix-remodelling associated 7
MXRA7	439921	NP_940932	6808	matrix-remodelling associated 7
MYADM	91663	NP_001018654	671	myeloid-associated differentiation marker
MYADM	91663	NP_001018655	672	myeloid-associated differentiation marker
MYADM	91663	NP_001018656	673	myeloid-associated differentiation marker
MYADM	91663	NP_001018657	674	myeloid-associated differentiation marker
MYADM	91663	NP_612382	5736	myeloid-associated differentiation marker
MYADML2	255275	NP_001138585	1815	myeloid-associated differentiation marker-like 2
MYCT1	80177	NP_079383	5120	myc target 1
MYOF	26509	NP_038479	3826	myoferlin
MYOF	26509	NP_579899	5688	myoferlin
MYRF	745	NP_001120864	1323	chromosome 11 open reading frame 9

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
MYRF	745	NP_037411	3794	chromosome 11 open reading frame 9
MYRFL	196446	XP_001716702	7054	chromosome 12 open reading frame 28
MYRFL	196446	XP_001718110	7059	chromosome 12 open reading frame 28
MYRFL	196446	XP_001718960	7063	chromosome 12 open reading frame 28
N4BP2L2	10443	NP_055702	4008	NEDD4 binding protein 2-like 2
N4BP2L2	10443	NP_149102	5451	NEDD4 binding protein 2-like 2
NAAA	27163	NP_001035861	885	N-acylethanolamine acid amidase
NAAA	27163	NP_055250	3937	N-acylethanolamine acid amidase
NAALAD2	10003	NP_005458	3289	N-acetylated alpha-linked acidic dipeptidase 2
NAALADL1	10004	NP_005459	3290	N-acetylated alpha-linked acidic dipeptidase-like 1
NAALADL2	254827	NP_996898	6997	N-acetylated alpha-linked acidic dipeptidase-like 2
NALCN	259232	NP_443099	5520	sodium leak channel, non-selective
NAT10	55226	NP_001137502	1759	N-acetyltransferase 10 (GCN5-related)
NAT10	55226	NP_078938	5066	N-acetyltransferase 10 (GCN5-related)
NAT14	57106	NP_065111	4640	N-acetyltransferase 14 (GCN5-related, putative)
NAT2	10	NP_000006	1	N-acetyltransferase 2 (arylamine N-acetyltransferase)
NAT8	9027	NP_003951	2930	N-acetyltransferase 8 (GCN5-related, putative)
NAT8	9027	NP_057431	4205	N-acetyltransferase 8 (GCN5-related, putative)
NAT8L	339983	NP_848652	6555	N-acetyltransferase 8-like (GCN5-related, putative)
NBAS	51594	NP_056993	4131	neuroblastoma amplified sequence
NCAM1	4684	NP_000606	173	neural cell adhesion molecule 1
NCAM1	4684	NP_001070150	949	neural cell adhesion molecule 1
NCAM1	4684	NP_851996	6595	neural cell adhesion molecule 1
NCAM2	4685	NP_004531.2	7327	neural cell adhesion molecule 2
NCEH1	57552	NP_001139748	1924	arylacetamide deacetylase-like 1
NCEH1	57552	NP_001139749	1925	arylacetamide deacetylase-like 1
NCEH1	57552	NP_001139750	1926	arylacetamide deacetylase-like 1
NCEH1	57552	NP_065843	4724	arylacetamide deacetylase-like 1
NCR1	9437	NP_001138929	1860	natural cytotoxicity triggering receptor 1
NCR1	9437	NP_001138930	1861	natural cytotoxicity triggering receptor 1
NCR1	9437	NP_004820	3144	natural cytotoxicity triggering receptor 1
NCR2	9436	NP_004819	3143	natural cytotoxicity triggering receptor 2
NCR3	259197	NP_001138938	1866	natural cytotoxicity triggering receptor 3
NCR3	259197	NP_001138939	1867	natural cytotoxicity triggering receptor 3
NCR3	259197	NP_667341	5989	natural cytotoxicity triggering receptor 3
NCR3LG1	374383	NP_001189368.1	7428	natural killer cell cytotoxicity receptor 3 ligand 1
NCSTN	23385	NP_056146	4062	nicastrin
NDC1	55706	NP_001162023	2234	transmembrane protein 48
NDC1	55706	NP_060557	4389	transmembrane protein 48
NDFIP2	54602	NP_001154879	2007	Nedd4 family interacting protein 2
NDFIP2	54602	NP_061953	4561	Nedd4 family interacting protein 2
NDRG1	10397	NP_001128714	1539	N-myc downstream regulated 1
NDRG1	10397	NP_006087	3434	N-myc downstream regulated 1
NDRG2	57447	NP_057334	4194	NDRG family member 2
NDRG2	57447	NP_963293	6902	NDRG family member 2
NDRG2	57447	NP_963294	6903	NDRG family member 2
NDRG2	57447	NP_963831	6904	NDRG family member 2
NDRG2	57447	NP_963832	6905	NDRG family member 2
NDRG2	57447	NP_963833	6906	NDRG family member 2
NDRG2	57447	NP_963834	6907	NDRG family member 2
NDRG2	57447	NP_963835	6908	NDRG family member 2
NDRG4	65009	NP_001123959	1430	NDRG family member 4
NDRG4	65009	NP_065198	4664	NDRG family member 4
NDRG4	65009	NP_075061	4961	NDRG family member 4
NDST1	3340	NP_001534	2409	N-deacetylase/N-sulfotransferase (heparan glucosaminyl) 1
NDST2	8509	NP_003626	2858	N-deacetylase/N-sulfotransferase (heparan glucosaminyl) 2
NDST3	9348	NP_004775	3126	N-deacetylase/N-sulfotransferase (heparan glucosaminyl) 3
NDST4	64579	NP_072091	4931	N-deacetylase/N-sulfotransferase (heparan glucosaminyl) 4
NDUFA1	4694	NP_004532	3071	NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 1, 7.5kDa
NDUFA11	126328	NP_783313	6446	NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 11, 14.7kDa
NDUFA13	51079	NP_057049	4143	NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 13
NDUFA3	4696	NP_004533	3072	NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 3, 9kDa
NDUFA4	4697	NP_002480	2658	NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 4, 9kDa
NDUFA4L2	56901	NP_064527	4603	NADH dehydrogenase (ubiquinone) 1 alpha subcomplex, 4-like 2
NDUFB1	4707	NP_004536	3073	NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 1, 7kDa
NDUFB11	54539	NP_001129470	1582	NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 11, 17.3kDa
NDUFB11	54539	NP_061929	4553	NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 11, 17.3kDa
NDUFB3	4709	NP_002482	2659	NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 3, 12kDa
NDUFB5	4711	NP_002483	2660	NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 5, 16kDa
NDUFB6	4712	NP_002484	2661	NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 6, 17kDa
NDUFB6	4712	NP_877416	6679	NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 6, 17kDa
NDUFB8	4714	NP_004995	3194	NADH dehydrogenase (ubiquinone) 1 beta subcomplex, 8, 19kDa
NDUFC2	4718	NP_004540	3074	NADH dehydrogenase (ubiquinone) 1, subcomplex unknown, 2, 14.5kDa
NELL2	4753	NP_001138579	1811	NEL-like 2 (chicken)
NELL2	4753	NP_001138580	1812	NEL-like 2 (chicken)
NELL2	4753	NP_001138581	1813	NEL-like 2 (chicken)
NELL2	4753	NP_001138582	1814	NEL-like 2 (chicken)

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
NELL2	4753	NP_006150	3448	NEL-like 2 (chicken)
NEMP1	23306	NP_001124435	1451	transmembrane protein 194A
NEMP1	23306	NP_056072	4053	transmembrane protein 194A
NEMP2	100131211	NP_001136117	1681	transmembrane protein 194B
NEO1	4756	NP_002490	2662	neogeninHomolog 1 (chicken)
NETO1	81832	NP_001188394.1	7427	neuropilin and tolloid like 1
NETO2	81831	NP_060562	4390	neuropilin (NRP) and tolloid (TLL)-like 2
NFAM1	150372	NP_666017	5986	NFAT activating protein with ITAM motif 1
NFASC	23114	NP_001005388	456	neurofascinHomolog (chicken)
NFASC	23114	NP_001005389	457	neurofascinHomolog (chicken)
NFASC	23114	NP_001153803	1989	neurofascinHomolog (chicken)
NFASC	23114	NP_001153804	1990	neurofascinHomolog (chicken)
NFASC	23114	NP_001153805	1991	neurofascinHomolog (chicken)
NFASC	23114	NP_055905	4031	neurofascinHomolog (chicken)
NFE2L3	9603	NP_004280	3009	nuclear factor (erythroid-derived 2)-like 3
NFXL1	152518	NP_694540	6142	nuclear transcription factor, X-box binding-like 1
NGFR	4804	NP_002498	2663	nerve growth factor receptor (TNFR superfamily, member 16)
NGLY1	55768	NP_001138765	1845	N-glycanase 1
NGLY1	55768	NP_001138766	1846	N-glycanase 1
NGLY1	55768	NP_001138767	1847	N-glycanase 1
NGLY1	55768	NP_060767	4416	N-glycanase 1
NINJ1	4814	NP_004139	2973	ninjurin 1
NINJ2	4815	NP_057617	4238	ninjurin 2
NIPA1	123606	NP_001135747	1630	non imprinted in Prader-Willi/Angelman syndrome 1
NIPA1	123606	NP_653200	5872	non imprinted in Prader-Willi/Angelman syndrome 1
NIPA2	81614	NP_001008860	555	non imprinted in Prader-Willi/Angelman syndrome 2
NIPA2	81614	NP_001008892	556	non imprinted in Prader-Willi/Angelman syndrome 2
NIPA2	81614	NP_001008894	557	non imprinted in Prader-Willi/Angelman syndrome 2
NIPA2	81614	NP_112184	5192	non imprinted in Prader-Willi/Angelman syndrome 2
NIPAL2	79815	NP_079035	5077	NIPA-like domain containing 2
NIPAL3	57185	NP_065181	4661	NIPA-like domain containing 3
NIPAL4	348938	NP_001092757	1154	NIPA-like domain containing 4
NIPAL4	348938	NP_001165763	2308	NIPA-like domain containing 4
NKAIN1	79570	NP_078798	5039	Na+/K+ transporting ATPase interacting 1
NKAIN2	154215	NP_001035304	864	Na+/K+ transporting ATPase interacting 2
NKAIN4	128414	NP_690603	6122	Na+/K+ transporting ATPase interacting 4
NKG7	4818	NP_005592	3321	natural killer cell group 7 sequence
NKPD1	284353	NP_940880.3	1469	NTPase, KAP family P-loop domain containing 1
NKPD1	284353	NP_940880	6799	NTPase, KAP family P-loop domain containing 1
NLGN1	22871	NP_055747	4016	neuroligin 1
NLGN2	57555	NP_065846.1	7325	neuroligin 2
NLGN3	54413	NP_001160132	2174	neuroligin 3
NLGN3	54413	NP_061850	4543	neuroligin 3
NLGN3	54413	NP_851820	6589	neuroligin 3
NLGN4X	57502	NP_001269074.1	7441	neuroligin 4, X-linked
NLGN4Y	22829	NP_001157710	2069	neuroligin 4, Y-linked
NLGN4Y	22829	NP_055708	4009	neuroligin 4, Y-linked
NMB	4828	NP_066563	4764	neuromedin B
NMB	4828	NP_995580	6961	neuromedin B
NMBR	4829	NP_002502	2665	neuromedin B receptor
NMU	10874	NP_006672	3544	neuromedin U
NMUR1	10316	NP_006047	3423	neuromedin U receptor 1
NMUR2	56923	NP_064552	4610	neuromedin U receptor 2
NNAT	4826	NP_005377	3279	neuronatin
NNAT	4826	NP_859017	6620	neuronatin
NNT	23530	NP_036475	3752	nicotinamide nucleotide transhydrogenase
NNT	23530	NP_892022	6698	nicotinamide nucleotide transhydrogenase
NOTCH2	4853	NP_077719	5027	NotchHomolog 2 (Drosophila)
NOTCH4	4855	NP_004548	3075	NotchHomolog 4 (Drosophila)
NOX1	27035	NP_008983	3623	NADPH oxidase 1
NOX1	27035	NP_039249	3833	NADPH oxidase 1
NOX3	50508	NP_056533.1	7294	NADPH oxidase 3
NOX4	50507	NP_001137308.1	7404	NADPH oxidase 4
NOX5	79400	NP_078781	5035	NADPH oxidase, EF-hand calcium binding domain 5
NPB	256933	NP_683694	6004	neuropeptide B
NPBWR1	2831	NP_005276	3253	neuropeptides B/W receptor 1
NPBWR2	2832	NP_005277	3254	neuropeptides B/W receptor 2
NPC1	4864	NP_000262	78	Niemann-Pick disease, type C1
NPC1L1	29881	NP_001095118	1203	NPC1 (Niemann-Pick disease, type C1, gene)-like 1
NPC1L1	29881	NP_037521	3817	NPC1 (Niemann-Pick disease, type C1, gene)-like 1
NPDC1	56654	NP_056207	4075	neural proliferation, differentiation and control, 1
NPFFR1	64106	NP_071429	4886	neuropeptide FF receptor 1
NPFFR2	10886	NP_001138228	1771	neuropeptide FF receptor 2
NPFFR2	10886	NP_004876	3160	neuropeptide FF receptor 2
NPFFR2	10886	NP_444264	5558	neuropeptide FF receptor 2
NPHP4	261734	NP_055917	4032	nephronophthisis 4
NPHS1	4868	NP_004637	3096	nephrosis 1, congenital, Finnish type (nephrin)

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
NPHS2	7827	NP_055440	3966	nephrosis 2, idiopathic, steroid-resistant (podocin)
NPPA	4878	NP_006163	3449	natriuretic peptide precursor A
NPR3	4883	NP_000899	322	natriuretic peptide receptor C/guanylate cyclase C (atrionatriuretic peptide receptor C)
NPTN	27020	NP_001154835	2004	neuroplastin
NPTN	27020	NP_001154836	2005	neuroplastin
NPTN	27020	NP_036560	3774	neuroplastin
NPTN	27020	NP_059429	4295	neuroplastin
NPTXR	23467	NP_055108	3907	neuronal pentraxin receptor
NPY	4852	NP_000896	321	neuropeptide Y
NPY1R	4886	NP_000900	323	neuropeptide Y receptor Y1
NPY2R	4887	NP_000901.1	7262	neuropeptide Y receptor Y2
NPY4R	5540	NP_005963	3409	pancreatic polypeptide receptor 1
NPY5R	4889	NP_006165	3450	neuropeptide Y receptor Y5
NRCAM	4897	NP_001032209	768	neuronal cell adhesion molecule
NRCAM	4897	NP_001032209.1	769	neuronal cell adhesion molecule
NRCAM	4897	NP_005001	3195	neuronal cell adhesion molecule
NRG1	3084	NP_001153467	1964	neuregulin 1
NRG1	3084	NP_001153468	1965	neuregulin 1
NRG1	3084	NP_001153471	1966	neuregulin 1
NRG1	3084	NP_001153473	1967	neuregulin 1
NRG1	3084	NP_001153474	1968	neuregulin 1
NRG1	3084	NP_001153476	1969	neuregulin 1
NRG1	3084	NP_001153477	1970	neuregulin 1
NRG1	3084	NP_001153479	1971	neuregulin 1
NRG1	3084	NP_001153480	1972	neuregulin 1
NRG1	3084	NP_004486	3065	neuregulin 1
NRG1	3084	NP_039250	3834	neuregulin 1
NRG1	3084	NP_039251	3835	neuregulin 1
NRG1	3084	NP_039252	3836	neuregulin 1
NRG1	3084	NP_039253	3837	neuregulin 1
NRG1	3084	NP_039254	3838	neuregulin 1
NRG1	3084	NP_001153467.1	3839	neuregulin 1
NRG1	3084	NP_039256	3840	neuregulin 1
NRG1	3084	NP_039258	3841	neuregulin 1
NRG2	9542	NP_004874	3158	neuregulin 2
NRG2	9542	NP_053584	3845	neuregulin 2
NRG2	9542	NP_053585	3846	neuregulin 2
NRG2	9542	NP_053586	3847	neuregulin 2
NRG3	10718	NP_001010848	569	neuregulin 3
NRG3	10718	NP_001159444	2135	neuregulin 3
NRG3	10718	NP_001159445	2136	neuregulin 3
NRG4	145957	NP_612640	5761	neuregulin 4
NRK	203447	NP_940867	6797	Nik related kinase
NRM	11270	NP_009174	3658	nurim (nuclear envelope membrane protein)
NRP1	8829	NP_001019799	680	neuropilin 1
NRP1	8829	NP_001019800	681	neuropilin 1
NRP1	8829	NP_003864	2923	neuropilin 1
NRP2	8828	NP_003863	2922	neuropilin 2
NRP2	8828	NP_061004	4464	neuropilin 2
NRP2	8828	NP_957716	6892	neuropilin 2
NRP2	8828	NP_957718	6893	neuropilin 2
NRP2	8828	NP_957719	6894	neuropilin 2
NRP2	8828	NP_958436	6895	neuropilin 2
NRROS	375387	NP_940967	6817	leucine rich repeat containing 33
NRSN1	140767	NP_542454	5624	neurensin 1
NRSN2	80023	NP_079234	5108	neurensin 2
NRXN1	9378	NP_001129131	1545	neurexin 1
NRXN1	9378	NP_004792	3131	neurexin 1
NRXN1	9378	NP_620072	5783	neurexin 1
NRXN2	9379	NP_055895	4030	neurexin 2
NRXN2	9379	NP_620060	5781	neurexin 2
NRXN2	9379	NP_620063	5782	neurexin 2
NRXN3	9369	NP_001098720	1222	neurexin 3
NRXN3	9369	NP_004787	3129	neurexin 3
NRXN3	9369	NP_620426	5810	neurexin 3
NSDHL	50814	NP_001123237	1383	NAD(P) dependent steroid dehydrogenase-like
NSDHL	50814	NP_057006	4136	NAD(P) dependent steroid dehydrogenase-like
NSG1	27065	NP_001035190	832	DNA segment on chromosome 4 (unique) 234 expressed sequence
NSG1	27065	NP_055207	3929	DNA segment on chromosome 4 (unique) 234 expressed sequence
NT5C3A	51251	NP_001002009	385	5'-nucleotidase, cytosolic III
NT5C3A	51251	NP_001002010	386	5'-nucleotidase, cytosolic III
NT5C3A	51251	NP_001159590	2150	5'-nucleotidase, cytosolic III
NT5C3A	51251	NP_057573	4227	5'-nucleotidase, cytosolic III
NT5E	4907	NP_002517	2666	5'-nucleotidase, ecto (CD73)
NTF3	4908	NP_001096124	1211	neurotrophin 3
NTF3	4908	NP_002518	2667	neurotrophin 3
NTM	50863	NP_001041674	929	neurotrimin

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
NTM	50863	NP_001137530	1766	neurotrimin
NTM	50863	NP_001137531	1767	neurotrimin
NTM	50863	NP_057606	4234	neurotrimin
NTN1	9423	NP_004813	3138	netrin 1
NTN4	59277	NP_067052	4798	netrin 4
NTRK1	4914	NP_001007793	525	neurotrophic tyrosine kinase, receptor, type 1
NTRK1	4914	NP_001012331	592	neurotrophic tyrosine kinase, receptor, type 1
NTRK1	4914	NP_002520	2668	neurotrophic tyrosine kinase, receptor, type 1
NTRK2	4915	NP_001007098	506	neurotrophic tyrosine kinase, receptor, type 2
NTRK2	4915	NP_001018074	665	neurotrophic tyrosine kinase, receptor, type 2
NTRK2	4915	NP_001018075	666	neurotrophic tyrosine kinase, receptor, type 2
NTRK2	4915	NP_001018076	667	neurotrophic tyrosine kinase, receptor, type 2
NTRK2	4915	NP_006171	3451	neurotrophic tyrosine kinase, receptor, type 2
NTRK3	4916	NP_001007157	507	neurotrophic tyrosine kinase, receptor, type 3
NTRK3	4916	NP_001012338	593	neurotrophic tyrosine kinase, receptor, type 3
NTRK3	4916	NP_002521	2669	neurotrophic tyrosine kinase, receptor, type 3
NTSR1	4923	NP_002522	2670	neurotensin receptor 1 (high affinity)
NTSR2	23620	NP_036476	3753	neurotensin receptor 2
NUCB1	4924	NP_006175	3453	nucleobindin 1
NUCB2	4925	NP_005004.1	7278	nucleobindin 2
NUDC	10726	NP_006591	3526	nuclear distribution gene CHomolog (A. nidulans)
NUP210	23225	NP_079199	5103	nucleoporin 210kDa
NUP210L	91181	NP_001152956	1943	nucleoporin 210kDa-like
NUP210L	91181	NP_997191	7005	nucleoporin 210kDa-like
NUP50	10762	NP_009103	3643	nucleoporin 50kDa
NUP50	10762	NP_705931	6208	nucleoporin 50kDa
NXPE1	120400	NP_689528	6032	family with sequence similarity 55, member A
NXPE3	91775	NP_001127928	1495	family with sequence similarity 55, member C
NXPE3	91775	NP_659474	5922	family with sequence similarity 55, member C
NXPE4	54827	NP_001071107	975	family with sequence similarity 55, member D
NXPE4	54827	NP_060148	4322	family with sequence similarity 55, member D
NYX	60506	NP_072089	4930	nyctalopin
OAS1	4938	NP_001027581.1	7354	2'-5'-oligoadenylate synthetase 1
OCA2	4948	NP_000266	80	oculocutaneous albinism II
OCEL1	79629	NP_078854	5051	occludin/ELL domain containing 1
OCIAD1	54940	NP_001073308	999	OCIA domain containing 1
OCIAD1	54940	NP_001073309	1000	OCIA domain containing 1
OCIAD1	54940	NP_001073310	1001	OCIA domain containing 1
OCIAD1	54940	NP_001073311	1002	OCIA domain containing 1
OCIAD1	54940	NP_001161726	2212	OCIA domain containing 1
OCIAD1	54940	NP_060300	4351	OCIA domain containing 1
OCLM	10896	NP_071770	4910	oculomedin
ODF4	146852	NP_694552	6148	outer dense fiber of sperm tails 4
OGFOD3	79701	NP_078924	5064	chromosome 17 open reading frame 101
OGFOD3	79701	NP_787098	6467	chromosome 17 open reading frame 101
OLFML2B	25903	NP_056256	4083	olfactomedin-like 2B
OLR1	4973	NP_002534	2671	oxidized low density lipoprotein (lectin-like) receptor 1
OMA1	115209	NP_660286	5938	OMA1Homolog, zinc metallopeptidase (S. cerevisiae)
OMG	4974	NP_002535	2672	oligodendrocyte myelin glycoprotein
OPALIN	93377	NP_001035191	833	oligodendrocytic myelin paranodal and inner loop protein
OPALIN	93377	NP_001035192	834	oligodendrocytic myelin paranodal and inner loop protein
OPALIN	93377	NP_149984	5469	oligodendrocytic myelin paranodal and inner loop protein
OPCML	4978	NP_001012393	594	opioid binding protein/cell adhesion molecule-like
OPCML	4978	NP_002536	2673	opioid binding protein/cell adhesion molecule-like
OPN1SW	611	NP_001699	2441	opsin 1 (cone pigments), short-wave-sensitive
OPN3	23596	NP_055137	3914	opsin 3
OPN4	94233	NP_001025186	713	opsin 4
OPN4	94233	NP_150598	5480	opsin 4
OPN5	221391	NP_859528.1	714	opsin 5
OPN5	221391	NP_859528	6627	opsin 5
OPRD1	4985	NP_000902	324	opioid receptor, delta 1
OPRK1	4986	NP_000903	325	opioid receptor, kappa 1
OPRL1	4987	NP_000904	326	opiate receptor-like 1
OPRL1	4987	NP_872588	6670	opiate receptor-like 1
OPRM1	4988	NP_000905	327	opioid receptor, mu 1
OPRM1	4988	NP_001008503	541	opioid receptor, mu 1
OPRM1	4988	NP_001008504	542	opioid receptor, mu 1
OPRM1	4988	NP_001008505	543	opioid receptor, mu 1
OPRM1	4988	NP_001138751	1833	opioid receptor, mu 1
OPRM1	4988	NP_001138752	1834	opioid receptor, mu 1
OPRM1	4988	NP_001138753	1835	opioid receptor, mu 1
OPRM1	4988	NP_001138754	1836	opioid receptor, mu 1
OPRM1	4988	NP_001138755	1837	opioid receptor, mu 1
OPRM1	4988	NP_001138756	1838	opioid receptor, mu 1
OPRM1	4988	NP_001138757	1839	opioid receptor, mu 1
OPRM1	4988	NP_001138758	1840	opioid receptor, mu 1
OPRM1	4988	NP_001138759	1841	opioid receptor, mu 1

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
OR10A3	26496	NP_001003745	422	olfactory receptor, family 10, subfamily A, member 3
OR10A4	283297	NP_997069	7002	olfactory receptor, family 10, subfamily A, member 4
OR10A5	144124	NP_835462	6525	olfactory receptor, family 10, subfamily A, member 5
OR10C1	442194	NP_039229	3832	olfactory receptor, family 10, subfamily C, member 1
OR10H1	26539	NP_039228	3831	olfactory receptor, family 10, subfamilyH, member 1
OR10H2	26538	NP_039227	3830	olfactory receptor, family 10, subfamilyH, member 2
OR10H3	26532	NP_039226	3829	olfactory receptor, family 10, subfamilyH, member 3
OR10J1	26476	NP_036483	3754	olfactory receptor, family 10, subfamily J, member 1
OR11A1	26531	NP_039225	3828	olfactory receptor, family 11, subfamily A, member 1
OR12D2	26529	NP_039224	3827	olfactory receptor, family 12, subfamily D, member 2
OR12D3	81797	NP_112221.1	5182	olfactory receptor, family 12, subfamily D, member 3
OR12D3	81797	NP_112221	5204	olfactory receptor, family 12, subfamily D, member 3
OR14J1	442191	NP_112208	5200	olfactory receptor, family 14, subfamily J, member 1
OR1A1	8383	NP_055380	3956	olfactory receptor, family 1, subfamily A, member 1
OR1A2	26189	NP_036484	3755	olfactory receptor, family 1, subfamily A, member 2
OR1C1	26188	NP_036485	3756	olfactory receptor, family 1, subfamily C, member 1
OR1D2	4991	NP_002539	2674	olfactory receptor, family 1, subfamily D, member 2
OR1F1	4992	NP_036492	3757	olfactory receptor, family 1, subfamily F, member 1
OR1G1	8390	NP_003546	2847	olfactory receptor, family 1, subfamily G, member 1
OR1I1	126370	NP_001004713	439	olfactory receptor, family 1, subfamily I, member 1
OR1J2	26740	NP_473448	5573	olfactory receptor, family 1, subfamily J, member 2
OR1J4	26219	NP_001004452	436	olfactory receptor, family 1, subfamily J, member 4
OR1Q1	158131	NP_036496	3758	olfactory receptor, family 1, subfamily Q, member 1
OR2B2	81697	NP_149046	5446	olfactory receptor, family 2, subfamily B, member 2
OR2B3	442184	NP_001005226	450	olfactory receptor, family 2, subfamily B, member 3
OR2B6	26212	NP_036499	3759	olfactory receptor, family 2, subfamily B, member 6
OR2C1	4993	NP_036500	3760	olfactory receptor, family 2, subfamily C, member 1
OR2C3	81472	NP_932340	6739	olfactory receptor, family 2, subfamily C, member 3
OR2F1	26211	NP_036501	3761	olfactory receptor, family 2, subfamily F, member 1
OR2F2	135948	NP_001004685	437	olfactory receptor, family 2, subfamily F, member 2
OR2H1	26716	NP_112145	5185	olfactory receptor, family 2, subfamilyH, member 1
OR2H2	7932	NP_009091	3636	olfactory receptor, family 2, subfamilyH, member 2
OR2J2	26707	NP_112167	5190	olfactory receptor, family 2, subfamily J, member 2
OR2K2	26248	NP_995581	6962	olfactory receptor, family 2, subfamily K, member 2
OR2L13	284521	NP_787107	6468	olfactory receptor, family 2, subfamily L, member 13
OR2L2	26246	NP_001004686	438	olfactory receptor, family 2, subfamily L, member 2
OR2M4	26245	NP_059974	4297	olfactory receptor, family 2, subfamily M, member 4
OR2S2	56656	NP_063950.2	7385	olfactory receptor family 2 subfamily S member 2 (gene/pseudogene)
OR2W1	26692	NP_112165	5189	olfactory receptor, family 2, subfamily W, member 1
OR3A1	4994	NP_002541	2675	olfactory receptor, family 3, subfamily A, member 1
OR3A2	4995	NP_002542	2676	olfactory receptor, family 3, subfamily A, member 2
OR3A3	8392	NP_036505	3762	olfactory receptor, family 3, subfamily A, member 3
OR4D1	26689	NP_036506	3763	olfactory receptor, family 4, subfamily D, member 1
OR4N4	283694	NP_001005241	451	olfactory receptor, family 4, subfamily N, member 4
OR51B2	79345	NP_149420	5463	olfactory receptor, family 51, subfamily B, member 2
OR51B4	79339	NP_149419	5462	olfactory receptor, family 51, subfamily B, member 4
OR51B5	282763	NP_001005567	463	olfactory receptor, family 51, subfamily B, member 5
OR51B6	390058	NP_001004750	441	olfactory receptor, family 51, subfamily B, member 6
OR51E1	143503	NP_689643.2	7395	olfactory receptor family 51 subfamily E member 1
OR51E2	81285	NP_110401	5168	olfactory receptor, family 51, subfamily E, member 2
OR51I1	390063	NP_001005288	453	olfactory receptor, family 51, subfamily I, member 1
OR51I2	390064	NP_001004754	442	olfactory receptor, family 51, subfamily I, member 2
OR51M1	390059	NP_001004756	443	olfactory receptor, family 51, subfamily M, member 1
OR52A1	23538	NP_036507	3764	olfactory receptor, family 52, subfamily A, member 1
OR52D1	390066	NP_001005163	444	olfactory receptor, family 52, subfamily D, member 1
OR5I1	10798	NP_006628	3533	olfactory receptor, family 5, subfamily I, member 1
OR5J2	282775	NP_001005492	460	olfactory receptor, family 5, subfamily J, member 2
OR5K1	26339	NP_001004736	440	olfactory receptor, family 5, subfamily K, member 1
OR5L2	26338	NP_001004739.1	7337	olfactory receptor family 5 subfamily L member 2
OR5P2	120065	NP_703145	6191	olfactory receptor, family 5, subfamily P, member 2
OR5P3	120066	NP_703146	6192	olfactory receptor, family 5, subfamily P, member 3
OR5V1	81696	NP_110503	5183	olfactory receptor, family 12, subfamily D, member 3
OR5V1	81696	NP_110503.3	5205	olfactory receptor, family 12, subfamily D, member 3
OR6A2	8590	NP_003687.2	7329	olfactory receptor family 6 subfamily A member 2
OR6B1	135946	NP_001005281	452	olfactory receptor, family 6, subfamily B, member 1
OR7A10	390892	NP_001005190	445	olfactory receptor, family 7, subfamily A, member 10
OR7A17	26333	NP_112163	5188	olfactory receptor, family 7, subfamily A, member 17
OR7A5	26659	NP_059976	4298	olfactory receptor, family 7, subfamily A, member 5
OR7C1	26664	NP_945182	6846	olfactory receptor, family 7, subfamily C, member 1
OR7C2	26658	NP_036509	3765	olfactory receptor, family 7, subfamily C, member 2
OR7D2	162998	NP_787079	6465	olfactory receptor, family 7, subfamily D, member 2
OR8B2	26595	NP_001005468	459	olfactory receptor, family 8, subfamily B, member 2
OR8B8	26493	NP_036510	3766	olfactory receptor, family 8, subfamily B, member 8
OR8D1	283159	NP_001002917	406	olfactory receptor, family 8, subfamily D, member 1
OR8D2	283160	NP_001002918	407	olfactory receptor, family 8, subfamily D, member 2
OR8G1	26494	NP_001002905	405	olfactory receptor, family 8, subfamily G, member 5, olfactory receptor, family 8, subfamily G, member 1
OR8G1	26494	NP_001005198	446	olfactory receptor, family 8, subfamily G, member 5, olfactory receptor, family 8, subfamily G, member 1

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
ORAI1	84876	NP_116179.2	7330	ORAI calcium release-activated calcium modulator 1
ORAI2	80228	NP_001119812	1306	ORAI calcium release-activated calcium modulator 2
ORAI2	80228	NP_116220	5413	ORAI calcium release-activated calcium modulator 2
ORAI3	93129	NP_689501	6026	ORAI calcium release-activated calcium modulator 3
ORMDL1	94101	NP_001121622	1363	ORM1-like 1 (S. cerevisiae)
ORMDL1	94101	NP_057551	4224	ORM1-like 1 (S. cerevisiae)
ORMDL2	29095	NP_054901	3876	ORM1-like 2 (S. cerevisiae)
ORMDL3	94103	NP_644809	5851	ORM1-like 3 (S. cerevisiae)
OS9	10956	NP_001017956	650	osteosarcoma amplified 9, endoplasmic reticulum associated protein
OS9	10956	NP_001017957	651	osteosarcoma amplified 9, endoplasmic reticulum associated protein
OS9	10956	NP_001017958	652	osteosarcoma amplified 9, endoplasmic reticulum associated protein
OS9	10956	NP_006803	3569	osteosarcoma amplified 9, endoplasmic reticulum associated protein
OSBPL8	114882	NP_001003712	419	oxysterol binding protein-like 8
OSBPL8	114882	NP_065892	4730	oxysterol binding protein-like 8
OSMR	9180	NP_001161827	2220	oncostatin M receptor
OSMR	9180	NP_003990	2943	oncostatin M receptor
OST4	100128731	NP_001128165	1503	dolichyl-diphosphooligosaccharide--protein glycosyltransferase subunit 4
OSTC	58505	NP_067050	4797	oligosaccharyltransferase complex subunit
OSTM1	28962	NP_054747	3857	osteopetrosis associated transmembrane protein 1
OTOF	9381	NP_004793	3132	otoferlin
OTOF	9381	NP_919224	6710	otoferlin
OTOF	9381	NP_919303	6719	otoferlin
OTOF	9381	NP_919304	6720	otoferlin
OTOR	56914	NP_064542	4607	otoraplin
OXA1L	5018	NP_005006	3197	oxidase (cytochrome c) assembly 1-like
OXER1	165140	NP_683765	6011	oxoeicosanoid (OXE) receptor 1
OXGR1	27199	NP_543008	5632	oxoglutarate (alpha-ketoglutarate) receptor 1
OXTR	5021	NP_000907.2	7326	oxytocin receptor
P2RX1	5023	NP_002549	2678	purinergic receptor P2X, ligand-gated ion channel, 1
P2RX2	22953	NP_036358	3724	purinergic receptor P2X, ligand-gated ion channel, 2
P2RX2	22953	NP_057402	4202	purinergic receptor P2X, ligand-gated ion channel, 2
P2RX2	22953	NP_733782	6246	purinergic receptor P2X, ligand-gated ion channel, 2
P2RX2	22953	NP_733783	6247	purinergic receptor P2X, ligand-gated ion channel, 2
P2RX2	22953	NP_777361	6400	purinergic receptor P2X, ligand-gated ion channel, 2
P2RX2	22953	NP_777362	6401	purinergic receptor P2X, ligand-gated ion channel, 2
P2RX3	5024	NP_002550.2	7323	purinergic receptor P2X 3
P2RX4	5025	NP_002551	2679	purinergic receptor P2X, ligand-gated ion channel, 4
P2RX5	5026	NP_002552	2680	purinergic receptor P2X, ligand-gated ion channel, 5
P2RX5	5026	NP_778255	6441	purinergic receptor P2X, ligand-gated ion channel, 5
P2RX5	5026	NP_778256	6442	purinergic receptor P2X, ligand-gated ion channel, 5
P2RX6	9127	NP_001153026	1945	purinergic receptor P2X, ligand-gated ion channel, 6
P2RX6	9127	NP_005437	3286	purinergic receptor P2X, ligand-gated ion channel, 6
P2RX7	5027	NP_002553	2681	purinergic receptor P2X, ligand-gated ion channel, 7
P2RY1	5028	NP_002554	2682	purinergic receptor P2Y, G-protein coupled, 1
P2RY10	27334	NP_055314	3948	purinergic receptor P2Y, G-protein coupled, 10
P2RY10	27334	NP_938147	6769	purinergic receptor P2Y, G-protein coupled, 10
P2RY12	64805	NP_073625	4946	purinergic receptor P2Y, G-protein coupled, 12
P2RY12	64805	NP_795345	6482	purinergic receptor P2Y, G-protein coupled, 12
P2RY13	53829	NP_795713.2	7394	purinergic receptor P2Y13
P2RY14	9934	NP_001074924	1049	purinergic receptor P2Y, G-protein coupled, 14
P2RY14	9934	NP_055694	4006	purinergic receptor P2Y, G-protein coupled, 14
P2RY2	5029	NP_002555	2683	purinergic receptor P2Y, G-protein coupled, 2
P2RY2	5029	NP_788085	6473	purinergic receptor P2Y, G-protein coupled, 2
P2RY2	5029	NP_788086	6474	purinergic receptor P2Y, G-protein coupled, 2
P2RY4	5030	NP_002556	2684	pyrimidinergic receptor P2Y, G-protein coupled, 4
P2RY6	5031	NP_001264133.1	2974	pyrimidinergic receptor P2Y, G-protein coupled, 6
P2RY6	5031	NP_789766	6477	pyrimidinergic receptor P2Y, G-protein coupled, 6
P2RY6	5031	NP_789767	6478	pyrimidinergic receptor P2Y, G-protein coupled, 6
P2RY6	5031	NP_789768	6479	pyrimidinergic receptor P2Y, G-protein coupled, 6
P2RY8	286530	NP_835230	6518	purinergic receptor P2Y, G-protein coupled, 8
P4HA2	8974	NP_001017973	656	prolyl 4-hydroxylase, alpha polypeptide II
P4HA2	8974	NP_001017974	657	prolyl 4-hydroxylase, alpha polypeptide II
P4HA2	8974	NP_001136070	1675	prolyl 4-hydroxylase, alpha polypeptide II
P4HA2	8974	NP_001136071	1676	prolyl 4-hydroxylase, alpha polypeptide II
P4HA2	8974	NP_004190	2985	prolyl 4-hydroxylase, alpha polypeptide II
P4HTM	54681	NP_808807	6503	prolyl 4-hydroxylase, transmembrane (endoplasmic reticulum)
P4HTM	54681	NP_808808	6504	prolyl 4-hydroxylase, transmembrane (endoplasmic reticulum)
PAAF1	80227	NP_079431	5128	proteasomal ATPase-associated factor 1
PACRG	135138	NP_001073847	1015	PARK2 co-regulated
PACRG	135138	NP_001073848	1016	PARK2 co-regulated
PACRG	135138	NP_689623	6055	PARK2 co-regulated
PAG1	55824	NP_060910	4450	phosphoprotein associated with glycosphingolipid microdomains 1
PAM	5066	NP_000910	328	peptidylglycine alpha-amidating monooxygenase
PAM	5066	NP_620121	5790	peptidylglycine alpha-amidating monooxygenase
PAM	5066	NP_620176	5803	peptidylglycine alpha-amidating monooxygenase
PAM	5066	NP_620177	5804	peptidylglycine alpha-amidating monooxygenase
PANX1	24145	NP_056183	4070	pannexin 1

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
PANX2	56666	NP_001153772	1988	pannexin 2
PANX2	56666	NP_443071	5517	pannexin 2
PANX3	116337	NP_443191	5551	pannexin 3
PAQR3	152559	NP_001035292	863	progestin and adipoQ receptor family member III
PAQR4	124222	NP_689554	6036	progestin and adipoQ receptor family member IV
PAQR5	54852	NP_001098024	1215	progestin and adipoQ receptor family member V
PAQR5	54852	NP_060175	4326	progestin and adipoQ receptor family member V
PAQR6	79957	NP_079173	5099	progestin and adipoQ receptor family member VI
PAQR6	79957	NP_940798	6793	progestin and adipoQ receptor family member VI
PAQR7	164091	NP_848509	6539	progestin and adipoQ receptor family member VII
PAQR8	85315	NP_588608	5689	progestin and adipoQ receptor family member VIII
PAQR9	344838	NP_940906	6804	progestin and adipoQ receptor family member IX
PARL	55486	NP_001032728	780	presenilin associated, rhomboid-like
PARL	55486	NP_061092	4471	presenilin associated, rhomboid-like
PARM1	25849	NP_056208	4076	DKFZP564O0823 protein
PARP16	54956	NP_060321	4357	poly (ADP-ribose) polymerase family, member 16
PCDH1	5097	NP_002578	2686	protocadherin 1
PCDH1	5097	NP_115796	5353	protocadherin 1
PCDH10	57575	NP_065866	4727	protocadherin 10
PCDH10	57575	NP_116586	5428	protocadherin 10
PCDH11X	27328	NP_001161832	2221	protocadherin 11 X-linked
PCDH11X	27328	NP_001161833	2222	protocadherin 11 X-linked
PCDH11X	27328	NP_001161834	2223	protocadherin 11 X-linked
PCDH11X	27328	NP_001161835	2224	protocadherin 11 X-linked
PCDH11X	27328	NP_001161832.1	3953	protocadherin 11 X-linked
PCDH11X	27328	NP_116750.1	5429	protocadherin 11 X-linked
PCDH11X	27328	NP_116750	5430	protocadherin 11 X-linked
PCDH11X	27328	NP_116751	5431	protocadherin 11 X-linked
PCDH11Y	83259	NP_001265548.1	7437	protocadherin 11 Y-linked
PCDH12	51294	NP_057664	4250	protocadherin 12
PCDH15	65217	NP_001136235	1688	protocadherin 15
PCDH15	65217	NP_001136236	1689	protocadherin 15
PCDH15	65217	NP_001136237	1690	protocadherin 15
PCDH15	65217	NP_001136238	1691	protocadherin 15
PCDH15	65217	NP_001136239	1692	protocadherin 15
PCDH15	65217	NP_001136240	1693	protocadherin 15
PCDH15	65217	NP_001136241	1694	protocadherin 15
PCDH15	65217	NP_001136242	1695	protocadherin 15
PCDH15	65217	NP_001136243	1696	protocadherin 15
PCDH15	65217	NP_001136244	1697	protocadherin 15
PCDH15	65217	NP_001136245	1698	protocadherin 15
PCDH15	65217	NP_149045	5445	protocadherin 15
PCDH17	27253	NP_001035519	866	protocadherin 17
PCDH18	54510	NP_061908	4552	protocadherin 18
PCDH19	57526	NP_001098713	1218	protocadherin 19
PCDH19	57526	NP_065817	4718	protocadherin 19
PCDH20	64881	NP_073754	4955	protocadherin 20
PCDH7	5099	NP_002580.2	7303	protocadherin 7
PCDH8	5100	NP_002581	2690	protocadherin 8
PCDH8	5100	NP_116567	5427	protocadherin 8
PCDH9	5101	NP_065136	4648	protocadherin 9
PCDH9	5101	NP_982354	6947	protocadherin 9
PCDHA10	56139	NP_061724.1	4497	protocadherin alpha 13, protocadherin alpha 10, protocadherin alpha subfamily C, 1, C, 2
PCDHA10	56139	NP_061724.1	4500	protocadherin alpha 13, protocadherin alpha 10, protocadherin alpha subfamily C, 1, C, 2
PCDHA10	56139	NP_061724.1	4503	protocadherin alpha 13, protocadherin alpha 10, protocadherin alpha subfamily C, 1, C, 2
PCDHA10	56139	NP_061727.1	4506	protocadherin alpha 13, protocadherin alpha 10, protocadherin alpha subfamily C, 1, C, 2
PCDHA10	56139	NP_114065	5243	protocadherin alpha 13, protocadherin alpha 10, protocadherin alpha subfamily C, 1, C, 2
PCDHA10	56139	NP_114065.1	5246	protocadherin alpha 13, protocadherin alpha 10, protocadherin alpha subfamily C, 1, C, 2
PCDHA10	56139	NP_114071	5249	protocadherin alpha 13, protocadherin alpha 10, protocadherin alpha subfamily C, 1, C, 2
PCDHA10	56139	NP_114088	5253	protocadherin alpha 13, protocadherin alpha 10, protocadherin alpha subfamily C, 1, C, 2
PCDHA10	56139	NP_114089	5256	protocadherin alpha 13, protocadherin alpha 10, protocadherin alpha subfamily C, 1, C, 2
PCDHA2	56146	NP_061728	4509	protocadherin alpha 2
PCDHA2	56146	NP_113683	5236	protocadherin alpha 2
PCDHA5	56143	NP_061731	4510	protocadherin alpha 5
PCDHA5	56143	NP_113689	5237	protocadherin alpha 5
PCDHA6	56142	NP_061732	4511	protocadherin alpha 8 protocadherin alpha 6
PCDHA6	56142	NP_061734	4512	protocadherin alpha 8 protocadherin alpha 6
PCDHA6	56142	NP_114036	5238	protocadherin alpha 8 protocadherin alpha 6
PCDHA6	56142	NP_114037	5239	protocadherin alpha 8 protocadherin alpha 6
PCDHA6	56142	NP_114062	5241	protocadherin alpha 8 protocadherin alpha 6
PCDHA9	9752	NP_054724	3852	protocadherin alpha 9
PCDHA9	9752	NP_114063	5242	protocadherin alpha 9
PCDHAC1	56135	NP_061721	4499	protocadherin alpha 13, protocadherin alpha 10, protocadherin alpha subfamily C, 1, C, 2
PCDHAC1	56135	NP_061722.1	4502	protocadherin alpha 13, protocadherin alpha 10, protocadherin alpha subfamily C, 1, C, 2
PCDHAC1	56135	NP_061721.2	4505	protocadherin alpha 13, protocadherin alpha 10, protocadherin alpha subfamily C, 1, C, 2
PCDHAC1	56135	NP_061721.2	4508	protocadherin alpha 13, protocadherin alpha 10, protocadherin alpha subfamily C, 1, C, 2
PCDHAC1	56135	NP_061721.2	5245	protocadherin alpha 13, protocadherin alpha 10, protocadherin alpha subfamily C, 1, C, 2

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
PCDHAC1	56135	NP_061721.2	5248	protocadherin alpha 13, protocadherin alpha 10, protocadherin alpha subfamily C, 1, C, 2
PCDHAC1	56135	NP_114088.2	5251	protocadherin alpha 13, protocadherin alpha 10, protocadherin alpha subfamily C, 1, C, 2
PCDHAC1	56135	NP_114088	5255	protocadherin alpha 13, protocadherin alpha 10, protocadherin alpha subfamily C, 1, C, 2
PCDHAC1	56135	NP_114089	5258	protocadherin alpha 13, protocadherin alpha 10, protocadherin alpha subfamily C, 1, C, 2
PCDHAC2	56134	NP_061722.1	4498	protocadherin alpha 13, protocadherin alpha 10, protocadherin alpha subfamily C, 1, C, 2
PCDHAC2	56134	NP_061722.1	4501	protocadherin alpha 13, protocadherin alpha 10, protocadherin alpha subfamily C, 1, C, 2
PCDHAC2	56134	NP_061722.1	4504	protocadherin alpha 13, protocadherin alpha 10, protocadherin alpha subfamily C, 1, C, 2
PCDHAC2	56134	NP_061724.1	4507	protocadherin alpha 13, protocadherin alpha 10, protocadherin alpha subfamily C, 1, C, 2
PCDHAC2	56134	NP_061722.1	5244	protocadherin alpha 13, protocadherin alpha 10, protocadherin alpha subfamily C, 1, C, 2
PCDHAC2	56134	NP_114089.1	5247	protocadherin alpha 13, protocadherin alpha 10, protocadherin alpha subfamily C, 1, C, 2
PCDHAC2	56134	NP_061722.1	5250	protocadherin alpha 13, protocadherin alpha 10, protocadherin alpha subfamily C, 1, C, 2
PCDHAC2	56134	NP_114088	5254	protocadherin alpha 13, protocadherin alpha 10, protocadherin alpha subfamily C, 1, C, 2
PCDHAC2	56134	NP_114089	5257	protocadherin alpha 13, protocadherin alpha 10, protocadherin alpha subfamily C, 1, C, 2
PCDHB1	29930	NP_037472	3808	protocadherin beta 1
PCDHB10	56126	NP_061753	4526	protocadherin beta 10 protocadherin beta 12
PCDHB10	56126	NP_061753.1	4570	protocadherin beta 10 protocadherin beta 11
PCDHB11	56125	NP_061754.1	4527	protocadherin beta 11
PCDHB12	56124	NP_061755	4528	protocadherin beta 12
PCDHB13	56123	NP_061756.1	7292	protocadherin beta 13
PCDHB14	56122	NP_061757	4529	protocadherin beta 14
PCDHB16	57717	NP_066008	4746	protocadherin beta 16
PCDHB2	56133	NP_061759.1	7286	protocadherin beta 2
PCDHB4	56131	NP_061761.1	7287	protocadherin beta 4
PCDHB5	26167	NP_056484	4111	protocadherin beta 5
PCDHB6	56130	NP_061762	4530	protocadherin beta 6
PCDHB7	56129	NP_061763	4531	protocadherin beta 7
PCDHB8	56128	NP_061993.3	7450	protocadherin beta 8
PCDHB9	56127	NP_061753	4525	protocadherin beta 10 protocadherin beta 10
PCDHB9	56127	NP_061992	4569	protocadherin beta 10 protocadherin beta 9
PCDHGA1	56114	NP_061735	4513	protocadherin gamma subfamily A, 1
PCDHGA1	56114	NP_114382	5276	protocadherin gamma subfamily A, 1
PCDHGA10	56106	NP_061736	4514	protocadherin gamma subfamily A, 10
PCDHGA10	56106	NP_114479	5289	protocadherin gamma subfamily A, 10
PCDHGA11	56105	NP_061737	4515	protocadherin gamma subfamily A, 11
PCDHGA11	56105	NP_114480	5290	protocadherin gamma subfamily A, 11
PCDHGA11	56105	NP_114481	5291	protocadherin gamma subfamily A, 11
PCDHGA9	56107	NP_061744	4516	protocadherin gamma subfamily A, 9
PCDHGA9	56107	NP_114478	5288	protocadherin gamma subfamily A, 9
PCDHGB5	56101	NP_061748	4517	protocadherin gamma subfamily B, 5
PCDHGB5	56101	NP_115270	5295	protocadherin gamma subfamily B, 5
PCDHGB6	56100	NP_061749	4518	protocadherin gamma subfamily B, 6
PCDHGB6	56100	NP_115271	5296	protocadherin gamma subfamily B, 6
PCDHGB7	56099	NP_061750.1	7293	protocadherin gamma subfamily B, 7
PCDHGC3	5098	NP_002579	2687	protocadherin gamma subfamily C, 3, C, 5, C, 4, A, 12
PCDHGC3	5098	NP_002579.2	2878	protocadherin gamma subfamily C, 3, C, 5, C, 4, A, 12
PCDHGC3	5098	NP_002579.2	4519	protocadherin gamma subfamily C, 3, C, 5, C, 4, A, 12
PCDHGC3	5098	NP_061752	4522	protocadherin gamma subfamily C, 3, C, 5, C, 4, A, 12
PCDHGC3	5098	NP_002579.2	5292	protocadherin gamma subfamily C, 3, C, 5, C, 4, A, 12
PCDHGC3	5098	NP_002579.2	5338	protocadherin gamma subfamily C, 3, C, 5, C, 4, A, 12
PCDHGC3	5098	NP_002579.2	5341	protocadherin gamma subfamily C, 3, C, 5, C, 4, A, 12
PCDHGC3	5098	NP_115782	5345	protocadherin gamma subfamily C, 3, C, 5, C, 4, A, 12
PCDHGC3	5098	NP_115783	5348	protocadherin gamma subfamily C, 3, C, 5, C, 4, A, 12
PCDHGC4	56098	NP_061751.1	2688	protocadherin gamma subfamily C, 3, C, 5, C, 4, A, 12
PCDHGC4	56098	NP_061751.1	2879	protocadherin gamma subfamily C, 3, C, 5, C, 4, A, 12
PCDHGC4	56098	NP_061751.1	4520	protocadherin gamma subfamily C, 3, C, 5, C, 4, A, 12
PCDHGC4	56098	NP_061752	4523	protocadherin gamma subfamily C, 3, C, 5, C, 4, A, 12
PCDHGC4	56098	NP_061751.1	5293	protocadherin gamma subfamily C, 3, C, 5, C, 4, A, 12
PCDHGC4	56098	NP_061751.1	5339	protocadherin gamma subfamily C, 3, C, 5, C, 4, A, 12
PCDHGC4	56098	NP_061751.1	5342	protocadherin gamma subfamily C, 3, C, 5, C, 4, A, 12
PCDHGC4	56098	NP_115782	5346	protocadherin gamma subfamily C, 3, C, 5, C, 4, A, 12
PCDHGC4	56098	NP_115783	5349	protocadherin gamma subfamily C, 3, C, 5, C, 4, A, 12
PCDHGC5	56097	NP_061752.1	2689	protocadherin gamma subfamily C, 3, C, 5, C, 4, A, 12
PCDHGC5	56097	NP_061752.1	2880	protocadherin gamma subfamily C, 3, C, 5, C, 4, A, 12
PCDHGC5	56097	NP_061751	4521	protocadherin gamma subfamily C, 3, C, 5, C, 4, A, 12
PCDHGC5	56097	NP_061752	4524	protocadherin gamma subfamily C, 3, C, 5, C, 4, A, 12
PCDHGC5	56097	NP_115783.1	5294	protocadherin gamma subfamily C, 3, C, 5, C, 4, A, 12
PCDHGC5	56097	NP_061752.1	5340	protocadherin gamma subfamily C, 3, C, 5, C, 4, A, 12
PCDHGC5	56097	NP_061752.1	5343	protocadherin gamma subfamily C, 3, C, 5, C, 4, A, 12
PCDHGC5	56097	NP_115782	5347	protocadherin gamma subfamily C, 3, C, 5, C, 4, A, 12
PCDHGC5	56097	NP_115783	5350	protocadherin gamma subfamily C, 3, C, 5, C, 4, A, 12
PCGF3	10336	NP_006306	3464	polycomb group ring finger 3
PCLO	27445	NP_055325	3951	piccolo (presynaptic cytomatrix protein)
PCLO	27445	NP_149015	5438	piccolo (presynaptic cytomatrix protein)
PCNX	22990	NP_055797	4023	pecanexHomolog (Drosophila)
PCNXL2	80003	NP_055616	3990	pecanex-like 2 (Drosophila)
PCNXL4	64430	NP_071940	4928	chromosome 14 open reading frame 135
PCSK1N	27344	NP_037403	3792	proprotein convertase subtilisin/kexin type 1 inhibitor

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
PCSK5	5125	NP_006191	3454	proprotein convertase subtilisin/kexin type 5
PCSK7	9159	NP_004707	3108	proprotein convertase subtilisin/kexin type 7 pseudogene
PCYT1A	5130	NP_001299602.1	7455	phosphate cytidylyltransferase 1, choline, alpha
PDCD1	5133	NP_005009	3198	programmed cell death 1
PDCD1LG2	80380	NP_079515	5139	programmed cell death 1 ligand 2
PDE3A	5139	NP_000912	329	phosphodiesterase 3A, cGMP-inhibited
PDE3B	5140	NP_000913	330	phosphodiesterase 3B, cGMP-inhibited
PDE6B	5158	NP_000274	81	phosphodiesterase 6B, cGMP-specific, rod, beta
PDE6B	5158	NP_001138763	1843	phosphodiesterase 6B, cGMP-specific, rod, beta
PDE6B	5158	NP_001138764	1844	phosphodiesterase 6B, cGMP-specific, rod, beta
PDGFRA	5156	NP_006197	3455	platelet-derived growth factor receptor, alpha polypeptide
PDGFRB	5159	NP_002600	2691	platelet-derived growth factor receptor, beta polypeptide
PDIA4	9601	NP_004902	3165	protein disulfide isomerase family A, member 4
PDPN	10630	NP_001006625	481	podoplanin
PDPN	10630	NP_001006626	482	podoplanin
PDPN	10630	NP_006465	3496	podoplanin
PDPN	10630	NP_938203	6786	podoplanin
PDZD8	118987	NP_776152	6384	PDZ domain containing 8
PDZK1IP1	10158	NP_005755	3359	PDZK1 interacting protein 1
PEAR1	375033	NP_001073940	1027	platelet endothelial aggregation receptor 1
PEBP4	157310	NP_659399	5906	phosphatidylethanolamine-binding protein 4
PECAM1	5175	NP_000433	128	platelet/endothelial cell adhesion molecule
PEMT	10400	NP_009100	3641	phosphatidylethanolamine N-methyltransferase
PEMT	10400	NP_680477	6001	phosphatidylethanolamine N-methyltransferase
PEMT	10400	NP_680478	6002	phosphatidylethanolamine N-methyltransferase
PERP	64065	NP_071404	4876	PERP, TP53 apoptosis effector
PET117	100303755	NP_001158283	7102	protein PET117 homolog, mitochondrial precursor
PEX10	5192	NP_002608	2692	peroxisomal biogenesis factor 10
PEX10	5192	NP_722540	6232	peroxisomal biogenesis factor 10
PEX11A	8800	NP_003838	2913	peroxisomal biogenesis factor 11 alpha
PEX11B	8799	NP_003837	2912	peroxisomal biogenesis factor 11 beta
PEX11G	92960	NP_542393	5613	peroxisomal biogenesis factor 11 gamma
PEX2	5828	NP_000309	91	peroxisomal membrane protein 3, 35kDa
PEX2	5828	NP_001073336	1009	peroxisomal membrane protein 3, 35kDa
PF4	5196	NP_002610	2693	platelet factor 4
PFKP	5214	NP_002618	2694	phosphofructokinase, platelet
PFN2	5217	NP_002619	2695	profilin 2
PFN2	5217	NP_444252	5556	profilin 2
PGAM5	192111	NP_001164014	2242	phosphoglycerate mutase family member 5
PGAM5	192111	NP_001164015	2243	phosphoglycerate mutase family member 5
PGAM5	192111	NP_612642	5762	phosphoglycerate mutase family member 5
PGAP1	80055	NP_079265	5112	post-GPI attachment to proteins 1
PGAP2	27315	NP_001138910	1858	post-GPI attachment to proteins 2
PGAP2	27315	NP_001138910	1859	post-GPI attachment to proteins 2
PGAP2	27315	NP_055304	3946	post-GPI attachment to proteins 2
PGAP3	93210	NP_219487	5495	post-GPI attachment to proteins 3
PGRMC1	10857	NP_006658	3536	progesterone receptor membrane component 1
PGRMC2	10424	NP_006311	3466	progesterone receptor membrane component 2
PHEX	5251	NP_000435	130	phosphate regulating endopeptidaseHomolog, X-linked
PHLDB2	90102	NP_001127909	1490	pleckstrinHomology-like domain, family B, member 2, phosphatidylinositol-specific phospholipase C, X domain containing 2
PHLDB2	90102	NP_001127910	1491	pleckstrinHomology-like domain, family B, member 2, phosphatidylinositol-specific phospholipase C, X domain containing 2
PHLDB2	90102	NP_001127911	1492	pleckstrinHomology-like domain, family B, member 2, phosphatidylinositol-specific phospholipase C, X domain containing 2
PHLDB2	90102	NP_001127909.1	1498	pleckstrinHomology-like domain, family B, member 2, phosphatidylinositol-specific phospholipase C, X domain containing 2
PHLDB2	90102	NP_665696	5974	pleckstrinHomology-like domain, family B, member 2, phosphatidylinositol-specific phospholipase C, X domain containing 2
PHLDB2	90102	NP_695000	6170	pleckstrinHomology-like domain, family B, member 2, phosphatidylinositol-specific phospholipase C, X domain containing 2
PHTF1	10745	NP_006599	3527	putativeHomeodomain transcription factor 1
PI16	221476	NP_699201	6187	peptidase inhibitor 16
PI3	5266	NP_002629	2697	peptidase inhibitor 3, skin-derived
PIANP	196500	NP_710152	6214	chromosome 12 open reading frame 53
PICK1	9463	NP_001034672	802	protein interacting with PRKCA 1
PICK1	9463	NP_001034673	803	protein interacting with PRKCA 1
PICK1	9463	NP_036539	3770	protein interacting with PRKCA 1
PIEZO1	9780	NP_001136336	1706	family with sequence similarity 38, member A
PIGA	5277	NP_002632	2698	phosphatidylinositol glycan anchor biosynthesis, class A
PIGA	5277	NP_065206	4668	phosphatidylinositol glycan anchor biosynthesis, class A
PIGB	9488	NP_004846	3149	phosphatidylinositol glycan anchor biosynthesis, class B
PIGC	5279	NP_002633	2699	phosphatidylinositol glycan anchor biosynthesis, class C
PIGC	5279	NP_714969	6222	phosphatidylinositol glycan anchor biosynthesis, class C
PIGF	5281	NP_002634	2700	phosphatidylinositol glycan anchor biosynthesis, class F
PIGF	5281	NP_775097	6331	phosphatidylinositol glycan anchor biosynthesis, class F
PIGG	54872	NP_001120650	1308	phosphatidylinositol glycan anchor biosynthesis, class G

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
PIGG	54872	NP_060203	4330	phosphatidylinositol glycan anchor biosynthesis, class G
PIGH	5283	NP_004560	3077	phosphatidylinositol glycan anchor biosynthesis, classH
PIGK	10026	NP_005473	3293	phosphatidylinositol glycan anchor biosynthesis, class K
PIGL	9487	NP_004269	3006	phosphatidylinositol glycan anchor biosynthesis, class L
PIGM	93183	NP_660150	5929	phosphatidylinositol glycan anchor biosynthesis, class M
PIGN	23556	NP_036459	3746	phosphatidylinositol glycan anchor biosynthesis, class N
PIGN	23556	NP_789744	6476	phosphatidylinositol glycan anchor biosynthesis, class N
PIGO	84720	NP_116023	5382	phosphatidylinositol glycan anchor biosynthesis, class O
PIGO	84720	NP_690577	6119	phosphatidylinositol glycan anchor biosynthesis, class O
PIGP	51227	NP_710148	6212	phosphatidylinositol glycan anchor biosynthesis, class P
PIGP	51227	NP_710149	6213	phosphatidylinositol glycan anchor biosynthesis, class P
PIGQ	9091	NP_004195	2986	phosphatidylinositol glycan anchor biosynthesis, class Q
PIGQ	9091	NP_683721	6007	phosphatidylinositol glycan anchor biosynthesis, class Q
PIGR	5284	NP_002635	2701	polymeric immunoglobulin receptor
PIGS	94005	NP_149975	5466	phosphatidylinositol glycan anchor biosynthesis, class S
PIGT	51604	NP_057021	4139	phosphatidylinositol glycan anchor biosynthesis, class T
PIGU	128869	NP_536724	5604	phosphatidylinositol glycan anchor biosynthesis, class U
PIGV	55650	NP_060307	4353	phosphatidylinositol glycan anchor biosynthesis, class V
PIGW	284098	NP_848612	6548	phosphatidylinositol glycan anchor biosynthesis, class W
PIGX	54965	NP_001159776	2159	phosphatidylinositol glycan anchor biosynthesis, class X
PIGX	54965	NP_060331	4359	phosphatidylinositol glycan anchor biosynthesis, class X
PIGZ	80235	NP_079439	5129	phosphatidylinositol glycan anchor biosynthesis, class Z
PIK3IP1	113791	NP_001129383	1574	phosphoinositide-3-kinase interacting protein 1
PIK3IP1	113791	NP_443112	5524	phosphoinositide-3-kinase interacting protein 1
PILRA	29992	NP_038467	3823	paired immunoglobin-like type 2 receptor alpha
PILRA	29992	NP_840056	6531	paired immunoglobin-like type 2 receptor alpha
PILRA	29992	NP_840057	6532	paired immunoglobin-like type 2 receptor alpha
PIPOX	51268	NP_057602	4233	pipecolic acid oxidase
PIRT	644139	NP_001094857	1201	phosphoinositide-interacting regulator of transient receptor potential channels
PKD1	5587	NP_002733.2	7364	protein kinase D1
PKD1L1	168507	NP_612152	5720	polycystic kidney disease 1 like 1
PKD1L2	114780	NP_001070248	950	polycystic kidney disease 1-like 2
PKD1L2	114780	NP_443124	5529	polycystic kidney disease 1-like 2
PKD2	5311	NP_000288	83	polycystic kidney disease 2 (autosomal dominant)
PKD2L1	9033	NP_057196	4171	polycystic kidney disease 2-like 1
PKD2L2	27039	NP_055201	3928	polycystic kidney disease 2-like 2
PKDCC	91461	NP_612379	5735	protein kinase-like protein SgK493
PKDREJ	10343	NP_006062	3428	polycystic kidney disease (polycystin) and REJHomolog (sperm receptor for egg jellyHomolog, sea urchin)
PKHD1	5314	NP_062565	4577	polycystic kidney andHepatic disease 1 (autosomal recessive)
PKHD1	5314	NP_619639	5771	polycystic kidney andHepatic disease 1 (autosomal recessive)
PKHD1	5314	NP_733842	6250	polycystic kidney andHepatic disease 1 (autosomal recessive)
PKN2	5586	NP_006247	3458	protein kinase N2
PLA2G10	8399	NP_003552	2848	phospholipase A2, group X
PLA2G16	11145	NP_001121675	1366	phospholipase A2, group XVI
PLA2G16	11145	NP_009000	3625	phospholipase A2, group XVI
PLA2G2A	5320	NP_000291	84	phospholipase A2, group IIA (platelets, synovial fluid)
PLA2G2A	5320	NP_001155199	2023	phospholipase A2, group IIA (platelets, synovial fluid)
PLA2G2A	5320	NP_001155200	2024	phospholipase A2, group IIA (platelets, synovial fluid)
PLA2G2A	5320	NP_001155201	2025	phospholipase A2, group IIA (platelets, synovial fluid)
PLA2G2D	26279	NP_036532	3768	phospholipase A2, group IID
PLA2G2E	30814	NP_055404	3962	phospholipase A2, group IIE
PLA2R1	22925	NP_001007268	512	phospholipase A2 receptor 1, 180kDa
PLA2R1	22925	NP_031392	3692	phospholipase A2 receptor 1, 180kDa
PLB1	151056	NP_001164056	2247	phospholipase B1
PLB1	151056	NP_694566	6150	phospholipase B1
PLBD2	196463	NP_001153199	1957	phospholipase B domain containing 2
PLBD2	196463	NP_775813	6355	phospholipase B domain containing 2
PLD3	23646	NP_001026866	723	phospholipase D family, member 3
PLD3	23646	NP_036400	3732	phospholipase D family, member 3
PLD4	122618	NP_620145	5796	phospholipase D family, member 4
PLD5	200150	NP_689879	6093	phospholipase D family, member 5
PLD6	201164	NP_849158	6569	phospholipase D family, member 6
PLGRKT	55848	NP_060935	4455	chromosome 9 open reading frame 46
PLIN1	5346	NP_001138783	1853	perilipin
PLIN1	5346	NP_002657	2702	perilipin
PLLP	51090	NP_057077	4148	plasma membrane proteolipid (plasmolipin)
PLN	5350	NP_002658	2703	phospholamban
PLOD2	5352	NP_000926	332	procollagen-lysine, 2-oxoglutarate 5-dioxygenase 2
PLOD2	5352	NP_891988	6696	procollagen-lysine, 2-oxoglutarate 5-dioxygenase 2
PLP1	5354	NP_000524	155	proteolipid protein 1
PLP1	5354	NP_001122306	1376	proteolipid protein 1
PLP1	5354	NP_955772	6888	proteolipid protein 1
PLP2	5355	NP_002659	2704	proteolipid protein 2 (colonic epithelium-enriched)
PLVAP	83483	NP_112600	5220	plasmalemma vesicle associated protein
PLXDC1	57125	NP_065138	4650	plexin domain containing 1
PLXDC2	84898	NP_116201	5408	plexin domain containing 2
PLXNA1	5361	NP_115618	5313	plexin A1

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
PLXNA2	5362	NP_079455	5130	plexin A2
PLXNA4	91584	NP_001099013	1226	plexin A4
PLXNA4	91584	NP_065962	4740	plexin A4
PLXNA4	91584	NP_861440	6631	plexin A4
PLXNC1	10154	NP_005752	3358	plexin C1
PLXND1	23129	NP_055918	4033	plexin D1
PM20D1	148811	NP_689704	6069	peptidase M20 domain containing 1
PMEL	6490	NP_008859	3597	silverHomolog (mouse)
PMEPA1	56937	NP_064567	4613	prostate transmembrane protein, androgen induced 1
PMEPA1	56937	NP_954638	6861	prostate transmembrane protein, androgen induced 1
PMEPA1	56937	NP_954639	6862	prostate transmembrane protein, androgen induced 1
PMEPA1	56937	NP_954640	6863	prostate transmembrane protein, androgen induced 1
PMP22	5376	NP_000295	85	peripheral myelin protein 22
PMP22	5376	NP_696996	6176	peripheral myelin protein 22
PMP22	5376	NP_696997	6177	peripheral myelin protein 22
PNKD	25953	NP_001070867	963	paroxysmal nonkinesigenic dyskinesia
PNKD	25953	NP_056303	4090	paroxysmal nonkinesigenic dyskinesia
PNKD	25953	NP_072094	4933	paroxysmal nonkinesigenic dyskinesia
PNPLA2	57104	NP_065109	4638	patatin-like phospholipase domain containing 2
PNPLA4	8228	NP_001135861	1651	patatin-like phospholipase domain containing 4
PNPLA4	8228	NP_004641	3097	patatin-like phospholipase domain containing 4
PNPLA6	10908	NP_001159583	2146	patatin-like phospholipase domain containing 6
PNPLA6	10908	NP_001159584	2147	patatin-like phospholipase domain containing 6
PNPLA6	10908	NP_001159585	2148	patatin-like phospholipase domain containing 6
PNPLA6	10908	NP_001159586	2149	patatin-like phospholipase domain containing 6
PNPLA6	10908	NP_006693	3551	patatin-like phospholipase domain containing 6
PNPLA7	375775	NP_001092007	1137	patatin-like phospholipase domain containing 7
PNPLA7	375775	NP_689499	6025	patatin-like phospholipase domain containing 7
PODXL	5420	NP_001018121	669	podocalyxin-like
PODXL	5420	NP_005388	3280	podocalyxin-like
PODXL2	50512	NP_056535	4118	podocalyxin-like 2
POLG2	11232	NP_009146	3651	polymerase (DNA directed), gamma 2, accessory subunit
POMGNT1	55624	NP_060209	4331	protein O-linked mannose beta1,2-N-acetylglucosaminyltransferase
POMK	84197	NP_115613	5312	protein kinase-like protein SgK196
POMT1	10585	NP_001070833	961	protein-O-mannosyltransferase 1
POMT1	10585	NP_001070834	962	protein-O-mannosyltransferase 1
POMT1	10585	NP_001129585	1599	protein-O-mannosyltransferase 1
POMT1	10585	NP_001129586	1600	protein-O-mannosyltransferase 1
POMT1	10585	NP_009102	3642	protein-O-mannosyltransferase 1
PON2	5445	NP_000296	86	paraoxonase 2
PON2	5445	NP_001018171	670	paraoxonase 2
PON3	5446	NP_000931	333	paraoxonase 3
POPDC2	64091	NP_071418	4881	popeye domain containing 2
POPDC3	64208	NP_071756	4904	popeye domain containing 3
POR	5447	NP_000932	334	P450 (cytochrome) oxidoreductase
PORCN	64840	NP_073736	4952	porcupineHomolog (Drosophila)
PORCN	64840	NP_982299	6942	porcupineHomolog (Drosophila)
PORCN	64840	NP_982300	6943	porcupineHomolog (Drosophila)
PORCN	64840	NP_982301	6944	porcupineHomolog (Drosophila)
PORCN	64840	NP_982302	6945	porcupineHomolog (Drosophila)
PPAP2A	8611	NP_003702	2873	phosphatidic acid phosphatase type 2A
PPAP2A	8611	NP_795714	6488	phosphatidic acid phosphatase type 2A
PPAP2B	8613	NP_003704	2875	phosphatidic acid phosphatase type 2B
PPAP2B	8613	NP_003704.3	6491	phosphatidic acid phosphatase type 2B
PPAP2C	8612	NP_003703	2874	phosphatidic acid phosphatase type 2C
PPAP2C	8612	NP_803545	6495	phosphatidic acid phosphatase type 2C
PPAP2C	8612	NP_808211	6500	phosphatidic acid phosphatase type 2C
PPAPDC1A	196051	NP_001025230	715	phosphatidic acid phosphatase type 2 domain containing 1A
PPAPDC1B	84513	NP_001096029	1207	phosphatidic acid phosphatase type 2 domain containing 1B
PPAPDC1B	84513	NP_001096030	1208	phosphatidic acid phosphatase type 2 domain containing 1B
PPAPDC1B	84513	NP_115872	5359	phosphatidic acid phosphatase type 2 domain containing 1B
PPAPDC2	403313	NP_982278	6938	phosphatidic acid phosphatase type 2 domain containing 2
PPAPDC3	84814	NP_116117	5390	phosphatidic acid phosphatase type 2 domain containing 3
PPBP	5473	NP_002695	2705	pro-platelet basic protein (chemokine (C-X-C motif) ligand 7)
PPIB	5479	NP_000933	335	peptidylprolyl isomerase B (cyclophilin B)
PPIC	5480	NP_000934	336	peptidylprolyl isomerase C (cyclophilin C)
PPM1L	151742	NP_640338	5848	protein phosphatase 1 (formerly 2C)-like
PPP1R3A	5506	NP_002702	2706	protein phosphatase 1, regulatory (inhibitor) subunit 3A
PPP1R3F	89801	NP_149992	5470	protein phosphatase 1, regulatory (inhibitor) subunit 3F
PPT2	9374	NP_005146	3220	palmitoyl-protein thioesterase 2
PPT2	9374	NP_619731	5776	palmitoyl-protein thioesterase 2
PQLC1	80148	NP_001139815	1933	PQ loop repeat containing 1
PQLC1	80148	NP_001139817	1934	PQ loop repeat containing 1
PQLC1	80148	NP_079354	5118	PQ loop repeat containing 1
PQLC2	54896	NP_001035214	838	PQ loop repeat containing 2
PQLC2	54896	NP_001035215	839	PQ loop repeat containing 2
PQLC2	54896	NP_060235	4338	PQ loop repeat containing 2

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
PQLC3	130814	NP_689604	6050	PQ loop repeat containing 3
PRAC2	360205	NP_001269204	7111	prostate cancer susceptibility candidate 2
PRADC1	84279	NP_115695	5326	chromosome 2 open reading frame 7
PRAF2	11230	NP_009144	3649	PRA1 domain family, member 2
PRCP	5547	NP_005031	3201	prolylcarboxypeptidase (angiotensinase C)
PRCP	5547	NP_955450	6883	prolylcarboxypeptidase (angiotensinase C)
PRDX4	10549	NP_006397	3482	peroxiredoxin 4
PREB	10113	NP_037520	3816	prolactin regulatory element binding
PRIMA1	145270	NP_821092	6513	proline rich membrane anchor 1
PRLH	51052	NP_056977	4130	prolactin releasingHormone
PRLHR	2834	NP_004239	2998	prolactin releasingHormone receptor
PRLR	5618	NP_000940	337	prolactin receptor
PRND	23627	NP_036541	3771	prion protein 2 (dublet)
PRNP	5621	NP_000302	88	prion protein
PRNP	5621	NP_001073590	1011	prion protein
PRNP	5621	NP_001073591	1012	prion protein
PRNP	5621	NP_001073592	1013	prion protein
PRNP	5621	NP_898902	6703	prion protein
PROCR	10544	NP_006395	3480	protein C receptor, endothelial (EPCR)
PROKR2	128674	NP_658986	5903	prokineticin receptor 2
PROM1	8842	NP_001139319	1878	prominin 1
PROM1	8842	NP_001139320	1879	prominin 1
PROM1	8842	NP_001139321	1880	prominin 1
PROM1	8842	NP_001139322	1881	prominin 1
PROM1	8842	NP_001139323	1882	prominin 1
PROM1	8842	NP_001139324	1883	prominin 1
PROM1	8842	NP_006008	3415	prominin 1
PROM2	150696	NP_001159449	2137	prominin 2
PROM2	150696	NP_001159450	2138	prominin 2
PROM2	150696	NP_653308	5895	prominin 2
PROS1	5627	NP_000304	89	protein S (alpha)
PRPF38B	55119	NP_060531	4386	PRP38 pre-mRNA processing factor 38 (yeast) domain containing B
PRPH2	5961	NP_000313	92	peripherin 2 (retinal degeneration, slow)
PRR4	11272	NP_001092008.2	7429	proline rich 4 (lacrimal)
PRR7	80758	NP_085044	5147	proline rich 7 (synaptic)
PRRG1	5638	NP_000941	338	proline rich Gla (G-carboxyglutamic acid) 1
PRRG1	5638	NP_001135867	1654	proline rich Gla (G-carboxyglutamic acid) 1
PRRG2	5639	NP_000942	339	proline rich Gla (G-carboxyglutamic acid) 2
PRRG3	79057	NP_076987	5006	proline rich Gla (G-carboxyglutamic acid) 3 (transmembrane)
PRRG4	79056	NP_076986	5005	proline rich Gla (G-carboxyglutamic acid) 4 (transmembrane)
PRRT1	80863	NP_085154	5155	proline-rich transmembrane protein 1
PRRT2	112476	NP_660282	5937	proline-rich transmembrane protein 2
PRRT3	285368	NP_997234	7014	proline-rich transmembrane protein 3
PRSS16	10279	NP_005856	3389	protease, serine, 16 (thymus)
PRUNE2	158471	NP_056040	4048	pruneHomolog 2 (Drosophila)
PSCA	8000	NP_005663	3340	prostate stem cell antigen
PSEN1	5663	NP_000012	3	presenilin 1
PSEN1	5663	NP_015557	3685	presenilin 1
PSEN2	5664	NP_000438	131	presenilin 2 (Alzheimer disease 4)
PSEN2	5664	NP_036618	3786	presenilin 2 (Alzheimer disease 4)
PSENEN	55851	NP_001268461.1	7440	presenilin enhancer gamma-secretase subunit
PSMB7	5695	NP_002790	2708	proteasome (prosome, macropain) subunit, beta type, 7
PTAFR	5724	NP_000943	340	platelet-activating factor receptor
PTAFR	5724	NP_001158193	2100	platelet-activating factor receptor
PTAFR	5724	NP_001158194	2101	platelet-activating factor receptor
PTAFR	5724	NP_001158195	2102	platelet-activating factor receptor
PTCH1	5727	NP_000255	77	patchedHomolog 1 (Drosophila)
PTCH1	5727	NP_001077071	1071	patchedHomolog 1 (Drosophila)
PTCH1	5727	NP_001077072	1072	patchedHomolog 1 (Drosophila)
PTCH1	5727	NP_001077073	1073	patchedHomolog 1 (Drosophila)
PTCH1	5727	NP_001077074	1074	patchedHomolog 1 (Drosophila)
PTCH1	5727	NP_001077075	1075	patchedHomolog 1 (Drosophila)
PTCH1	5727	NP_001077076	1076	patchedHomolog 1 (Drosophila)
PTCH2	8643	NP_001159764	2158	patchedHomolog 2 (Drosophila)
PTCH2	8643	NP_003729	2882	patchedHomolog 2 (Drosophila)
PTCHD1	139411	NP_775766	6347	patched domain containing 1
PTCHD2	57540	NP_065831	4721	patched domain containing 2
PTCHD4	442213	NP_001013754	619	chromosome 6 open reading frame 138
PTCRA	171558	NP_612153	5721	pre T-cell antigen receptor alpha
PTDSS1	9791	NP_055569	3983	phosphatidylserine synthase 1
PTDSS2	81490	NP_110410	5172	phosphatidylserine synthase 2
PTGDR	5729	NP_000944	341	prostaglandin D2 receptor (DP)
PTGDR2	11251	NP_004769	3123	G protein-coupled receptor 44
PTGER1	5731	NP_000946	342	prostaglandin E receptor 1 (subtype EP1), 42kDa
PTGER2	5732	NP_000947	343	prostaglandin E receptor 2 (subtype EP2), 53kDa
PTGER3	5733	NP_000948	344	prostaglandin E receptor 3 (subtype EP3)
PTGER3	5733	NP_001119516	1300	prostaglandin E receptor 3 (subtype EP3)

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
PTGER3	5733	NP_942005	6827	prostaglandin E receptor 3 (subtype EP3)
PTGER3	5733	NP_942006	6828	prostaglandin E receptor 3 (subtype EP3)
PTGER3	5733	NP_942007	6829	prostaglandin E receptor 3 (subtype EP3)
PTGER3	5733	NP_942008	6830	prostaglandin E receptor 3 (subtype EP3)
PTGER3	5733	NP_942009	6831	prostaglandin E receptor 3 (subtype EP3)
PTGER3	5733	NP_942010	6832	prostaglandin E receptor 3 (subtype EP3)
PTGER3	5733	NP_942011	6833	prostaglandin E receptor 3 (subtype EP3)
PTGER3	5733	NP_942012	6834	prostaglandin E receptor 3 (subtype EP3)
PTGER4	5734	NP_000949	345	prostaglandin E receptor 4 (subtype EP4)
PTGFR	5737	NP_000950	346	prostaglandin F receptor (FP)
PTGFR	5737	NP_001034674	804	prostaglandin F receptor (FP)
PTGFRN	5738	NP_065173	4660	prostaglandin F2 receptor negative regulator
PTGIR	5739	NP_000951.1	7263	prostaglandin I2 (prostacyclin) receptor (IP)
PTGIS	5740	NP_000952	347	prostaglandin I2 (prostacyclin) synthase
PTGS1	5742	NP_000953	348	prostaglandin-endoperoxide synthase 1 (prostaglandin G/H synthase and cyclooxygenase)
PTGS1	5742	NP_542158	5607	prostaglandin-endoperoxide synthase 1 (prostaglandin G/H synthase and cyclooxygenase)
PTH1R	5745	NP_000307	90	parathyroidHormone 1 receptor
PTH2	113091	NP_848544	6540	parathyroidHormone 2
PTK7	5754	NP_002812	2709	PTK7 protein tyrosine kinase 7
PTK7	5754	NP_690619	6125	PTK7 protein tyrosine kinase 7
PTK7	5754	NP_690620	6126	PTK7 protein tyrosine kinase 7
PTK7	5754	NP_690621	6127	PTK7 protein tyrosine kinase 7
PTN	5764	NP_002816	2710	pleiotrophin
PTPN1	5770	NP_002818	2712	protein tyrosine phosphatase, non-receptor type 1
PTPN2	5771	NP_002819	2713	protein tyrosine phosphatase, non-receptor type 2
PTPN2	5771	NP_536347	5601	protein tyrosine phosphatase, non-receptor type 2
PTPN2	5771	NP_536348	5602	protein tyrosine phosphatase, non-receptor type 2
PTPN5	84867	NP_001035059	820	protein tyrosine phosphatase, non-receptor type 5 (striatum-enriched)
PTPN5	84867	NP_008837	3592	protein tyrosine phosphatase, non-receptor type 5 (striatum-enriched)
PTPN5	84867	NP_116170	5400	protein tyrosine phosphatase, non-receptor type 5 (striatum-enriched)
PTPRA	5786	NP_002827	2714	protein tyrosine phosphatase, receptor type, A
PTPRA	5786	NP_543030	5637	protein tyrosine phosphatase, receptor type, A
PTPRA	5786	NP_543031	5638	protein tyrosine phosphatase, receptor type, A
PTPRC	5788	NP_002829	2715	protein tyrosine phosphatase, receptor type, C
PTPRC	5788	NP_563578	5645	protein tyrosine phosphatase, receptor type, C
PTPRC	5788	NP_563579	5646	protein tyrosine phosphatase, receptor type, C
PTPRC	5788	NP_563580	5647	protein tyrosine phosphatase, receptor type, C
PTPRCAP	5790	NP_005599	3324	protein tyrosine phosphatase, receptor type, C-associated protein
PTPRD	5789	NP_001035802	883	protein tyrosine phosphatase, receptor type, D
PTPRD	5789	NP_001164496	2263	protein tyrosine phosphatase, receptor type, D
PTPRD	5789	NP_002830	2716	protein tyrosine phosphatase, receptor type, D
PTPRD	5789	NP_569075	5651	protein tyrosine phosphatase, receptor type, D
PTPRD	5789	NP_569076	5652	protein tyrosine phosphatase, receptor type, D
PTPRD	5789	NP_569077	5653	protein tyrosine phosphatase, receptor type, D
PTPRE	5791	NP_006495	3501	protein tyrosine phosphatase, receptor type, E
PTPRE	5791	NP_569119	5655	protein tyrosine phosphatase, receptor type, E
PTPRF	5792	NP_002831	2717	protein tyrosine phosphatase, receptor type, F
PTPRF	5792	NP_569707	5656	protein tyrosine phosphatase, receptor type, F
PTPRG	5793	NP_002832	2718	protein tyrosine phosphatase, receptor type, G
PTPRH	5794	NP_001154912	2011	protein tyrosine phosphatase, receptor type,H
PTPRH	5794	NP_002833	2719	protein tyrosine phosphatase, receptor type,H
PTPRJ	5795	NP_001091973	1127	protein tyrosine phosphatase, receptor type, J
PTPRJ	5795	NP_002834	2720	protein tyrosine phosphatase, receptor type, J
PTPRK	5796	NP_001129120	1543	protein tyrosine phosphatase, receptor type, K
PTPRK	5796	NP_002835	2721	protein tyrosine phosphatase, receptor type, K
PTPRM	5797	NP_001098714	1219	protein tyrosine phosphatase, receptor type, M
PTPRM	5797	NP_002836	2722	protein tyrosine phosphatase, receptor type, M
PTPRN	5798	NP_001186692.1	7426	protein tyrosine phosphatase, receptor type N
PTPRO	5800	NP_002839	2723	protein tyrosine phosphatase, receptor type, O
PTPRO	5800	NP_109592	5157	protein tyrosine phosphatase, receptor type, O
PTPRO	5800	NP_109593	5158	protein tyrosine phosphatase, receptor type, O
PTPRO	5800	NP_109594	5159	protein tyrosine phosphatase, receptor type, O
PTPRO	5800	NP_109595	5160	protein tyrosine phosphatase, receptor type, O
PTPRO	5800	NP_109596	5161	protein tyrosine phosphatase, receptor type, O
PTPRR	5801	NP_002840	2724	protein tyrosine phosphatase, receptor type, R
PTPRR	5801	NP_570897	5671	protein tyrosine phosphatase, receptor type, R
PTPRS	5802	NP_002841	2725	protein tyrosine phosphatase, receptor type, S
PTPRS	5802	NP_570923	5673	protein tyrosine phosphatase, receptor type, S
PTPRS	5802	NP_570924	5674	protein tyrosine phosphatase, receptor type, S
PTPRS	5802	NP_570925	5675	protein tyrosine phosphatase, receptor type, S
PTPRT	11122	NP_008981	3622	protein tyrosine phosphatase, receptor type, T
PTPRT	11122	NP_573400	5676	protein tyrosine phosphatase, receptor type, T
PTPRU	10076	NP_005695	3348	protein tyrosine phosphatase, receptor type, U
PTPRU	10076	NP_573438	5677	protein tyrosine phosphatase, receptor type, U
PTPRU	10076	NP_573439	5678	protein tyrosine phosphatase, receptor type, U
PTPRZ1	5803	NP_002842	2726	protein tyrosine phosphatase, receptor-type, Z polypeptide 1
PTRH2	51651	NP_057161	4162	peptidyl-tRNAHydrolase 2

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
PTTG1IP	754	NP_004330	3026	pituitary tumor-transforming 1 interacting protein
PVR	5817	NP_001129240	1562	poliovirus receptor
PVR	5817	NP_001129241	1563	poliovirus receptor
PVR	5817	NP_001129242	1564	poliovirus receptor
PVR	5817	NP_006496	3502	poliovirus receptor
PVRIG	79037	NP_076975	5002	poliovirus receptor related immunoglobulin domain containing
PVRL1	5818	NP_002846	2727	poliovirus receptor-related 1 (herpesvirus entry mediator C)
PVRL1	5818	NP_976030	6921	poliovirus receptor-related 1 (herpesvirus entry mediator C)
PVRL1	5818	NP_976031	6922	poliovirus receptor-related 1 (herpesvirus entry mediator C)
PVRL2	5819	NP_001036189	918	poliovirus receptor-related 2 (herpesvirus entry mediator B)
PVRL2	5819	NP_002847	2728	poliovirus receptor-related 2 (herpesvirus entry mediator B)
PVRL3	25945	NP_056295	4088	poliovirus receptor-related 3
PVRL4	81607	NP_112178	5191	poliovirus receptor-related 4
PXDN	7837	NP_036425	3739	peroxidasinHomolog (Drosophila)
PXYLP1	92370	NP_001032249	770	acid phosphatase-like 2
PXYLP1	92370	NP_689495	6024	acid phosphatase-like 2
PYY	5697	NP_004151	2975	peptide YY
QPCTL	54814	NP_001156849	2052	glutaminyl-peptide cyclotransferase-like
QPCTL	54814	NP_060129	4318	glutaminyl-peptide cyclotransferase-like
QRFPR	84109	NP_937822	6751	pyroglutamylated RFamide peptide receptor
QSOX1	5768	NP_001004128	427	quiescin Q6 sulfhydryl oxidase 1
QSOX1	5768	NP_002817	2711	quiescin Q6 sulfhydryl oxidase 1
QSOX2	169714	NP_859052	6621	quiescin Q6 sulfhydryl oxidase 2
RABAC1	10567	NP_006414	3489	Rab acceptor 1 (prenylated)
RABGAP1	23637	NP_036329	3719	RAB GTPase activating protein 1
RAD51B	5890	NP_002868	2729	RAD51-like 1 (S. cerevisiae)
RAD51B	5890	NP_598193	5699	RAD51-like 1 (S. cerevisiae)
RAD51B	5890	NP_598194	5700	RAD51-like 1 (S. cerevisiae)
RAET1E	135250	NP_631904	5840	retinoic acid early transcript 1E
RAET1G	353091	NP_001001788.2	7369	retinoic acid early transcript 1G
RAET1L	154064	NP_570970.2	7368	retinoic acid early transcript 1L
RAMP1	10267	NP_005846	3386	receptor (G protein-coupled) activity modifying protein 1
RAMP2	10266	NP_005845	3385	receptor (G protein-coupled) activity modifying protein 2
RAMP3	10268	NP_005847	3387	receptor (G protein-coupled) activity modifying protein 3
RARRES3	5920	NP_004576	3078	retinoic acid receptor responder (tazarotene induced) 3
RBFOX3	146713	NP_001076044	1061	hexaribonucleotide binding protein 3
RBM3	5935	NP_006734	3555	RNA binding motif (RNP1, RRM) protein 3
RDH10	157506	NP_742034	6270	retinol dehydrogenase 10 (all-trans)
RDH11	51109	NP_057110	4154	retinol dehydrogenase 11 (all-trans/9-cis/11-cis)
RDM1	201299	NP_001030008	760	RAD52 motif 1
RDM1	201299	NP_001156592	2042	RAD52 motif 1
RDM1	201299	NP_001156593	2043	RAD52 motif 1
RDM1	201299	NP_001156594	2044	RAD52 motif 1
RDM1	201299	NP_001156596	2045	RAD52 motif 1
RDM1	201299	NP_001156597	2046	RAD52 motif 1
RDM1	201299	NP_001156602	2047	RAD52 motif 1
RDM1	201299	NP_663629	5967	RAD52 motif 1
RECK	8434	NP_066934	4774	reversion-inducing-cysteine-rich protein with kazal motifs
REEP2	51308	NP_057690	4259	receptor accessory protein 2
REEP3	221035	NP_001001330	352	receptor accessory protein 3
REEP4	80346	NP_079508	5138	receptor accessory protein 4
REEP5	7905	NP_005660	3338	receptor accessory protein 5
REEP6	92840	NP_612402	5742	receptor accessory protein 6
RELL1	768211	NP_001078868	1091	RELT-like 1
RELL1	768211	NP_001078869	1092	RELT-like 1
RELL2	285613	NP_001123501	1411	RELT-like 2
RELL2	285613	NP_776189	6391	RELT-like 2
RELT	84957	NP_116260	5419	RELT tumor necrosis factor receptor
RELT	84957	NP_689408	6019	RELT tumor necrosis factor receptor
RER1	11079	NP_008964	3618	RER1 retention in endoplasmic reticulum 1Homolog (S. cerevisiae)
RET	5979	NP_065681	4685	ret proto-oncogene
RET	5979	NP_066124	4750	ret proto-oncogene
RETSAT	54884	NP_060220	4335	retinol saturase (all-trans-retinol 13,14-reductase)
RFT1	91869	NP_443091	5519	RFT1Homolog (S. cerevisiae)
RGR	5995	NP_001012738	604	retinal G protein coupled receptor
RGR	5995	NP_001012740	605	retinal G protein coupled receptor
RGR	5995	NP_002912	2730	retinal G protein coupled receptor
RGS9BP	388531	NP_997274	7020	regulator of G protein signaling 9 binding protein
RGSL1	353299	NP_001131141	1615	regulator of G-protein signaling like 1
RHAG	6005	NP_000315	93	Rh-associated glycoprotein
RHBDD1	84236	NP_001161080	2187	rhomboid domain containing 1
RHBDD1	84236	NP_115652	5317	rhomboid domain containing 1
RHBDD2	57414	NP_001035546	872	rhomboid domain containing 2
RHBDD2	57414	NP_001035547	873	rhomboid domain containing 2
RHBDD3	25807	NP_036397	3731	rhomboid domain containing 3
RHBDF1	64285	NP_071895	4919	rhomboid 5Homolog 1 (Drosophila)
RHBDF2	79651	NP_001005498	461	rhomboid 5Homolog 2 (Drosophila)

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
RHBDF2	79651	NP_078875	5054	rhomboid 5Homolog 2 (Drosophila)
RHBDL1	9028	NP_001265649.1	2931	rhomboid, veinlet-like 1 (Drosophila)
RHBDL2	54933	NP_060291	4349	rhomboid, veinlet-like 2 (Drosophila)
RHBDL3	162494	NP_612201	5727	rhomboid, veinlet-like 3 (Drosophila)
RHBG	57127	NP_065140	4651	Rh family, B glycoprotein (gene/pseudogene)
RHCE	6006	NP_065231	4676	Rh blood group, CcEe antigens
RHCE	6006	NP_619522	5765	Rh blood group, CcEe antigens
RHCE	6006	NP_619523	5766	Rh blood group, CcEe antigens
RHCE	6006	NP_619524	5767	Rh blood group, CcEe antigens
RHCG	51458	NP_057405	4203	Rh family, C glycoprotein
RHD	6007	NP_001121163	1347	Rh blood group, D antigen
RHD	6007	NP_057208	4173	Rh blood group, D antigen
RHO	6010	NP_000530	156	rhodopsin
RHOT1	55288	NP_001028738.1	7355	ras homolog family member T1
RHOT2	89941	NP_620124	5791	rasHomolog gene family, member T2
RIC3	79608	NP_001128581	1525	resistance to inhibitors of cholinesterase 3Homolog (C. elegans)
RIC3	79608	NP_078833	5047	resistance to inhibitors of cholinesterase 3Homolog (C. elegans)
RMDN2	151393	NP_001164262	2253	family with sequence similarity 82, member A1
RMDN2	151393	NP_001164263	2254	family with sequence similarity 82, member A1
RMDN2	151393	NP_001164264	2255	family with sequence similarity 82, member A1
RMDN2	151393	NP_653314	5897	family with sequence similarity 82, member A1
RMDN3	55177	NP_060615	4398	family with sequence similarity 82, member A2
RNASE1	6035	NP_002924	2731	ribonuclease, RNase A family, 1 (pancreatic)
RNASE1	6035	NP_937875	6760	ribonuclease, RNase A family, 1 (pancreatic)
RNASE1	6035	NP_937877	6761	ribonuclease, RNase A family, 1 (pancreatic)
RNASE1	6035	NP_937878	6762	ribonuclease, RNase A family, 1 (pancreatic)
RNASE4	6038	NP_002928	2732	ribonuclease, RNase A family, 4
RNASE4	6038	NP_919412	6722	ribonuclease, RNase A family, 4
RNASE6	6039	NP_005606	3325	ribonuclease, RNase A family, k6
RNASE7	84659	NP_115961	5373	ribonuclease, RNase A family, 7
RNF103	7844	NP_001005753	471	vacuolar protein sorting 24Homolog (S. cerevisiae) ring finger protein 103
RNF103	7844	NP_005658	3337	vacuolar protein sorting 24Homolog (S. cerevisiae) ring finger protein 103
RNF103	7844	NP_057163	4164	vacuolar protein sorting 24Homolog (S. cerevisiae) ring finger protein 103
RNF112	7732	NP_009079	3634	ring finger protein 112
RNF121	55298	NP_060790	4420	ring finger protein 121
RNF122	79845	NP_079063	5082	ring finger protein 122
RNF128	79589	NP_078815	5043	ring finger protein 128
RNF128	79589	NP_919445	6732	ring finger protein 128
RNF13	11342	NP_009213	3668	ring finger protein 13
RNF13	11342	NP_899237	6708	ring finger protein 13
RNF130	55819	NP_060904	4449	ring finger protein 130
RNF133	168433	NP_631914	5844	ring finger protein 133
RNF139	11236	NP_009149	3652	ring finger protein 139
RNF144A	9781	NP_055561	3981	ring finger protein 144A
RNF144B	255488	NP_877434	6680	ring finger protein 144B
RNF145	153830	NP_653327	5901	ring finger protein 145
RNF149	284996	NP_775918	6370	ring finger protein 149
RNF150	57484	NP_065775	4708	ring finger protein 150
RNF152	220441	NP_775828	6357	ring finger protein 152
RNF167	26001	NP_056343	4095	ring finger protein 167
RNF170	81790	NP_001153695	1979	ring finger protein 170
RNF170	81790	NP_001153696	1980	ring finger protein 170
RNF170	81790	NP_001153697	1981	ring finger protein 170
RNF170	81790	NP_112216	5201	ring finger protein 170
RNF175	285533	NP_775933	6376	ring finger protein 175
RNF180	285671	NP_001107033	1264	ring finger protein 180
RNF180	285671	NP_848627	6550	ring finger protein 180
RNF182	221687	NP_001158504	2122	ring finger protein 182
RNF182	221687	NP_001158505	2123	ring finger protein 182
RNF182	221687	NP_001158506	2124	ring finger protein 182
RNF182	221687	NP_689950	6108	ring finger protein 182
RNF183	138065	NP_659488	5924	ring finger protein 183
RNF186	54546	NP_061935	4555	ring finger protein 186
RNF19A	25897	NP_056250	4082	ring finger protein 19A
RNF19A	25897	NP_904355	6709	ring finger protein 19A
RNF19B	127544	NP_001120833	1318	ring finger protein 19B
RNF19B	127544	NP_699172	6178	ring finger protein 19B
RNF217	154214	NP_689766	6081	ring finger protein 217
RNF24	11237	NP_001127809	1475	ring finger protein 24
RNF24	11237	NP_001127810	1476	ring finger protein 24
RNF24	11237	NP_009150	3653	ring finger protein 24
RNF26	79102	NP_114404	5278	ring finger protein 26
RNF43	54894	NP_060233	4337	ring finger protein 43
ROBO2	6092	NP_001122401	1380	roundabout, axon guidance receptor,Homolog 2 (Drosophila)
ROBO2	6092	NP_002933	2733	roundabout, axon guidance receptor,Homolog 2 (Drosophila)
ROBO3	64221	NP_071765	4907	roundabout, axon guidance receptor,Homolog 3 (Drosophila)
ROM1	6094	NP_000318	94	retinal outer segment membrane protein 1

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
ROMO1	140823	NP_542786	5626	reactive oxygen species modulator 1
ROR1	4919	NP_001077061	1070	receptor tyrosine kinase-like orphan receptor 1
ROR1	4919	NP_005003	3196	receptor tyrosine kinase-like orphan receptor 1
ROR2	4920	NP_004551	3076	receptor tyrosine kinase-like orphan receptor 2
ROS1	6098	NP_002935	2734	c-ros oncogene 1 , receptor tyrosine kinase
RPL37A	6168	NP_000989	349	ribosomal protein L37a
RPN1	6184	NP_002941	2735	ribophorin I
RPN2	6185	NP_001129243	1565	ribophorin II
RPN2	6185	NP_002942	2736	ribophorin II
RPRM	56475	NP_062819	4583	reprimo, TP53 dependent G2 arrest mediator candidate
RPRML	388394	NP_981945	6931	reprimo-like
RPS23	6228	NP_001016	638	ribosomal protein S23
RRBP1	6238	NP_001036041	905	ribosome binding protein 1Homolog 180kDa (dog)
RRBP1	6238	NP_004578	3079	ribosome binding protein 1Homolog 180kDa (dog)
RRH	10692	NP_006574	3522	retinal pigment epithelium-derived rhodopsinHomolog
RRM2B	50484	NP_056528	4116	ribonucleotide reductase M2 B (TP53 inducible)
RTN1	6252	NP_066959	4779	reticulon 1
RTN1	6252	NP_996734	6974	reticulon 1
RTN1	6252	NP_996739	6975	reticulon 1
RTN2	6253	NP_005610	3327	reticulon 2
RTN2	6253	NP_996783	6978	reticulon 2
RTN2	6253	NP_996784	6979	reticulon 2
RTN3	10313	NP_006045	3422	reticulon 3
RTN3	10313	NP_958831	6899	reticulon 3
RTN3	10313	NP_958832	6900	reticulon 3
RTN3	10313	NP_958833	6901	reticulon 3
RTN4	57142	NP_008939	3613	reticulon 4
RTN4	57142	NP_065393	4680	reticulon 4
RTN4	57142	NP_722550	6235	reticulon 4
RTN4	57142	NP_997403	7023	reticulon 4
RTN4	57142	NP_997404	7024	reticulon 4
RTP1	132112	NP_714919	6218	receptor (chemosensory) transporter protein 1
RTP3	83597	NP_113628	5224	receptor (chemosensory) transporter protein 3
RXFP1	59350	NP_067647	4812	relaxin/insulin-like family peptide receptor 1
RXFP2	122042	NP_001159530	2142	relaxin/insulin-like family peptide receptor 2
RXFP2	122042	NP_570718	5667	relaxin/insulin-like family peptide receptor 2
RXFP3	51289	NP_057652	4246	relaxin/insulin-like family peptide receptor 3
RYK	6259	NP_001005861	473	RYK receptor-like tyrosine kinase
RYK	6259	NP_002949	2737	RYK receptor-like tyrosine kinase
RYR1	6261	NP_000531	157	ryanodine receptor 1 (skeletal)
RYR1	6261	NP_001036188	917	ryanodine receptor 1 (skeletal)
RYR2	6262	NP_001026	718	ryanodine receptor 2 (cardiac)
RYR3	6263	NP_001027	736	ryanodine receptor 3
S100A10	6281	NP_002957.1	7264	S100 calcium binding protein A10
S1PR1	1901	NP_001391	2360	sphingosine-1-phosphate receptor 1
S1PR2	9294	NP_004221	2994	sphingosine-1-phosphate receptor 2
S1PR3	1903	NP_005217	3234	sphingosine-1-phosphate receptor 3
S1PR4	8698	NP_003766	2889	sphingosine-1-phosphate receptor 4
S1PR5	53637	NP_001159687	2154	sphingosine-1-phosphate receptor 5
S1PR5	53637	NP_110387	5164	sphingosine-1-phosphate receptor 5
SACM1L	22908	NP_054735	3855	SAC1 suppressor of actin mutations 1-like (yeast)
SAMD5	389432	NP_001025231	716	sterile alpha motif domain containing 5
SAMD8	142891	NP_653261	5885	sterile alpha motif domain containing 8
SAYSD1	55776	NP_060792	4421	chromosome 6 open reading frame 64
SC5D	6309	NP_001020127	692	sterol-C5-desaturase (ERG3 delta-5-desaturaseHomolog, S. cerevisiae)-like
SC5D	6309	NP_008849	3593	sterol-C5-desaturase (ERG3 delta-5-desaturaseHomolog, S. cerevisiae)-like
SCAMP1	9522	NP_004857	3152	secretory carrier membrane protein 1
SCAMP2	10066	NP_005688	3346	secretory carrier membrane protein 2
SCAMP3	10067	NP_005689	3347	secretory carrier membrane protein 3
SCAMP3	10067	NP_443069	5516	secretory carrier membrane protein 3
SCAMP4	113178	NP_524558	5596	secretory carrier membrane protein 4
SCAMP5	192683	NP_620417	5809	secretory carrier membrane protein 5
SCAP	22937	NP_036367	3725	SREBF chaperone
SCARA3	51435	NP_057324	4190	scavenger receptor class A, member 3
SCARA3	51435	NP_878185	6686	scavenger receptor class A, member 3
SCARA5	286133	NP_776194	6392	scavenger receptor class A, member 5 (putative)
SCARB1	949	NP_001076428	1063	scavenger receptor class B, member 1
SCARB1	949	NP_005496	3299	scavenger receptor class B, member 1
SCARB2	950	NP_005497	3300	scavenger receptor class B, member 2
SCCPDH	51097	NP_057086	4150	saccharopine dehydrogenase (putative)
SCD	6319	NP_005054	3203	stearoyl-CoA desaturase (delta-9-desaturase)
SCG2	7857	NP_003460	2838	secretogranin II (chromogranin C)
SCIMP	388325	NP_996986	6998	chromosome 17 open reading frame 87
SCN10A	6336	NP_006505	3503	sodium channel, voltage-gated, type X, alpha subunit
SCN11A	11280	NP_054858	3870	sodium channel, voltage-gated, type XI, alpha subunit
SCN1A	6323	NP_001159435	2133	sodium channel, voltage-gated, type I, alpha subunit
SCN1A	6323	NP_001159436	2134	sodium channel, voltage-gated, type I, alpha subunit

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
SCN1A	6323	NP_008851	3594	sodium channel, voltage-gated, type I, alpha subunit
SCN1B	6324	NP_001028	748	sodium channel, voltage-gated, type I, beta
SCN1B	6324	NP_950238	6851	sodium channel, voltage-gated, type I, beta
SCN2A	6326	NP_001035232	842	sodium channel, voltage-gated, type II, alpha subunit
SCN2A	6326	NP_001035233	843	sodium channel, voltage-gated, type II, alpha subunit
SCN2A	6326	NP_066287	4757	sodium channel, voltage-gated, type II, alpha subunit
SCN2B	6327	NP_004579	3080	sodium channel, voltage-gated, type II, beta
SCN3A	6328	NP_001075145	1053	sodium channel, voltage-gated, type III, alpha subunit
SCN3A	6328	NP_001075146	1054	sodium channel, voltage-gated, type III, alpha subunit
SCN3A	6328	NP_008853	3595	sodium channel, voltage-gated, type III, alpha subunit
SCN3B	55800	NP_001035241	844	sodium channel, voltage-gated, type III, beta
SCN3B	55800	NP_060870	4439	sodium channel, voltage-gated, type III, beta
SCN4A	6329	NP_000325	95	sodium channel, voltage-gated, type IV, alpha subunit
SCN4B	6330	NP_001135820	1646	sodium channel, voltage-gated, type IV, beta
SCN4B	6330	NP_001135821	1647	sodium channel, voltage-gated, type IV, beta
SCN4B	6330	NP_777594	6413	sodium channel, voltage-gated, type IV, beta
SCN5A	6331	NP_000326.2	7324	sodium voltage-gated channel alpha subunit 5
SCN7A	6332	NP_002967	2739	sodium channel, voltage-gated, type VII, alpha
SCN8A	6334	NP_055006	3879	sodium channel, voltage gated, type VIII, alpha subunit
SCN9A	6335	NP_002968	2740	sodium channel, voltage-gated, type IX, alpha subunit
SCNN1A	6337	NP_001029	756	sodium channel, nonvoltage-gated 1 alpha
SCNN1A	6337	NP_001153047	1946	sodium channel, nonvoltage-gated 1 alpha
SCNN1A	6337	NP_001153048	1947	sodium channel, nonvoltage-gated 1 alpha
SCNN1B	6338	NP_000327	96	sodium channel, nonvoltage-gated 1, beta
SCNN1D	6339	NP_001123885	1422	sodium channel, nonvoltage-gated 1, delta
SCNN1D	6339	NP_001123885.2	2741	sodium channel, nonvoltage-gated 1, delta
SCNN1G	6340	NP_001030	759	sodium channel, nonvoltage-gated 1, gamma
SCRG1	11341	NP_009212	3667	stimulator of chondrogenesis 1
SCTR	6344	NP_002971	2742	secretin receptor
SCUBE2	57758	NP_001164161	2251	signal peptide, CUB domain, EGF-like 2
SCUBE2	57758	NP_066025	4749	signal peptide, CUB domain, EGF-like 2
SDC1	6382	NP_001006947	496	syndecan 1
SDC1	6382	NP_002988	2747	syndecan 1
SDC2	6383	NP_002989	2748	syndecan 2
SDC3	9672	NP_055469	3969	syndecan 3
SDC4	6385	NP_002990	2749	syndecan 4
SDCBP	6386	NP_001007068	501	syndecan binding protein (syntenin)
SDCBP	6386	NP_001007069	502	syndecan binding protein (syntenin)
SDCBP	6386	NP_001007070	503	syndecan binding protein (syntenin)
SDCBP	6386	NP_001007071	504	syndecan binding protein (syntenin)
SDCBP	6386	NP_005616	3328	syndecan binding protein (syntenin)
SDF2L1	23753	NP_071327	4867	stromal cell-derived factor 2-like 1
SDF4	51150	NP_057260.2	7307	stromal cell derived factor 4
SDHC	6391	NP_001030588	764	succinate dehydrogenase complex, subunit C, integral membrane protein, 15kDa
SDHC	6391	NP_001030589	765	succinate dehydrogenase complex, subunit C, integral membrane protein, 15kDa
SDHC	6391	NP_001030590	766	succinate dehydrogenase complex, subunit C, integral membrane protein, 15kDa
SDHC	6391	NP_002992	2750	succinate dehydrogenase complex, subunit C, integral membrane protein, 15kDa
SDK1	221935	NP_001073121	989	sidekickHomolog 1, cell adhesion molecule (chicken)Hypothetical LOC730351
SDK1	221935	NP_689957	6109	sidekickHomolog 1, cell adhesion molecule (chicken)Hypothetical LOC730351
SDK2	54549	NP_001138424	1799	sidekickHomolog 2 (chicken)
SDR42E1	93517	NP_660151	5930	short chain dehydrogenase/reductase family 42E, member 1
SDSL	113675	NP_612441	5751	serine dehydratase-like
SEC11A	23478	NP_055115	3909	SEC11Homolog A (S. cerevisiae)
SEC11C	90701	NP_150596	5479	SEC11Homolog C (S. cerevisiae)
SEC22A	26984	NP_036562	3775	SEC22 vesicle trafficking proteinHomolog A (S. cerevisiae)
SEC61A1	29927	NP_037468	3804	Sec61 alpha 1 subunit (S. cerevisiae)
SEC61A2	55176	NP_001136099	1679	Sec61 alpha 2 subunit (S. cerevisiae)
SEC61A2	55176	NP_001136100	1680	Sec61 alpha 2 subunit (S. cerevisiae)
SEC61A2	55176	NP_060614	4397	Sec61 alpha 2 subunit (S. cerevisiae)
SEC61G	23480	NP_001012474	596	Sec61 gamma subunit
SEC61G	23480	NP_055117	3910	Sec61 gamma subunit
SEC62	7095	NP_003253	2805	SEC62Homolog (S. cerevisiae)
SEC63	11231	NP_009145	3650	SEC63Homolog (S. cerevisiae)
SECTM1	6398	NP_002995.1	7265	secreted and transmembrane 1
SEL1L	6400	NP_005056	3204	sel-1 suppressor of lin-12-like (C. elegans)
SEL1L3	23231	NP_056002	4043	KIAA0746 protein
SELE	6401	NP_000441	132	selectin E
SELK	58515	NP_067060	4800	selenoprotein K
SELL	6402	NP_000646	187	selectin L
SELM	140606	NP_536355	5603	selenoprotein M
SELP	6403	NP_002996	2751	selectin P (granule membrane protein 140kDa, antigen CD62)
SELPLG	6404	NP_002997	2752	selectin P ligand
SELT	51714	NP_057359	4197	selenoprotein T
SEMA3B	7869	NP_001005914	475	sema domain, immunoglobulin domain (Ig), short basic domain, secreted, (semaphorin) 3B
SEMA3B	7869	NP_004627	3094	sema domain, immunoglobulin domain (Ig), short basic domain, secreted, (semaphorin) 3B
SEMA3D	223117	NP_689967	6111	sema domain, immunoglobulin domain (Ig), short basic domain, secreted, (semaphorin) 3D

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
SEMA4A	64218	NP_071762	4906	sema domain, immunoglobulin domain (Ig), transmembrane domain (TM) and short cytoplasmic domain, (semaphorin) 4A
SEMA4B	10509	NP_064595	4618	sema domain, immunoglobulin domain (Ig), transmembrane domain (TM) and short cytoplasmic domain, (semaphorin) 4B
SEMA4B	10509	NP_945119	6844	sema domain, immunoglobulin domain (Ig), transmembrane domain (TM) and short cytoplasmic domain, (semaphorin) 4B
SEMA4C	54910	NP_060259	4343	sema domain, immunoglobulin domain (Ig), transmembrane domain (TM) and short cytoplasmic domain, (semaphorin) 4C
SEMA4D	10507	NP_001135759	1634	sema domain, immunoglobulin domain (Ig), transmembrane domain (TM) and short cytoplasmic domain, (semaphorin) 4D
SEMA4D	10507	NP_006369	3478	sema domain, immunoglobulin domain (Ig), transmembrane domain (TM) and short cytoplasmic domain, (semaphorin) 4D
SEMA4F	10505	NP_004254	3003	sema domain, immunoglobulin domain (Ig), transmembrane domain (TM) and short cytoplasmic domain, (semaphorin) 4F
SEMA4G	57715	NP_060363	4365	sema domain, immunoglobulin domain (Ig), transmembrane domain (TM) and short cytoplasmic domain, (semaphorin) 4G
SEMA5A	9037	NP_003957	2934	sema domain, seven thrombospondin repeats (type 1 and type 1-like), transmembrane domain (TM) and short cytoplasmic domain, (semaphorin) 5A
SEMA5B	54437	NP_001026872.2	7357	semaphorin 5B
SEMA6A	57556	NP_065847	4725	sema domain, transmembrane domain (TM), and cytoplasmic domain, (semaphorin) 6A
SEMA6B	10501	NP_115484	5297	sema domain, transmembrane domain (TM), and cytoplasmic domain, (semaphorin) 6B
SEMA6D	80031	NP_065909	4733	sema domain, transmembrane domain (TM), and cytoplasmic domain, (semaphorin) 6D
SEMA6D	80031	NP_079242	5110	sema domain, transmembrane domain (TM), and cytoplasmic domain, (semaphorin) 6D
SEMA6D	80031	NP_705869	6204	sema domain, transmembrane domain (TM), and cytoplasmic domain, (semaphorin) 6D
SEMA6D	80031	NP_705870	6205	sema domain, transmembrane domain (TM), and cytoplasmic domain, (semaphorin) 6D
SEMA6D	80031	NP_705871	6206	sema domain, transmembrane domain (TM), and cytoplasmic domain, (semaphorin) 6D
SEMA6D	80031	NP_705872	6207	sema domain, transmembrane domain (TM), and cytoplasmic domain, (semaphorin) 6D
SEPP1	6414	NP_001078955	1096	selenoprotein P, plasma, 1
SEPP1	6414	NP_001087195	1100	selenoprotein P, plasma, 1
SEPP1	6414	NP_005401	3281	selenoprotein P, plasma, 1
SERINC1	57515	NP_065806	4717	serine incorporator 1
SERINC2	347735	NP_849196	6575	serine incorporator 2
SERINC3	10955	NP_006802	3568	serine incorporator 3
SERINC3	10955	NP_945179	6845	serine incorporator 3
SERINC5	256987	NP_840060	6533	serine incorporator 5
SERP1	27230	NP_055260	3941	stress-associated endoplasmic reticulum protein 1
SERP2	387923	NP_001010897	575	stress-associated endoplasmic reticulum protein family member 2
SERPINA1	5265	NP_000286	82	serpin peptidase inhibitor, clade A (alpha-1 antiproteinase, antitrypsin), member 1
SERPINA1	5265	NP_001002235	389	serpin peptidase inhibitor, clade A (alpha-1 antiproteinase, antitrypsin), member 1
SERPINA1	5265	NP_001002236	390	serpin peptidase inhibitor, clade A (alpha-1 antiproteinase, antitrypsin), member 1
SERPINA1	5265	NP_001121172	1349	serpin peptidase inhibitor, clade A (alpha-1 antiproteinase, antitrypsin), member 1
SERPINA1	5265	NP_001121173	1350	serpin peptidase inhibitor, clade A (alpha-1 antiproteinase, antitrypsin), member 1
SERPINA1	5265	NP_001121174	1351	serpin peptidase inhibitor, clade A (alpha-1 antiproteinase, antitrypsin), member 1
SERPINA1	5265	NP_001121175	1352	serpin peptidase inhibitor, clade A (alpha-1 antiproteinase, antitrypsin), member 1
SERPINA1	5265	NP_001121176	1353	serpin peptidase inhibitor, clade A (alpha-1 antiproteinase, antitrypsin), member 1
SERPINA1	5265	NP_001121177	1354	serpin peptidase inhibitor, clade A (alpha-1 antiproteinase, antitrypsin), member 1
SERPINA1	5265	NP_001121178	1355	serpin peptidase inhibitor, clade A (alpha-1 antiproteinase, antitrypsin), member 1
SERPINA1	5265	NP_001121179	1356	serpin peptidase inhibitor, clade A (alpha-1 antiproteinase, antitrypsin), member 1
SERPINA9	327657	NP_001035983	901	serpin peptidase inhibitor, clade A (alpha-1 antiproteinase, antitrypsin), member 9
SERPINA9	327657	NP_783866	6458	serpin peptidase inhibitor, clade A (alpha-1 antiproteinase, antitrypsin), member 9
SERPINB13	5275	NP_036529	3767	serpin peptidase inhibitor, clade B (ovalbumin), member 13
SERTM1	400120	NP_982276	6937	chromosome 13 open reading frame 36
SEZ6	124925	NP_001092105	1143	seizure related 6Homolog (mouse)
SEZ6	124925	NP_849191	6574	seizure related 6Homolog (mouse)
SEZ6L	23544	NP_066938	4776	seizure related 6Homolog (mouse)-like
SEZ6L2	26470	NP_001107571	1266	seizure related 6Homolog (mouse)-like 2
SEZ6L2	26470	NP_001107572	1267	seizure related 6Homolog (mouse)-like 2
SEZ6L2	26470	NP_036542	3772	seizure related 6Homolog (mouse)-like 2
SEZ6L2	26470	NP_963869	6911	seizure related 6Homolog (mouse)-like 2
SFRP1	6422	NP_003003	2753	secreted frizzled-related protein 1
SFT2D1	113402	NP_660152	5931	SFT2 domain containing 1
SFT2D2	375035	NP_955376	6879	SFT2 domain containing 2
SFT2D3	84826	NP_116129	5394	SFT2 domain containing 3
SFTPC	6440	NP_003009	2754	surfactant protein C
SFTPD	6441	NP_003010	2755	surfactant protein D
SFXN1	94081	NP_073591	4943	sideroflexin 1
SFXN2	118980	NP_849189	6572	sideroflexin 2
SFXN3	81855	NP_112233	5209	sideroflexin 3
SFXN4	119559	NP_998814	7046	sideroflexin 4
SFXN5	94097	NP_653180	5865	sideroflexin 5
SGCA	6442	NP_000014	4	sarcoglycan, alpha (50kDa dystrophin-associated glycoprotein)
SGCA	6442	NP_001129169	1554	sarcoglycan, alpha (50kDa dystrophin-associated glycoprotein)
SGCB	6443	NP_000223	73	sarcoglycan, beta (43kDa dystrophin-associated glycoprotein)
SGCD	6444	NP_000328	97	sarcoglycan, delta (35kDa dystrophin-associated glycoprotein)
SGCD	6444	NP_001121681	1368	sarcoglycan, delta (35kDa dystrophin-associated glycoprotein)
SGCD	6444	NP_758447	6301	sarcoglycan, delta (35kDa dystrophin-associated glycoprotein)
SGCE	8910	NP_001092870	1156	sarcoglycan, epsilon

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
SGCE	8910	NP_001092871	1157	sarcoglycan, epsilon
SGCE	8910	NP_003910	2927	sarcoglycan, epsilon
SGCG	6445	NP_000222	72	sarcoglycan, gamma (35kDa dystrophin-associated glycoprotein)
SGCZ	137868	NP_631906	5841	sarcoglycan zeta
SGMS1	259230	NP_671512	5991	sphingomyelin synthase 1
SGMS2	166929	NP_001129729	1604	sphingomyelin synthase 2
SGMS2	166929	NP_001129730	1605	sphingomyelin synthase 2
SGMS2	166929	NP_689834	6090	sphingomyelin synthase 2
SGPL1	8879	NP_003892	2926	sphingosine-1-phosphate lyase 1
SGPP1	81537	NP_110418	5176	sphingosine-1-phosphate phosphatase 1
SGPP2	130367	NP_689599	6046	sphingosine-1-phosphate phosphotase 2
SHH	6469	NP_000184	58	sonicHedgehogHomolog (Drosophila)
SHISA2	387914	NP_001007539	522	shisaHomolog 2 (Xenopus laevis)
SHISA3	152573	NP_001073974	1032	shisaHomolog 3 (Xenopus laevis)
SHISA4	149345	NP_937792	6746	shisaHomolog 4 (Xenopus laevis)
SHISA5	51246	NP_057563	4225	shisaHomolog 5 (Xenopus laevis)
SHISA6	388336	NP_997269	7019	FLJ45455 protein
SHISA7	729956	NP_001138648.1	7406	shisa family member 7
SHISA8	440829	XP_496530	7089	transmembrane protein 46-like
SHISA8	440829	XP_945572	7094	transmembrane protein 46-like
SHISA9	729993	NP_001138676.2	7418	shisa family member 9
SI	6476	NP_001032	767	sucrase-isomaltase (alpha-glucosidase)
SIDT1	54847	NP_060169	4325	SID1 transmembrane family, member 1
SIDT2	51092	NP_001035545	871	SID1 transmembrane family, member 2
SIGIRR	59307	NP_001128525	1515	single immunoglobulin and toll-interleukin 1 receptor (TIR) domain
SIGIRR	59307	NP_001128526	1516	single immunoglobulin and toll-interleukin 1 receptor (TIR) domain
SIGIRR	59307	NP_068577	4832	single immunoglobulin and toll-interleukin 1 receptor (TIR) domain
SIGLEC1	6614	NP_075556	4971	sialic acid binding Ig-like lectin 1, sialoadhesin
SIGLEC10	89790	NP_001164627.1	7416	sialic acid binding Ig like lectin 10
SIGLEC11	114132	NP_001128635	1533	sialic acid binding Ig-like lectin 11
SIGLEC11	114132	NP_443116	5525	sialic acid binding Ig-like lectin 11
SIGLEC6	946	NP_001236	2325	sialic acid binding Ig-like lectin 6
SIGLEC6	946	NP_942142	6839	sialic acid binding Ig-like lectin 6
SIGLEC6	946	NP_942143	6840	sialic acid binding Ig-like lectin 6
SIGLEC7	27036	NP_055200	3927	sialic acid binding Ig-like lectin 7
SIGLEC7	27036	NP_057627	4240	sialic acid binding Ig-like lectin 7
SIGLEC8	27181	NP_055257	3940	sialic acid binding Ig-like lectin 8
SIGLEC9	27180	NP_055256	3939	sialic acid binding Ig-like lectin 9
SIGLECL1	284369	NP_775906	6368	hypothetical protein FLJ40235
SIGMAR1	10280	NP_005857	3390	sigma non-opioid intracellular receptor 1
SIGMAR1	10280	NP_671513	5992	sigma non-opioid intracellular receptor 1
SIRPA	140885	NP_001035111	821	signal-regulatory protein alpha
SIRPA	140885	NP_001035112	822	signal-regulatory protein alpha
SIRPA	140885	NP_542970	5629	signal-regulatory protein alpha
SIRPB1	10326	NP_001077379	1082	signal-regulatory protein beta 1
SIRPB1	10326	NP_001129316	1573	signal-regulatory protein beta 1
SIRPB1	10326	NP_006056	3426	signal-regulatory protein beta 1
SIRPB2	284759	NP_001116434	1294	signal-regulatory protein beta 2
SIRPB2	284759	NP_001128308	1507	signal-regulatory protein beta 2
SIRPG	55423	NP_001034597	797	signal-regulatory protein gamma
SIRPG	55423	NP_061026	4466	signal-regulatory protein gamma
SIRPG	55423	NP_543006	5630	signal-regulatory protein gamma
SIT1	27240	NP_055265	3943	signaling threshold regulating transmembrane adaptor 1
SLAMF1	6504	NP_003028	2759	signaling lymphocytic activation molecule family member 1
SLAMF6	114836	NP_443163	5541	SLAM family member 6
SLAMF7	57823	NP_067004	4788	SLAM family member 7
SLAMF8	56833	NP_064510	4599	SLAM family member 8
SLAMF9	89886	NP_001139644	1910	SLAM family member 9
SLAMF9	89886	NP_001139645	1911	SLAM family member 9
SLAMF9	89886	NP_254273	5498	SLAM family member 9
SLC10A1	6554	NP_003040	2770	solute carrier family 10 (sodium/bile acid cotransporter family), member 1
SLC10A2	6555	NP_000443	133	solute carrier family 10 (sodium/bile acid cotransporter family), member 2
SLC10A3	8273	NP_001135863	1652	solute carrier family 10 (sodium/bile acid cotransporter family), member 3
SLC10A3	8273	NP_001135864	1653	solute carrier family 10 (sodium/bile acid cotransporter family), member 3
SLC10A3	8273	NP_062822	4584	solute carrier family 10 (sodium/bile acid cotransporter family), member 3
SLC10A4	201780	NP_689892	6095	solute carrier family 10 (sodium/bile acid cotransporter family), member 4
SLC10A7	84068	NP_001025169	712	solute carrier family 10 (sodium/bile acid cotransporter family), member 7
SLC10A7	84068	NP_115504	5301	solute carrier family 10 (sodium/bile acid cotransporter family), member 7
SLC11A1	6556	NP_000569	166	solute carrier family 11 (proton-coupled divalent metal ion transporters), member 1
SLC11A2	4891	NP_000608	175	solute carrier family 11 (proton-coupled divalent metal ion transporters), member 2
SLC12A1	6557	NP_000329	98	solute carrier family 12 (sodium/potassium/chloride transporters), member 1
SLC12A2	6558	NP_001037	919	solute carrier family 12 (sodium/potassium/chloride transporters), member 2
SLC12A3	6559	NP_000330	99	solute carrier family 12 (sodium/chloride transporters), member 3
SLC12A3	6559	NP_001119579	1303	solute carrier family 12 (sodium/chloride transporters), member 3
SLC12A3	6559	NP_001119580	1304	solute carrier family 12 (sodium/chloride transporters), member 3
SLC12A4	6560	NP_001139433	1888	solute carrier family 12 (potassium/chloride transporters), member 4
SLC12A4	6560	NP_001139434	1889	solute carrier family 12 (potassium/chloride transporters), member 4

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
SLC12A4	6560	NP_001139435	1890	solute carrier family 12 (potassium/chloride transporters), member 4
SLC12A4	6560	NP_001139436	1891	solute carrier family 12 (potassium/chloride transporters), member 4
SLC12A4	6560	NP_005063	3207	solute carrier family 12 (potassium/chloride transporters), member 4
SLC12A5	57468	NP_001128243	1506	solute carrier family 12 (potassium-chloride transporter), member 5
SLC12A5	57468	NP_065759	4706	solute carrier family 12 (potassium-chloride transporter), member 5
SLC12A6	9990	NP_001035959	896	solute carrier family 12 (potassium/chloride transporters), member 6
SLC12A6	9990	NP_001035960	897	solute carrier family 12 (potassium/chloride transporters), member 6
SLC12A6	9990	NP_001035961	898	solute carrier family 12 (potassium/chloride transporters), member 6
SLC12A6	9990	NP_001035962	899	solute carrier family 12 (potassium/chloride transporters), member 6
SLC12A6	9990	NP_005126	3218	solute carrier family 12 (potassium/chloride transporters), member 6
SLC12A6	9990	NP_598408	5702	solute carrier family 12 (potassium/chloride transporters), member 6
SLC12A7	10723	NP_006589	3525	solute carrier family 12 (potassium/chloride transporters), member 7
SLC12A8	84561	NP_078904	5058	solute carrier family 12 (potassium/chloride transporters), member 8
SLC12A9	56996	NP_064631	4623	solute carrier family 12 (potassium/chloride transporters), member 9
SLC13A1	6561	NP_071889	4918	solute carrier family 13 (sodium/sulfate symporters), member 1
SLC13A2	9058	NP_001139447	1892	solute carrier family 13 (sodium-dependent dicarboxylate transporter), member 2
SLC13A2	9058	NP_001139447.1	1893	solute carrier family 13 (sodium-dependent dicarboxylate transporter), member 2
SLC13A2	9058	NP_003975	2940	solute carrier family 13 (sodium-dependent dicarboxylate transporter), member 2
SLC13A3	64849	NP_001011554	589	solute carrier family 13 (sodium-dependent dicarboxylate transporter), member 3
SLC13A3	64849	NP_073740	4953	solute carrier family 13 (sodium-dependent dicarboxylate transporter), member 3
SLC13A4	26266	NP_036582	3778	solute carrier family 13 (sodium/sulfate symporters), member 4
SLC13A5	284111	NP_001137310	1732	solute carrier family 13 (sodium-dependent citrate transporter), member 5
SLC13A5	284111	NP_808218	6501	solute carrier family 13 (sodium-dependent citrate transporter), member 5
SLC14A1	6563	NP_001122060	1374	solute carrier family 14 (urea transporter), member 1 (Kidd blood group)
SLC14A1	6563	NP_001139508	1897	solute carrier family 14 (urea transporter), member 1 (Kidd blood group)
SLC14A1	6563	NP_001139509	1898	solute carrier family 14 (urea transporter), member 1 (Kidd blood group)
SLC14A1	6563	NP_056949	4125	solute carrier family 14 (urea transporter), member 1 (Kidd blood group)
SLC14A2	8170	NP_009094	3638	solute carrier family 14 (urea transporter), member 2
SLC15A1	6564	NP_005064	3208	solute carrier family 15 (oligopeptide transporter), member 1
SLC15A2	6565	NP_001139470	1894	solute carrier family 15 (H+/peptide transporter), member 2
SLC15A2	6565	NP_066568	4765	solute carrier family 15 (H+/peptide transporter), member 2
SLC15A3	51296	NP_057666	4251	solute carrier family 15, member 3
SLC15A4	121260	NP_663623	5964	solute carrier family 15, member 4
SLC16A1	6566	NP_001159968	2171	solute carrier family 16, member 1 (monocarboxylic acid transporter 1)
SLC16A1	6566	NP_003042	2771	solute carrier family 16, member 1 (monocarboxylic acid transporter 1)
SLC16A10	117247	NP_061063	4470	solute carrier family 16, member 10 (aromatic amino acid transporter)
SLC16A11	162515	NP_699188	6183	solute carrier family 16, member 11 (monocarboxylic acid transporter 11)
SLC16A14	151473	NP_689740	6075	solute carrier family 16, member 14 (monocarboxylic acid transporter 14)
SLC16A2	6567	NP_006508	3505	solute carrier family 16, member 2 (monocarboxylic acid transporter 8)
SLC16A3	9123	NP_001035887.1	7361	solute carrier family 16 member 3
SLC16A4	9122	NP_004687	3103	solute carrier family 16, member 4 (monocarboxylic acid transporter 5)
SLC16A5	9121	NP_004686	3102	similar to MCT solute carrier family 16, member 5 (monocarboxylic acid transporter 6)
SLC16A6	9120	NP_004685	3101	solute carrier family 16, member 6 (monocarboxylic acid transporter 7)
SLC16A7	9194	NP_004722	3110	solute carrier family 16, member 7 (monocarboxylic acid transporter 2)
SLC16A8	23539	NP_037488	3811	solute carrier family 16, member 8 (monocarboxylic acid transporter 3)
SLC16A9	220963	NP_919274	6715	solute carrier family 16, member 9 (monocarboxylic acid transporter 9)
SLC17A1	6568	NP_005065.2	7376	solute carrier family 17 member 1
SLC17A2	10246	NP_005826	3381	solute carrier family 17 (sodium phosphate), member 2
SLC17A3	10786	NP_001091956	1126	solute carrier family 17 (sodium phosphate), member 3
SLC17A3	10786	NP_006623	3531	solute carrier family 17 (sodium phosphate), member 3
SLC17A4	10050	NP_005486	3294	solute carrier family 17 (sodium phosphate), member 4
SLC17A5	26503	NP_036566	3776	solute carrier family 17 (anion/sugar transporter), member 5
SLC17A7	57030	NP_064705	4627	solute carrier family 17 (sodium-dependent inorganic phosphate cotransporter), member 7
SLC17A8	246213	NP_001138760	1842	solute carrier family 17 (sodium-dependent inorganic phosphate cotransporter), member 8
SLC17A8	246213	NP_647480	5854	solute carrier family 17 (sodium-dependent inorganic phosphate cotransporter), member 8
SLC18A1	6570	NP_001129163	1553	solute carrier family 18 (vesicular monoamine), member 1
SLC18A1	6570	NP_001135796	1641	solute carrier family 18 (vesicular monoamine), member 1
SLC18A1	6570	NP_001135797	1642	solute carrier family 18 (vesicular monoamine), member 1
SLC18A1	6570	NP_003044	2773	solute carrier family 18 (vesicular monoamine), member 1
SLC18A2	6571	NP_003045	2774	solute carrier family 18 (vesicular monoamine), member 2
SLC18A3	6572	NP_003046	2775	solute carrier family 18 (vesicular acetylcholine), member 3
SLC18B1	116843	NP_439896	5513	chromosome 6 open reading frame 192
SLC19A1	6573	NP_919231	6711	solute carrier family 19 (folate transporter), member 1
SLC19A2	10560	NP_008927	3610	solute carrier family 19 (thiamine transporter), member 2
SLC19A3	80704	NP_079519.1	7300	solute carrier family 19 member 3
SLC1A1	6505	NP_004161	2976	solute carrier family 1 (neuronal/epithelialHigh affinity glutamate transporter, system Xag), member 1
SLC1A2	6506	NP_004162	2977	solute carrier family 1 (glialHigh affinity glutamate transporter), member 2
SLC1A3	6507	NP_001160167	2178	solute carrier family 1 (glialHigh affinity glutamate transporter), member 3
SLC1A3	6507	NP_001160168	2179	solute carrier family 1 (glialHigh affinity glutamate transporter), member 3
SLC1A3	6507	NP_004163	2978	solute carrier family 1 (glialHigh affinity glutamate transporter), member 3
SLC1A4	6509	NP_001180422.1	1541	solute carrier family 1 (glutamate/neutral amino acid transporter), member 4
SLC1A4	6509	NP_003029	2760	solute carrier family 1 (glutamate/neutral amino acid transporter), member 4
SLC1A5	6510	NP_001138616	1816	solute carrier family 1 (neutral amino acid transporter), member 5
SLC1A5	6510	NP_001138617	1817	solute carrier family 1 (neutral amino acid transporter), member 5
SLC1A5	6510	NP_005619	3329	solute carrier family 1 (neutral amino acid transporter), member 5
SLC1A6	6511	NP_005062	3206	solute carrier family 1 (high affinity aspartate/glutamate transporter), member 6
SLC1A7	6512	NP_006662	3539	solute carrier family 1 (glutamate transporter), member 7

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
SLC20A1	6574	NP_005406	3282	solute carrier family 20 (phosphate transporter), member 1
SLC20A2	6575	NP_006740	3556	solute carrier family 20 (phosphate transporter), member 2
SLC22A1	6580	NP_003048	2776	solute carrier family 22 (organic cation transporter), member 1
SLC22A1	6580	NP_694857	6155	solute carrier family 22 (organic cation transporter), member 1
SLC22A11	55867	NP_060954	4460	solute carrier family 22 (organic anion/urate transporter), member 11
SLC22A12	116085	NP_653186	5868	solute carrier family 22 (organic anion/urate transporter), member 12
SLC22A12	116085	NP_700357	6189	solute carrier family 22 (organic anion/urate transporter), member 12
SLC22A13	9390	NP_004247	3000	solute carrier family 22 (organic anion transporter), member 13
SLC22A14	9389	NP_004794	3133	solute carrier family 22, member 14
SLC22A15	55356	NP_060890	4446	solute carrier family 22, member 15
SLC22A16	85413	NP_149116	5452	solute carrier family 22 (organic cation/carnitine transporter), member 16
SLC22A17	51310	NP_057693	4262	solute carrier family 22, member 17
SLC22A17	51310	NP_065105	4636	solute carrier family 22, member 17
SLC22A18	5002	NP_002546	2677	solute carrier family 22, member 18
SLC22A18	5002	NP_899056	6704	solute carrier family 22, member 18
SLC22A2	6582	NP_003049	2777	solute carrier family 22 (organic cation transporter), member 2
SLC22A23	63027	NP_056297	4089	solute carrier family 22, member 23
SLC22A23	63027	NP_068764	4850	solute carrier family 22, member 23
SLC22A24	283238	NP_001129978	1609	solute carrier family 22, member 24
SLC22A25	387601	NP_955384	6881	solute carrier family 22, member 25
SLC22A3	6581	NP_068812	4857	solute carrier family 22 (extraneuronal monoamine transporter), member 3
SLC22A31	146429	XP_001714620	7053	Putative solute carrier family 22 member ENSG00000182157
SLC22A31	146429	XP_370997	7088	Putative solute carrier family 22 member ENSG00000182157
SLC22A31	146429	XP_951429	7097	Putative solute carrier family 22 member ENSG00000182157
SLC22A4	6583	NP_003050	2778	solute carrier family 22 (organic cation/ergothioneine transporter), member 4
SLC22A5	6584	NP_003051	2779	solute carrier family 22 (organic cation/carnitine transporter), member 5
SLC22A6	9356	NP_004781	3127	solute carrier family 22 (organic anion transporter), member 6
SLC22A6	9356	NP_695008	6172	solute carrier family 22 (organic anion transporter), member 6
SLC22A6	9356	NP_695009	6173	solute carrier family 22 (organic anion transporter), member 6
SLC22A6	9356	NP_695010	6174	solute carrier family 22 (organic anion transporter), member 6
SLC22A7	10864	NP_006663	3540	solute carrier family 22 (organic anion transporter), member 7
SLC22A7	10864	NP_696961	6175	solute carrier family 22 (organic anion transporter), member 7
SLC22A8	9376	NP_004245	2999	solute carrier family 22 (organic anion transporter), member 8
SLC22A9	114571	NP_543142	5640	solute carrier family 22 (organic anion transporter), member 9
SLC23A1	9963	NP_005838	3383	solute carrier family 23 (nucleobase transporters), member 1
SLC23A1	9963	NP_689898	6097	solute carrier family 23 (nucleobase transporters), member 1
SLC23A2	9962	NP_005107	3214	solute carrier family 23 (nucleobase transporters), member 2
SLC23A2	9962	NP_976072	6923	solute carrier family 23 (nucleobase transporters), member 2
SLC23A3	151295	NP_001138361	1780	solute carrier family 23 (nucleobase transporters), member 3
SLC23A3	151295	NP_001138362	1781	solute carrier family 23 (nucleobase transporters), member 3
SLC23A3	151295	NP_653313	5896	solute carrier family 23 (nucleobase transporters), member 3
SLC24A1	9187	NP_001287960.1	7448	solute carrier family 24 member 1
SLC24A2	25769	NP_065077	4632	solute carrier family 24 (sodium/potassium/calcium exchanger), member 2
SLC24A3	57419	NP_065740	4699	solute carrier family 24 (sodium/potassium/calcium exchanger), member 3
SLC24A4	123041	NP_705932	6209	solute carrier family 24 (sodium/potassium/calcium exchanger), member 4
SLC24A4	123041	NP_705933	6210	solute carrier family 24 (sodium/potassium/calcium exchanger), member 4
SLC24A4	123041	NP_705934	6211	solute carrier family 24 (sodium/potassium/calcium exchanger), member 4
SLC25A17	10478	NP_006349	3474	solute carrier family 25 (mitochondrial carrier peroxisomal membrane protein, 34kDa), member 17
SLC25A18	83733	NP_113669	5233	solute carrier family 25 (mitochondrial carrier), member 18
SLC25A2	83884	NP_114153	5274	solute carrier family 25 (mitochondrial carrier ornithine transporter) member 2
SLC25A20	788	NP_000378	115	solute carrier family 25 (carnitine/acylcarnitine translocase), member 20
SLC25A23	79085	NP_077008	5010	solute carrier family 25 (mitochondrial carrier phosphate carrier), member 23
SLC25A25	114789	NP_001006642	490	solute carrier family 25 (mitochondrial carrier phosphate carrier), member 25
SLC25A25	114789	NP_001006643	491	solute carrier family 25 (mitochondrial carrier phosphate carrier), member 25
SLC25A25	114789	NP_001006644	492	solute carrier family 25 (mitochondrial carrier phosphate carrier), member 25
SLC25A25	114789	NP_443133	5533	solute carrier family 25 (mitochondrial carrier phosphate carrier), member 25
SLC25A26	115286	NP_001158268	2113	solute carrier family 25, member 26
SLC25A26	115286	NP_775742	6343	solute carrier family 25, member 26
SLC25A3	5250	NP_002626	2696	solute carrier family 25 (mitochondrial carrier phosphate carrier), member 3
SLC25A3	5250	NP_005879	3394	solute carrier family 25 (mitochondrial carrier phosphate carrier), member 3
SLC25A3	5250	NP_998776	7042	solute carrier family 25 (mitochondrial carrier phosphate carrier), member 3
SLC25A31	83447	NP_112581	5216	solute carrier family 25 (mitochondrial carrier adenine nucleotide translocator), member 31
SLC25A33	84275	NP_115691	5324	solute carrier family 25, member 33
SLC25A34	284723	NP_997231	7013	solute carrier family 25, member 34
SLC25A39	51629	NP_001137252	1714	solute carrier family 25, member 39
SLC25A39	51629	NP_057100	4151	solute carrier family 25, member 39
SLC25A4	291	NP_001142	1937	solute carrier family 25 (mitochondrial carrier adenine nucleotide translocator), member 5
SLC25A43	203427	NP_660348	5958	solute carrier family 25, member 43
SLC25A44	9673	NP_001273113.1	1547	solute carrier family 25, member 44
SLC25A44	9673	NP_055470	3970	solute carrier family 25, member 44
SLC25A45	283130	NP_001070709	957	solute carrier family 25, member 45
SLC25A45	283130	NP_872362	6659	solute carrier family 25, member 45
SLC25A48	153328	NP_660325	5952	mitochondrial carrier protein-like
SLC25A5	292	NP_001143	1938	solute carrier family 25 (mitochondrial carrier adenine nucleotide translocator), member 5
SLC26A1	10861	NP_071325	4866	solute carrier family 26 (sulfate transporter), member 1
SLC26A1	10861	NP_602297	5707	solute carrier family 26 (sulfate transporter), member 1
SLC26A1	10861	NP_998778	7043	solute carrier family 26 (sulfate transporter), member 1

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
SLC26A10	65012	NP_597996.2	7350	solute carrier family 26 member 10
SLC26A11	284129	NP_001159819	2160	solute carrier family 26, member 11
SLC26A11	284129	NP_001159820	2161	solute carrier family 26, member 11
SLC26A11	284129	NP_001159821	2162	solute carrier family 26, member 11
SLC26A11	284129	NP_775897	6366	solute carrier family 26, member 11
SLC26A2	1836	NP_000103	32	solute carrier family 26 (sulfate transporter), member 2
SLC26A3	1811	NP_000102	31	solute carrier family 26, member 3
SLC26A4	5172	NP_000432	127	solute carrier family 26, member 4
SLC26A6	65010	NP_001035544	859	solute carrier family 26, member 6 cadherin, EGF LAG seven-pass G-type receptor 3 (flamingoHomolog, Drosophila)
SLC26A6	65010	NP_001035544.1	2363	solute carrier family 26, member 6 cadherin, EGF LAG seven-pass G-type receptor 3 (flamingoHomolog, Drosophila)
SLC26A6	65010	NP_001035544.1	4962	solute carrier family 26, member 6 cadherin, EGF LAG seven-pass G-type receptor 3 (flamingoHomolog, Drosophila)
SLC26A6	65010	NP_599025	5703	solute carrier family 26, member 6 cadherin, EGF LAG seven-pass G-type receptor 3 (flamingoHomolog, Drosophila)
SLC26A6	65010	NP_602298	5708	solute carrier family 26, member 6 cadherin, EGF LAG seven-pass G-type receptor 3 (flamingoHomolog, Drosophila)
SLC26A7	115111	NP_439897	5514	solute carrier family 26, member 7
SLC26A7	115111	NP_599028	5705	solute carrier family 26, member 7
SLC26A8	116369	NP_443193	5552	solute carrier family 26, member 8
SLC26A8	116369	NP_619732	5777	solute carrier family 26, member 8
SLC26A9	115019	NP_443166.1	1677	solute carrier family 26, member 9
SLC26A9	115019	NP_443166	5543	solute carrier family 26, member 9
SLC26A9	115019	NP_599152	5706	solute carrier family 26, member 9
SLC27A1	376497	NP_940982	6820	solute carrier family 27 (fatty acid transporter), member 1
SLC27A2	11001	NP_001153101	1948	solute carrier family 27 (fatty acid transporter), member 2
SLC27A2	11001	NP_003636	2862	solute carrier family 27 (fatty acid transporter), member 2
SLC27A3	11000	NP_077306	5025	solute carrier family 27 (fatty acid transporter), member 3
SLC27A4	10999	NP_005085	3211	solute carrier family 27 (fatty acid transporter), member 4
SLC27A5	10998	NP_036386	3728	solute carrier family 27 (fatty acid transporter), member 5
SLC28A1	9154	NP_004204	2992	solute carrier family 28 (sodium-coupled nucleoside transporter), member 1
SLC28A1	9154	NP_964014	6918	solute carrier family 28 (sodium-coupled nucleoside transporter), member 1
SLC28A2	9153	NP_004203	2991	solute carrier family 28 (sodium-coupled nucleoside transporter), member 2
SLC28A3	64078	NP_071410	4878	solute carrier family 28 (sodium-coupled nucleoside transporter), member 3
SLC29A1	2030	NP_001071643.1	980	solute carrier family 29 (nucleoside transporters), member 1
SLC29A1	2030	NP_001071643	981	solute carrier family 29 (nucleoside transporters), member 1
SLC29A1	2030	NP_001071643.1	982	solute carrier family 29 (nucleoside transporters), member 1
SLC29A1	2030	NP_001071645	983	solute carrier family 29 (nucleoside transporters), member 1
SLC29A1	2030	NP_001071643.1	3177	solute carrier family 29 (nucleoside transporters), member 1
SLC29A2	3177	NP_001523	2407	solute carrier family 29 (nucleoside transporters), member 2
SLC29A3	55315	NP_001138722	1830	solute carrier family 29 (nucleoside transporters), member 3
SLC29A3	55315	NP_060814	4426	solute carrier family 29 (nucleoside transporters), member 3
SLC29A4	222962	NP_001035751	878	solute carrier family 29 (nucleoside transporters), member 4
SLC29A4	222962	NP_694979	6165	solute carrier family 29 (nucleoside transporters), member 4
SLC2A1	6513	NP_006507	3504	solute carrier family 2 (facilitated glucose transporter), member 1
SLC2A10	81031	NP_110404	5169	solute carrier family 2 (facilitated glucose transporter), member 10
SLC2A12	154091	NP_660159	5933	solute carrier family 2 (facilitated glucose transporter), member 12
SLC2A13	114134	NP_443117	5526	solute carrier family 2 (facilitated glucose transporter), member 13
SLC2A2	6514	NP_000331.1	7272	solute carrier family 2 member 2
SLC2A3	6515	NP_008862	3598	solute carrier family 2 (facilitated glucose transporter), member 3
SLC2A4	6517	NP_001033	781	solute carrier family 2 (facilitated glucose transporter), member 4
SLC2A5	6518	NP_001129057	1542	solute carrier family 2 (facilitated glucose/fructose transporter), member 5
SLC2A5	6518	NP_003030	2761	solute carrier family 2 (facilitated glucose/fructose transporter), member 5
SLC2A6	11182	NP_001138571	1809	solute carrier family 2 (facilitated glucose transporter), member 6
SLC2A6	11182	NP_060055	4309	solute carrier family 2 (facilitated glucose transporter), member 6
SLC2A8	29988	NP_055395	3960	solute carrier family 2 (facilitated glucose transporter), member 8
SLC2A9	56606	NP_001001290.1	7335	solute carrier family 2 member 9
SLC30A1	7779	NP_067017	4791	solute carrier family 30 (zinc transporter), member 1
SLC30A10	55532	NP_061183	4487	solute carrier family 30, member 10
SLC30A2	7780	NP_001004434	434	solute carrier family 30 (zinc transporter), member 2
SLC30A2	7780	NP_115902	5364	solute carrier family 30 (zinc transporter), member 2
SLC30A3	7781	NP_003450	2834	solute carrier family 30 (zinc transporter), member 3
SLC30A4	7782	NP_037441	3801	solute carrier family 30 (zinc transporter), member 4
SLC30A5	64924	NP_075053	4960	solute carrier family 30 (zinc transporter), member 5
SLC30A5	64924	NP_076960	5000	solute carrier family 30 (zinc transporter), member 5
SLC30A6	55676	NP_060434	4375	solute carrier family 30 (zinc transporter), member 6
SLC30A7	148867	NP_001138356	1779	solute carrier family 30 (zinc transporter), member 7
SLC30A7	148867	NP_598003	5697	solute carrier family 30 (zinc transporter), member 7
SLC30A8	169026	NP_776250	6394	solute carrier family 30 (zinc transporter), member 8
SLC30A9	10463	NP_006336	3473	solute carrier family 30 (zinc transporter), member 9
SLC31A1	1317	NP_001850	2489	solute carrier family 31 (copper transporters), member 1
SLC31A2	1318	NP_001851	2490	solute carrier family 31 (copper transporters), member 2
SLC32A1	140679	NP_542119	5606	solute carrier family 32 (GABA vesicular transporter), member 1
SLC33A1	9197	NP_004724	3111	solute carrier family 33 (acetyl-CoA transporter), member 1
SLC34A1	6569	NP_001161051	2184	solute carrier family 34 (sodium phosphate), member 1
SLC34A1	6569	NP_003043	2772	solute carrier family 34 (sodium phosphate), member 1

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
SLC34A2	10568	NP_006415	3490	solute carrier family 34 (sodium phosphate), member 2
SLC34A3	142680	NP_543153	5644	solute carrier family 34 (sodium phosphate), member 3
SLC35A1	10559	NP_001161870	2231	solute carrier family 35 (CMP-sialic acid transporter), member A1
SLC35A1	10559	NP_006407	3488	solute carrier family 35 (CMP-sialic acid transporter), member A1
SLC35A2	7355	NP_001027460	742	solute carrier family 35 (UDP-galactose transporter), member A2
SLC35A2	7355	NP_001035963	900	solute carrier family 35 (UDP-galactose transporter), member A2
SLC35A2	7355	NP_005651	3336	solute carrier family 35 (UDP-galactose transporter), member A2
SLC35A3	23443	NP_036375	3726	solute carrier family 35 (UDP-N-acetylglucosamine (UDP-GlcNAc) transporter), member A3
SLC35A4	113829	NP_542401	5615	solute carrier family 35, member A4
SLC35A5	55032	NP_060415	4374	solute carrier family 35, member A5
SLC35B1	10237	NP_005818	3378	solute carrier family 35, member B1
SLC35B2	347734	NP_835361	6520	solute carrier family 35, member B2
SLC35B3	51000	NP_001136013	1670	solute carrier family 35, member B3
SLC35B3	51000	NP_001136013	1671	solute carrier family 35, member B3
SLC35B3	51000	NP_057032	4141	solute carrier family 35, member B3
SLC35B4	84912	NP_116215	5411	solute carrier family 35, member B4
SLC35C1	55343	NP_001138737	1831	solute carrier family 35, member C1
SLC35C1	55343	NP_001138738	1832	solute carrier family 35, member C1
SLC35C1	55343	NP_060859	4437	solute carrier family 35, member C1
SLC35C2	51006	NP_057029	4140	solute carrier family 35, member C2
SLC35C2	51006	NP_775096	6330	solute carrier family 35, member C2
SLC35C2	51006	NP_775271	6338	solute carrier family 35, member C2
SLC35D1	23169	NP_055954	4040	solute carrier family 35 (UDP-glucuronic acid/UDP-N-acetylgalactosamine dual transporter), member D1
SLC35D2	11046	NP_008932	3612	solute carrier family 35, member D2
SLC35D3	340146	NP_001008783	554	solute carrier family 35, member D3
SLC35E1	79939	NP_079157	5095	solute carrier family 35, member E1
SLC35E2	9906	NP_001104251	1242	solute carrier family 35, member E2
SLC35E2	9906	NP_878258	6688	solute carrier family 35, member E2
SLC35E3	55508	NP_061126	4480	solute carrier family 35, member E3
SLC35E4	339665	NP_001001479	361	solute carrier family 35, member E4
SLC35F1	222553	NP_001025029	710	solute carrier family 35, member F1
SLC35F2	54733	NP_059985	4299	solute carrier family 35, member F2
SLC35F4	341880	NP_001193849.1	1025	solute carrier family 35, member F4
SLC35F5	80255	NP_079457	5131	solute carrier family 35, member F5
SLC35F6	54978	NP_060347	4362	chromosome 2 open reading frame 18
SLC35G1	159371	NP_001128130	1500	transmembrane protein 20
SLC35G1	159371	NP_694958	6162	transmembrane protein 20
SLC35G3	146861	NP_001096084	1210	acyl-malonyl condensing enzyme 1-like 3, 1, 2, enzyme 1
SLC35G3	146861	NP_473369	5569	acyl-malonyl condensing enzyme 1-like 3, 1, 2, enzyme 1
SLC35G3	146861	NP_689675	6063	acyl-malonyl condensing enzyme 1-like 3, 1, 2, enzyme 1
SLC36A1	206358	NP_510968	5593	solute carrier family 36 (proton/amino acid symporter), member 1
SLC36A4	120103	NP_689526	6031	solute carrier family 36 (proton/amino acid symporter), member 4
SLC37A1	54020	NP_061837	4536	solute carrier family 37 (glycerol-3-phosphate transporter), member 1
SLC37A3	84255	NP_115671	5320	solute carrier family 37 (glycerol-3-phosphate transporter), member 3
SLC37A3	84255	NP_996996	7000	solute carrier family 37 (glycerol-3-phosphate transporter), member 3
SLC37A4	2542	NP_001157749	2073	solute carrier family 37 (glucose-6-phosphate transporter), member 4
SLC37A4	2542	NP_001157750	2074	solute carrier family 37 (glucose-6-phosphate transporter), member 4
SLC37A4	2542	NP_001157751	2075	solute carrier family 37 (glucose-6-phosphate transporter), member 4
SLC37A4	2542	NP_001157752	2076	solute carrier family 37 (glucose-6-phosphate transporter), member 4
SLC37A4	2542	NP_001458	2383	solute carrier family 37 (glucose-6-phosphate transporter), member 4
SLC38A1	81539	NP_001070952.1	7367	solute carrier family 38 member 1
SLC38A10	124565	NP_001033073	782	solute carrier family 38, member 10
SLC38A10	124565	NP_612637	5760	solute carrier family 38, member 10
SLC38A11	151258	NP_775783	6350	solute carrier family 38, member 11
SLC38A2	54407	NP_061849	4542	solute carrier family 38, member 2
SLC38A3	10991	NP_006832	3577	solute carrier family 38, member 3
SLC38A4	55089	NP_001137296.1	7402	solute carrier family 38 member 4
SLC38A5	92745	NP_277053	5504	solute carrier family 38, member 5
SLC38A6	145389	NP_722518	6230	solute carrier family 38, member 6
SLC38A7	55238	NP_060701	4405	solute carrier family 38, member 7
SLC38A9	153129	NP_775785	6351	solute carrier family 38, member 9
SLC39A1	27173	NP_055252	3938	solute carrier family 39 (zinc transporter), member 1
SLC39A10	57181	NP_001120729.1	7389	solute carrier family 39 member 10
SLC39A11	201266	NP_001153242	1959	solute carrier family 39 (metal ion transporter), member 11
SLC39A11	201266	NP_631916	5845	solute carrier family 39 (metal ion transporter), member 11
SLC39A12	221074	NP_001138667	1826	solute carrier family 39 (zinc transporter), member 12
SLC39A12	221074	NP_689938	6106	solute carrier family 39 (zinc transporter), member 12
SLC39A13	91252	NP_001121697	1370	solute carrier family 39 (zinc transporter), member 13
SLC39A13	91252	NP_689477	6022	solute carrier family 39 (zinc transporter), member 13
SLC39A14	23516	NP_001121903	1373	solute carrier family 39 (zinc transporter), member 14
SLC39A14	23516	NP_001128625	1529	solute carrier family 39 (zinc transporter), member 14
SLC39A14	23516	NP_001128626	1530	solute carrier family 39 (zinc transporter), member 14
SLC39A14	23516	NP_056174	4068	solute carrier family 39 (zinc transporter), member 14
SLC39A2	29986	NP_055394	3959	solute carrier family 39 (zinc transporter), member 2
SLC39A3	29985	NP_653165	5861	solute carrier family 39 (zinc transporter), member 3
SLC39A3	29985	NP_998733	7037	solute carrier family 39 (zinc transporter), member 3
SLC39A4	55630	NP_060237	4339	solute carrier family 39 (zinc transporter), member 4

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
SLC39A4	55630	NP_570901	5672	solute carrier family 39 (zinc transporter), member 4
SLC39A5	283375	NP_001128667	1537	solute carrier family 39 (metal ion transporter), member 5
SLC39A5	283375	NP_775867	6362	solute carrier family 39 (metal ion transporter), member 5
SLC39A6	25800	NP_001092876	1158	solute carrier family 39 (zinc transporter), member 6
SLC39A6	25800	NP_036451	3745	solute carrier family 39 (zinc transporter), member 6
SLC39A7	7922	NP_001070984	973	solute carrier family 39 (zinc transporter), member 7
SLC39A7	7922	NP_008910	3605	solute carrier family 39 (zinc transporter), member 7
SLC39A8	64116	NP_001128618	1526	solute carrier family 39 (zinc transporter), member 8
SLC39A8	64116	NP_001128619	1527	solute carrier family 39 (zinc transporter), member 8
SLC39A8	64116	NP_001128620	1528	solute carrier family 39 (zinc transporter), member 8
SLC39A8	64116	NP_071437	4889	solute carrier family 39 (zinc transporter), member 8
SLC39A9	55334	NP_060845	4435	solute carrier family 39 (zinc transporter), member 9
SLC3A1	6519	NP_000332	100	solute carrier family 3 (cystine, dibasic and neutral amino acid transporters, activator of cystine, dibasic and neutral amino acid transport), member 1
SLC3A2	6520	NP_001012680.1	600	solute carrier family 3 (activators of dibasic and neutral amino acid transport), member 2
SLC3A2	6520	NP_001012680	601	solute carrier family 3 (activators of dibasic and neutral amino acid transport), member 2
SLC3A2	6520	NP_001012680.1	602	solute carrier family 3 (activators of dibasic and neutral amino acid transport), member 2
SLC3A2	6520	NP_001012682	603	solute carrier family 3 (activators of dibasic and neutral amino acid transport), member 2
SLC3A2	6520	NP_001013269	610	solute carrier family 3 (activators of dibasic and neutral amino acid transport), member 2
SLC3A2	6520	NP_002385	2641	solute carrier family 3 (activators of dibasic and neutral amino acid transport), member 2
SLC40A1	30061	NP_055400	3961	solute carrier family 40 (iron-regulated transporter), member 1
SLC41A1	254428	NP_776253	6397	solute carrier family 41, member 1
SLC41A2	84102	NP_115524	5302	solute carrier family 41, member 2
SLC41A3	54946	NP_001008485	537	solute carrier family 41, member 3
SLC41A3	54946	NP_001008486	538	solute carrier family 41, member 3
SLC41A3	54946	NP_001008487	539	solute carrier family 41, member 3
SLC41A3	54946	NP_001157947	2083	solute carrier family 41, member 3
SLC41A3	54946	NP_060306	4352	solute carrier family 41, member 3
SLC43A1	8501	NP_003618	2856	solute carrier family 43, member 1
SLC43A2	124935	NP_689559	6037	solute carrier family 43, member 2
SLC43A3	29015	NP_054815	3869	solute carrier family 43, member 3
SLC43A3	29015	NP_060081	4312	solute carrier family 43, member 3
SLC43A3	29015	NP_955361	6875	solute carrier family 43, member 3
SLC44A1	23446	NP_536856	5605	solute carrier family 44, member 1
SLC44A2	57153	NP_001138528	1807	solute carrier family 44, member 2
SLC44A2	57153	NP_065161	4655	solute carrier family 44, member 2
SLC44A3	126969	NP_001107578	1268	solute carrier family 44, member 3
SLC44A3	126969	NP_689582	6044	solute carrier family 44, member 3
SLC44A4	80736	NP_079533	5142	solute carrier family 44, member 4
SLC44A5	204962	NP_001123530	1412	solute carrier family 44, member 5
SLC44A5	204962	NP_689910	6101	solute carrier family 44, member 5
SLC45A1	50651	NP_001073866	1019	solute carrier family 45, member 1
SLC45A2	51151	NP_001012527	597	solute carrier family 45, member 2
SLC45A2	51151	NP_057264	4181	solute carrier family 45, member 2
SLC45A3	85414	NP_149093	5450	solute carrier family 45, member 3
SLC45A4	57210	NP_001073900	1021	solute carrier family 45, member 4
SLC46A1	113235	NP_542400	5614	solute carrier family 46 (folate transporter), member 1
SLC46A2	57864	NP_149040	5444	solute carrier family 46, member 2
SLC46A3	283537	NP_001129391	1576	solute carrier family 46, member 3
SLC46A3	283537	NP_861450	6634	solute carrier family 46, member 3
SLC47A1	55244	NP_060712	4407	solute carrier family 47, member 1
SLC47A2	146802	NP_001093116	1162	solute carrier family 47, member 2
SLC47A2	146802	NP_690872	6131	solute carrier family 47, member 2
SLC4A1	6521	NP_000333	101	solute carrier family 4, anion exchanger, member 1 (erythrocyte membrane protein band 3, Diego blood group)
SLC4A10	57282	NP_001171486.1	7421	solute carrier family 4 member 10
SLC4A11	83959	NP_114423	5282	solute carrier family 4, sodium borate transporter, member 11
SLC4A2	6522	NP_003031	2762	solute carrier family 4, anion exchanger, member 2 (erythrocyte membrane protein band 3-like 1)
SLC4A3	6508	NP_005061	3205	solute carrier family 4, anion exchanger, member 3
SLC4A3	6508	NP_963868	6910	solute carrier family 4, anion exchanger, member 3
SLC4A4	8671	NP_001091954	1125	solute carrier family 4, sodium bicarbonate cotransporter, member 4
SLC4A4	8671	NP_001128214	1504	solute carrier family 4, sodium bicarbonate cotransporter, member 4
SLC4A4	8671	NP_003750	2885	solute carrier family 4, sodium bicarbonate cotransporter, member 4
SLC4A5	57835	NP_067019	4793	solute carrier family 4, sodium bicarbonate cotransporter, member 5
SLC4A5	57835	NP_597812	5695	solute carrier family 4, sodium bicarbonate cotransporter, member 5
SLC4A7	9497	NP_003606	2855	solute carrier family 4, sodium bicarbonate cotransporter, member 7
SLC4A8	9498	NP_001035049	818	solute carrier family 4, sodium bicarbonate cotransporter, member 8
SLC4A8	9498	NP_001035049.1	3150	solute carrier family 4, sodium bicarbonate cotransporter, member 8
SLC4A9	83697	NP_113655	5229	solute carrier family 4, sodium bicarbonate cotransporter, member 9
SLC50A1	55974	NP_001116309	1288	recombination activating gene 1 activating protein 1
SLC50A1	55974	NP_001116311	1289	recombination activating gene 1 activating protein 1
SLC50A1	55974	NP_061333	4493	recombination activating gene 1 activating protein 1
SLC51A	200931	NP_689885	6094	organic solute transporter alpha
SLC51B	123264	NP_849190	6573	organic solute transporter beta
SLC52A1	55065	NP_001098047	1217	G protein-coupled receptor 172B
SLC52A1	55065	NP_060456	4378	G protein-coupled receptor 172B
SLC52A2	79581	NP_078807	5042	G protein-coupled receptor 172A
SLC52A3	113278	NP_212134	5494	chromosome 20 open reading frame 54

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
SLC5A1	6523	NP_000334	102	solute carrier family 5 (sodium/glucose cotransporter), member 1
SLC5A10	125206	NP_001035915	888	solute carrier family 5 (sodium/glucose cotransporter), member 10
SLC5A10	125206	NP_689564	6038	solute carrier family 5 (sodium/glucose cotransporter), member 10
SLC5A11	115584	NP_443176	5546	solute carrier family 5 (sodium/glucose cotransporter), member 11
SLC5A12	159963	NP_848593	6545	solute carrier family 5 (sodium/glucose cotransporter), member 12
SLC5A2	6524	NP_003032	2763	solute carrier family 5 (sodium/glucose cotransporter), member 2
SLC5A3	6526	NP_008864	3599	solute carrier family 5 (sodium/myo-inositol cotransporter), member 3
SLC5A4	6527	NP_055042	3889	solute carrier family 5 (low affinity glucose cotransporter), member 4
SLC5A5	6528	NP_000444	134	solute carrier family 5 (sodium iodide symporter), member 5
SLC5A6	8884	NP_066918	4767	solute carrier family 5 (sodium-dependent vitamin transporter), member 6
SLC5A7	60482	NP_068587	4837	solute carrier family 5 (choline transporter), member 7
SLC5A8	160728	NP_666018	5987	solute carrier family 5 (iodide transporter), member 8
SLC5A9	200010	NP_001011547	587	solute carrier family 5 (sodium/glucose cotransporter), member 9
SLC5A9	200010	NP_001128653	1536	solute carrier family 5 (sodium/glucose cotransporter), member 9
SLC6A1	6529	NP_003033	2764	solute carrier family 6 (neurotransmitter transporter, GABA), member 1
SLC6A11	6538	NP_055044	3890	solute carrier family 6 (neurotransmitter transporter, GABA), member 11
SLC6A12	6539	NP_001116319	1290	solute carrier family 6 (neurotransmitter transporter, betaine/GABA), member 12
SLC6A12	6539	NP_001116320	1291	solute carrier family 6 (neurotransmitter transporter, betaine/GABA), member 12
SLC6A12	6539	NP_003035	2766	solute carrier family 6 (neurotransmitter transporter, betaine/GABA), member 12
SLC6A13	6540	NP_057699	4264	solute carrier family 6 (neurotransmitter transporter, GABA), member 13
SLC6A14	11254	NP_009162	3656	solute carrier family 6 (amino acid transporter), member 14
SLC6A15	55117	NP_001139807	1932	solute carrier family 6 (neutral amino acid transporter), member 15
SLC6A15	55117	NP_060527	4385	solute carrier family 6 (neutral amino acid transporter), member 15
SLC6A15	55117	NP_877499	6682	solute carrier family 6 (neutral amino acid transporter), member 15
SLC6A16	28968	NP_054756	3859	solute carrier family 6, member 16
SLC6A18	348932	NP_872438	6668	solute carrier family 6, member 18
SLC6A19	340024	NP_001003841	424	solute carrier family 6 (neutral amino acid transporter), member 19
SLC6A2	6530	NP_001034	785	solute carrier family 6 (neurotransmitter transporter, noradrenalin), member 2
SLC6A20	54716	NP_064593	4617	solute carrier family 6 (proline IMINO transporter), member 20
SLC6A20	54716	NP_071800	4912	solute carrier family 6 (proline IMINO transporter), member 20
SLC6A3	6531	NP_001035	816	solute carrier family 6 (neurotransmitter transporter, dopamine), member 3
SLC6A4	6532	NP_001036	902	solute carrier family 6 (neurotransmitter transporter, serotonin), member 4
SLC6A5	9152	NP_004202	2990	solute carrier family 6 (neurotransmitter transporter, glycine), member 5
SLC6A6	6533	NP_001127839	1477	solute carrier family 6 (neurotransmitter transporter, taurine), member 6
SLC6A6	6533	NP_001127840	1478	solute carrier family 6 (neurotransmitter transporter, taurine), member 6
SLC6A6	6533	NP_003034	2765	solute carrier family 6 (neurotransmitter transporter, taurine), member 6
SLC6A7	6534	NP_055043.2	7370	solute carrier family 6 member 7
SLC6A8	6535	NP_001136277	1704	solute carrier family 6 (neurotransmitter transporter, creatine), member 8
SLC6A8	6535	NP_001136278	1705	solute carrier family 6 (neurotransmitter transporter, creatine), member 8
SLC6A8	6535	NP_005620	3330	solute carrier family 6 (neurotransmitter transporter, creatine), member 8
SLC6A9	6536	NP_001020016	688	solute carrier family 6 (neurotransmitter transporter, glycine), member 9
SLC6A9	6536	NP_008865	3600	solute carrier family 6 (neurotransmitter transporter, glycine), member 9
SLC6A9	6536	NP_964012	6917	solute carrier family 6 (neurotransmitter transporter, glycine), member 9
SLC7A1	6541	NP_003036	2767	solute carrier family 7 (cationic amino acid transporter, y+ system), member 1
SLC7A10	56301	NP_062823	4585	solute carrier family 7, (neutral amino acid transporter, y+ system) member 10
SLC7A11	23657	NP_055146	3915	solute carrier family 7, (cationic amino acid transporter, y+ system) member 11
SLC7A13	157724	NP_620172.2	7353	solute carrier family 7 member 13
SLC7A14	57709	NP_066000	4744	solute carrier family 7 (cationic amino acid transporter, y+ system), member 14
SLC7A2	6542	NP_001008539	546	solute carrier family 7 (cationic amino acid transporter, y+ system), member 2
SLC7A2	6542	NP_001158243	2112	solute carrier family 7 (cationic amino acid transporter, y+ system), member 2
SLC7A2	6542	NP_003037	2768	solute carrier family 7 (cationic amino acid transporter, y+ system), member 2
SLC7A4	6545	NP_004164	2979	solute carrier family 7 (cationic amino acid transporter, y+ system), member 4
SLC7A5	8140	NP_003477.4	7352	solute carrier family 7 member 5
SLC7A6	9057	NP_001070253	951	solute carrier family 7 (cationic amino acid transporter, y+ system), member 6
SLC7A6	9057	NP_003974	2939	solute carrier family 7 (cationic amino acid transporter, y+ system), member 6
SLC7A7	9056	NP_001119577	1301	solute carrier family 7 (cationic amino acid transporter, y+ system), member 7
SLC7A7	9056	NP_001119578	1302	solute carrier family 7 (cationic amino acid transporter, y+ system), member 7
SLC7A7	9056	NP_003973	2938	solute carrier family 7 (cationic amino acid transporter, y+ system), member 7
SLC7A8	23428	NP_036376	3727	solute carrier family 7 (cationic amino acid transporter, y+ system), member 8
SLC7A8	23428	NP_877392	6678	solute carrier family 7 (cationic amino acid transporter, y+ system), member 8
SLC7A9	11136	NP_001119807	1305	solute carrier family 7 (cationic amino acid transporter, y+ system), member 9
SLC7A9	11136	NP_055085	3904	solute carrier family 7 (cationic amino acid transporter, y+ system), member 9
SLC8A1	6546	NP_001106271	1250	solute carrier family 8 (sodium/calcium exchanger), member 1
SLC8A1	6546	NP_001106272	1251	solute carrier family 8 (sodium/calcium exchanger), member 1
SLC8A1	6546	NP_001106273	1252	solute carrier family 8 (sodium/calcium exchanger), member 1
SLC8A1	6546	NP_066920	4769	solute carrier family 8 (sodium/calcium exchanger), member 1
SLC8A2	6543	NP_055878	4028	solute carrier family 8 (sodium/calcium exchanger), member 2
SLC8A3	6547	NP_001123889	1424	solute carrier family 8 (sodium/calcium exchanger), member 3
SLC8A3	6547	NP_150287	5476	solute carrier family 8 (sodium/calcium exchanger), member 3
SLC8A3	6547	NP_489479	5587	solute carrier family 8 (sodium/calcium exchanger), member 3
SLC8A3	6547	NP_891977	6694	solute carrier family 8 (sodium/calcium exchanger), member 3
SLC8A3	6547	NP_891981	6695	solute carrier family 8 (sodium/calcium exchanger), member 3
SLC8A3	6547	NP_892114	6700	solute carrier family 8 (sodium/calcium exchanger), member 3
SLC8B1	80024	NP_079235	5109	solute carrier family 24 (sodium/potassium/calcium exchanger), member 6
SLC9A1	6548	NP_003038.2	7322	solute carrier family 9 member A1
SLC9A2	6549	NP_003039	2769	solute carrier family 9 (sodium/hydrogen exchanger), member 2
SLC9A3	6550	NP_004165	2980	solute carrier family 9 (sodium/hydrogen exchanger), member 3

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
SLC9A5	6553	NP_004585	3081	solute carrier family 9 (sodium/hydrogen exchanger), member 5
SLC9A6	10479	NP_001036002	903	solute carrier family 9 (sodium/hydrogen exchanger), member 6
SLC9A6	10479	NP_006350	3475	solute carrier family 9 (sodium/hydrogen exchanger), member 6
SLC9A7	84679	NP_115980	5375	solute carrier family 9 (sodium/hydrogen exchanger), member 7
SLC9A8	23315	NP_056081	4055	solute carrier family 9 (sodium/hydrogen exchanger), member 8
SLC9A9	285195	NP_775924	6373	solute carrier family 9 (sodium/hydrogen exchanger), member 9
SLC9B1	150159	NP_001094344	1196	similar to CG10806-like, similar to Na+/H+ exchanger domain containing 1
SLC9B1	150159	NP_631912	5843	similar to CG10806-like, similar to Na+/H+ exchanger domain containing 1
SLC9B2	133308	NP_849155	6567	Na+/H+ exchanger domain containing 2
SLC9C2	284525	NP_848622.2	7407	solute carrier family 9 member C2 (putative)
SLCO1A2	6579	NP_066580.1	7290	solute carrier organic anion transporter family member 1A2
SLCO1B1	10599	NP_006437	3492	solute carrier organic anion transporter family, member 1B1
SLCO1B3	28234	NP_062818	4582	solute carrier organic anion transporter family, member 1B3
SLCO1C1	53919	NP_001139416	1885	solute carrier organic anion transporter family, member 1C1
SLCO1C1	53919	NP_001139417	1886	solute carrier organic anion transporter family, member 1C1
SLCO1C1	53919	NP_001139418	1887	solute carrier organic anion transporter family, member 1C1
SLCO1C1	53919	NP_059131	4290	solute carrier organic anion transporter family, member 1C1
SLCO2A1	6578	NP_005621	3331	solute carrier organic anion transporter family, member 2A1
SLCO2B1	11309	NP_001138683	1828	solute carrier organic anion transporter family, member 2B1
SLCO2B1	11309	NP_001138684	1829	solute carrier organic anion transporter family, member 2B1
SLCO2B1	11309	NP_009187	3662	solute carrier organic anion transporter family, member 2B1
SLCO3A1	28232	NP_001138516	1806	solute carrier organic anion transporter family, member 3A1
SLCO3A1	28232	NP_037404	3793	solute carrier organic anion transporter family, member 3A1
SLCO4A1	28231	NP_057438	4209	solute carrier organic anion transporter family, member 4A1
SLCO4C1	353189	NP_851322	6579	solute carrier organic anion transporter family, member 4C1
SLCO5A1	81796	NP_001139480	1895	solute carrier organic anion transporter family, member 5A1
SLCO5A1	81796	NP_001139481	1896	solute carrier organic anion transporter family, member 5A1
SLCO5A1	81796	NP_112220	5203	solute carrier organic anion transporter family, member 5A1
SLCO6A1	133482	NP_775759	6345	solute carrier organic anion transporter family, member 6A1
SLITRK1	114798	NP_443142	5536	SLIT and NTRK-like family, member 1
SLITRK2	84631	NP_001137475	1750	similar to CXorf2 protein SLIT and NTRK-like family, member 2
SLITRK2	84631	NP_001137476	1751	similar to CXorf2 protein SLIT and NTRK-like family, member 2
SLITRK2	84631	NP_001137477	1752	similar to CXorf2 protein SLIT and NTRK-like family, member 2
SLITRK2	84631	NP_001137478	1753	similar to CXorf2 protein SLIT and NTRK-like family, member 2
SLITRK2	84631	NP_001137479	1754	similar to CXorf2 protein SLIT and NTRK-like family, member 2
SLITRK2	84631	NP_001137480	1755	similar to CXorf2 protein SLIT and NTRK-like family, member 2
SLITRK2	84631	NP_001137481	1756	similar to CXorf2 protein SLIT and NTRK-like family, member 2
SLITRK2	84631	NP_001137482	1757	similar to CXorf2 protein SLIT and NTRK-like family, member 2
SLITRK2	84631	NP_115928	5368	similar to CXorf2 protein SLIT and NTRK-like family, member 2
SLITRK3	22865	NP_055741	4015	SLIT and NTRK-like family, member 3
SLITRK4	139065	NP_775101	6333	SLIT and NTRK-like family, member 4
SLITRK5	26050	NP_056382	4103	SLIT and NTRK-like family, member 5
SLITRK6	84189	NP_115605	5310	SLIT and NTRK-like family, member 6
SLMAP	7871	NP_009090	3635	sarcolemma associated protein
SLN	6588	NP_003054	2780	sarcolipin
SMAD2	4087	NP_001003652	409	SMAD family member 2
SMAD2	4087	NP_001129409	1577	SMAD family member 2
SMAD2	4087	NP_005892	3396	SMAD family member 2
SMAGP	57228	NP_001026798	719	small trans-membrane and glycosylated protein
SMAGP	57228	NP_001029045	757	small trans-membrane and glycosylated protein
SMCO3	440087	NP_001013720	618	chromosome 12 open reading frame 69
SMDT1	91689	NP_201575	5487	chromosome 22 open reading frame 32
SMIM1	388588	NP_001157196	2056	hypothetical LOC388588
SMIM10	644538	NP_001156910	2053	hypothetical protein LOC644538
SMIM11	54065	NP_478062	5583	family with sequence similarity 165, member B
SMIM12	113444	NP_001158296	2114	chromosome 1 open reading frame 212
SMIM12	113444	NP_001158297	2115	chromosome 1 open reading frame 212
SMIM12	113444	NP_612437	5749	chromosome 1 open reading frame 212
SMIM14	201895	NP_777581	6410	chromosome 4 open reading frame 34
SMIM15	643155	NP_001041714	932	chromosome 5 open reading frame 43
SMIM17	147670	XP_001718525	7060	hypothetical protein LOC147670
SMIM17	147670	XP_001718558	7061	hypothetical protein LOC147670
SMIM17	147670	XP_001719239	7064	hypothetical protein LOC147670
SMIM19	114926	NP_001129146	1549	chromosome 8 open reading frame 40
SMIM19	114926	NP_001129147	1550	chromosome 8 open reading frame 40
SMIM19	114926	NP_001129148	1551	chromosome 8 open reading frame 40
SMIM19	114926	NP_612445	5752	chromosome 8 open reading frame 40
SMIM2	79024	XP_935153	7090	hypothetical protein MGC5590
SMIM2	79024	XP_946545	7095	hypothetical protein MGC5590
SMIM20	389203	NP_001138904	1857	hypothetical protein LOC389203
SMIM22	440335	XP_001722243	7071	hypothetical protein LOC440335
SMIM22	440335	XP_002343454	7077	hypothetical protein LOC440335
SMIM22	440335	XP_002343455	7078	hypothetical protein LOC440335
SMIM22	440335	XP_002344988	7079	hypothetical protein LOC440335
SMIM22	440335	XP_002344989	7080	hypothetical protein LOC440335
SMIM22	440335	XP_002347626	7084	hypothetical protein LOC440335
SMIM22	440335	XP_002347627	7085	hypothetical protein LOC440335

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
SMIM22	440335	XP_940484	7092	hypothetical protein LOC440335
SMIM22	440335	XP_950835	7096	hypothetical protein LOC440335
SMIM24	284422	NP_001129975	1607	similar toHSPC323
SMIM3	85027	NP_116565	5426	MSTP150
SMIM4	440957	NP_001118239	1299	similar to CG32736-PA
SMIM5	643008	NP_001156467	2039	PP12104
SMIM6	100130933	NP_001156469	2040	hypothetical LOC100130933
SMIM7	79086	NP_077009	5011	chromosome 19 open reading frame 42
SMIM8	57150	NP_001035958	895	chromosome 6 open reading frame 162
SMIM8	57150	NP_065158	4654	chromosome 6 open reading frame 162
SMLR1	100507203	NP_001182526.1	7107	small leucine-rich protein 1
SMO	6608	NP_005622	3332	smoothenedHomolog (Drosophila)
SMOC2	64094	NP_001159884	2167	SPARC related modular calcium binding 2
SMOC2	64094	NP_071421	4882	SPARC related modular calcium binding 2
SMPD1	6609	NP_000534	158	sphingomyelin phosphodiesterase 1, acid lysosomal
SMPD1	6609	NP_001007594	524	sphingomyelin phosphodiesterase 1, acid lysosomal
SMPD2	6610	NP_003071	2781	sphingomyelin phosphodiesterase 2, neutral membrane (neutral sphingomyelinase)
SMPD3	55512	NP_061137	4482	sphingomyelin phosphodiesterase 3, neutral membrane (neutral sphingomyelinase II)
SNAP23	8773	NP_003816.2	7308	synaptosome associated protein 23
SNN	8303	NP_003489	2843	stannin
SNRNP40	9410	NP_004805	3135	small nuclear ribonucleoprotein 40kDa (U5)
SNX13	23161	NP_055947	4038	sorting nexin 13
SNX14	57231	NP_065201	4665	sorting nexin 14
SNX14	57231	NP_722523	6231	sorting nexin 14
SNX19	399979	NP_055573	3984	sorting nexin 19
SOAT1	6646	NP_003092	2782	sterol O-acyltransferase 1
SOAT2	8435	NP_003569	2851	sterol O-acyltransferase 2
SORCS1	114815	NP_001013049	609	sortilin-related VPS10 domain containing receptor 1
SORCS1	114815	NP_443150	5540	sortilin-related VPS10 domain containing receptor 1
SORCS2	57537	NP_065828	4720	sortilin-related VPS10 domain containing receptor 2
SORCS3	22986	NP_055793	4021	sortilin-related VPS10 domain containing receptor 3
SORL1	6653	NP_003096	2783	sortilin-related receptor, L(DLR class) A repeats-containing
SORT1	6272	NP_002950	2738	sortilin 1
SOX1	6656	NP_005977	3411	SRY (sex determining region Y)-box 1
SP4	6671	NP_003103	2784	Sp4 transcription factor
SPACA1	81833	NP_112222	5206	sperm acrosome associated 1
SPAG4	6676	NP_003107	2785	sperm associated antigen 4
SPAG9	9043	NP_001124000	1432	sperm associated antigen 9
SPAG9	9043	NP_003962	2936	sperm associated antigen 9
SPAM1	6677	NP_003108	2786	sperm adhesion molecule 1 (PH-20Hyaluronidase, zona pellucida binding)
SPAM1	6677	NP_694859	6156	sperm adhesion molecule 1 (PH-20Hyaluronidase, zona pellucida binding)
SPAST	6683	NP_055761	4019	spastin
SPAST	6683	NP_955468	6884	spastin
SPATA3	130560	NP_620712	5834	spermatogenesis associated 3
SPATA31D1	389763	NP_001001670	373	FAM75-like protein FLJ46321
SPATA9	83890	NP_114158	5275	spermatogenesis associated 9
SPCS1	28972	NP_054760	3860	signal peptidase complex subunit 1Homolog (S. cerevisiae)
SPCS2	9789	NP_055567	3982	signal peptidase complex subunit 2Homolog (S. cerevisiae) signal peptidase complex subunit 2Homolog pseudogene
SPEM1	374768	NP_955371	6877	spermatid maturation 1
SPG7	6687	NP_003110	2787	spastic paraplegia 7 (pure and complicated autosomal recessive)
SPG7	6687	NP_955399	6882	spastic paraplegia 7 (pure and complicated autosomal recessive)
SPI1	6688	NP_001074016	1037	spleen focus forming virus (SFFV) proviral integration oncogene spi1
SPI1	6688	NP_003111	2788	spleen focus forming virus (SFFV) proviral integration oncogene spi1
SPINK13	153218	NP_001035218	840	serine PI Kazal type 5-like 3
SPINK2	6691	NP_066937	4775	serine peptidase inhibitor, Kazal type 2 (acrosin-trypsin inhibitor)
SPINK6	404203	NP_995313	6956	serine peptidase inhibitor, Kazal type 6
SPINT1	6692	NP_001027539	744	serine peptidase inhibitor, Kunitz type 1
SPINT1	6692	NP_003701	2872	serine peptidase inhibitor, Kunitz type 1
SPINT1	6692	NP_857593	6616	serine peptidase inhibitor, Kunitz type 1
SPINT2	10653	NP_001159575	2145	serine peptidase inhibitor, Kunitz type, 2
SPINT2	10653	NP_066925	4772	serine peptidase inhibitor, Kunitz type, 2
SPN	6693	NP_001025459	717	sialophorin
SPN	6693	NP_003114	2789	sialophorin
SPNS1	83985	NP_001135920	1662	spinsterHomolog 1 (Drosophila)
SPNS1	83985	NP_001135921	1663	spinsterHomolog 1 (Drosophila)
SPNS1	83985	NP_001135922	1664	spinsterHomolog 1 (Drosophila)
SPNS1	83985	NP_001135923	1665	spinsterHomolog 1 (Drosophila)
SPNS1	83985	NP_114427	5283	spinsterHomolog 1 (Drosophila)
SPNS2	124976	NP_001118230	1298	spinsterHomolog 2 (Drosophila)
SPNS3	201305	NP_872344	6653	spinsterHomolog 3 (Drosophila)
SPOCK3	50859	NP_001035249	845	sparc/osteonectin, cwcv and kazal-like domains proteoglycan (testican) 3
SPOCK3	50859	NP_058646	4279	sparc/osteonectin, cwcv and kazal-like domains proteoglycan (testican) 3
SPON2	10417	NP_001121797.1	7392	spondin 2
SPPL2A	84888	NP_116191	5405	signal peptide peptidase-like 2A
SPPL2B	56928	NP_001070706	956	signal peptide peptidase-like 2B
SPPL2B	56928	NP_694533	6141	signal peptide peptidase-like 2B

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
SPPL2C	162540	NP_787078	6464	intramembrane protease 5
SPPL3	121665	NP_620584	5825	signal peptide peptidase 3
SPTLC1	10558	NP_006406	3487	serine palmitoyltransferase, long chain base subunit 1
SPTLC1	10558	NP_847894	6536	serine palmitoyltransferase, long chain base subunit 1
SPTLC2	9517	NP_004854	3151	serine palmitoyltransferase, long chain base subunit 2
SPTLC3	55304	NP_060797	4422	serine palmitoyltransferase, long chain base subunit 3
SPTSSA	171546	NP_612145	5719	chromosome 14 open reading frame 147
SPTSSB	165679	NP_001035189	831	chromosome 3 open reading frame 57
SPX	80763	NP_085049	5149	chromosome 12 open reading frame 39
SQLE	6713	NP_003120	2790	squalene epoxidase
SRD5A1	6715	NP_001038	926	steroid-5-alpha-reductase, alpha polypeptide 1 (3-oxo-5 alpha-steroid delta 4-dehydrogenase alpha 1)
SRD5A2	6716	NP_000339	103	steroid-5-alpha-reductase, alpha polypeptide 2 (3-oxo-5 alpha-steroid delta 4-dehydrogenase alpha 2)
SRD5A3	79644	NP_078868	5053	steroid 5 alpha-reductase 3
SRPRB	58477	NP_067026	4795	signal recognition particle receptor, B subunit
SRR	63826	NP_068766	4851	serine racemase
SSBP4	170463	NP_001009998	568	single stranded DNA binding protein 4
SSBP4	170463	NP_116016	5381	single stranded DNA binding protein 4
SSMEM1	136263	NP_660311	5947	chromosome 7 open reading frame 45
SSPN	8082	NP_001129295	1572	sarcospan (Kras oncogene-associated gene)
SSPN	8082	NP_005077	3209	sarcospan (Kras oncogene-associated gene)
SSR1	6745	NP_003135	2791	signal sequence receptor, alpha
SSR2	6746	NP_003136	2792	signal sequence receptor, beta (translocon-associated protein beta)
SSR3	6747	NP_009038	3631	signal sequence receptor, gamma (translocon-associated protein gamma)
SST	6750	NP_001039.1	7266	somatostatin
SSTR1	6751	NP_001040	927	somatostatin receptor 1
SSTR2	6752	NP_001041	928	somatostatin receptor 2
SSTR3	6753	NP_001042	935	somatostatin receptor 3
SSTR4	6754	NP_001043	936	somatostatin receptor 4
SSTR5	6755	NP_001044	937	somatostatin receptor 5
SSX1	6756	NP_001265620.1	7438	SSX family member 1
ST14	6768	NP_068813	4858	suppression of tumorigenicity 14 (colon carcinoma)
ST3GAL1	6482	NP_003024	2757	ST3 beta-galactoside alpha-2,3-sialyltransferase 1
ST3GAL1	6482	NP_775479	6341	ST3 beta-galactoside alpha-2,3-sialyltransferase 1
ST3GAL2	6483	NP_008858	3596	ST3 beta-galactoside alpha-2,3-sialyltransferase 2
ST3GAL3	6487	NP_006270	3460	ST3 beta-galactoside alpha-2,3-sialyltransferase 3
ST3GAL3	6487	NP_777623	6424	ST3 beta-galactoside alpha-2,3-sialyltransferase 3
ST3GAL3	6487	NP_777624	6425	ST3 beta-galactoside alpha-2,3-sialyltransferase 3
ST3GAL3	6487	NP_777625	6426	ST3 beta-galactoside alpha-2,3-sialyltransferase 3
ST3GAL3	6487	NP_777626	6427	ST3 beta-galactoside alpha-2,3-sialyltransferase 3
ST3GAL3	6487	NP_777627	6428	ST3 beta-galactoside alpha-2,3-sialyltransferase 3
ST3GAL3	6487	NP_777628	6429	ST3 beta-galactoside alpha-2,3-sialyltransferase 3
ST3GAL3	6487	NP_777629	6430	ST3 beta-galactoside alpha-2,3-sialyltransferase 3
ST3GAL3	6487	NP_777630	6431	ST3 beta-galactoside alpha-2,3-sialyltransferase 3
ST3GAL3	6487	NP_777631	6432	ST3 beta-galactoside alpha-2,3-sialyltransferase 3
ST3GAL4	6484	NP_006269	3459	ST3 beta-galactoside alpha-2,3-sialyltransferase 4
ST3GAL5	8869	NP_001035902	887	ST3 beta-galactoside alpha-2,3-sialyltransferase 5
ST3GAL5	8869	NP_003887	2925	ST3 beta-galactoside alpha-2,3-sialyltransferase 5
ST3GAL6	10402	NP_006091	3435	ST3 beta-galactoside alpha-2,3-sialyltransferase 6
ST6GAL2	84620	NP_001135823	1648	ST6 beta-galactosamide alpha-2,6-sialyltranferase 2
ST6GAL2	84620	NP_001135824	1649	ST6 beta-galactosamide alpha-2,6-sialyltranferase 2
ST6GAL2	84620	NP_115917	5366	ST6 beta-galactosamide alpha-2,6-sialyltranferase 2
ST6GALNAC1	55808	NP_060884	4444	ST6 (alpha-N-acetyl-neuraminyl-2,3-beta-galactosyl-1,3)-N-acetylgalactosaminide alpha-2,6-sialyltransferase 1
ST6GALNAC2	10610	NP_006447	3493	ST6 (alpha-N-acetyl-neuraminyl-2,3-beta-galactosyl-1,3)-N-acetylgalactosaminide alpha-2,6-sialyltransferase 2
ST6GALNAC3	256435	NP_001153483	1973	ST6 (alpha-N-acetyl-neuraminyl-2,3-beta-galactosyl-1,3)-N-acetylgalactosaminide alpha-2,6-sialyltransferase 3
ST6GALNAC3	256435	NP_694541	6143	ST6 (alpha-N-acetyl-neuraminyl-2,3-beta-galactosyl-1,3)-N-acetylgalactosaminide alpha-2,6-sialyltransferase 3
ST6GALNAC4	27090	NP_778204	6436	ST6 (alpha-N-acetyl-neuraminyl-2,3-beta-galactosyl-1,3)-N-acetylgalactosaminide alpha-2,6-sialyltransferase 4
ST6GALNAC4	27090	NP_778205	6437	ST6 (alpha-N-acetyl-neuraminyl-2,3-beta-galactosyl-1,3)-N-acetylgalactosaminide alpha-2,6-sialyltransferase 4
ST6GALNAC5	81849	NP_112227	5207	ST6 (alpha-N-acetyl-neuraminyl-2,3-beta-galactosyl-1,3)-N-acetylgalactosaminide alpha-2,6-sialyltransferase 5
ST6GALNAC6	30815	NP_038471	3824	ST6 (alpha-N-acetyl-neuraminyl-2,3-beta-galactosyl-1,3)-N-acetylgalactosaminide alpha-2,6-sialyltransferase 6
ST7	7982	NP_060882	4442	suppression of tumorigenicity 7
ST7	7982	NP_068708	4843	suppression of tumorigenicity 7
ST7L	54879	NP_060214	4334	suppression of tumorigenicity 7 like
ST7L	54879	NP_620055	5778	suppression of tumorigenicity 7 like
ST7L	54879	NP_620056	5779	suppression of tumorigenicity 7 like
ST7L	54879	NP_620057	5780	suppression of tumorigenicity 7 like
ST8SIA1	6489	NP_003025	2758	ST8 alpha-N-acetyl-neuraminide alpha-2,8-sialyltransferase 1
ST8SIA2	8128	NP_006002	3413	ST8 alpha-N-acetyl-neuraminide alpha-2,8-sialyltransferase 2
ST8SIA3	51046	NP_056963	4127	ST8 alpha-N-acetyl-neuraminide alpha-2,8-sialyltransferase 3
ST8SIA5	29906	NP_037437	3799	ST8 alpha-N-acetyl-neuraminide alpha-2,8-sialyltransferase 5
STAB1	23166	NP_055951	4039	stabilin 1
STAB2	55576	NP_060034.9	7345	stabilin 2
STARD3	10948	NP_001159409	2129	StAR-related lipid transfer (START) domain containing 3
STARD3	10948	NP_001159410	2130	StAR-related lipid transfer (START) domain containing 3
STARD3	10948	NP_006795	3567	StAR-related lipid transfer (START) domain containing 3
STARD3NL	83930	NP_114405	5279	STARD3 N-terminal like
STEAP1	26872	NP_036581	3777	six transmembrane epithelial antigen of the prostate 1
STEAP1B	256227	NP_001157932	2079	similar to Six transmembrane epithelial antigen of prostate

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
STEAP1B	256227	NP_997225	7012	similar to Six transmembrane epithelial antigen of prostate
STEAP2	261729	NP_001035755	879	six transmembrane epithelial antigen of the prostate 2
STEAP2	261729	NP_001035756	880	six transmembrane epithelial antigen of the prostate 2
STEAP2	261729	NP_694544	6145	six transmembrane epithelial antigen of the prostate 2
STEAP3	55240	NP_001008410.1	7340	STEAP3 metalloreductase
STEAP4	79689	NP_078912	5060	STEAP family member 4
STIM1	6786	NP_003147	2793	stromal interaction molecule 1
STIM2	57620	NP_001162588	2236	stromal interaction molecule 2
STIM2	57620	NP_001162589	2237	stromal interaction molecule 2
STIM2	57620	NP_065911	4734	stromal interaction molecule 2
STOM	2040	NP_004090	2962	stomatin
STOM	2040	NP_937837	6756	stomatin
STOML1	9399	NP_004800	3134	stomatin (EPB72)-like 1
STOML3	161003	NP_001137505	1760	stomatin (EPB72)-like 3
STOML3	161003	NP_660329	5953	stomatin (EPB72)-like 3
STRA6	64220	NP_001136089.1	7401	stimulated by retinoic acid 6
STS	412	NP_000342	105	steroid sulfatase (microsomal), isozyme S
STT3A	3703	NP_689926	6103	STT3, subunit of the oligosaccharyltransferase complex,Homolog A (S. cerevisiae)
STX10	8677	NP_003756	2888	syntaxin 10
STX12	23673	NP_803173	6492	syntaxin 12
STX17	55014	NP_060389	4371	syntaxin 17
STX18	53407	NP_058626	4274	syntaxin 18
STX1B	112755	NP_443106	5523	syntaxin 1B
STX2	2054	NP_001971	2516	syntaxin 2
STX2	2054	NP_919337	6721	syntaxin 2
STX3	6809	NP_004168	2981	syntaxin 3
STX4	6810	NP_004595	3082	syntaxin 4
STX6	10228	NP_005810	3376	syntaxin 6
STX7	8417	NP_003560	2849	syntaxin 7
STX8	9482	NP_004844	3147	syntaxin 8
STXBP2	6813	NP_001120868	1326	syntaxin binding protein 2
STXBP2	6813	NP_008880	3602	syntaxin binding protein 2
STYK1	55359	NP_060893	4447	serine/threonine/tyrosine kinase 1
SUCNR1	56670	NP_149039	5443	succinate receptor 1
SULF2	55959	NP_001155313.1	7411	sulfatase 2
SUMF2	25870	NP_001035934	890	sulfatase modifying factor 2
SUMF2	25870	NP_001035935	891	sulfatase modifying factor 2
SUMF2	25870	NP_001123541	1416	sulfatase modifying factor 2
SUMF2	25870	NP_001139805	1931	sulfatase modifying factor 2
SUMF2	25870	NP_056226	4078	sulfatase modifying factor 2
SUN1	23353	NP_001124437	1452	unc-84Homolog A (C. elegans)
SUN1	23353	NP_001165415	2301	unc-84Homolog A (C. elegans)
SUN1	23353	NP_001165416	2302	unc-84Homolog A (C. elegans)
SUN1	23353	NP_001165417	2303	unc-84Homolog A (C. elegans)
SUN1	23353	NP_079430	5127	unc-84Homolog A (C. elegans)
SUN2	25777	NP_056189	4071	unc-84Homolog B (C. elegans)
SURF4	6836	NP_149351	5454	surfeit 4
SUSD1	64420	NP_071931	4926	sushi domain containing 1
SUSD2	56241	NP_062547	4574	sushi domain containing 2
SUSD3	203328	NP_659443	5916	sushi domain containing 3
SUSD4	55061	NP_001032252	771	sushi domain containing 4
SUSD4	55061	NP_060452	4377	sushi domain containing 4
SUSD5	26032	NP_056366	4100	sushi domain containing 5
SUSD6	9766	NP_055549	3979	KIAA0247
SV2A	9900	NP_055664	3999	synaptic vesicle glycoprotein 2A
SV2B	9899	NP_001161052	2185	synaptic vesicle glycoprotein 2BHypothetical protein LOC100128403
SV2B	9899	NP_055663	3998	synaptic vesicle glycoprotein 2BHypothetical protein LOC100128403
SV2C	22987	NP_055794	4022	synaptic vesicle glycoprotein 2C
SVOP	55530	NP_061181	4486	SV2 related proteinHomolog (rat)
SVOPL	136306	NP_001132928	1618	SVOP-like
SVOPL	136306	NP_777619	6423	SVOP-like
SYNDIG1	79953	NP_079169	5097	chromosome 20 open reading frame 39
SYNDIG1L	646658	NP_001099049	1230	transmembrane protein 90A
SYNE3	161176	NP_689805	6086	chromosome 14 open reading frame 49
SYNE4	163183	NP_001034965	814	chromosome 19 open reading frame 46
SYNGR1	9145	NP_004702	3107	synaptogyrin 1
SYNGR1	9145	NP_663783	5972	synaptogyrin 1
SYNGR1	9145	NP_663791	5973	synaptogyrin 1
SYNGR2	9144	NP_004701	3106	synaptogyrin 2
SYNGR3	9143	NP_004200	2989	synaptogyrin 3
SYNGR4	23546	NP_036583	3779	synaptogyrin 4
SYNJ2BP	55333	NP_060843	4433	synaptojanin 2 binding protein
SYNPR	132204	NP_001123475	1408	synaptoporin
SYNPR	132204	NP_653243	5882	synaptoporin
SYP	6855	NP_003170	2794	synaptophysin
SYPL1	6856	NP_006745	3557	synaptophysin-like 1
SYPL1	6856	NP_874384	6677	synaptophysin-like 1

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
SYPL2	284612	NP_001035799	882	synaptophysin-like 2
SYS1	90196	NP_291020	5505	SYS1 Golgi-localized integral membrane proteinHomolog (S. cerevisiae)
SYT1	6857	NP_001129277	1567	synaptotagmin I
SYT1	6857	NP_001129278	1568	synaptotagmin I
SYT1	6857	NP_005630	3334	synaptotagmin I
SYT11	23208	NP_689493	6023	synaptotagmin XI
SYT12	91683	NP_808878	6508	synaptotagmin XII
SYT13	57586	NP_065877	4729	synaptotagmin XIII
SYT14	255928	NP_001139733	1917	synaptotagmin XIV
SYT14	255928	NP_001139734	1918	synaptotagmin XIV
SYT14	255928	NP_001139736	1919	synaptotagmin XIV
SYT14	255928	NP_694994	6169	synaptotagmin XIV
SYT15	83849	NP_114118	5268	synaptotagmin XV
SYT15	83849	NP_852660	6605	synaptotagmin XV
SYT2	127833	NP_001129976	1608	synaptotagmin II
SYT2	127833	NP_796376	6489	synaptotagmin II
SYT3	84258	NP_001153800.1	7410	synaptotagmin 3
SYT4	6860	NP_065834	4722	synaptotagmin IV
SYT5	6861	NP_003171	2795	synaptotagmin V
SYT6	148281	NP_995320	6957	synaptotagmin VI
SYT7	9066	NP_001238994.1	7431	synaptotagmin 7
SYT8	90019	NP_612634.3	7358	synaptotagmin 8
SYT9	143425	NP_783860	6455	synaptotagmin IX
TAAR1	134864	NP_612200	5726	trace amine associated receptor 1
TAAR2	9287	NP_001028252	753	trace amine associated receptor 2
TAAR2	9287	NP_055441	3967	trace amine associated receptor 2
TAAR5	9038	NP_003958	2935	trace amine associated receptor 5
TAAR8	83551	NP_444508	5561	trace amine associated receptor 8
TAAR9	134860	NP_778227	6439	trace amine associated receptor 9
TACR1	6869	NP_001049	939	tachykinin receptor 1
TACR1	6869	NP_056542	4120	tachykinin receptor 1
TACR2	6865	NP_001048	938	tachykinin receptor 2
TACR3	6870	NP_001050	940	tachykinin receptor 3
TACSTD2	4070	NP_002344	2631	tumor-associated calcium signal transducer 2
TAOK2	9344	NP_004774	3125	TAO kinase 2
TAOK2	9344	NP_057235	4178	TAO kinase 2
TAPBP	6892	NP_003181	2797	TAP binding protein (tapasin)
TAPBP	6892	NP_757345	6290	TAP binding protein (tapasin)
TAPBP	6892	NP_757346	6291	TAP binding protein (tapasin)
TAPBPL	55080	NP_060479	4381	TAP binding protein-like
TAPT1	202018	NP_699196	6184	transmembrane anterior posterior transformation 1
TAS1R1	80835	NP_619642	5772	taste receptor, type 1, member 1
TAS1R1	80835	NP_803883	6497	taste receptor, type 1, member 1
TAS1R1	80835	NP_803884	6498	taste receptor, type 1, member 1
TAS1R1	80835	NP_803885	6499	taste receptor, type 1, member 1
TAS1R2	80834	NP_689418	6020	taste receptor, type 1, member 2
TAS2R1	50834	NP_062545	4573	taste receptor, type 2, member 1
TAS2R10	50839	NP_076410.1	7297	taste 2 receptor member 10
TAS2R13	50838	NP_076409	4983	taste receptor, type 2, member 13
TAS2R14	50840	NP_076411	4984	taste receptor, type 2, member 14
TAS2R16	50833	NP_058641	4277	taste receptor, type 2, member 16
TAS2R3	50831	NP_058639	4276	taste receptor, type 2, member 3
TAS2R38	5726	NP_789787	6480	taste receptor, type 2, member 38
TAS2R39	259285	NP_795362	6483	taste receptor, type 2, member 39
TAS2R4	50832	NP_058640.1	7285	taste 2 receptor member 4
TAS2R40	259286	NP_795363	6484	taste receptor, type 2, member 40
TAS2R41	259287	NP_795364	6485	taste receptor, type 2, member 41
TAS2R5	54429	NP_061853	4544	taste receptor, type 2, member 5
TAS2R50	259296	NP_795371	6486	taste receptor, type 2, member 50
TAS2R7	50837	NP_076408	4982	taste receptor, type 2, member 7
TAS2R8	50836	NP_076407	4981	taste receptor, type 2, member 8
TAS2R9	50835	NP_076406	4980	taste receptor, type 2, member 9
TAZ	6901	NP_000107	34	tafazzin
TAZ	6901	NP_851828	6590	tafazzin
TAZ	6901	NP_851829	6591	tafazzin
TAZ	6901	NP_851830	6592	tafazzin
TBC1D20	128637	NP_653229	5878	TBC1 domain family, member 20
TBC1D2B	23102	NP_055894	4029	TBC1 domain family, member 2B
TBC1D2B	23102	NP_653173	5864	TBC1 domain family, member 2B
TBC1D7	51256	NP_001137436	1744	TBC1 domain family, member 7
TBC1D7	51256	NP_001137437	1745	TBC1 domain family, member 7
TBC1D7	51256	NP_001137438	1746	TBC1 domain family, member 7
TBC1D7	51256	NP_057579	4228	TBC1 domain family, member 7
TBL2	26608	NP_036585	3781	transducin (beta)-like 2
TCIRG1	10312	NP_006010	3416	T-cell, immune regulator 1, ATPase,H+ transporting, lysosomal V0 subunit A3
TCIRG1	10312	NP_006044	3421	T-cell, immune regulator 1, ATPase,H+ transporting, lysosomal V0 subunit A3
TCTA	6988	NP_071503	4894	T-cell leukemia translocation altered gene

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
TCTN3	26123	NP_001137445	1749	tectonic family member 3
TCTN3	26123	NP_056446	4108	tectonic family member 3
TDGF1	6997	NP_001167607.1	7419	teratocarcinoma-derived growth factor 1
TDGF1P3	6998	AAA61135.1	7466	teratocarcinoma-derived growth factor 3
TDRKH	11022	NP_001077432	1086	tudor and KH domain containing
TDRKH	11022	NP_001077433	1087	tudor and KH domain containing
TDRKH	11022	NP_001077434	1088	tudor and KH domain containing
TDRKH	11022	NP_006853	3585	tudor and KH domain containing
TECRL	253017	NP_001010874	573	steroid 5 alpha-reductase 2-like 2
TEDDM1	127670	NP_741997	6266	transmembrane epididymal protein 1
TEK	7010	NP_000450	135	TEK tyrosine kinase, endothelial
TENM1	10178	NP_001156750	2049	odz, odd Oz/ten-mHomolog 1(Drosophila)
TENM1	10178	NP_001156751	2050	odz, odd Oz/ten-mHomolog 1(Drosophila)
TENM1	10178	NP_055068	3896	odz, odd Oz/ten-mHomolog 1(Drosophila)
TENM2	57451	NP_001073897.2	7373	teneurin transmembrane protein 2
TENM3	55714	NP_001073946	1029	odz, odd Oz/ten-mHomolog 3 (Drosophila)
TENM4	26011	NP_001092286	1147	odz, odd Oz/ten-mHomolog 4 (Drosophila)
TEX2	55852	NP_060939	4456	testis expressed 2
TEX261	113419	NP_653183	5867	testis expressed 261
TEX264	51368	NP_001123356	1402	testis expressed 264
TEX264	51368	NP_057010	4138	testis expressed 264
TEX29	121793	NP_689537	6033	chromosome 13 open reading frame 16
TEX38	374973	NP_001138946	1868	chromosome 1 open reading frame 223
TFR2	7036	NP_003218	2799	transferrin receptor 2
TGFA	7039	NP_001093161	1170	transforming growth factor, alpha
TGFA	7039	NP_003227	2800	transforming growth factor, alpha
TGFB3	7043	NP_003230	2801	transforming growth factor, beta 3
TGFBI	7045	NP_000349	107	transforming growth factor, beta-induced, 68kDa
TGFBR1	7046	NP_001124388	1447	transforming growth factor, beta receptor 1
TGFBR1	7046	NP_004603	3083	transforming growth factor, beta receptor 1
TGFBR2	7048	NP_001020018	689	transforming growth factor, beta receptor II (70/80kDa)
TGFBR2	7048	NP_003233	2802	transforming growth factor, beta receptor II (70/80kDa)
TGFBR3	7049	NP_003234	2803	transforming growth factor, beta receptor III
TGOLN2	10618	NP_006455	3495	trans-golgi network protein 2
THADA	63892	NP_001077422	1085	thyroid adenoma associated
THADA	63892	NP_071348	4871	thyroid adenoma associated
THAP6	152815	NP_653322	5899	THAP domain containing 6
THBD	7056	NP_000352	108	thrombomodulin
THSD4	79875	NP_079093	5086	thrombospondin, type I, domain containing 4
THSD7A	221981	NP_056019	4044	thrombospondin, type I, domain containing 7A
THSD7B	80731	NP_001303278.1	1020	thrombospondin, type I, domain containing 7B
THY1	7070	NP_001298089.1	7454	Thy-1 cell surface antigen
TIE1	7075	NP_005415	3283	tyrosine kinase with immunoglobulin-like and EGF-like domains 1
TIGIT	201633	NP_776160	6385	T cell immunoreceptor with Ig and ITIM domains
TIMD4	91937	NP_001140198	1935	T-cell immunoglobulin and mucin domain containing 4
TIMD4	91937	NP_612388	5737	T-cell immunoglobulin and mucin domain containing 4
TIMM17A	10440	NP_006326	3469	translocase of inner mitochondrial membrane 17Homolog A (yeast)
TIMM17B	10245	NP_001161419	2208	translocase of inner mitochondrial membrane 17Homolog B (yeast)
TIMM17B	10245	NP_005825	3380	translocase of inner mitochondrial membrane 17Homolog B (yeast)
TIMM21	29090	NP_054896	3875	chromosome 18 open reading frame 55
TIMM22	29928	NP_037469	3805	translocase of inner mitochondrial membrane 22Homolog (yeast)
TIMM23	100287932	NP_006318	3467	similar to translocase of inner mitochondrial membrane 23 (yeast)Homolog
TIMM50	92609	NP_001001563	370	translocase of inner mitochondrial membrane 50Homolog (S. cerevisiae)
TIMMDC1	51300	NP_057673	4252	chromosome 3 open reading frame 1
TKFC	26007	NP_056348	4096	dihydroxyacetone kinase 2Homolog (S. cerevisiae)
TLCD1	116238	NP_001153879	1995	TLC domain containing 1
TLCD1	116238	NP_612472	5756	TLC domain containing 1
TLCD2	727910	NP_001157879	2077	TLC domain containing 2
TLR1	7096	NP_003254	2806	toll-like receptor 1
TLR10	81793	NP_001017388	639	toll-like receptor 10
TLR10	81793	NP_112218	5202	toll-like receptor 10
TLR2	7097	NP_003255	2807	toll-like receptor 2
TLR3	7098	NP_003256	2808	toll-like receptor 3
TLR4	7099	NP_612564	5759	toll-like receptor 4
TLR5	7100	NP_003259	2809	toll-like receptor 5
TLR6	10333	NP_006059	3427	toll-like receptor 6
TLR7	51284	NP_057646	4244	toll-like receptor 7
TLR9	54106	NP_059138	4293	toll-like receptor 9
TM2D1	83941	NP_114416	5281	TM2 domain containing 1
TM2D2	83877	NP_001019551	676	TM2 domain containing 2
TM2D2	83877	NP_001019552	677	TM2 domain containing 2
TM2D2	83877	NP_114146	5272	TM2 domain containing 2
TM2D2	83877	NP_510882	5590	TM2 domain containing 2
TM4SF1	4071	NP_055035	3888	transmembrane 4 L six family member 1
TM4SF18	116441	NP_620141	5794	transmembrane 4 L six family member 18
TM4SF20	79853	NP_079071	5085	transmembrane 4 L six family member 20
TM4SF4	7104	NP_004608	3086	transmembrane 4 L six family member 4

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
TM4SF5	9032	NP_003954	2932	transmembrane 4 L six family member 5
TM6SF1	53346	NP_001138375	1783	transmembrane 6 superfamily member 1
TM6SF1	53346	NP_075379	4969	transmembrane 6 superfamily member 1
TM6SF2	53345	NP_001001524	366	transmembrane 6 superfamily member 2
TM7SF2	7108	NP_003264	2813	transmembrane 7 superfamily member 2
TM7SF3	51768	NP_057635	4242	transmembrane 7 superfamily member 3
TM9SF1	10548	NP_001014842	628	transmembrane 9 superfamily member 1
TM9SF1	10548	NP_006396	3481	transmembrane 9 superfamily member 1
TM9SF2	9375	NP_004791	3130	transmembrane 9 superfamily member 2
TM9SF3	56889	NP_064508	4598	transmembrane 9 superfamily member 3
TM9SF4	9777	NP_055557	3980	transmembrane 9 superfamily protein member 4
TMBIM1	64114	NP_071435	4887	transmembrane BAX inhibitor motif containing 1
TMBIM4	51643	NP_057140	4159	transmembrane BAX inhibitor motif containing 4
TMBIM6	7009	NP_001092046	1138	transmembrane BAX inhibitor motif containing 6
TMBIM6	7009	NP_003208	2798	transmembrane BAX inhibitor motif containing 6
TMC1	117531	NP_619636	5770	transmembrane channel-like 1
TMC2	117532	NP_542789	5627	transmembrane channel-like 2
TMC4	147798	NP_001138775	1848	transmembrane channel-like 4
TMC4	147798	NP_653287	5892	transmembrane channel-like 4
TMC5	79838	NP_001098718	1220	transmembrane channel-like 5
TMC5	79838	NP_001098719	1221	transmembrane channel-like 5
TMC5	79838	NP_079056	5080	transmembrane channel-like 5
TMC6	11322	NP_001120670	1309	transmembrane channel-like 6
TMC6	11322	NP_009198	3665	transmembrane channel-like 6
TMC7	79905	NP_001153836	1994	transmembrane channel-like 7
TMC7	79905	NP_079123	5092	transmembrane channel-like 7
TMC8	147138	NP_689681	6065	transmembrane channel-like 8
TMCC1	23023	NP_001017395	640	transmembrane and coiled-coil domain family 1
TMCC1	23023	NP_001121696	1369	transmembrane and coiled-coil domain family 1
TMCC1	23023	NP_001017395.2	4025	transmembrane and coiled-coil domain family 1
TMCC2	9911	NP_055673	4000	transmembrane and coiled-coil domain family 2
TMCC3	57458	NP_065749	4703	transmembrane and coiled-coil domain family 3
TMCO1	54499	NP_061899	4550	transmembrane and coiled-coil domains 1
TMCO2	127391	NP_001008740	552	transmembrane and coiled-coil domains 2
TMCO3	55002	NP_060375	4368	transmembrane and coiled-coil domains 3
TMCO4	255104	NP_859070	6624	transmembrane and coiled-coil domains 4
TMCO5A	145942	NP_689666	6062	transmembrane and coiled-coil domains 5A
TMED1	11018	NP_006849	3584	transmembrane emp24 protein transport domain containing 1
TMED10	10972	NP_006818	3574	transmembrane emp24-like trafficking protein 10 (yeast)
TMED2	10959	NP_006806	3570	transmembrane emp24 domain trafficking protein 2
TMED3	23423	NP_031390	3691	transmembrane emp24 protein transport domain containing 3
TMED4	222068	NP_872353	6655	transmembrane emp24 protein transport domain containing 4
TMED5	50999	NP_001161302	2202	transmembrane emp24 protein transport domain containing 5
TMED5	50999	NP_057124	4157	transmembrane emp24 protein transport domain containing 5
TMED6	146456	NP_653277	5889	transmembrane emp24 protein transport domain containing 6
TMED7	51014	NP_001157941	2081	transmembrane emp24 protein transport domain containing 7 toll-like receptor adaptor molecule 2
TMED7	51014	NP_067681	4817	transmembrane emp24 protein transport domain containing 7 toll-like receptor adaptor molecule 2
TMED7	51014	NP_861974	6639	transmembrane emp24 protein transport domain containing 7 toll-like receptor adaptor molecule 2
TMEFF2	23671	NP_057276	4183	transmembrane protein with EGF-like and two follistatin-like domains 2
TMEM100	55273	NP_001093110	1161	transmembrane protein 100
TMEM100	55273	NP_060756	4414	transmembrane protein 100
TMEM101	84336	NP_115752	5336	transmembrane protein 101
TMEM104	54868	NP_060198	4329	transmembrane protein 104
TMEM105	284186	NP_848615	6549	transmembrane protein 105
TMEM106A	113277	NP_659478	5923	hypothetical LOC728772 transmembrane protein 106A
TMEM106B	54664	NP_001127704	1474	transmembrane protein 106B
TMEM106B	54664	NP_060844	4434	transmembrane protein 106B
TMEM106C	79022	NP_001137313	1733	transmembrane protein 106C
TMEM106C	79022	NP_001137314	1734	transmembrane protein 106C
TMEM106C	79022	NP_001137315	1735	transmembrane protein 106C
TMEM106C	79022	NP_076961	5001	transmembrane protein 106C
TMEM107	84314	NP_115730	5333	transmembrane protein 107
TMEM107	84314	NP_898888	6701	transmembrane protein 107
TMEM109	79073	NP_076997	5009	transmembrane protein 109
TMEM110	375346	NP_940965	6816	transmembrane protein 110 musculoskeletal, embryonic nuclear protein 1
TMEM110	375346	NP_995325	6960	transmembrane protein 110 musculoskeletal, embryonic nuclear protein 1
TMEM115	11070	NP_008955	3617	transmembrane protein 115
TMEM116	89894	NP_612350	5731	transmembrane protein 116
TMEM117	84216	NP_115632	5314	transmembrane protein 117
TMEM119	338773	NP_859075	6626	transmembrane protein 119
TMEM120A	83862	NP_114131	5270	transmembrane protein 120A
TMEM120B	144404	NP_001074294	1042	transmembrane protein 120B
TMEM121	80757	NP_079544	5146	transmembrane protein 121
TMEM123	114908	NP_443164	5542	transmembrane protein 123
TMEM125	128218	NP_653227	5877	transmembrane protein 125
TMEM126A	84233	NP_115649	5316	transmembrane protein 126A
TMEM126B	55863	NP_060950	4459	transmembrane protein 126B

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
TMEM127	55654	NP_060319	4356	transmembrane protein 127
TMEM128	85013	NP_116316	5423	transmembrane protein 128
TMEM129	92305	NP_001120738	1317	transmembrane protein 129
TMEM129	92305	NP_612394	5738	transmembrane protein 129
TMEM130	222865	NP_001127922	1493	transmembrane protein 130
TMEM130	222865	NP_001127923	1494	transmembrane protein 130
TMEM130	222865	NP_690877	6132	transmembrane protein 130
TMEM131	23505	NP_056163	4066	transmembrane protein 131
TMEM132B	114795	NP_443139	5535	transmembrane protein 132BHypothetical LOC121296
TMEM132C	92293	NP_001129575	1598	transmembrane protein 132C
TMEM132D	121256	NP_597705	5692	transmembrane protein 132D
TMEM132E	124842	NP_001291367.1	7006	transmembrane protein 132E
TMEM133	83935	NP_114410	5280	transmembrane protein 133
TMEM134	80194	NP_001072118	984	transmembrane protein 134
TMEM134	80194	NP_001072119	985	transmembrane protein 134
TMEM134	80194	NP_001072118.1	5121	transmembrane protein 134
TMEM135	65084	NP_001162195	2235	transmembrane protein 135
TMEM135	65084	NP_075069	4964	transmembrane protein 135
TMEM136	219902	NP_777586	6412	transmembrane protein 136
TMEM138	51524	NP_057548	4223	transmembrane protein 138
TMEM139	135932	NP_699176	6180	transmembrane protein 139
TMEM140	55281	NP_060765	4415	transmembrane protein 140
TMEM141	85014	NP_001034463	791	KIAA1984 transmembrane protein 141
TMEM141	85014	NP_116317	5424	KIAA1984 transmembrane protein 141
TMEM143	55260	NP_060743	4412	transmembrane protein 143
TMEM144	55314	NP_060812	4425	transmembrane protein 144
TMEM145	284339	NP_775904	6367	transmembrane protein 145
TMEM147	10430	NP_116024	5383	transmembrane protein 147
TMEM14A	28978	NP_054770	3863	transmembrane protein 14A
TMEM14C	51522	NP_001158730	2126	transmembrane protein 14C
TMEM14C	51522	NP_057546	4222	transmembrane protein 14C
TMEM150A	129303	NP_001026908	731	transmembrane protein 150A
TMEM150A	129303	NP_001026908.1	6179	transmembrane protein 150A
TMEM150B	284417	NP_001078957	1097	transmembrane protein 150B
TMEM150C	441027	NP_001073975	1033	hypothetical LOC441027
TMEM151B	441151	NP_001131032	1614	transmembrane protein 151B
TMEM154	201799	NP_689893	6096	transmembrane protein 154
TMEM156	80008	NP_079219	5105	transmembrane protein 156
TMEM158	25907	NP_056259	4084	transmembrane protein 158
TMEM159	57146	NP_065155	4653	transmembrane protein 159
TMEM160	54958	NP_060324	4358	transmembrane protein 160
TMEM161A	54929	NP_060284	4348	transmembrane protein 161A
TMEM161B	153396	NP_699185	6182	transmembrane protein 161B
TMEM163	81615	NP_112185	5193	transmembrane protein 163
TMEM165	55858	NP_060945	4457	transmembrane protein 165
TMEM167A	153339	NP_777569	6407	similar to transmembrane protein 167A
TMEM168	64418	NP_071929	4925	transmembrane protein 168
TMEM169	92691	NP_001135782	1638	transmembrane protein 169
TMEM169	92691	NP_001135783	1639	transmembrane protein 169
TMEM169	92691	NP_001135784	1640	transmembrane protein 169
TMEM169	92691	NP_612399	5740	transmembrane protein 169
TMEM17	200728	NP_938017	6766	transmembrane protein 17
TMEM170A	124491	NP_660297	5942	transmembrane protein 170A
TMEM170B	100113407	NP_001094299	1195	transmembrane protein 170B
TMEM171	134285	NP_001154814	2001	transmembrane protein 171
TMEM171	134285	NP_775761	6346	transmembrane protein 171
TMEM174	134288	NP_694949	6161	transmembrane protein 174
TMEM175	84286	NP_115702	5328	transmembrane protein 175
TMEM176B	28959	NP_001094781	1197	transmembrane protein 176B
TMEM176B	28959	NP_001094782	1198	transmembrane protein 176B
TMEM176B	28959	NP_001094781.1	1199	transmembrane protein 176B
TMEM176B	28959	NP_001094784	1200	transmembrane protein 176B
TMEM176B	28959	NP_054739	3856	transmembrane protein 176B
TMEM178A	130733	NP_001161431	2209	transmembrane protein 178
TMEM178A	130733	NP_689603	6049	transmembrane protein 178
TMEM178B	100507421	XP_011514007	7108	transmembrane protein 178B
TMEM179B	374395	NP_955369	6876	transmembrane protein 179B
TMEM18	129787	NP_690047	6118	transmembrane protein 18
TMEM181	57583	NP_065874	4728	transmembrane protein 181
TMEM182	130827	NP_653233	5879	transmembrane protein 182
TMEM183A	92703	NP_001073277	996	transmembrane protein 183A transmembrane protein 183B
TMEM183A	92703	NP_612400	5741	transmembrane protein 183A transmembrane protein 183B
TMEM184B	25829	NP_036396	3730	transmembrane protein 184B
TMEM185B	79134	NP_077026	7100	transmembrane protein 185B
TMEM186	25880	NP_056236	4080	transmembrane protein 186
TMEM187	8269	NP_003483	2841	transmembrane protein 187
TMEM189	387521	NP_001027459	741	ubiquitin-conjugating enzyme E2 variant 1

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
TMEM189	387521	NP_001155977	2032	ubiquitin-conjugating enzyme E2 variant 1
TMEM189	387521	NP_001155977.1	2824	ubiquitin-conjugating enzyme E2 variant 1
TMEM189	387521	NP_068823	4860	ubiquitin-conjugating enzyme E2 variant 1
TMEM189	387521	NP_071887	4917	ubiquitin-conjugating enzyme E2 variant 1
TMEM189	387521	NP_954580	6857	ubiquitin-conjugating enzyme E2 variant 1
TMEM189	387521	NP_954595	6859	ubiquitin-conjugating enzyme E2 variant 1
TMEM189	387521	NP_954673	6864	ubiquitin-conjugating enzyme E2 variant 1
TMEM19	55266	NP_060749	4413	transmembrane protein 19
TMEM190	147744	NP_631911	5842	transmembrane protein 190
TMEM192	201931	NP_001093859	1185	transmembrane protein 192
TMEM196	256130	NP_689987	6114	transmembrane protein 196
TMEM198	130612	NP_001005209	447	transmembrane protein 198
TMEM199	147007	NP_689677	6064	transmembrane protein 199
TMEM2	23670	NP_001129292	1569	transmembrane protein 2
TMEM2	23670	NP_037522	3818	transmembrane protein 2
TMEM200A	114801	NP_443145	5537	transmembrane protein 200A
TMEM200B	399474	NP_001003682	415	transmembrane protein 200B
TMEM200B	399474	NP_001165339	2289	transmembrane protein 200B
TMEM200C	645369	NP_001073678	1014	two transmembrane domain family member A
TMEM201	199953	NP_001010866	571	transmembrane protein 201
TMEM201	199953	NP_001124396	1448	transmembrane protein 201
TMEM203	94107	NP_444273	5559	transmembrane protein 203
TMEM204	79652	NP_078876	5055	transmembrane protein 204
TMEM205	374882	NP_001138888	1855	transmembrane protein 205
TMEM205	374882	NP_212133	5493	transmembrane protein 205
TMEM205	374882	NP_940938	6810	transmembrane protein 205
TMEM206	55248	NP_060722	4409	transmembrane protein 206
TMEM207	131920	NP_997199	7007	transmembrane protein 207
TMEM208	29100	NP_054906	3878	transmembrane protein 208
TMEM209	84928	NP_116231	5416	transmembrane protein 209
TMEM212	389177	NP_001157908	2078	hypothetical LOC389177
TMEM213	155006	NP_001078898	1094	transmembrane protein 213
TMEM216	51259	NP_057583	4230	transmembrane protein 216
TMEM217	221468	NP_001156372	2038	transmembrane protein 217
TMEM217	221468	NP_660359	5961	transmembrane protein 217
TMEM218	219854	NP_001074015	1036	transmembrane protein 218
TMEM219	124446	NP_001077082	1077	transmembrane protein 219
TMEM219	124446	NP_919256	6712	transmembrane protein 219
TMEM220	388335	NP_001004313	429	transmembrane protein 220
TMEM221	100130519	XP_001718804	7062	transmembrane protein 221
TMEM221	100130519	XP_001719748	7067	transmembrane protein 221
TMEM222	84065	NP_115501	5300	transmembrane protein 222
TMEM225	338661	NP_001013765	620	transmembrane protein 225
TMEM229B	161145	NP_872332	6650	chromosome 14 open reading frame 83
TMEM230	29058	NP_001009923	565	hypothetical LOC642975 chromosome 20 open reading frame 30
TMEM230	29058	NP_001009924	566	hypothetical LOC642975 chromosome 20 open reading frame 30
TMEM230	29058	NP_001009925	567	hypothetical LOC642975 chromosome 20 open reading frame 30
TMEM230	29058	NP_054864	3873	hypothetical LOC642975 chromosome 20 open reading frame 30
TMEM231	79583	NP_001070884	967	hypothetical protein FLJ22167
TMEM231	79583	NP_001070886	968	hypothetical protein FLJ22167
TMEM231	79583	NP_001070884.2	969	hypothetical protein FLJ22167
TMEM234	56063	NP_061991	4568	chromosome 1 open reading frame 91
TMEM235	283999	NP_001191139.1	1869	hypothetical protein LOC283999
TMEM237	65062	NP_001037850	921	amyotrophic lateral sclerosis 2 (juvenile) chromosome region, candidate 4
TMEM237	65062	NP_689601	6047	amyotrophic lateral sclerosis 2 (juvenile) chromosome region, candidate 4
TMEM239	100288797	NP_001161142	2193	hypothetical protein LOC100288797
TMEM242	729515	NP_001138890	1856	chromosome 6 open reading frame 35HCG1820764 tetratricopeptide repeat domain 28
TMEM242	729515	NP_060922	4454	chromosome 6 open reading frame 35HCG1820764 tetratricopeptide repeat domain 28
TMEM243	79161	NP_077291	5024	chromosome 7 open reading frame 23
TMEM245	23731	NP_114401.2	1171	chromosome 9 open reading frame 5
TMEM245	23731	NP_114401	5277	chromosome 9 open reading frame 5
TMEM246	84302	NP_115718	5332	chromosome 9 open reading frame 125
TMEM248	55069	NP_060464	4379	chromosome 7 open reading frame 42
TMEM25	84866	NP_001137506	1761	transmembrane protein 25
TMEM25	84866	NP_001137507	1762	transmembrane protein 25
TMEM25	84866	NP_001137508	1763	transmembrane protein 25
TMEM25	84866	NP_001137509	1764	transmembrane protein 25
TMEM25	84866	NP_001137510	1765	transmembrane protein 25
TMEM25	84866	NP_116169	5399	transmembrane protein 25
TMEM251	26175	NP_001092091	1141	chromosome 14 open reading frame 109
TMEM251	26175	NP_056491	4112	chromosome 14 open reading frame 109
TMEM252	169693	NP_694969	6164	chromosome 9 open reading frame 71
TMEM254	80195	NP_079401	5122	chromosome 10 open reading frame 58 chromosome 10 open reading frame 57
TMEM254	80195	NP_001257296.1	5330	chromosome 10 open reading frame 58 chromosome 10 open reading frame 57
TMEM255A	55026	NP_001098014	1213	family with sequence similarity 70, member A
TMEM255A	55026	NP_001098015	1214	family with sequence similarity 70, member A
TMEM255A	55026	NP_060408	4373	family with sequence similarity 70, member A

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
TMEM255B	348013	NP_872420	6667	family with sequence similarity 70, member B
TMEM256	254863	NP_689979	6112	chromosome 17 open reading frame 61
TMEM257	9142	NP_004700	3105	chromosome X open reading frame 1
TMEM258	746	NP_055021	3880	chromosome 11 open reading frame 10
TMEM259	91304	NP_001028198	750	chromosome 19 open reading frame 6
TMEM259	91304	NP_219488	5496	chromosome 19 open reading frame 6
TMEM26	219623	NP_848600	6546	transmembrane protein 26
TMEM260	54916	NP_060269	4345	chromosome 14 open reading frame 101
TMEM261	90871	NP_219500	5497	chromosome 9 open reading frame 123
TMEM263	90488	NP_689474	6021	chromosome 12 open reading frame 23
TMEM265	100862671	NP_001243758	7113	transmembrane protein 265
TMEM27	57393	NP_065716	4695	transmembrane protein 27
TMEM30A	55754	NP_001137430	1740	transmembrane protein 30A
TMEM30A	55754	NP_060717	4408	transmembrane protein 30A
TMEM30B	161291	NP_001017970	654	transmembrane protein 30B
TMEM31	203562	NP_872347	6654	transmembrane protein 31
TMEM33	55161	NP_060596	4396	transmembrane protein 33
TMEM35	59353	NP_067650	4814	transmembrane protein 35
TMEM37	140738	NP_899063	6705	transmembrane protein 37
TMEM38B	55151	NP_060582	4394	transmembrane protein 38B
TMEM39A	55254	NP_060736	4411	transmembrane protein 39A
TMEM40	55287	NP_060776	4418	transmembrane protein 40
TMEM41A	90407	NP_542383	5612	transmembrane protein 41A
TMEM41B	440026	NP_001158502	2121	transmembrane protein 41B
TMEM41B	440026	NP_055827	4026	transmembrane protein 41B
TMEM42	131616	NP_653239	5881	transmembrane protein 42
TMEM43	79188	NP_077310	5026	transmembrane protein 43
TMEM45A	55076	NP_060474	4380	transmembrane protein 45A
TMEM45B	120224	NP_620143	5795	transmembrane protein 45B
TMEM47	83604	NP_113630	5225	transmembrane protein 47
TMEM5	10329	NP_055069	3897	transmembrane protein 5
TMEM50A	23585	NP_055128	3912	transmembrane protein 50A
TMEM50B	757	NP_006125	3440	transmembrane protein 50B
TMEM51	55092	NP_001129688	1601	transmembrane protein 51
TMEM51	55092	NP_001129689	1602	transmembrane protein 51
TMEM51	55092	NP_001129690	1603	transmembrane protein 51
TMEM51	55092	NP_060492	4382	transmembrane protein 51
TMEM52	339456	NP_848640	6554	transmembrane protein 52
TMEM52B	120939	NP_694567	6151	chromosome 12 open reading frame 59
TMEM54	113452	NP_277039	5502	transmembrane protein 54
TMEM55B	90809	NP_001094284	1194	transmembrane protein 55B
TMEM55B	90809	NP_653169	5863	transmembrane protein 55B
TMEM56	148534	NP_689700	6068	transmembrane protein 56
TMEM57	55219	NP_060672	4402	transmembrane protein 57
TMEM59	9528	NP_004863	3156	transmembrane protein 59
TMEM59L	25789	NP_036241	3706	transmembrane protein 59-like
TMEM60	85025	NP_116325	5425	transmembrane protein 60
TMEM61	199964	NP_872338	6652	transmembrane protein 61
TMEM62	80021	NP_079232	5107	transmembrane protein 62
TMEM63A	9725	NP_055513	3975	transmembrane protein 63A
TMEM63B	55362	NP_060896	4448	transmembrane protein 63B
TMEM63C	57156	NP_065164	4656	transmembrane protein 63C
TMEM64	169200	NP_001008495	540	transmembrane protein 64
TMEM64	169200	NP_001139745	1923	transmembrane protein 64
TMEM65	157378	NP_919267	6714	transmembrane protein 65
TMEM67	91147	NP_001135773	1636	transmembrane protein 67
TMEM67	91147	NP_714915	6217	transmembrane protein 67
TMEM68	137695	NP_689630	6056	transmembrane protein 68
TMEM69	51249	NP_057570	4226	transmembrane protein 69
TMEM70	54968	NP_001035703	875	transmembrane protein 70
TMEM70	54968	NP_060336	4360	transmembrane protein 70
TMEM71	137835	NP_001138625	1822	transmembrane protein 71
TMEM71	137835	NP_653250	5883	transmembrane protein 71
TMEM74	157753	NP_694560	6149	transmembrane protein 74
TMEM74B	55321	NP_060824	4428	chromosome 20 open reading frame 46
TMEM79	84283	NP_115699	5327	transmembrane protein 79
TMEM80	283232	NP_001035928	889	transmembrane protein 80
TMEM80	283232	NP_777600	6415	transmembrane protein 80
TMEM86A	144110	NP_699178	6181	transmembrane protein 86A
TMEM86B	255043	NP_776165	6387	transmembrane protein 86B
TMEM87A	25963	NP_001103973	1241	transmembrane protein 87A
TMEM87A	25963	NP_056312	4091	transmembrane protein 87A
TMEM87B	84910	NP_116213	5410	transmembrane protein 87B
TMEM88	92162	NP_981956	6933	transmembrane protein 88
TMEM8A	58986	NP_067082.2	7380	transmembrane protein 8A
TMEM8B	51754	NP_001036054	911	transmembrane protein 8B
TMEM8B	51754	NP_001036055	912	transmembrane protein 8B

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
TMEM8B	51754	NP_057530	4219	transmembrane protein 8B
TMEM9	252839	NP_057540	4221	transmembrane protein 9
TMEM91	641649	NP_001036060	913	transmembrane protein 91
TMEM91	641649	NP_001092291	1148	transmembrane protein 91
TMEM91	641649	NP_001092292	1149	transmembrane protein 91
TMEM91	641649	NP_001092293	1150	transmembrane protein 91
TMEM91	641649	NP_001092294	1151	transmembrane protein 91
TMEM91	641649	NP_001092295	1152	transmembrane protein 91
TMEM92	162461	NP_001161687	2211	transmembrane protein 92
TMEM92	162461	NP_694961	6163	transmembrane protein 92
TMEM95	339168	NP_937797	6750	transmembrane protein 95
TMEM97	27346	NP_055388	3958	transmembrane protein 97
TMEM98	26022	NP_001028676	754	similar to transmembrane protein 98
TMEM98	26022	NP_056359	4098	similar to transmembrane protein 98
TMEM99	147184	NP_660317	5949	transmembrane protein 99
TMEM9B	56674	NP_065695	4688	TMEM9 domain family, member B
TMIE	259236	NP_671729	5996	transmembrane inner ear
TMIGD2	126259	NP_001162597	2238	transmembrane and immunoglobulin domain containing 2
TMIGD2	126259	NP_653216	5875	transmembrane and immunoglobulin domain containing 2
TMPO	7112	NP_001027454	739	thymopoietin
TMPO	7112	NP_001027455	740	thymopoietin
TMPO	7112	NP_003267	2814	thymopoietin
TMPRSS11B	132724	NP_872308	6645	transmembrane protease, serine 11B
TMPRSS11D	9407	NP_004253	3002	transmembrane protease, serine 11D
TMPRSS11E	28983	NP_054777.2	1167	transmembrane protease, serine 11E
TMPRSS11E	28983	NP_054777	3866	transmembrane protease, serine 11E
TMPRSS12	283471	NP_872365	6660	transmembrane protease, serine 12
TMPRSS13	84000	NP_001070731	960	transmembrane protease, serine 13
TMPRSS15	5651	NP_002763	2707	protease, serine, 7 (enterokinase)
TMPRSS2	7113	NP_001128571	1521	transmembrane protease, serine 2
TMPRSS2	7113	NP_005647	3335	transmembrane protease, serine 2
TMPRSS3	64699	NP_076927	4995	transmembrane protease, serine 3
TMPRSS3	64699	NP_115781	5344	transmembrane protease, serine 3
TMPRSS4	56649	NP_001077416	1084	transmembrane protease, serine 4
TMPRSS4	56649	NP_063947	4592	transmembrane protease, serine 4
TMPRSS5	80975	NP_110397	5167	transmembrane protease, serine 5
TMPRSS6	164656	NP_705837	6200	transmembrane protease, serine 6
TMTC1	83857	NP_787057	6462	transmembrane and tetratricopeptide repeat containing 1
TMTC2	160335	NP_689801	6085	transmembrane and tetratricopeptide repeat containing 2
TMTC3	160418	NP_861448	6633	transmembrane and tetratricopeptide repeat containing 3
TMTC4	84899	NP_001073137	990	transmembrane and tetratricopeptide repeat containing 4
TMTC4	84899	NP_116202	5409	transmembrane and tetratricopeptide repeat containing 4
TMUB1	83590	NP_001129516	1593	transmembrane and ubiquitin-like domain containing 1
TMUB1	83590	NP_113622	5223	transmembrane and ubiquitin-like domain containing 1
TMUB2	79089	NP_001070142	948	transmembrane and ubiquitin-like domain containing 2
TMUB2	79089	NP_077012	5013	transmembrane and ubiquitin-like domain containing 2
TMUB2	79089	NP_803190	6493	transmembrane and ubiquitin-like domain containing 2
TMX1	81542	NP_110382	5163	thioredoxin-related transmembrane protein 1
TMX2	51075	NP_001137484	1758	thioredoxin-related transmembrane protein 2
TMX2	51075	NP_057043	4142	thioredoxin-related transmembrane protein 2
TMX3	54495	NP_061895	4549	thioredoxin-related transmembrane protein 3
TNC	3371	NP_002151	2562	tenascin C
TNF	7124	NP_000585	169	tumor necrosis factor (TNF superfamily, member 2)
TNFRSF10A	8797	NP_003835	2911	tumor necrosis factor receptor superfamily, member 10a
TNFRSF10B	8795	NP_003833	2910	tumor necrosis factor receptor superfamily, member 10b
TNFRSF10B	8795	NP_671716	5994	tumor necrosis factor receptor superfamily, member 10b
TNFRSF10C	8794	NP_003832	2909	tumor necrosis factor receptor superfamily, member 10c, decoy without an intracellular domain
TNFRSF10D	8793	NP_003831.2	7317	TNF receptor superfamily member 10d
TNFRSF11A	8792	NP_003830	2908	tumor necrosis factor receptor superfamily, member 11a, NFKB activator
TNFRSF12A	51330	NP_057723.1	7284	TNF receptor superfamily member 12A
TNFRSF13B	23495	NP_036584	3780	tumor necrosis factor receptor superfamily, member 13B
TNFRSF13C	115650	NP_443177	5547	tumor necrosis factor receptor superfamily, member 13C
TNFRSF14	8764	NP_003811	2907	tumor necrosis factor receptor superfamily, member 14 (herpesvirus entry mediator)
TNFRSF17	608	NP_001183	2312	tumor necrosis factor receptor superfamily, member 17
TNFRSF18	8784	NP_004186	2984	tumor necrosis factor receptor superfamily, member 18
TNFRSF18	8784	NP_683699	6005	tumor necrosis factor receptor superfamily, member 18
TNFRSF18	8784	NP_683700	6006	tumor necrosis factor receptor superfamily, member 18
TNFRSF19	55504	NP_061117	4477	tumor necrosis factor receptor superfamily, member 19
TNFRSF19	55504	NP_683760	6009	tumor necrosis factor receptor superfamily, member 19
TNFRSF1A	7132	NP_001056	941	tumor necrosis factor receptor superfamily, member 1A
TNFRSF1B	7133	NP_001057	942	tumor necrosis factor receptor superfamily, member 1B
TNFRSF21	27242	NP_055267	3944	tumor necrosis factor receptor superfamily, member 21
TNFRSF25	8718	NP_001034753	806	tumor necrosis factor receptor superfamily, member 25
TNFRSF25	8718	NP_003781	2896	tumor necrosis factor receptor superfamily, member 25
TNFRSF25	8718	NP_683866	6013	tumor necrosis factor receptor superfamily, member 25
TNFRSF25	8718	NP_683867	6014	tumor necrosis factor receptor superfamily, member 25
TNFRSF25	8718	NP_683868	6015	tumor necrosis factor receptor superfamily, member 25

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
TNFRSF25	8718	NP_683871	6016	tumor necrosis factor receptor superfamily, member 25
TNFRSF4	7293	NP_003318	2822	tumor necrosis factor receptor superfamily, member 4
TNFRSF8	943	NP_001234	2323	tumor necrosis factor receptor superfamily, member 8
TNFRSF8	943	NP_001234.3	6140	tumor necrosis factor receptor superfamily, member 8
TNFRSF9	3604	NP_001552.2	7280	TNF receptor superfamily member 9
TNFSF10	8743	NP_003801	2899	tumor necrosis factor (ligand) superfamily, member 10
TNFSF11	8600	NP_003692	2871	tumor necrosis factor (ligand) superfamily, member 11
TNFSF11	8600	NP_143026	5437	tumor necrosis factor (ligand) superfamily, member 11
TNFSF12	407977	NP_003800.1	7268	TNFSF12-TNFSF13 readthrough
TNFSF13B	10673	NP_001139117	1871	tumor necrosis factor (ligand) superfamily, member 13b
TNFSF13B	10673	NP_006564	3515	tumor necrosis factor (ligand) superfamily, member 13b
TNFSF14	8740	NP_003798	2898	tumor necrosis factor (ligand) superfamily, member 14
TNFSF14	8740	NP_742011	6269	tumor necrosis factor (ligand) superfamily, member 14
TNFSF15	9966	NP_005109	3216	tumor necrosis factor (ligand) superfamily, member 15
TNFSF18	8995	NP_005083.2	7379	TNF superfamily member 18
TNFSF4	7292	NP_003317	2821	tumor necrosis factor (ligand) superfamily, member 4
TNFSF8	944	NP_001235	2324	tumor necrosis factor (ligand) superfamily, member 8
TNFSF9	8744	NP_003802	2900	tumor necrosis factor (ligand) superfamily, member 9
TNMD	64102	NP_071427	4885	tenomodulin
TOMM20	9804	NP_055580	3986	similar to translocase of outer mitochondrial membrane 20Homolog
TOMM7	54543	NP_061932	4554	translocase of outer mitochondrial membrane 7Homolog (yeast)
TOMM70A	9868	NP_055635	3995	translocase of outer mitochondrial membrane 70Homolog A (S. cerevisiae)
TOR1AIP2	163590	NP_071742	4899	torsin A interacting protein 2
TOR1AIP2	163590	NP_659471	5921	torsin A interacting protein 2
TOR1B	27348	NP_055321	3950	torsin family 1, member B (torsin B)
TOR2A	27433	NP_001078816	1089	torsin family 2, member A
TOR2A	27433	NP_001127902	1488	torsin family 2, member A
TOR2A	27433	NP_001127903	1489	torsin family 2, member A
TOR2A	27433	NP_569726	5658	torsin family 2, member A
TP53I13	90313	NP_612358.3	7363	tumor protein p53 inducible protein 13
TPBG	7162	NP_001159864	2164	trophoblast glycoprotein
TPBG	7162	NP_006661	3538	trophoblast glycoprotein
TPBGL	100507050	NP_001182457	7106	trophoblast glycoprotein-like
TPCN1	53373	NP_001137291	1729	two pore segment channel 1
TPCN1	53373	NP_060371	4367	two pore segment channel 1
TPCN2	219931	NP_620714	5835	two pore segment channel 2
TPM1	7168	NP_000357.3	7321	tropomyosin 1 (alpha)
TPO	7173	NP_000538	159	thyroid peroxidase
TPO	7173	NP_783650	6447	thyroid peroxidase
TPO	7173	NP_783652	6448	thyroid peroxidase
TPO	7173	NP_783653	6449	thyroid peroxidase
TPRA1	131601	NP_001129525	1597	G protein-coupled receptor 175
TPRA1	131601	NP_001136118	1682	G protein-coupled receptor 175
TPRA1	131601	NP_001129525.1	4214	G protein-coupled receptor 175
TPSG1	25823	NP_036599	3784	tryptase gamma 1
TPST1	8460	NP_003587	2853	tyrosylprotein sulfotransferase 1
TPST2	8459	NP_001008566	548	tyrosylprotein sulfotransferase 2
TPST2	8459	NP_003586	2852	tyrosylprotein sulfotransferase 2
TPTE	7179	NP_954868	6870	transmembrane phosphatase with tensinHomology
TPTE	7179	NP_954869	6871	transmembrane phosphatase with tensinHomology
TPTE	7179	NP_954870	6872	transmembrane phosphatase with tensinHomology
TRAF3IP3	80342	NP_079504	5137	TRAF3 interacting protein 3
TRAM1	23471	NP_055109	3908	translocation associated membrane protein 1
TRAM1L1	133022	NP_689615	6052	translocation associated membrane protein 1-like 1
TRAM2	9697	NP_036420	3738	translocation associated membrane protein 2
TRAT1	50852	NP_057472	4216	T cell receptor associated transmembrane adaptor 1
TRDN	10345	NP_006064	3429	triadin
TREM1	54210	NP_061113	4474	triggering receptor expressed on myeloid cells 1
TREM2	54209	NP_061838	4537	triggering receptor expressed on myeloid cells 2
TREML1	340205	NP_835468	6526	triggering receptor expressed on myeloid cells-like 1
TREML2	79865	NP_079083.2	7387	triggering receptor expressed on myeloid cells like 2
TRHDE	29953	NP_037513	3815	thyrotropin-releasingHormone degrading enzyme
TRHR	7201	NP_003292	2816	thyrotropin-releasingHormone receptor
TRIL	9865	NP_055632	3994	KIAA0644 gene product
TRIM13	10206	NP_001007279	517	tripartite motif-containing 13
TRIM13	10206	NP_005789	3372	tripartite motif-containing 13
TRIM13	10206	NP_434698	5512	tripartite motif-containing 13
TRIM13	10206	NP_775876	6364	tripartite motif-containing 13
TRIM13	10206	NP_955751	6887	tripartite motif-containing 13
TRIM13	10206	NP_998755	7038	tripartite motif-containing 13
TRIM59	286827	NP_775107	6334	tripartite motif-containing 59
TRIQK	286144	NP_001165266	2284	chromosome 8 open reading frame 83
TRIQK	286144	NP_001165267	2285	chromosome 8 open reading frame 83
TRIQK	286144	NP_001165268	2286	chromosome 8 open reading frame 83
TRIQK	286144	NP_001165269	2287	chromosome 8 open reading frame 83
TRIQK	286144	NP_001165270	2288	chromosome 8 open reading frame 83
TRPA1	8989	NP_015628	3688	transient receptor potential cation channel, subfamily A, member 1

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
TRPC1	7220	NP_003295	2817	transient receptor potential cation channel, subfamily C, member 1
TRPC3	7222	NP_001124170	1434	transient receptor potential cation channel, subfamily C, member 3
TRPC3	7222	NP_003296	2818	transient receptor potential cation channel, subfamily C, member 3
TRPC4	7223	NP_001129427	1578	transient receptor potential cation channel, subfamily C, member 4
TRPC4	7223	NP_001129428	1579	transient receptor potential cation channel, subfamily C, member 4
TRPC4	7223	NP_001129429	1580	transient receptor potential cation channel, subfamily C, member 4
TRPC4	7223	NP_001129430	1581	transient receptor potential cation channel, subfamily C, member 4
TRPC4	7223	NP_003297	2819	transient receptor potential cation channel, subfamily C, member 4
TRPC4	7223	NP_057263	4180	transient receptor potential cation channel, subfamily C, member 4
TRPC5	7224	NP_036603	3785	transient receptor potential cation channel, subfamily C, member 5
TRPC6	7225	NP_004612	3087	transient receptor potential cation channel, subfamily C, member 6
TRPC7	57113	NP_001161048	2182	transient receptor potential cation channel, subfamily C, member 7
TRPC7	57113	NP_001161049	2183	transient receptor potential cation channel, subfamily C, member 7
TRPC7	57113	NP_065122	4644	transient receptor potential cation channel, subfamily C, member 7
TRPM1	4308	NP_002411	2650	transient receptor potential cation channel, subfamily M, member 1
TRPM2	7226	NP_003298	2820	transient receptor potential cation channel, subfamily M, member 2
TRPM3	80036	NP_001007471	519	transient receptor potential cation channel, subfamily M, member 3
TRPM3	80036	NP_001007472	520	transient receptor potential cation channel, subfamily M, member 3
TRPM3	80036	NP_066003	4745	transient receptor potential cation channel, subfamily M, member 3
TRPM3	80036	NP_079247	5111	transient receptor potential cation channel, subfamily M, member 3
TRPM3	80036	NP_996827	6986	transient receptor potential cation channel, subfamily M, member 3
TRPM3	80036	NP_996828	6987	transient receptor potential cation channel, subfamily M, member 3
TRPM3	80036	NP_996829	6988	transient receptor potential cation channel, subfamily M, member 3
TRPM3	80036	NP_996830	6989	transient receptor potential cation channel, subfamily M, member 3
TRPM3	80036	NP_996831	6990	transient receptor potential cation channel, subfamily M, member 3
TRPM4	54795	NP_060106	4315	transient receptor potential cation channel, subfamily M, member 4
TRPM5	29850	NP_055370	3955	transient receptor potential cation channel, subfamily M, member 5
TRPM6	140803	NP_060132	4319	transient receptor potential cation channel, subfamily M, member 6
TRPM7	54822	NP_060142	4320	transient receptor potential cation channel, subfamily M, member 7
TRPM8	79054	NP_076985	5004	transient receptor potential cation channel, subfamily M, member 8
TRPV1	7442	NP_061197	4490	transient receptor potential cation channel, subfamily V, member 1
TRPV1	7442	NP_542435	5621	transient receptor potential cation channel, subfamily V, member 1
TRPV1	7442	NP_542436	5622	transient receptor potential cation channel, subfamily V, member 1
TRPV1	7442	NP_542437	5623	transient receptor potential cation channel, subfamily V, member 1
TRPV2	51393	NP_057197	4172	transient receptor potential cation channel, subfamily V, member 2
TRPV3	162514	NP_659505	5925	transient receptor potential cation channel, subfamily V, member 3
TRPV4	59341	NP_067638	4811	transient receptor potential cation channel, subfamily V, member 4
TRPV4	59341	NP_671737	6000	transient receptor potential cation channel, subfamily V, member 4
TRPV5	56302	NP_062815	4581	transient receptor potential cation channel, subfamily V, member 5
TRPV6	55503	NP_061116	4476	transient receptor potential cation channel, subfamily V, member 6
TSHR	7253	NP_000360	109	thyroid stimulatingHormone receptor
TSHR	7253	NP_001018046	663	thyroid stimulatingHormone receptor
TSHR	7253	NP_001136098	1678	thyroid stimulatingHormone receptor
TSNARE1	203062	NP_659440	5914	t-SNARE domain containing 1
TSPAN1	10103	NP_005718	3354	tetraspanin 1
TSPAN10	83882	NP_114151	5273	tetraspanin 10
TSPAN11	441631	NP_001073978	1034	tetraspanin 11
TSPAN12	23554	NP_036470	3749	tetraspanin 12
TSPAN13	27075	NP_055214	3932	tetraspanin 13
TSPAN14	81619	NP_001121781	1371	tetraspanin 14
TSPAN14	81619	NP_112189	5195	tetraspanin 14
TSPAN15	23555	NP_036471	3750	tetraspanin 15
TSPAN16	26526	NP_036598	3783	tetraspanin 16
TSPAN17	26262	NP_001006617.2	7423	tetraspanin 17
TSPAN18	90139	NP_570139.3	730	tetraspanin 18
TSPAN18	90139	NP_570139	5664	tetraspanin 18
TSPAN2	10100	NP_005716	3353	tetraspanin 2
TSPAN3	10099	NP_001161884	2232	tetraspanin 3
TSPAN3	10099	NP_005715	3352	tetraspanin 3
TSPAN3	10099	NP_944492	6841	tetraspanin 3
TSPAN31	6302	NP_005972	3410	tetraspanin 31
TSPAN32	10077	NP_620591	5828	tetraspanin 32
TSPAN33	340348	NP_848657	6556	tetraspanin 33
TSPAN4	7106	NP_001020405	702	tetraspanin 4
TSPAN4	7106	NP_001020406	703	tetraspanin 4
TSPAN4	7106	NP_001020407	704	tetraspanin 4
TSPAN4	7106	NP_001020408	705	tetraspanin 4
TSPAN4	7106	NP_001020409	706	tetraspanin 4
TSPAN4	7106	NP_001020410	707	tetraspanin 4
TSPAN4	7106	NP_003262	2811	tetraspanin 4
TSPAN5	10098	NP_005714	3351	tetraspanin 5
TSPAN6	7105	NP_003261	2810	tetraspanin 6
TSPAN7	7102	NP_004606	3084	tetraspanin 7
TSPAN8	7103	NP_004607	3085	tetraspanin 8
TSPAN9	10867	NP_001161792	2216	tetraspanin 9
TSPAN9	10867	NP_006666	3541	tetraspanin 9
TSPO	706	NP_000705	209	translocator protein (18kDa)

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
TSPO	706	NP_000705.2	3684	translocator protein (18kDa)
TSPO2	222642	NP_001010873	572	benzodiazapine receptor (peripheral)-like 1
TSPO2	222642	NP_001153198	1956	benzodiazapine receptor (peripheral)-like 1
TTC13	79573	NP_001116307	1287	tetratricopeptide repeat domain 13
TTC13	79573	NP_078801	5041	tetratricopeptide repeat domain 13
TTYH1	57348	NP_001005367	455	tweetyHomolog 1 (Drosophila)
TTYH1	57348	NP_065710	4692	tweetyHomolog 1 (Drosophila)
TTYH2	94015	NP_116035	5384	tweetyHomolog 2 (Drosophila)
TTYH2	94015	NP_443101	5522	tweetyHomolog 2 (Drosophila)
TTYH3	80727	NP_079526	5141	tweetyHomolog 3 (Drosophila)
TUBA8	51807	NP_061816	4533	tubulin, alpha 8
TUBD1	51174	NP_057345	4196	tubulin, delta 1
TUSC3	7991	NP_006756	3559	tumor suppressor candidate 3
TUSC3	7991	NP_839952	6530	tumor suppressor candidate 3
TUSC5	286753	NP_758955	6322	tumor suppressor candidate 5
TVP23A	780776	NP_001072980	986	family with sequence similarity 18, member A
TVP23C	201158	NP_001128508	1513	family with sequence similarity 18, member B2 family with sequence similarity 18, member B
TVP23C	201158	NP_057162	4163	family with sequence similarity 18, member B2 family with sequence similarity 18, member B
TVP23C	201158	NP_660344	5957	family with sequence similarity 18, member B2 family with sequence similarity 18, member B
TXNDC12	51060	NP_056997.1	2987	thioredoxin domain containing 12 (endoplasmic reticulum)
TXNDC12	51060	NP_056997.1	4132	thioredoxin domain containing 12 (endoplasmic reticulum)
TXNDC12	51060	NP_056997.1	5979	thioredoxin domain containing 12 (endoplasmic reticulum)
TXNDC12	51060	NP_056997.1	5981	thioredoxin domain containing 12 (endoplasmic reticulum)
TXNDC15	79770	NP_078991	5069	thioredoxin domain containing 15
TXNDC16	57544	NP_001153519	1974	thioredoxin domain containing 16
TXNDC16	57544	NP_065835	4723	thioredoxin domain containing 16
TYR	7299	NP_000363	110	tyrosinase-like (pseudogene)
TYRO3	7301	NP_006284	3461	TYRO3 protein tyrosine kinase
TYROBP	7305	NP_003323	2823	TYRO protein tyrosine kinase binding protein
TYROBP	7305	NP_937758	6743	TYRO protein tyrosine kinase binding protein
TYRP1	7306	NP_000541	160	tyrosinase-related protein 1
TYW1	55253	NP_060734	4410	tRNA-yW synthesizing protein 1Homolog (S. cerevisiae)
UBA2	10054	NP_005490	3296	ubiquitin-like modifier activating enzyme 2
UBA5	79876	NP_079094	5087	ubiquitin-like modifier activating enzyme 5
UBA5	79876	NP_938143	6768	ubiquitin-like modifier activating enzyme 5
UBAC2	337867	NP_001137544	1769	UBA domain containing 2
UBAC2	337867	NP_808882	6510	UBA domain containing 2
UBE2J1	51465	NP_057105	4152	ubiquitin-conjugating enzyme E2, J1 (UBC6Homolog, yeast)
UBE2J2	118424	NP_477515	5581	ubiquitin-conjugating enzyme E2, J2 (UBC6Homolog, yeast)
UBE2J2	118424	NP_919296	6717	ubiquitin-conjugating enzyme E2, J2 (UBC6Homolog, yeast)
UBE2J2	118424	NP_919439	6730	ubiquitin-conjugating enzyme E2, J2 (UBC6Homolog, yeast)
UBE2J2	118424	NP_919440	6731	ubiquitin-conjugating enzyme E2, J2 (UBC6Homolog, yeast)
UBIAD1	29914	NP_037451	3802	UbiA prenyltransferase domain containing 1
UBXN8	7993	NP_005662	3339	UBX domain protein 8
UCN2	90226	NP_149976	5467	urocortin 2
UGGT1	56886	NP_064505.1	709	UDP-glucose ceramide glucosyltransferase-like 1
UGGT1	56886	NP_064505	4597	UDP-glucose ceramide glucosyltransferase-like 1
UGT1A8	54576	NP_000454	136	UDP glucuronosyltransferase 1 family, polypeptide A3
UGT1A8	54576	NP_001063	943	UDP glucuronosyltransferase 1 family, polypeptide A3
UGT1A8	54576	NP_009051	3632	UDP glucuronosyltransferase 1 family, polypeptide A3
UGT1A8	54576	NP_061948	4557	UDP glucuronosyltransferase 1 family, polypeptide A3
UGT1A8	54576	NP_061949	4558	UDP glucuronosyltransferase 1 family, polypeptide A3
UGT1A8	54576	NP_061950	4559	UDP glucuronosyltransferase 1 family, polypeptide A3
UGT1A8	54576	NP_061951	4560	UDP glucuronosyltransferase 1 family, polypeptide A1, A3, A4, A5, A6, A7, A8, A9, A10
UGT1A8	54576	NP_061966	4563	UDP glucuronosyltransferase 1 family, polypeptide A3
UGT1A8	54576	NP_066307	4760	UDP glucuronosyltransferase 1 family, polypeptide A3
UGT1A8	54576	NP_995584	6963	UDP glucuronosyltransferase 1 family, polypeptide A3
UGT2A3	79799	NP_079019	5075	UDP glucuronosyltransferase 2 family, polypeptide A3
UGT2B15	7366	NP_001067	944	UDP glucuronosyltransferase 2 family, polypeptide B15
UGT2B17	7367	NP_001068	945	UDP glucuronosyltransferase 2 family, polypeptide B17
UGT2B4	7363	NP_066962	4780	UDP glucuronosyltransferase 2 family, polypeptide B4
UGT3A1	133688	NP_001165344	2290	UDP glycosyltransferase 3 family, polypeptide A1
UGT3A1	133688	NP_689617	6054	UDP glycosyltransferase 3 family, polypeptide A1
UGT8	7368	NP_001121646	1364	UDP glycosyltransferase 8
UGT8	7368	NP_003351	2825	UDP glycosyltransferase 8
ULBP1	80329	NP_001304018.1	7458	UL16 binding protein 1
ULBP2	80328	NP_079493	5133	UL16 binding protein 2
ULBP3	79465	NP_078794.1	7298	UL16 binding protein 3
UMOD	7369	NP_001008390	531	uromodulin
UMOD	7369	NP_003352	2826	uromodulin
UMODL1	89766	NP_001004416.2	7371	uromodulin like 1
UNC50	25972	NP_054763	3861	unc-50Homolog (C. elegans)
UNC5A	90249	NP_588610	5690	unc-5Homolog A (C. elegans)
UNC5B	219699	NP_734465	6261	unc-5Homolog B (C. elegans)
UNC5C	8633	NP_003719	2876	unc-5Homolog C (C. elegans)
UNC5CL	222643	NP_775832	6358	unc-5Homolog C (C. elegans)-like
UNC5D	137970	NP_001309747.1	7461	unc-5 netrin receptor D

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
UNC93B1	81622	NP_112192	5196	unc-93Homolog B1 (C. elegans), unc-93Homolog B6 (C. elegans), unc-93Homolog B3 pseudogene (C. elegans), similar to unc-93Homolog B1 (C. elegans)
UPK1A	11045	NP_008931	3611	uroplakin 1A
UPK1B	7348	NP_008883	3603	uroplakin 1B
UPK2	7379	NP_006751.1	7281	uroplakin 2
UPK3A	7380	NP_001161046	2181	uroplakin 3A
UPK3A	7380	NP_008884	3604	uroplakin 3A
UPK3B	80761	NP_085047	5148	uroplakin 3B
UPK3B	80761	NP_872625	6675	uroplakin 3B
UQCR11	10975	NP_006821	3575	ubiquinol-cytochrome c reductase, 6.4kDa subunit
USH2A	7399	NP_009054.5	7403	usherin
USP19	10869	NP_006668	3542	ubiquitin specific peptidase 19
USP27X	389856	NP_001138545	1808	ubiquitin specific peptidase 27, X-linked
USP30	84749	NP_116052	5385	ubiquitin specific peptidase 30
USP48	84196	NP_001027902	747	ubiquitin specific peptidase 48
USP48	84196	NP_115612	5311	ubiquitin specific peptidase 48
UST	10090	NP_005706	3350	uronyl-2-sulfotransferase
UTS2B	257313	NP_937795	6749	urotensin 2 domain containing
UTS2R	2837	NP_061822	4534	urotensin 2 receptor
UTY	7404	NP_009056	3633	ubiquitously transcribed tetratricopeptide repeat gene, Y-linked
UTY	7404	NP_872600	6671	ubiquitously transcribed tetratricopeptide repeat gene, Y-linked
UTY	7404	NP_872601	6672	ubiquitously transcribed tetratricopeptide repeat gene, Y-linked
UXS1	80146	NP_079352	5117	UDP-glucuronate decarboxylase 1
VAMP1	6843	NP_055046	3891	vesicle-associated membrane protein 1 (synaptobrevin 1)
VAMP1	6843	NP_058439	4273	vesicle-associated membrane protein 1 (synaptobrevin 1)
VAMP1	6843	NP_954740	6869	vesicle-associated membrane protein 1 (synaptobrevin 1)
VAMP2	6844	NP_055047	3892	vesicle-associated membrane protein 2 (synaptobrevin 2)
VAMP3	9341	NP_004772	3124	vesicle-associated membrane protein 3 (cellubrevin)
VAMP4	8674	NP_003753	2887	vesicle-associated membrane protein 4
VAMP5	10791	NP_006625	3532	vesicle-associated membrane protein 5 (myobrevin)
VAMP7	6845	NP_001138621	1821	vesicle-associated membrane protein 7
VAMP7	6845	NP_005629	3333	vesicle-associated membrane protein 7
VAMP8	8673	NP_003752	2886	vesicle-associated membrane protein 8 (endobrevin)
VANGL1	81839	NP_620409	5808	vang-like 1 (van gogh, Drosophila)
VANGL2	57216	NP_065068	4631	vang-like 2 (van gogh, Drosophila)
VAPA	9218	NP_003565	2850	VAMP (vesicle-associated membrane protein)-associated protein A, 33kDa
VAPA	9218	NP_919415	6723	VAMP (vesicle-associated membrane protein)-associated protein A, 33kDa
VAPB	9217	NP_004729	3114	VAMP (vesicle-associated membrane protein)-associated protein B and C
VASN	114990	NP_612449	5753	vasorin
VCAM1	7412	NP_001069	946	vascular cell adhesion molecule 1
VCAM1	7412	NP_542413	5620	vascular cell adhesion molecule 1
VEZT	55591	NP_060069	4311	vezatin, adherens junctions transmembrane protein
VIMP	55829	NP_060915	4451	selenoprotein S
VIMP	55829	NP_982298	6941	selenoprotein S
VIP	7432	NP_003372	2827	vasoactive intestinal peptide
VIP	7432	NP_919416	6724	vasoactive intestinal peptide
VIPR1	7433	NP_004615	3088	vasoactive intestinal peptide receptor 1
VIPR2	7434	NP_003373	2828	vasoactive intestinal peptide receptor 2
VKORC1	79001	NP_076869	4991	vitamin K epoxide reductase complex, subunit 1
VKORC1	79001	NP_996560	6972	vitamin K epoxide reductase complex, subunit 1
VKORC1L1	154807	NP_775788	6352	vitamin K epoxide reductase complex, subunit 1-like 1
VMA21	203547	NP_001017980	659	VMA21 vacuolarH+-ATPaseHomolog (S. cerevisiae)
VMP1	81671	NP_001316323.1	7462	vacuole membrane protein 1
VN1R1	57191	NP_065684.1	7289	vomeronasal 1 receptor 1
VN1R2	317701	NP_776255	6398	vomeronasal 1 receptor 2
VNN2	8875	NP_004656	3098	vanin 2
VNN2	8875	NP_511043	5594	vanin 2
VNN3	55350	NP_001278631.1	678	vanin 3
VNN3	55350	NP_001278631.1	4438	vanin 3
VNN3	55350	NP_001278631.1	5595	vanin 3
VRK1	7443	NP_003375	2829	vaccinia related kinase 1
VRK2	7444	NP_001123952	1426	vaccinia related kinase 2
VRK2	7444	NP_001123953	1427	vaccinia related kinase 2
VRK2	7444	NP_001123954	1428	vaccinia related kinase 2
VRK2	7444	NP_001123955	1429	vaccinia related kinase 2
VRK2	7444	NP_001123952.1	1591	vaccinia related kinase 2
VRK2	7444	NP_006287	3462	vaccinia related kinase 2
VSIG1	340547	NP_001164024	2244	V-set and immunoglobulin domain containing 1
VSIG1	340547	NP_872413	6666	V-set and immunoglobulin domain containing 1
VSIG10	54621	NP_061959	4562	hypothetical protein FLJ20674
VSIG10L	147645	NP_001157394	2059	hypothetical protein LOC147645
VSIG2	23584	NP_055127	3911	V-set and immunoglobulin domain containing 2
VSIG4	11326	NP_001093901	1188	V-set and immunoglobulin domain containing 4
VSIG4	11326	NP_009199	3666	V-set and immunoglobulin domain containing 4
VSTM1	284415	NP_940883	6800	V-set and transmembrane domain containing 1
VSTM2A	222008	NP_001304772.1	7459	V-set and transmembrane domain containing 2A
VSTM4	196740	NP_001026916	733	chromosome 10 open reading frame 72

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
VSTM4	196740	NP_659421	5912	chromosome 10 open reading frame 72
VTCN1	79679	NP_078902	5057	V-set domain containing T cell activation inhibitor 1
VTI1A	143187	NP_660207	5935	vesicle transport through interaction with t-SNAREsHomolog 1A (yeast)
VTI1B	10490	NP_006361	3476	vesicle transport through interaction with t-SNAREsHomolog 1B (yeast)
VWA9	81556	NP_001129515	1592	chromosome 15 open reading frame 44
VWA9	81556	NP_001129515.1	5179	chromosome 15 open reading frame 44
WBP1L	54838	NP_001077382	1083	chromosome 10 open reading frame 26
WBP1L	54838	NP_060257	4342	chromosome 10 open reading frame 26
WBSCR17	64409	NP_071924	4923	Williams-Beuren syndrome chromosome region 17
WBSCR28	135886	NP_872310	6646	Williams-Beuren syndrome chromosome region 28
WDR19	57728	NP_079408	5124	WD repeat domain 19
WDR33	55339	NP_001006623	479	WD repeat domain 33
WDR33	55339	NP_001006624	480	WD repeat domain 33
WDR33	55339	NP_060853	4436	WD repeat domain 33
WDR83OS	51398	NP_057229	4177	chromosome 19 open reading frame 56
WFDC10A	140832	NP_542791	5628	WAP four-disulfide core domain 10A
WFDC10B	280664	NP_742003	6268	WAP four-disulfide core domain 10B
WFDC10B	280664	NP_742143	6282	WAP four-disulfide core domain 10B
WFDC11	259239	NP_671730	5997	WAP four-disulfide core domain 11
WFDC2	10406	NP_006094	3436	WAP four-disulfide core domain 2
WFDC9	259240	NP_671731	5998	WAP four-disulfide core domain 9
WFIKKN2	124857	NP_783165	6444	WAP, follistatin/kazal, immunoglobulin, kunitz and netrin domain containing 2
WFS1	7466	NP_001139325	1884	Wolfram syndrome 1 (wolframin)
WFS1	7466	NP_005996	3412	Wolfram syndrome 1 (wolframin)
WLS	79971	NP_001002292	401	G protein-coupled receptor 177
WLS	79971	NP_079187	5101	G protein-coupled receptor 177
WNT1	7471	NP_005421	3284	wingless-type MMTV integration site family, member 1
WNT11	7481	NP_004617	3090	wingless-type MMTV integration site family, member 11
WNT16	51384	NP_057171	4167	wingless-type MMTV integration site family, member 16
WNT16	51384	NP_476509	5578	wingless-type MMTV integration site family, member 16
WNT3	7473	NP_110380	5162	wingless-type MMTV integration site family, member 3
WNT4	54361	NP_110388	5165	wingless-type MMTV integration site family, member 4
WNT5A	7474	NP_003383	2831	wingless-type MMTV integration site family, member 5A
WNT6	7475	NP_006513	3506	wingless-type MMTV integration site family, member 6
WNT7A	7476	NP_004616	3089	wingless-type MMTV integration site family, member 7A
WNT7B	7477	NP_478679	5586	wingless-type MMTV integration site family, member 7B
WNT9B	7484	NP_003387	2832	wingless-type MMTV integration site family, member 9B
WRB	7485	NP_001139690	1914	tryptophan rich basic protein
WRB	7485	NP_004618	3091	tryptophan rich basic protein
WSCD1	23302	NP_056068	4051	WSC domain containing 1
WSCD2	9671	NP_055468	3968	WSC domain containing 2
WT1	7490	NP_000369	111	Wilms tumor 1
WT1	7490	NP_077742	5031	Wilms tumor 1
WT1	7490	NP_077743	5032	Wilms tumor 1
WT1	7490	NP_077744	5033	Wilms tumor 1
XCR1	2829	NP_001019815	682	chemokine (C motif) receptor 1
XCR1	2829	NP_005274	3251	chemokine (C motif) receptor 1
XG	7499	NP_001135391	1625	Xg blood group
XG	7499	NP_001135392	1626	Xg blood group
XG	7499	NP_780778	6443	Xg blood group
XK	7504	NP_066569	4766	X-linked Kx blood group (McLeod syndrome)
XKR4	114786	NP_443130	5531	XK, Kell blood group complex subunit-related family, member 4
XKR6	286046	NP_775954	6380	XK, Kell blood group complex subunit-related family, member 6
XKRX	402415	NP_997724	7036	XK, Kell blood group complex subunit-related, X-linked
XPR1	9213	NP_001129141	1546	xenotropic and polytropic retrovirus receptor
XPR1	9213	NP_004727	3112	xenotropic and polytropic retrovirus receptor
XXYLT1	152002	NP_689744	6077	chromosome 3 open reading frame 21
XYLT1	64131	NP_071449	4891	xylosyltransferase I
XYLT2	64132	NP_071450	4892	xylosyltransferase II
YIF1A	10897	NP_065203	4667	Yip1 interacting factorHomolog A (S. cerevisiae)
YIF1B	90522	NP_001034760	807	Yip1 interacting factorHomolog B (S. cerevisiae)
YIF1B	90522	NP_001034761	808	Yip1 interacting factorHomolog B (S. cerevisiae)
YIF1B	90522	NP_001034762	809	Yip1 interacting factorHomolog B (S. cerevisiae)
YIF1B	90522	NP_001138933	1862	Yip1 interacting factorHomolog B (S. cerevisiae)
YIF1B	90522	NP_001138934	1863	Yip1 interacting factorHomolog B (S. cerevisiae)
YIF1B	90522	NP_001138935	1864	Yip1 interacting factorHomolog B (S. cerevisiae)
YIF1B	90522	NP_001034760.1	5508	Yip1 interacting factorHomolog B (S. cerevisiae)
YIPF1	54432	NP_061855	4546	Yip1 domain family, member 1
YIPF2	78992	NP_076934	4997	Yip1 domain family, member 2
YIPF3	25844	NP_056203	4074	Yip1 domain family, member 3
YIPF4	84272	NP_115688	5323	Yip1 domain family, member 4
YIPF5	81555	NP_001020118	691	Yip1 domain family, member 5
YIPF5	81555	NP_110426	5178	Yip1 domain family, member 5
YIPF6	286451	NP_776195	6393	Yip1 domain family, member 6
YME1L1	10730	NP_055078	3900	YME1-like 1 (S. cerevisiae)
YME1L1	10730	NP_647473	5852	YME1-like 1 (S. cerevisiae)
ZACN	353174	NP_851321	6578	zinc activated ligand-gated ion channel

FIG. 2 (CONT.)

Antigen	GeneID	Accession No.	SEQ ID NO:	RefSeq Gene Name
ZAN	7455	NP_003377	2830	zonadhesin
ZAN	7455	NP_775082	6327	zonadhesin
ZDHHC1	29800	NP_037436	3798	zinc finger, DHHC-type containing 1
ZDHHC11	79844	NP_079062	5081	zinc finger, DHHC-type containing 11
ZDHHC12	84885	NP_116188	5404	zinc finger, DHHC-type containing 12
ZDHHC13	54503	NP_001001483	362	zinc finger, DHHC-type containing 13
ZDHHC13	54503	NP_061901	4551	zinc finger, DHHC-type containing 13
ZDHHC14	79683	NP_078906	5059	zinc finger, DHHC-type containing 14
ZDHHC14	79683	NP_714968	6221	zinc finger, DHHC-type containing 14
ZDHHC15	158866	NP_001139728	1915	zinc finger, DHHC-type containing 15
ZDHHC15	158866	NP_001139729	1916	zinc finger, DHHC-type containing 15
ZDHHC15	158866	NP_659406	5909	zinc finger, DHHC-type containing 15
ZDHHC17	23390	NP_056151	4063	zinc finger, DHHC-type containing 17
ZDHHC18	84243	NP_115659	5319	zinc finger, DHHC-type containing 18
ZDHHC19	131540	NP_001034706	805	zinc finger, DHHC-type containing 19
ZDHHC2	51201	NP_057437	4208	zinc finger, DHHC-type containing 2
ZDHHC20	253832	NP_694983	6166	zinc finger, DHHC-type containing 20
ZDHHC21	340481	NP_848661	6557	zinc finger, DHHC-type containing 21
ZDHHC22	283576	NP_777636	6433	zinc finger, DHHC-type containing 22
ZDHHC23	254887	NP_775841	6360	zinc finger, DHHC-type containing 23
ZDHHC24	254359	NP_997223	7010	zinc finger, DHHC-type containing 24
ZDHHC3	51304	NP_001128651	1534	zinc finger, DHHC-type containing 3
ZDHHC3	51304	NP_001128651.1	1535	zinc finger, DHHC-type containing 3
ZDHHC3	51304	NP_057682	4256	zinc finger, DHHC-type containing 3
ZDHHC4	55146	NP_001127859	1479	zinc finger, DHHC-type containing 4
ZDHHC4	55146	NP_001127860	1480	zinc finger, DHHC-type containing 4
ZDHHC4	55146	NP_001127861	1481	zinc finger, DHHC-type containing 4
ZDHHC4	55146	NP_060576	4393	zinc finger, DHHC-type containing 4
ZDHHC5	25921	NP_056272	4086	zinc finger, DHHC-type containing 5
ZDHHC6	64429	NP_071939	4927	zinc finger, DHHC-type containing 6
ZDHHC7	55625	NP_001139020	1870	zinc finger, DHHC-type containing 7
ZDHHC7	55625	NP_060210	4332	zinc finger, DHHC-type containing 7
ZDHHC8	29801	NP_037505	3813	zinc finger, DHHC-type containing 8
ZDHHC9	51114	NP_001008223	527	zinc finger, DHHC-type containing 9
ZDHHC9	51114	NP_057116	4156	zinc finger, DHHC-type containing 9
ZFPL1	7542	NP_006773	3562	zinc finger protein-like 1
ZFYVE27	118813	NP_001002261	393	zinc finger, FYVE domain containing 27
ZFYVE27	118813	NP_001002262	394	zinc finger, FYVE domain containing 27
ZFYVE27	118813	NP_653189	5870	zinc finger, FYVE domain containing 27
ZG16B	124220	NP_660295	5941	zymogen granule protein 16Homolog B (rat)
ZMPSTE24	10269	NP_005848	3388	zinc metallopeptidase (STE24Homolog, S. cerevisiae)
ZMYM6	9204	NP_009098	3639	hypothetical LOC100130633 zinc finger, MYM-type 6
ZMYM6NB	100506144	NP_001182085	7112	ZMYM6 neighbor
ZMYND11	10771	NP_001189393.1	2014	zinc finger, MYND domain containing 11
ZMYND11	10771	NP_006615	3530	zinc finger, MYND domain containing 11
ZMYND11	10771	NP_997644	7033	zinc finger, MYND domain containing 11
ZNF138	7697	NP_001153655	1978	zinc finger protein 138
ZNF138	7697	NP_006515	3507	zinc finger protein 138
ZNF304	57343	NP_065708	4691	zinc finger protein 304
ZNF546	339327	NP_848639	6553	zinc finger protein 546
ZNF559	84527	NP_115886	5361	zinc finger protein 559
ZNF7	7553	NP_003407	2833	zinc finger protein 7
ZNF816	125893	NP_001026835	720	zinc finger protein 816A
ZNRF3	84133	NP_115549	5304	zinc and ring finger 3
ZNRF4	148066	NP_859061	6623	zinc and ring finger 4
ZP1	22917	NP_997224	7011	zona pellucida glycoprotein 1 (sperm receptor)
ZP2	7783	NP_001277033.1	7443	zona pellucida glycoprotein 2
ZP4	57829	NP_067009.1	7291	zona pellucida glycoprotein 4
ZPBP	11055	NP_001153350	1962	zona pellucida binding protein
ZPBP	11055	NP_008940	3614	zona pellucida binding protein
ZPLD1	131368	NP_778226	6438	zona pellucida-like domain containing 1

FIG. 3A

2 INPUT "AND" GATE with antigens A **AND** X

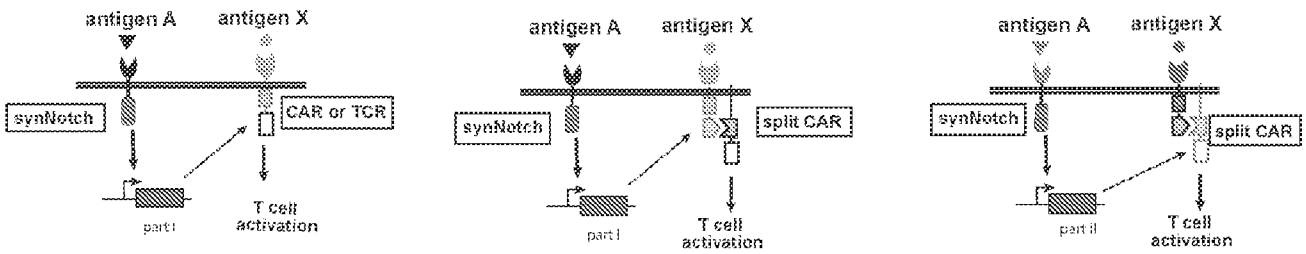

FIG. 3B

3 INPUT "AND-GATE" with antigens A **AND** B **AND** X

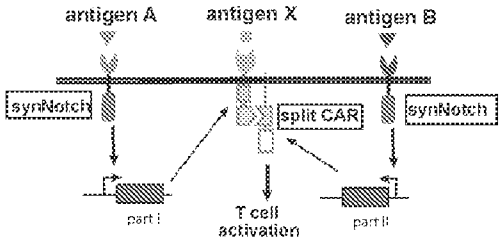

FIG. 3C

2 INPUT "AND-NOT" GATE with antigens X **AND-NOT** A

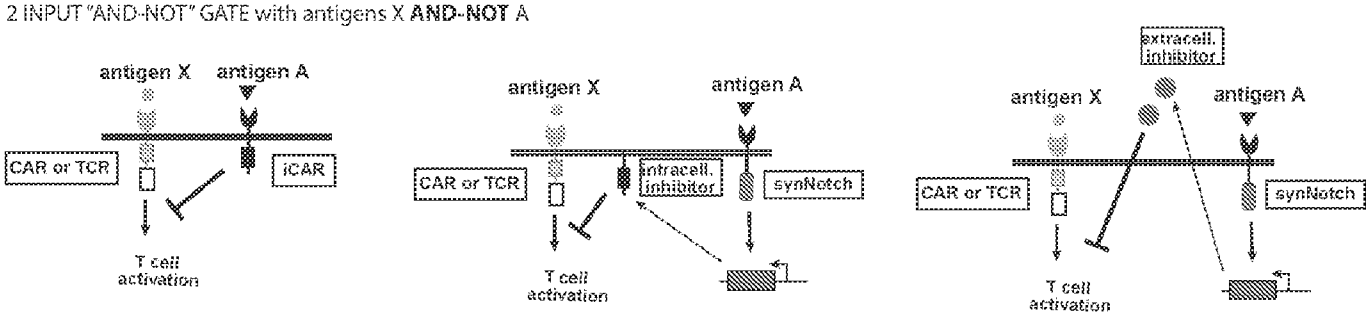

FIG. 3D

3 INPUT "AND-NOT" GATE with antigens A **AND** X **AND-NOT** B

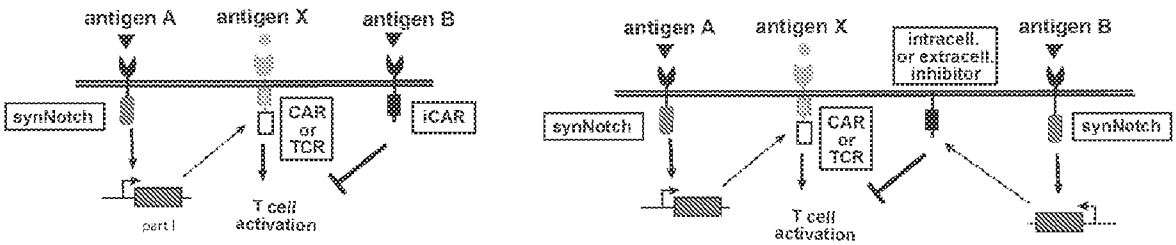

FIG. 4

Antigen	Gene Symbol	Entrez GeneID	Exemplary Cancers	Accession No.	SEQ ID NOs:
ALK	ALK	238	Hallmark, large cell lymphomas, neuroblastoma, and non small cell lung cancer	NP_004295	7119
Axl	AXL	558	Lung cancer	NP_068713, NP_001690	7120, 7121
B7H3	CD276	80381	Sarcoma, glioma	NP_079516, NP_001019907	7122, 7123
B7H6	NCR3LG1	374383	leukemia, lymphoma and gastrointestinal stromal tumors	NP_001189368	7124
BCMA	TNFRSF17	608	multiple myeloma	NP_001183	7125
CAIX	CA9	768	Kidney	NP_001207	7126
CD123	IL3RA	3563	Myeloid	NP_002174	7127
CD138	SDC1	6382	multiple myeloma	NP_002988, NP_001006947	7128, 7129
CD160	CD160	11126	Leukemia	NP_008984	7130
CD171	L1CAM	3897	Neuroblastoma	NP_076493, NP_000416, NP_001137435	7131, 7132, 7133
CD19	CD19	930	B-cell	NP_001761	7134
CD20	MS4A1	931	B-cell	NP_690605, NP_068769	7135, 7136
CD22	CD22	933	B-cell	NP_001762	7137
CD30	TNFRSF8	943	B-cell	NP_694421, NP_001234	7138, 7139
CD33	CD33	945	Myeloid	NP_001076087, NP_001763	7140, 7141
CD38	CD38	952	multiple myeloma	NP_001766	7142
CD44 v6/v7	CD44	960	Cervical	NP_001001389, NP_001001392, NP_001001391, NP_001001390, NP_000601	7143, 7144, 7145, 7146, 7147
CD70	CD70	970	B-cell/T-cell	NP_001243	7148
CS1	SLAMF7	57823	multiple myeloma	NP_067004	7149
EGP2 (EGP40)	EPCAM	4072	Colon, Carcinomas, breast, lung	NP_002345	7150
EphA2	EPHA2	1969	Glioma, lung	NP_004422	7151
EphA3	EPHA3	2042	Lung, kidney, melanoma, glioma, hematological malignancies	NP_872585, NP_005224	7152, 7153
ErbB3	ERBB3	2065	Breast, ovarian	NP_001005915, NP_001973	7154, 7155
ErbB4	ERBB4	2066	Breast, ovarian	NP_001036064, NP_005226	7156, 7157
FAP	FAP	2191	Cancer associated fibroblasts	NP_004451	7158
Folate receptor alpha	FOLR1	2348	Ovarian	NP_057937, NP_057941, NP_057943, NP_057942, NP_000793, NP_057936	7159, 7160, 7161, 7162, 7163, 7164
Folate receptor beta	FOLR2	2350	Acute Myeloid Leukemia	NP_000794, NP_001107007, NP_001107006, NP_001107008	7165, 7166, 7167, 7168
GD2	B4GALNT1	2583	Neuroblastoma, sarcoma, melanoma	NP_001469	7169
GD3	ST8SIA1	6489	Melanoma, lung cancer	NP_003025	7170
gpA33	GPA33	10223	Colorectal carcinoma	NP_005805	7171
GPC3	GPC3	2719	Hepatocellular Carcinoma	NP_004475, NP_001158089, NP_001158090, NP_001158091	7172, 7173, 7174, 7175
Her2 (NEU)	ERBB2	2064	Breast, lung, prostate, glioma	NP_001005862, NP_004439, NP_001005915, NP_001973, NP_001036064, NP_005226	7176, 7177, 7178, 7179, 7180, 7181

FIG. 4 (CONT.)

Antigen	Gene Symbol	Entrez GeneID	Exemplary Cancers	Accession No.	SEQ ID NOs:
HLA-A2/NY-ESO-1	CTAG1B	1485	Sarcoma, melanoma	NP_640343, NP_001318	7182, 7183
HMW-MAA	CSPG4	1464	Melanoma	NP_001888	7184
IGF1R	IGF1R	3480	Lung, breast, head and neck, prostate, thyroid, glioma	NP_000866	7185
IL11Ra	IL11RA	3590	Osteosarcoma	NP_671518, NP_004503, NP_001136256	7186, 7187, 7188
IL13Ra2	IL13RA2	3598	Glioma	NP_000631	7189
Kappa	IGK	50802	B-cell, multiple myeloma	NG_000834, NG_000833	7190, 7191
MAGE-A1	MAGEA1	4100	Melanoma	NP_004979	7192
MART-1	MLANA	2315	Melanoma	NP_005502	7193
Mesothelin	MSLN	10232	Mesothelioma, breast, pancreas	NP_005814, NP_037536	7194, 7195
Met	MET	4233	Lung cancer	NP_001120972, NP_000236	7196, 7197
MUC13	MUC13	56667	Adenocarcinoma	NP_149038	7198
MUC17	MUC17	140453	Adenocarcinoma	NP_001035194	7199
Mucin-1	MUC1	4582	Ovarian, breast, prostate	NP_002447, NP_001018017, NP_001018016, NP_001037857, NP_001037858, NP_001037855, NP_001037856	7200, 7201, 7202, 7203, 7204, 7205, 7206
Mucin-16	MUC16	94025	Adenocarcinoma	NP_078966, XP_002345182	7207, 7208
NCAM	NCAM1	4684	Neuroblastoma, colorectal	NP_000606, NP_851996, NP_001070150	7209 7210, 7211
PSCA	PSCA	8000	Prostate, pancreatic	NP_005663	7212
PSMA	FOLH1	2346	Prostate	NP_004467, NP_001014986	7213, 7214
RANKL	TNFSF11	8600	Prostate cancer and bone metastases	NP_003692, NP_143026	7215, 7216
ROR1	ROR1	4919	B-cell	NP_001077061, NP_005003	7217, 7218
TRAIL-R1	TNFRSF10A	8797	Solid tumors (colon, lung, pancreas) and hematological malignancies	NP_003835	7219
TRAIL-R2	TNFRSF10B	8795	Solid tumors (colon, lung, pancreas) and hematological malignancies	NP_671716, NP_003833	7220, 7221
VEGFR-2	KDR	3791	Tumor vasculature	NP_002244	7222
WT1	WT1	7490	Leukemia, solid tumors (e.g., lung cancer)	NP_077743, NP_077742, NP_000369, NP_077744	7223, 7224, 7225, 7226
α5β1	ITGB1	3688	Tumor vasculature	NP_391989, NP_391988, NP_391987, NP_389647, NP_002202, NP_596867	7227, 7228, 7229, 7230, 7231, 7232
α5β3	ITGB3	3690	Tumor vasculature	NP_000203	7233
NKG2D ligands	KLRK1	22914	Myeloid, Ovarian, Sarcoma	NP_031386	7234
Lewis Y (CD174)	FUT3	2525	Breast/ovarian/pancreatic , multiple myeloma	NP_001091109, NP_001091108, NP_000140, NP_001091110	7235, 7236, 7237, 7238,
CEA	CEACAM5	1048	Epithelial tumors (breast, colon, lung)	NP_004354	7239

FIG. 5A

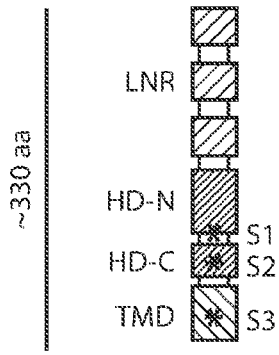

FIG. 5B

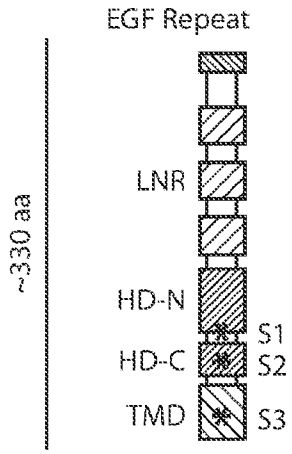

FIG. 5C

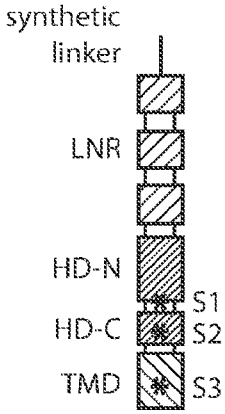

FIG. 5D

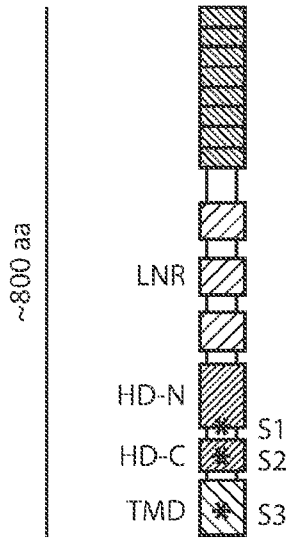

FIG. 5E

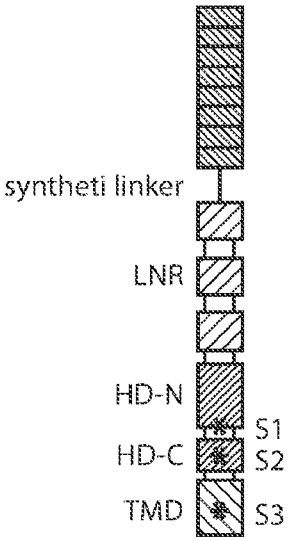

FIG. 5F

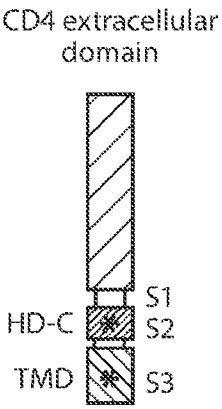

FIG. 5G

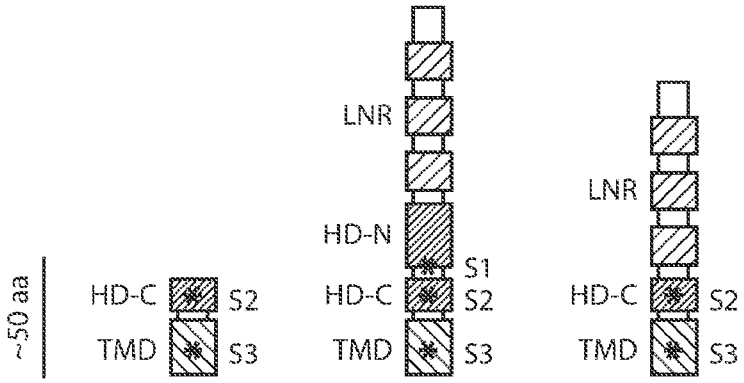

FIG. 6A

FIG. 6B

FIG. 6C

FIG. 6D

FIG. 6E

FIG. 6F

FIG. 7

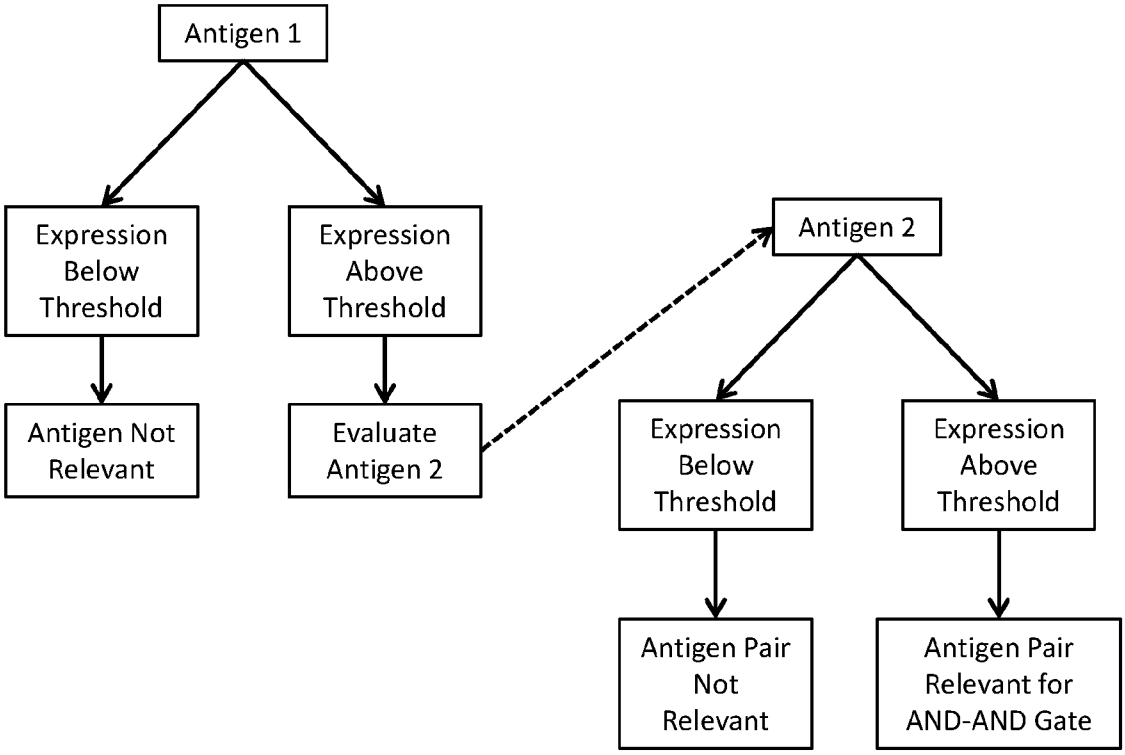
Antigen 1
Expression Below Threshold
Expression Above Threshold
Antigen Not Relevant
Evaluate Antigen 2
Antigen 2
Expression Below Threshold
Expression Above Threshold
Antigen Pair Not Relevant
Antigen Pair Relevant for AND-AND Gate

FIG. 8

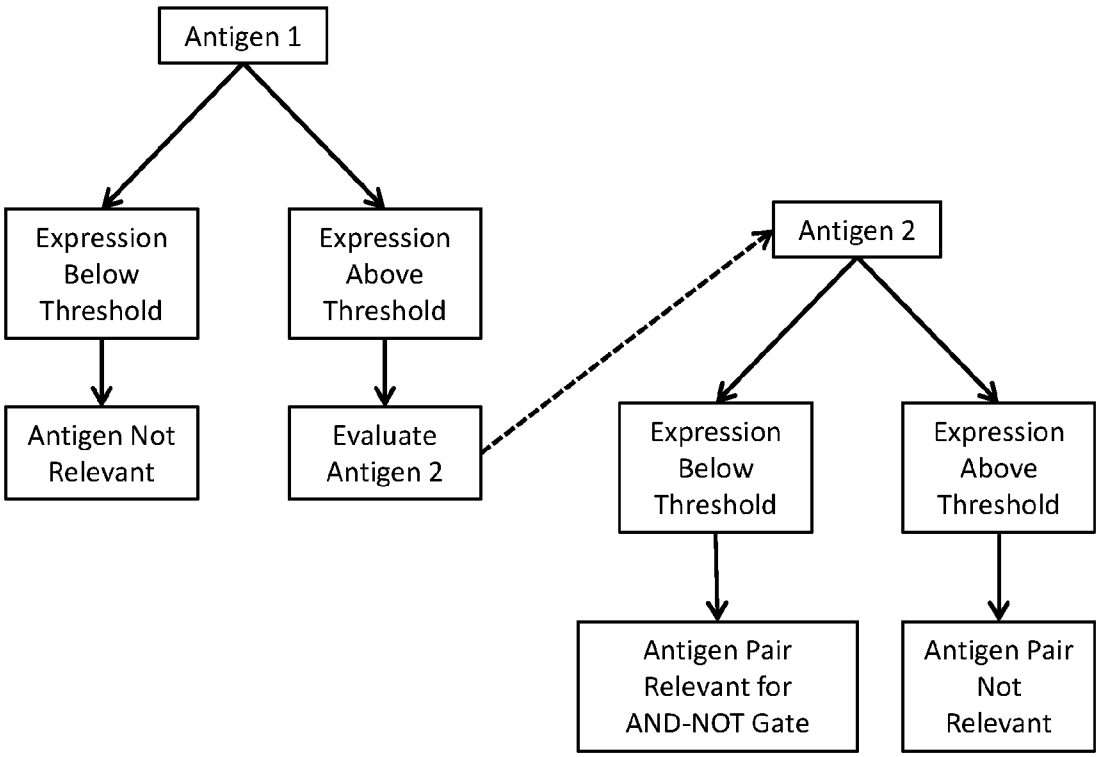
Antigen 1
Expression Below Threshold
Expression Above Threshold
Antigen Not Relevant
Evaluate Antigen 2
Antigen 2
Expression Below Threshold
Expression Above Threshold
Antigen Pair Relevant for AND-NOT Gate
Antigen Pair Not Relevant

FIG. 9

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
ADAM12 AND EPCAM	Breast	0.868687	0.955556	0.796296
NOX4 AND EPCAM	Breast	0.757282	0.795918	0.722222
SLC5A6 AND EPCAM	Breast	0.75	0.970588	0.611111
EPCAM AND CALHM2	Breast	0.821053	0.95122	0.722222
PLXND1 AND EPCAM	Breast	0.786517	1	0.648148
TGFB3 AND EPCAM	Breast	0.741573	0.942857	0.611111
PMEPA1 AND EPCAM	Breast	0.733333	0.916667	0.611111
SEMA4C AND EPCAM	Breast	0.791209	0.972973	0.666667
ADAM12 AND ERBB2	Breast	0.714286	1	0.555556
EPCAM AND APLNR	Breast	0.772277	0.829787	0.722222
PRLR AND EPCAM	Breast	0.707071	0.777778	0.648148
EPCAM AND NTM	Breast	0.705882	0.75	0.666667
UNC5B AND EPCAM	Breast	0.719101	0.914286	0.592593
EPCAM AND CELSR2	Breast	0.699029	0.734694	0.666667
SLC4A2 AND EPCAM	Breast	0.698795	1	0.537037
MFSD10 AND EPCAM	Breast	0.727273	0.941176	0.592593
FZD2 AND EPCAM	Breast	0.681818	0.882353	0.555556
EPCAM AND TNFSF4	Breast	0.828283	0.911111	0.759259
F2RL2 AND EPCAM	Breast	0.682927	1	0.518519
EPCAM AND SLC1A3	Breast	0.792453	0.807692	0.777778
EPCAM AND LRRC32	Breast	0.72549	0.770833	0.685185
EPCAM AND FGFR1	Breast	0.666667	0.708333	0.62963
EPCAM AND FZD7	Breast	0.666667	0.794872	0.574074
EPCAM AND CD84	Breast	0.674699	0.965517	0.518519
EPCAM AND SLC39A10	Breast	0.754717	0.769231	0.740741
CXCL9 AND EPCAM	Breast	0.817204	0.974359	0.703704
EPCAM AND GJA1	Breast	0.75	0.724138	0.777778
EPCAM AND S1PR3	Breast	0.660377	0.673077	0.648148
EPCAM AND SLC3A2	Breast	0.747664	0.754717	0.740741
SLC52A2 AND EPCAM	Breast	0.659794	0.744186	0.592593
EPCAM AND SLC10A3	Breast	0.6875	0.785714	0.611111
EPCAM AND TPCN2	Breast	0.707965	0.677966	0.740741
EPCAM AND PCDHB10	Breast	0.652632	0.756098	0.574074
EPCAM AND APOLD1	Breast	0.65625	0.567568	0.777778
EPCAM AND SLC9A7	Breast	0.651163	0.56	0.777778
EPCAM AND JAG2	Breast	0.752475	0.808511	0.703704
EPCAM AND CXCR4	Breast	0.649123	0.616667	0.685185
EPCAM AND ITGAM	Breast	0.848485	0.933333	0.777778
EPCAM AND LPAR2	Breast	0.673684	0.780488	0.592593
EPCAM AND ESYT1	Breast	0.658537	0.964286	0.5
EPCAM AND STX6	Breast	0.772277	0.829787	0.722222
EPCAM AND TSPAN5	Breast	0.719298	0.683333	0.759259
EPCAM AND PMP22	Breast	0.780952	0.803922	0.759259
EPCAM AND CD81	Breast	0.727273	0.714286	0.740741
EPCAM AND ITSN1	Breast	0.719298	0.683333	0.759259
EPCAM AND GJA5	Breast	0.711864	0.65625	0.777778
BAMBI AND EPCAM	Breast	0.641975	0.962963	0.481481
EPCAM AND TMEM119	Breast	0.651163	0.56	0.777778
EPCAM AND PCDHB16	Breast	0.650794	0.569444	0.759259
EPCAM AND PTPRS	Breast	0.666667	0.761905	0.592593
EPCAM AND ADAM8	Breast	0.638298	0.75	0.555556
EPCAM AND SLC16A6	Breast	0.772277	0.829787	0.722222
FGG AND MSLN	Liver	0.8	0.666667	1
HPN AND TNFRSF10A	Liver	0.666667	0.666667	0.666667
ASGR1 AND MSLN	Liver	0.75	0.6	1
FGG AND TNFRSF10A	Liver	0.666667	0.666667	0.666667
GJB1 AND MSLN	Liver	0.8	1	0.666667
ABCC6 AND MSLN	Liver	0.75	0.6	1
SLC27A5 AND MSLN	Liver	0.75	0.6	1
F10 AND TNFRSF10A	Liver	0.8	1	0.666667
ABCB4 AND MSLN	Liver	1	1	1
ABCB4 AND TNFRSF10A	Liver	0.8	1	0.666667
SLC27A5 AND TNFRSF10A	Liver	0.615385	0.571429	0.666667
ABCB11 AND MSLN	Liver	0.727273	0.8	0.666667
ABCC2 AND MSLN	Liver	1	1	1
ABCC2 AND TNFRSF10A	Liver	0.8	1	0.666667
F10 AND MSLN	Liver	1	1	1
APOB AND MSLN	Liver	0.833333	0.833333	0.833333
KLB AND MSLN	Liver	0.666667	0.666667	0.666667
APOB AND TNFRSF10A	Liver	0.6	0.75	0.5
KLB AND TNFRSF10A	Liver	0.6	0.75	0.5
SLC4A2 AND MSLN	Liver	0.666667	1	0.5
ABCC6 AND MUC1	Liver	0.666667	0.555556	0.833333
GJB1 AND MUC1	Liver	0.666667	1	0.5
TM4SF5 AND MSLN	Liver	0.8	1	0.666667
TM4SF5 AND TNFRSF10A	Liver	0.6	0.75	0.5
F10 AND MUC1	Liver	0.909091	1	0.833333

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
FOLH1 AND FAM57A	Prostate	0.857143	1	0.75
FOLH1 AND CDH3	Prostate	0.8	0.666667	1
FOLH1 AND DHRS3	Prostate	1	1	1
FOLH1 AND CD69	Prostate	0.888889	0.8	1
FOLH1 AND CTNS	Prostate	0.75	0.75	0.75
FOLH1 AND STYK1	Prostate	0.888889	0.8	1
DDR2 AND B4GALNT1	Sarcoma	0.648649	0.6	0.705882
CDH11 AND EPCAM	Breast	0.712644	0.939394	0.574074
PPAPDC1A AND EPCAM	Breast	0.711111	0.888889	0.592593
EPCAM AND BST2	Breast	0.817204	0.974359	0.703704
ENPP1 AND EPCAM	Breast	0.672566	0.644068	0.703704
EPCAM AND LAMP5	Breast	0.666667	0.794872	0.574074
EPCAM AND SLC38A1	Breast	0.647619	0.666667	0.62963
EPCAM AND IL17RA	Breast	0.666667	0.733333	0.611111
EPCAM AND LRRC8E	Breast	0.712871	0.765957	0.666667
EPCAM AND TGFBI	Breast	0.672	0.591549	0.777778
EPCAM AND SEMA4D	Breast	0.638655	0.584615	0.703704
EPCAM AND DYSF	Breast	0.618557	0.697674	0.555556
EPCAM AND LAPTM5	Breast	0.696629	0.885714	0.574074
EPCAM AND TTYH2	Breast	0.605042	0.553846	0.666667
PPAPDC1A AND ERBB2	Breast	0.60241	0.862069	0.462963
EPCAM AND TNFRSF12A	Breast	0.639175	0.72093	0.574074
EPCAM AND GPR137B	Breast	0.688172	0.820513	0.592593
EPCAM AND VANGL1	Breast	0.632479	0.587302	0.685185
EPCAM AND LAT	Breast	0.672414	0.629032	0.722222
EPCAM AND SLC6A6	Breast	0.643478	0.606557	0.685185
EPCAM AND FLVCR1	Breast	0.650407	0.57971	0.740741
EPCAM AND IFI6	Breast	0.638655	0.584615	0.703704
EPCAM AND CD163	Breast	0.792453	0.807692	0.777778
SLC2A2 AND MSLN	Liver	0.857143	0.75	1
SLC2A2 AND TNFRSF10A	Liver	0.727273	0.8	0.666667
SLCO1B1 AND MSLN	Liver	0.75	0.6	1
SLC17A2 AND MSLN	Liver	0.666667	0.555556	0.833333
ABCG8 AND MSLN	Liver	0.8	1	0.666667
SLC13A5 AND MSLN	Liver	0.666667	0.5	1
SLCO1B1 AND TNFRSF10A	Liver	0.615385	0.571429	0.666667
SLC17A2 AND TNFRSF10A	Liver	0.615385	0.571429	0.666667
ABCG8 AND TNFRSF10A	Liver	0.666667	1	0.5
SLC10A1 AND MSLN	Liver	0.666667	0.555556	0.833333
SLC43A1 AND TNFRSF10A	Liver	0.8	1	0.666667
ABCG5 AND MSLN	Liver	0.75	0.6	1
SLC30A8 AND CD19	Pancreas	0.705882	1	0.545455
PPAPDC1A AND MUC1	Pancreas	0.666667	0.857143	0.545455
KCND2 AND MUC1	Pancreas	0.736842	0.875	0.636364
PPAPDC1A AND ERBB2	Pancreas	0.666667	0.857143	0.545455
PPAPDC1A AND MSLN	Pancreas	0.705882	1	0.545455
CLDN10 AND MSLN	Pancreas	0.631579	0.75	0.545455
TNFRSF10B AND MUC1	Pancreas	0.631579	0.75	0.545455
MUC1 AND SYT11	Pancreas	0.615385	0.533333	0.727273
CD163 AND MUC1	Pancreas	0.666667	0.857143	0.545455
MSLN AND TGFBI	Pancreas	0.6	0.666667	0.545455
IFI6 AND MUC1	Colon	1	1	1
IFI6 AND ERBB2	Colon	0.909091	0.833333	1
SYT11 AND ERBB2	Glioma	0.964539	0.944444	0.985507
SYT11 AND MUC1	Glioma	0.94964	0.942857	0.956522
SYT11 AND EPHA2	Glioma	0.957143	0.943662	0.971014
GPR19 AND ERBB2	Glioma	0.816327	0.769231	0.869565
LYPD1 AND MUC1	Glioma	0.876923	0.934426	0.826087
GPR19 AND EPHA2	Glioma	0.802817	0.780822	0.826087
GPR19 AND MUC1	Glioma	0.805556	0.773333	0.84058
CSMD2 AND ERBB2	Glioma	0.691176	0.701493	0.681159
BEST3 AND MUC1	Glioma	0.682171	0.733333	0.637681
CSMD2 AND EPHA2	Glioma	0.666667	0.681818	0.652174
SLCO1C1 AND ERBB2	Glioma	0.653333	0.604938	0.710145
DSCAM AND ERBB2	Glioma	0.640777	0.970588	0.478261
ASTN1 AND ERBB2	Glioma	0.629834	0.508929	0.826087
SLCO1C1 AND MUC1	Glioma	0.613333	0.567901	0.666667
CDH10 AND ERBB2	Glioma	0.604396	0.486726	0.797101
FXYD6 AND ERBB2	Glioma	0.601156	0.5	0.753623
CD163 AND ERBB2	Glioma	0.666667	0.759259	0.594203
SLC22A16 AND CD33	AML	0.89505	0.889764	0.900398
CD33 AND ADAM29	AML	0.819085	0.81746	0.820717
CD33 AND SLC22A14	AML	0.834846	0.766667	0.916335
CD33 AND SLC43A1	AML	0.8	0.945652	0.693227
CD33 AND GJD2	AML	0.791252	0.789683	0.792829
CD33 AND SLCO1B3	AML	0.812834	0.735484	0.908367
CD33 AND P2RX4	AML	0.775862	0.683891	0.896414

FIG. 9 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
SDC2 AND MSLN	Liver	0.6	0.75	0.5
SLC12A7 AND MSLN	Liver	0.833333	0.833333	0.833333
KLB AND MUC1	Liver	0.666667	0.666667	0.666667
EVA1A AND TNFRSF10A	Liver	0.666667	1	0.5
NOX4 AND MSLN	Pancreas	0.705882	1	0.545455
TNFSF4 AND MUC1	Pancreas	0.777778	1	0.636364
MMP14 AND MUC1	Pancreas	0.705882	1	0.545455
VSIG1 AND MSLN	Pancreas	0.625	1	0.454545
TNFSF4 AND MSLN	Pancreas	0.631579	0.75	0.545455
ADAM12 AND MUC1	Pancreas	0.705882	1	0.545455
TNFSF13B AND MUC1	Pancreas	0.625	1	0.454545
MMP14 AND CD19	Pancreas	0.7	0.777778	0.636364
OSMR AND MUC1	Pancreas	0.777778	1	0.636364
RNF144A AND MSLN	Pancreas	0.631579	0.75	0.545455
ABCC8 AND MUC1	Pancreas	0.705882	1	0.545455
ABCC8 AND ERBB2	Pancreas	0.705882	1	0.545455
MUC1 AND CD84	Pancreas	0.625	1	0.454545
TNFSF4 AND ERBB2	Pancreas	0.636364	0.636364	0.636364
SLC4A7 AND MUC1	Pancreas	0.705882	1	0.545455
IFITM2 AND MSLN	Pancreas	0.631579	0.75	0.545455
MMP14 AND MSLN	Pancreas	0.705882	1	0.545455
TREM2 AND MUC1	Pancreas	0.625	1	0.454545
ABCC8 AND CD19	Pancreas	0.705882	1	0.545455
NALCN AND MUC1	Pancreas	0.631579	0.75	0.545455
ABCC8 AND MSLN	Pancreas	0.705882	1	0.545455
FLRT2 AND ERBB2	Pancreas	0.631579	0.75	0.545455
PTPN2 AND MUC1	Pancreas	0.8	0.888889	0.727273
SLC2A3 AND MUC1	Pancreas	0.631579	0.75	0.545455
SLC12A2 AND MUC1	Colon	0.75	1	0.6
GDE1 AND MSLN	Colon	0.75	1	0.6
SPINT1 AND MSLN	Colon	0.75	1	0.6
PMEPA1 AND MSLN	Colon	0.6	0.6	0.6
PMEPA1 AND MUC1	Colon	0.888889	1	0.8
TNFSF15 AND MUC1	Colon	0.6	0.6	0.6
PMEPA1 AND ERBB2	Colon	0.666667	0.571429	0.8
SLC39A10 AND MUC1	Colon	0.888889	1	0.8
SMAGP AND MSLN	Colon	0.75	1	0.6
NRCAM AND MUC1	Glioma	0.909091	0.952381	0.869565
NRCAM AND ERBB2	Glioma	0.917293	0.953125	0.884058
NLGN1 AND ERBB2	Glioma	0.938462	1	0.884058
SGCB AND MUC1	Glioma	0.859259	0.878788	0.84058
SCN3A AND ERBB2	Glioma	0.789116	0.74359	0.84058
SCN3A AND EPHA2	Glioma	0.808511	0.791667	0.826087
SCN3A AND MUC1	Glioma	0.820896	0.846154	0.797101
GRIK3 AND ERBB2	Glioma	0.754967	0.695122	0.826087
EPHB1 AND ERBB2	Glioma	0.75	0.882353	0.652174
FZD3 AND ERBB2	Glioma	0.736842	0.674699	0.811594
NLGN4X AND ERBB2	Glioma	0.73494	0.628866	0.884058
CPT1C AND ERBB2	Glioma	0.722892	0.618557	0.869565
CPT1C AND EPHA2	Glioma	0.714286	0.606061	0.869565
CPT1C AND MUC1	Glioma	0.693642	0.576923	0.869565
GPR34 AND MUC1	Glioma	0.787402	0.862069	0.724638
SLC39A10 AND ERBB2	Glioma	0.688889	0.558559	0.898551
RAMP1 AND MUC1	Glioma	0.683544	0.606742	0.782609
NCAM2 AND ERBB2	Glioma	0.675159	0.602273	0.768116
CNR1 AND ERBB2	Glioma	0.697368	0.638554	0.768116
NCAM2 AND EPHA2	Glioma	0.666667	0.588889	0.768116
SLC39A10 AND EPHA2	Glioma	0.666667	0.552381	0.84058
CD33 AND SLC2A5	AML	0.855319	0.917808	0.800797
P2RX1 AND CD33	AML	0.839858	0.758842	0.940239
MR1 AND MUC1	Lung Adenocarcinoma	0.60177	0.790698	0.485714
CXCL9 AND CD19	B-Cell Diffuse	0.774194	0.96	0.648649
CXCL9 AND MS4A1	B-Cell Diffuse	0.754098	0.958333	0.621622
CXCL9 AND CD22	B-Cell Diffuse	0.754098	0.958333	0.621622
MS4A1 AND MCOLN2	B-Cell Diffuse	0.709677	0.88	0.594595
MS4A1 AND GJA1	B-Cell Diffuse	0.705882	0.774194	0.648649
MS4A1 AND TNFRSF9	B-Cell Diffuse	0.705882	0.774194	0.648649
MS4A1 AND SLCO2B1	B-Cell Diffuse	0.705882	0.774194	0.648649
CD19 AND SLC2A5	B-Cell Diffuse	0.698413	0.846154	0.594595
MS4A1 AND SLC1A3	B-Cell Diffuse	0.69697	0.793103	0.621622
MS4A1 AND PPAP2B	B-Cell Diffuse	0.695652	0.75	0.648649
MS4A1 AND AQP7	B-Cell Diffuse	0.69697	0.793103	0.621622
CXCL10 AND CD22	B-Cell Diffuse	0.6875	0.814815	0.594595
MS4A1 AND PDGFRA	B-Cell Diffuse	0.686567	0.766667	0.621622
MS4A1 AND DDR2	B-Cell Diffuse	0.686567	0.766667	0.621622
MS4A1 AND ANTXR1	B-Cell Diffuse	0.686567	0.766667	0.621622
MS4A1 AND OSMR	B-Cell Diffuse	0.686567	0.766667	0.621622

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
CD33 AND KCND2	AML	0.720764	0.89881	0.601594
CD33 AND OTOF	AML	0.715294	0.873563	0.605578
CD33 AND SLC39A8	AML	0.710145	0.90184	0.585657
CD33 AND CSPG5	AML	0.709251	0.793103	0.641434
MLC1 AND CD33	AML	0.700405	0.711934	0.689243
CD33 AND SLC5A10	AML	0.693446	0.738739	0.653386
CD33 AND STAB1	AML	0.67803	0.646209	0.713147
CD33 AND KCNJ9	AML	0.664962	0.928571	0.517928
SLC22A16 AND IL3RA	AML	0.610966	0.886364	0.466135
CD33 AND SLC2A1	AML	0.608696	0.85	0.474104
CD33 AND SPAM1	AML	0.604396	0.973451	0.438247
MS4A1 AND KCNJ10	B-Cell Diffuse	0.69697	0.793103	0.621622
MS4A1 AND CDH11	B-Cell Diffuse	0.686567	0.766667	0.621622
CD19 AND KCNJ10	B-Cell Diffuse	0.656716	0.733333	0.594595
MS4A1 AND EFNB2	B-Cell Diffuse	0.656716	0.733333	0.594595
MS4A1 AND FAT1	B-Cell Diffuse	0.647059	0.709677	0.594595
MS4A1 AND PLVAP	B-Cell Diffuse	0.65625	0.777778	0.567568
CD19 AND CSPG5	B-Cell Diffuse	0.628571	0.666667	0.594595
CD19 AND CDH11	B-Cell Diffuse	0.628571	0.666667	0.594595
MS4A1 AND AOC3	B-Cell Diffuse	0.626866	0.7	0.567568
MS4A1 AND PROCR	B-Cell Diffuse	0.606061	0.689655	0.540541
CD19 AND ADCY10	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND JPH3	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND SLC22A18	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND SCN8A	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND HCN2	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND PCDHAC2	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND KCNA1	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND SLC22A16	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND MC2R	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND LCT	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND CLDN17	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND DCC	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND CLDN20	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND CATSPERD	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND PTGIR	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND ADAM20	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND CNTNAP4	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND MPL	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND MRGPRX2	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND GHSR	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND KCNC1	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND CCKBR	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND TMEM235	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND HTR3C	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND PRND	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND NTSR2	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND TAS1R1	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND GPR156	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND SLC30A8	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND GNRHR	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND DTNA	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND CDH22	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND OR3A2	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND CACNG1	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND SLC22A12	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND OR52D1	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND OR7C1	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND GPR50	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND FGF6	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND GPR158	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND DRD5	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND OR10J1	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND USH2A	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND MUC17	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND KCNV2	Mantle-Cell Lymphoma	0.898551	1	0.815789
CD19 AND EPHA10	Mantle-Cell Lymphoma	0.898551	1	0.815789
CD19 AND CALN1	Mantle-Cell Lymphoma	0.898551	1	0.815789
CD19 AND SLC4A5	Mantle-Cell Lymphoma	0.898551	1	0.815789
CD19 AND CLDN15	Mantle-Cell Lymphoma	0.898551	1	0.815789
CD19 AND TRPC7	Mantle-Cell Lymphoma	0.898551	1	0.815789
CD19 AND CDH18	Mantle-Cell Lymphoma	0.898551	1	0.815789
CD19 AND CRB1	Mantle-Cell Lymphoma	0.898551	1	0.815789
CD19 AND HTR6	Mantle-Cell Lymphoma	0.898551	1	0.815789
CD19 AND CACNG2	Mantle-Cell Lymphoma	0.898551	1	0.815789
CD19 AND STAB1	Mantle-Cell Lymphoma	0.898551	1	0.815789
CD19 AND SLC22A2	Mantle-Cell Lymphoma	0.898551	1	0.815789

FIG. 9 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
MS4A1 AND TMEM119	B-Cell Diffuse	0.686567	0.766667	0.621622
MS4A1 AND CDH5	B-Cell Diffuse	0.686567	0.766667	0.621622
MCOLN2 AND CD22	B-Cell Diffuse	0.760563	0.794118	0.72973
MS4A1 AND SLC2A5	B-Cell Diffuse	0.685714	0.727273	0.648649
CD80 AND CD22	B-Cell Diffuse	0.712329	0.722222	0.702703
CD19 AND MCOLN2	B-Cell Diffuse	0.677419	0.84	0.567568
CD19 AND BRCA1	B-Cell Diffuse	0.676923	0.785714	0.594595
MS4A1 AND KCNMA1	B-Cell Diffuse	0.676471	0.741935	0.621622
MS4A1 AND GJA4	B-Cell Diffuse	0.676923	0.785714	0.594595
MS4A1 AND MCAM	B-Cell Diffuse	0.666667	0.71875	0.621622
CD19 AND ATP2A1	B-Cell Diffuse	0.666667	0.758621	0.594595
MS4A1 AND UNC5B	B-Cell Diffuse	0.666667	0.71875	0.621622
CD22 AND SLC2A5	B-Cell Diffuse	0.698795	0.630435	0.783784
MS4A1 AND TM4SF1	B-Cell Diffuse	0.666667	0.758621	0.594595
MS4A1 AND TSPAN7	B-Cell Diffuse	0.666667	0.758621	0.594595
MS4A1 AND CSF1	B-Cell Diffuse	0.666667	0.758621	0.594595
MS4A1 AND TMEM47	B-Cell Diffuse	0.666667	0.758621	0.594595
MS4A1 AND AQP1	B-Cell Diffuse	0.666667	0.758621	0.594595
MS4A1 AND PMP22	B-Cell Diffuse	0.657534	0.666667	0.648649
MS4A1 AND ENPP2	B-Cell Diffuse	0.657534	0.666667	0.648649
MS4A1 AND SDC2	B-Cell Diffuse	0.657143	0.69697	0.621622
MS4A1 AND CAV1	B-Cell Diffuse	0.657143	0.69697	0.621622
MS4A1 AND FZD4	B-Cell Diffuse	0.666667	0.758621	0.594595
MS4A1 AND ECSCR	B-Cell Diffuse	0.647887	0.676471	0.621622
MS4A1 AND SLC25A4	B-Cell Diffuse	0.647059	0.709677	0.594595
CD19 AND TNFRSF9	B-Cell Diffuse	0.647059	0.709677	0.594595
MS4A1 AND CMKLR1	B-Cell Diffuse	0.647059	0.709677	0.594595
MS4A1 AND CD80	B-Cell Diffuse	0.646154	0.75	0.567568
CD19 AND SLC1A3	B-Cell Diffuse	0.646154	0.75	0.567568
CD19 AND NDRG4	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND DPP4	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND SLC13A3	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND P2RY2	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND KCNH5	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND LPAR3	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND SLC7A2	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND MARVELD2	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND LY6D	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND NOX4	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND TRHDE	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND MRGPRX4	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND STRA6	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND APOB	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND PCDHB11	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND GDE1	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND PDE6B	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND CLECL1	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND LRRC32	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND TMEM30B	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND ITGA2	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND RHBDL1	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND IL5RA	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND DCHS2	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND ADAM22	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND CORIN	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND TMEM150B	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND KNCN	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND TAS2R39	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND ANO1	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND CXADR	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND CCR8	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND CHRM3	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND FRAS1	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND EVA1A	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND CDH19	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND NPBWR2	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND OR1G1	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND PROKR2	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND SLC24A4	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND OR51B5	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND FCRL4	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND PCDHB12	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND VN1R1	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND SLC24A3	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND ADORA2B	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND OXGR1	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND AOC2	Mantle-Cell Lymphoma	0.914286	1	0.842105

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
CD19 AND SLC39A2	Mantle-Cell Lymphoma	0.898551	1	0.815789
CD19 AND CEACAM7	Mantle-Cell Lymphoma	0.898551	1	0.815789
CD19 AND TSHR	Mantle-Cell Lymphoma	0.898551	1	0.815789
CD19 AND SLC22A11	Mantle-Cell Lymphoma	0.898551	1	0.815789
CD19 AND PCDHGC4	Mantle-Cell Lymphoma	0.898551	1	0.815789
CD19 AND GRIN2B	Mantle-Cell Lymphoma	0.898551	1	0.815789
CD19 AND TRPM3	Mantle-Cell Lymphoma	0.898551	1	0.815789
CD19 AND ADAM7	Mantle-Cell Lymphoma	0.898551	1	0.815789
CD19 AND MRAP	Mantle-Cell Lymphoma	0.898551	1	0.815789
CD19 AND GRIN1	Mantle-Cell Lymphoma	0.898551	1	0.815789
CD19 AND GABRB2	Mantle-Cell Lymphoma	0.898551	1	0.815789
CD19 AND SLC17A1	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND PPAPDC1A	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND KCNJ9	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND CNGA4	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND OPALIN	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND GRM6	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND P2RX3	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND BEST2	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND DISP2	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND ZACN	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND GPR12	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND CABP7	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND SLC17A3	Mantle-Cell Lymphoma	0.898551	1	0.815789
CD19 AND CRB2	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND AJAP1	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND ZP4	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND OR5P3	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND KCNK10	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND EPHA8	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND HCN1	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND ATP6V0A4	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND SLC1A6	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND CHRNA1	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND SLC22A14	Mantle-Cell Lymphoma	0.914286	1	0.842105
MLANA AND GPR137B	Melanoma	1	1	1
GPR19 AND B4GALNT1	Neuroblastoma	0.624	0.46988	0.928571
FLVCR1 AND B4GALNT1	Neuroblastoma	0.643836	0.758065	0.559524
B4GALNT1 AND ATP7A	Neuroblastoma	0.675159	0.726027	0.630952
FOLH1 AND STEAP4	Prostate	0.857143	1	0.75
FOLH1 AND ANTXR2	Prostate	0.727273	0.571429	1
FOLH1 AND PTGIS	Prostate	0.8	0.666667	1
FOLH1 AND SLC43A1	Prostate	1	1	1
FOLH1 AND GGTLC1	Prostate	1	1	1
UPK3A AND FOLH1	Prostate	0.666667	0.6	0.75
FOLH1 AND GHR	Prostate	0.857143	1	0.75
FOLH1 AND ATP7A	Prostate	0.888889	0.8	1
FOLH1 AND GUCY2D	Prostate	0.666667	0.6	0.75
FOLH1 AND ATP8B2	Prostate	0.666667	0.5	1
FOLH1 AND FAT1	Prostate	0.857143	1	0.75
FOLH1 AND GABRG3	Prostate	0.666667	0.6	0.75
FOLH1 AND ATP8B1	Prostate	0.615385	0.444444	1
FOLH1 AND EMP3	Prostate	0.727273	0.571429	1
FOLH1 AND FLVCR1	Prostate	0.727273	0.571429	1
FOLH1 AND VANGL1	Prostate	0.727273	0.571429	1
FOLH1 AND MRGPRF	Prostate	0.6	0.5	0.75
FOLH1 AND SEMA4B	Prostate	0.727273	0.571429	1
FOLH1 AND PPAPDC1B	Prostate	0.615385	0.444444	1
FOLH1 AND CD93	Prostate	0.857143	1	0.75
FOLH1 AND SLC1A5	Prostate	0.615385	0.444444	1
FOLH1 AND PAQR7	Prostate	0.857143	1	0.75
FOLH1 AND ENPP1	Prostate	0.857143	1	0.75
FOLH1 AND SLC22A5	Prostate	0.857143	1	0.75
FOLH1 AND ZACN	Prostate	0.666667	1	0.5
FOLH1 AND TTYH3	Prostate	0.666667	1	0.5
FOLH1 AND CD44	Prostate	0.615385	0.444444	1
FOLH1 AND MPL	Prostate	0.8	0.666667	1
FOLH1 AND CD163	Prostate	1	1	1
FOLH1 AND F2R	Prostate	0.888889	0.8	1
FOLH1 AND AOC3	Prostate	0.727273	0.571429	1
FOLH1 AND PHLDB2	Prostate	0.615385	0.444444	1
FOLH1 AND ANPEP	Prostate	0.727273	0.571429	1
FOLH1 AND DGKE	Prostate	0.857143	1	0.75
FOLH1 AND KCNK6	Prostate	0.615385	0.444444	1
FOLH1 AND BTN3A1	Prostate	0.8	0.666667	1
FOLH1 AND PTGER4	Prostate	0.857143	1	0.75
FAP AND EPCAM	Breast	0.875	1	0.777778

FIG. 9 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
CD19 AND PCDH10	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND SLC26A7	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND HTR1B	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND OR51E1	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND HTR2A	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND IL1RL1	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND OR7A5	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND CHRM5	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND PKHD1	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND AQP4	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND EPHB1	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND EFNB3	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND TIE1	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND SI	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND KCNK1	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND AGTR1	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND P2RX2	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND OR51M1	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND UPK1B	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND SFRP1	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND SLC5A9	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND ATP4A	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND ABCC8	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND MAS1	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND CDH6	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND KCNG3	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND EMCN	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND LRP4	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND OPN5	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND KIR2DL4	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND KCNE1	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND EDNRA	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND TSPAN1	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND SLC8A1	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND EPHA1	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND FLRT3	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND ERVFRD-1	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND KIR2DL3	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND SLC16A1	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND NOX3	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND SLC26A10	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND OR1J4	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND SLC12A4	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND MFSD2A	Mantle-Cell Lymphoma	0.914286	1	0.842105
MLANA AND KCNK15	Melanoma	1	1	1
MLANA AND SLC20A1	Melanoma	0.952381	1	0.909091
SLC6A2 AND B4GALNT1	Neuroblastoma	0.810127	0.864865	0.761905
SCN9A AND B4GALNT1	Neuroblastoma	0.766467	0.771084	0.761905
THSD7A AND B4GALNT1	Neuroblastoma	0.695187	0.631068	0.77381
SLC29A4 AND B4GALNT1	Neuroblastoma	0.69281	0.768116	0.630952
TUSC3 AND B4GALNT1	Neuroblastoma	0.69281	0.768116	0.630952
XPR1 AND B4GALNT1	Neuroblastoma	0.6625	0.697368	0.630952
CADM1 AND B4GALNT1	Neuroblastoma	0.641791	0.86	0.511905
HTRA2 AND B4GALNT1	Neuroblastoma	0.626866	0.84	0.5
SLC26A11 AND B4GALNT1	Neuroblastoma	0.609375	0.886364	0.464286
B4GALNT1 AND PEMT	Neuroblastoma	0.666667	0.824561	0.559524
B4GALNT1 AND YIPF3	Neuroblastoma	0.657895	0.735294	0.595238
B4GALNT1 AND SLC38A2	Neuroblastoma	0.679739	0.753623	0.619048
LDLRAD3 AND B4GALNT1	Neuroblastoma	0.630872	0.723077	0.559524
B4GALNT1 AND VIMP	Neuroblastoma	0.601504	0.816327	0.47619
B4GALNT1 AND JTB	Neuroblastoma	0.607595	0.648649	0.571429
DCHS1 AND B4GALNT1	Neuroblastoma	0.714286	0.892857	0.595238
FOLH1 AND CACNA1D	Prostate	0.857143	1	0.75
FOLH1 AND BCAM	Prostate	0.727273	0.571429	1
OR51E1 AND FOLH1	Prostate	0.857143	1	0.75
FOLH1 AND ADRB2	Prostate	1	1	1
TSPAN1 AND FOLH1	Prostate	0.666667	1	0.5
FOLH1 AND PCDHB14	Prostate	0.75	0.75	0.75
FOLH1 AND VIPR1	Prostate	0.857143	1	0.75
FOLH1 AND SERINC5	Prostate	0.666667	1	0.5
FOLH1 AND GOLM1	Prostate	1	1	1
FOLH1 AND EDNRA	Prostate	0.727273	0.571429	1
FOLH1 AND SELE	Prostate	0.666667	0.5	1
FOLH1 AND HPN	Prostate	1	1	1
FOLH1 AND CD320	Prostate	0.727273	0.571429	1
FOLH1 AND COLEC12	Prostate	0.8	0.666667	1
FOLH1 AND ATP2C1	Prostate	0.666667	0.6	0.75

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
TPBG AND EPCAM	Breast	0.712644	0.939394	0.574074
EPCAM AND SLC7A5	Breast	0.744681	0.875	0.648148
FAP AND MSLN	Breast	0.762887	0.860465	0.685185
FAP AND ERBB2	Breast	0.682927	1	0.518519
EPCAM AND KDR	Breast	0.66129	0.585714	0.759259
EPCAM AND SDC1	Breast	0.764045	0.971429	0.62963
EPCAM AND AXL	Breast	0.701754	0.666667	0.740741
EPCAM AND GPNMB	Breast	0.8	0.823529	0.777778
EPCAM AND TNC	Breast	0.632479	0.587302	0.685185
TPBG AND MSLN	Breast	0.606061	0.666667	0.555556
EPCAM AND PCYT1A	Breast	0.740741	0.740741	0.740741
EPCAM AND ITGAV	Breast	0.65625	0.567568	0.777778
EPCAM AND CBX3	Breast	0.722689	0.661538	0.796296
FAP AND MUC1	Pancreas	0.777778	1	0.636364
FAP AND CD19	Pancreas	0.705882	1	0.545455
FAP AND MSLN	Pancreas	0.705882	1	0.545455
CLDN2 AND MSLN	Pancreas	0.615385	0.533333	0.727273
FAP AND ERBB2	Pancreas	0.705882	1	0.545455
VCAM1 AND MUC1	Pancreas	0.666667	0.7	0.636364
TNFSF11 AND MUC1	Pancreas	0.608696	0.583333	0.636364
VCAM1 AND ERBB2	Pancreas	0.692308	0.6	0.818182
MUC1 AND ITGAV	Pancreas	0.727273	0.727273	0.727273
PTK7 AND MSLN	Pancreas	0.625	1	0.454545
STEAP2 AND MUC1	Pancreas	0.642857	0.529412	0.818182
CBX3 AND MUC1	Colon	0.666667	0.571429	0.8
CBX3 AND ERBB2	Colon	0.666667	0.571429	0.8
BIRC5 AND MUC1	Colon	0.666667	0.5	1
ITGAV AND MUC1	Colon	0.909091	0.833333	1
CD33 AND CD70	AML	0.872659	0.823322	0.928287
CD33 AND SLC7A5	AML	0.857678	0.809187	0.912351
CD33 AND PROM1	AML	0.757576	0.829384	0.697211
CD33 AND CD38	AML	0.756661	0.682692	0.848606
CD33 AND TPBG	AML	0.64486	0.779661	0.549801
CD33 AND CD34	AML	0.631068	0.807453	0.517928
CD22 AND GPNMB	B-Cell Diffuse	0.8125	0.962963	0.702703
MS4A1 AND GPNMB	B-Cell Diffuse	0.716418	0.8	0.648649
MS4A1 AND VCAM1	B-Cell Diffuse	0.705882	0.774194	0.648649
CD19 AND GPNMB	B-Cell Diffuse	0.689655	0.952381	0.540541
MS4A1 AND SLC7A5	B-Cell Diffuse	0.686567	0.766667	0.621622
MS4A1 AND TPBG	B-Cell Diffuse	0.676471	0.741935	0.621622
MS4A1 AND EDNRB	B-Cell Diffuse	0.657143	0.69697	0.621622
CD22 AND VCAM1	B-Cell Diffuse	0.637363	0.537037	0.783784
CD19 AND VCAM1	B-Cell Diffuse	0.628571	0.666667	0.594595
MS4A1 AND CLDN12	B-Cell Diffuse	0.621622	0.621622	0.621622
CD19 AND TPBG	B-Cell Diffuse	0.619718	0.647059	0.594595
CD19 AND IL2RA	B-Cell Diffuse	0.60274	0.611111	0.594595
CD19 AND STEAP2	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND ERBB2	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND SDC1	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND ERBB3	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND SSTR1	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND CD34	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND MUC16	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND CLDN2	Mantle-Cell Lymphoma	0.898551	1	0.815789
CD19 AND NCAM1	Mantle-Cell Lymphoma	0.898551	1	0.815789
CD19 AND ITGB6	Mantle-Cell Lymphoma	0.898551	1	0.815789
CD19 AND EDNRB	Mantle-Cell Lymphoma	0.898551	1	0.815789
CD19 AND GUCY2C	Mantle-Cell Lymphoma	0.898551	1	0.815789
CD19 AND ENPP3	Mantle-Cell Lymphoma	0.898551	1	0.815789
CD19 AND LGR5	Mantle-Cell Lymphoma	0.898551	1	0.815789
CD19 AND ERBB4	Mantle-Cell Lymphoma	0.898551	1	0.815789
CD19 AND B4GALNT1	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND FOLH1	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND TNFRSF8	Mantle-Cell Lymphoma	0.898551	1	0.815789
CD19 AND SLC39A6	Mantle-Cell Lymphoma	0.882353	1	0.789474
CD19 AND TNFRSF10A	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND PMEL	Mantle-Cell Lymphoma	0.914286	1	0.842105
CD19 AND FOLR1	Mantle-Cell Lymphoma	0.882353	1	0.789474
CD19 AND CLDN12	Mantle-Cell Lymphoma	0.865672	1	0.763158
CD19 AND ABCB5	Mantle-Cell Lymphoma	0.865672	1	0.763158
CD19 AND ITGB3	Mantle-Cell Lymphoma	0.865672	1	0.763158
CD19 AND CD52	Mantle-Cell Lymphoma	0.898551	1	0.815789
CD19 AND IL11RA	Mantle-Cell Lymphoma	0.898551	1	0.815789
CD19 AND SSTR5	Mantle-Cell Lymphoma	0.830769	1	0.710526
CD19 AND CLDN18	Mantle-Cell Lymphoma	0.830769	1	0.710526
CD19 AND RNF43	Mantle-Cell Lymphoma	0.8125	1	0.684211
CD19 AND CA9	Mantle-Cell Lymphoma	0.793651	1	0.657895

FIG. 9 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
ABCC4 AND FOLH1	Prostate	0.857143	1	0.75
FOLH1 AND CDH26	Prostate	0.727273	0.571429	1
FOLH1 AND CD99	Prostate	0.666667	0.6	0.75
FOLH1 AND SLC2A10	Prostate	0.666667	1	0.5
FOLH1 AND EPHA7	Prostate	0.666667	0.5	1
FOLH1 AND AGTRAP	Prostate	0.727273	0.571429	1
FOLH1 AND IL1R1	Prostate	0.888889	0.8	1
FOLH1 AND ITGA5	Prostate	0.888889	0.8	1
FOLH1 AND SDK1	Prostate	0.727273	0.571429	1
FOLH1 AND POPDC2	Prostate	0.888889	0.8	1
FOLH1 AND BBS4	Prostate	0.727273	0.571429	1
FOLH1 AND PPAP2A	Prostate	0.6	0.5	0.75
FOLH1 AND KCNMB1	Prostate	0.666667	0.5	1
FOLH1 AND SLC26A4	Prostate	0.857143	1	0.75
NDRG1 AND FOLH1	Prostate	0.666667	1	0.5
FOLH1 AND KIAA0319L	Prostate	0.75	0.75	0.75
FOLH1 AND TNFSF4	Prostate	0.6	0.5	0.75
SLC2A12 AND FOLH1	Prostate	0.857143	1	0.75
FOLH1 AND ANKH	Prostate	0.6	0.5	0.75
FOLH1 AND OSMR	Prostate	0.857143	1	0.75
FOLH1 AND TLCD1	Prostate	1	1	1
FOLH1 AND TUSC3	Prostate	0.727273	0.571429	1
FOLH1 AND SLC10A3	Prostate	0.857143	1	0.75
FOLH1 AND SLC26A6	Prostate	0.75	0.75	0.75
FOLH1 AND FZD4	Prostate	1	1	1
FOLH1 AND SLC7A2	Prostate	0.888889	0.8	1
FOLH1 AND FREM2	Prostate	0.888889	0.8	1
FOLH1 AND MARVELD1	Prostate	0.8	0.666667	1
FOLH1 AND VMP1	Prostate	1	1	1
FOLH1 AND NPR3	Prostate	0.75	0.75	0.75
FOLH1 AND FZD10	Prostate	0.727273	0.571429	1
FOLH1 AND FUT1	Prostate	0.75	0.75	0.75
FOLH1 AND GPR183	Prostate	0.666667	1	0.5
FOLH1 AND C5AR1	Prostate	0.75	0.75	0.75
FOLH1 AND KCNG3	Prostate	0.666667	1	0.5
FOLH1 AND TSPAN2	Prostate	0.727273	0.571429	1
FOLH1 AND CYB5R1	Prostate	0.8	0.666667	1
FOLH1 AND KCNS3	Prostate	0.888889	0.8	1
FOLH1 AND TPM1	Prostate	1	1	1
FOLH1 AND TMPRSS2	Prostate	0.727273	0.571429	1
FOLH1 AND ITGB5	Prostate	0.857143	1	0.75
FOLH1 AND CATSPER2	Prostate	0.857143	1	0.75
FOLH1 AND TRPV1	Prostate	0.857143	1	0.75
FOLH1 AND NOX4	Prostate	1	1	1
FOLH1 AND MYADM	Prostate	0.857143	1	0.75
FOLH1 AND OPN1SW	Prostate	0.857143	1	0.75
FOLH1 AND VANGL2	Prostate	0.666667	0.6	0.75
FOLH1 AND LMAN2	Prostate	0.857143	1	0.75
FOLH1 AND EPHB4	Prostate	0.8	0.666667	1

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
CD19 AND SLC7A5	Mantle-Cell Lymphoma	0.793651	1	0.657895
CD19 AND ROR1	Mantle-Cell Lymphoma	0.793651	1	0.657895
CD19 AND CSPG4	Mantle-Cell Lymphoma	0.793651	1	0.657895
CD19 AND IL2RA	Mantle-Cell Lymphoma	0.793651	1	0.657895
CD19 AND EPCAM	Mantle-Cell Lymphoma	0.774194	1	0.631579
FCRL2 AND CD19	Mantle-Cell Lymphoma	0.754098	1	0.605263
CD19 AND CLDN9	Mantle-Cell Lymphoma	0.754098	1	0.605263
CD19 AND SLC34A2	Mantle-Cell Lymphoma	0.733333	1	0.578947
CD19 AND VTCN1	Mantle-Cell Lymphoma	0.733333	1	0.578947
CD19 AND GPA33	Mantle-Cell Lymphoma	0.711864	1	0.552632
CD19 AND MUC4	Mantle-Cell Lymphoma	0.711864	1	0.552632
CD19 AND STEAP1	Mantle-Cell Lymphoma	0.689655	1	0.526316
CD19 AND CLDN23	Mantle-Cell Lymphoma	0.666667	1	0.5
CD19 AND CLDN7	Mantle-Cell Lymphoma	0.642857	1	0.473684
CD19 AND IL3RA	Mantle-Cell Lymphoma	0.642857	1	0.473684
CD19 AND PROM1	Mantle-Cell Lymphoma	0.618182	1	0.447368
CD19 AND L1CAM	Mantle-Cell Lymphoma	0.618182	1	0.447368
MLANA AND TRPM4	Melanoma	0.952381	1	0.909091
MLANA AND IL11RA	Melanoma	1	1	1
MLANA AND L1CAM	Melanoma	0.952381	1	0.909091
EDNRB AND PMEL	Melanoma	0.9	1	0.818182
EDNRB AND MLANA	Melanoma	0.9	1	0.818182
PMEL AND CD22	Melanoma	0.9	1	0.818182
MLANA AND MET	Melanoma	0.842105	1	0.727273
MLANA AND IL13RA1	Melanoma	0.842105	1	0.727273
PMEL AND ITGAV	Melanoma	0.842105	1	0.727273
CBX3 AND B4GALNT1	Neuroblastoma	0.625954	0.87234	0.488095
B4GALNT1 AND CD276	Neuroblastoma	0.629371	0.762712	0.535714
STEAP2 AND FOLH1	Prostate	0.727273	0.571429	1
STEAP1 AND FOLH1	Prostate	0.75	0.75	0.75
FOLH1 AND EPHA3	Prostate	0.8	0.666667	1
FOLH1 AND GPNMB	Prostate	0.666667	1	0.5
FOLH1 AND FLOT2	Prostate	0.666667	0.5	1
FOLH1 AND ROR1	Prostate	0.888889	0.8	1
FOLH1 AND TNC	Prostate	0.888889	0.8	1
FOLH1 AND EPCAM	Prostate	0.857143	1	0.75
FOLH1 AND CLDN8	Prostate	0.8	0.666667	1
FOLH1 AND TPBG	Prostate	0.6	0.5	0.75
FOLH1 AND CLDN12	Prostate	0.666667	1	0.5
FOLH1 AND HSPA5	Prostate	0.666667	1	0.5
FOLH1 AND CD38	Prostate	0.727273	0.571429	1
FOLH1 AND IL20RA	Prostate	0.8	0.666667	1
FOLH1 AND DKK1	Prostate	0.727273	0.571429	1
FOLH1 AND PTK7	Prostate	0.615385	0.444444	1
FOLH1 AND MOK	Prostate	0.666667	1	0.5
FOLH1 AND ERBB2	Prostate	0.615385	0.444444	1
FOLH1 AND CBX3	Prostate	0.857143	1	0.75
FOLH1 AND MUC1	Prostate	0.888889	0.8	1
FOLH1 AND CD52	Prostate	0.75	0.75	0.75
FOLH1 AND ABCA5	Prostate	0.666667	1	0.5

FIG. 10

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
EPCAM AND NOT-SLC4A4	Breast	0.84848	0.9333	0.7778
ADAM12 AND NOT-MET	Breast	0.77358	0.7885	0.7593
EPCAM AND NOT-VSIG2	Breast	0.71795	0.6667	0.7778
EPCAM AND NOT-MRAP2	Breast	0.71429	0.7955	0.6481
EPCAM AND NOT-F3	Breast	0.71074	0.6418	0.7963
NOX4 AND NOT-ROR1	Breast	0.73171	0.6522	0.8333
NOX4 AND NOT-TYR	Breast	0.69173	0.5823	0.8519
EPCAM AND NOT-ABHD6	Breast	0.68908	0.6308	0.7593
PMEPA1 AND NOT-ROR1	Breast	0.68627	0.7292	0.6481
EPCAM AND NOT-SCNN1B	Breast	0.68333	0.6212	0.7593
SLC52A2 AND NOT-TYR	Breast	0.74725	0.9189	0.6296
EPCAM AND NOT-SLC16A9	Breast	0.68627	0.7292	0.6481
EPCAM AND NOT-CLCA4	Breast	0.67742	0.6	0.7778
F2RL2 AND NOT-TYR	Breast	0.67442	0.9063	0.537
PRLR AND NOT-MET	Breast	0.67327	0.7234	0.6296
EPCAM AND NOT-TSPAN8	Breast	0.67308	0.7	0.6481
PRLR AND NOT-TYR	Breast	0.67308	0.7	0.6481
EPCAM AND NOT-KCNQ1	Breast	0.71186	0.6563	0.7778
EPCAM AND NOT-KCNE3	Breast	0.672	0.5915	0.7778
F2RL2 AND NOT-ROR1	Breast	0.67442	0.9063	0.537
EPCAM AND NOT-CA4	Breast	0.76364	0.75	0.7778
EPCAM AND NOT-ATP10B	Breast	0.68421	0.65	0.7222
EPCAM AND NOT-PVRL3	Breast	0.66667	0.6491	0.6852
EPCAM AND NOT-BEST4	Breast	0.66142	0.5753	0.7778
EPCAM AND NOT-PIGR	Breast	0.66116	0.597	0.7407
EPCAM AND NOT-BMP2	Breast	0.66116	0.597	0.7407
EPCAM AND NOT-CYP4F12	Breast	0.71186	0.6563	0.7778
EPCAM AND NOT-TRHDE	Breast	0.66071	0.6379	0.6852
PMEPA1 AND NOT-TYR	Breast	0.66055	0.6545	0.6667
EPCAM AND NOT-UPK1B	Breast	0.66129	0.5857	0.7593
SLC4A2 AND NOT-TYR	Breast	0.65217	0.7895	0.5556
EPCAM AND NOT-SMPD3	Breast	0.65152	0.5513	0.7963
EPCAM AND NOT-DUOX2	Breast	0.7	0.6364	0.7778
EPCAM AND NOT-IL22RA1	Breast	0.64662	0.5443	0.7963
EPCAM AND NOT-IL1R2	Breast	0.64407	0.5938	0.7037
EPCAM AND NOT-SLC23A1	Breast	0.64407	0.5938	0.7037
EPCAM AND NOT-TSPAN7	Breast	0.66071	0.6379	0.6852
EPCAM AND NOT-ADORA2B	Breast	0.64122	0.5455	0.7778
EPCAM AND NOT-TMPRSS2	Breast	0.64078	0.6735	0.6111
MFSD10 AND NOT-MET	Breast	0.64078	0.6735	0.6111
EPCAM AND NOT-SLC17A4	Breast	0.65625	0.5676	0.7778
EPCAM AND NOT-GJB1	Breast	0.64078	0.6735	0.6111
EPCAM AND NOT-SLC3A1	Breast	0.64348	0.6066	0.6852
SLC4A2 AND NOT-ROR1	Breast	0.63736	0.7838	0.537
EPCAM AND NOT-TSPAN1	Breast	0.63704	0.5309	0.7963
SLC5A6 AND NOT-TYR	Breast	0.63636	0.625	0.6481
EPCAM AND NOT-RHBDL2	Breast	0.65649	0.5584	0.7963
EPCAM AND NOT-SLC5A1	Breast	0.65487	0.6271	0.6852
FGFR4 AND NOT-MUC1	Liver	0.8	1	0.6667
FGFR4 AND NOT-MSLN	Liver	0.8	1	0.6667
FGFR4 AND NOT-TNFRSF10A	Liver	0.8	1	0.6667
SLC4A2 AND NOT-MUC1	Liver	0.8	1	0.6667
LRP5 AND NOT-MUC1	Liver	0.83333	0.8333	0.8333
CLPTM1 AND NOT-MUC1	Liver	0.90909	1	0.8333
SEMA4G AND NOT-MUC1	Liver	0.625	0.5	0.8333
TMPRSS6 AND NOT-MUC1	Liver	0.66667	1	0.5
TMPRSS6 AND NOT-MSLN	Liver	0.66667	1	0.5
ITPR2 AND NOT-MUC1	Liver	0.83333	0.8333	0.8333
ABCB11 AND NOT-MUC1	Liver	0.8	1	0.6667
ABCC2 AND NOT-MUC1	Liver	1	1	1
TMPRSS6 AND NOT-TNFRSF10A	Liver	0.66667	1	0.5
APOB AND NOT-MUC1	Liver	0.83333	0.8333	0.8333
SLC4A2 AND NOT-TNFRSF10A	Liver	0.8	1	0.6667
SLC5A9 AND NOT-MUC1	Liver	0.72727	0.8	0.6667
RHBG AND NOT-MUC1	Liver	0.66667	1	0.5
LSR AND NOT-MUC1	Liver	0.90909	1	0.8333
SMO AND NOT-MUC1	Liver	0.66667	1	0.5
TM4SF5 AND NOT-MUC1	Liver	0.8	1	0.6667
CLPTM1 AND NOT-TNFRSF10A	Liver	0.90909	1	0.8333
TSPAN6 AND NOT-MUC1	Liver	0.90909	1	0.8333
LRP5 AND NOT-TNFRSF10A	Liver	0.83333	0.8333	0.8333
SLC16A11 AND NOT-MSLN	Liver	0.66667	1	0.5
SLC16A11 AND NOT-MUC1	Liver	0.66667	1	0.5
TMUB1 AND NOT-TNFRSF10A	Liver	0.8	1	0.6667
SLC16A11 AND NOT-TNFRSF10A	Liver	0.66667	1	0.5
RHBG AND NOT-MSLN	Liver	0.66667	1	0.5
ITPR2 AND NOT-TNFRSF10A	Liver	0.8	1	0.6667

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
EPCAM AND NOT-SLC13A2	Breast	0.6179	0.5507	0.7037
EPCAM AND NOT-CLDN16	Breast	0.6074	0.5062	0.7593
EPCAM AND NOT-PCDHA6	Breast	0.688	0.6056	0.7963
EPCAM AND NOT-C8B	Breast	0.6056	0.4886	0.7963
ENPP1 AND NOT-ROR1	Breast	0.6066	0.5441	0.6852
EPCAM AND NOT-SLC6A13	Breast	0.6167	0.5606	0.6852
EPCAM AND NOT-KCNK10	Breast	0.6176	0.5122	0.7778
EPCAM AND NOT-CHRNA9	Breast	0.6061	0.5128	0.7407
EPCAM AND NOT-SLC22A18	Breast	0.6	0.4884	0.7778
EPCAM AND NOT-HTR6	Breast	0.6043	0.4941	0.7778
EPCAM AND NOT-SLCO1C1	Breast	0.6116	0.5522	0.6852
EPCAM AND NOT-GABRA2	Breast	0.6074	0.5062	0.7593
EPCAM AND NOT-SYT4	Breast	0.7167	0.6515	0.7963
EPCAM AND NOT-ANPEP	Breast	0.6232	0.5119	0.7963
EPCAM AND NOT-HCN4	Breast	0.6043	0.4941	0.7778
TNFRSF12A AND NOT-ROR1	Breast	0.6271	0.5781	0.6852
EPCAM AND NOT-SLC28A1	Breast	0.6232	0.5119	0.7963
EPCAM AND NOT-OR1Q1	Breast	0.6087	0.5	0.7778
EPCAM AND NOT-TAS2R1	Breast	0.6232	0.5119	0.7963
EPCAM AND NOT-ADAM20	Breast	0.6299	0.5479	0.7407
ALCAM AND NOT-MUC1	Liver	0.7143	0.625	0.8333
SLC30A10 AND NOT-MUC1	Liver	0.6667	1	0.5
ABCG8 AND NOT-MUC1	Liver	0.8	1	0.6667
CLDN15 AND NOT-MUC1	Liver	0.8333	0.8333	0.8333
SLC22A18 AND NOT-MUC1	Liver	1	1	1
SLC43A1 AND NOT-MUC1	Liver	1	1	1
ALCAM AND NOT-MSLN	Liver	0.7143	0.625	0.8333
DIO1 AND NOT-MUC1	Liver	0.6	0.4286	1
ABCG5 AND NOT-MUC1	Liver	0.75	0.6	1
CLDN15 AND NOT-TNFRSF10A	Liver	0.6667	0.5556	0.8333
ENPP1 AND NOT-MUC1	Liver	0.6154	0.5714	0.6667
IFI6 AND NOT-GPC3	Colon	0.8333	0.7143	1
MUC1 AND NOT-AGTR2	Colon	0.75	1	0.6
PROCR AND NOT-GPC3	Colon	0.75	1	0.6
MUC1 AND NOT-EPHA10	Colon	0.75	1	0.6
MUC1 AND NOT-KIRREL3	Colon	0.75	1	0.6
MUC1 AND NOT-GUCY2D	Colon	0.75	1	0.6
MUC1 AND NOT-ATP1B4	Colon	0.75	1	0.6
MUC1 AND NOT-ROS1	Colon	0.75	1	0.6
MUC1 AND NOT-GABRA4	Colon	0.75	1	0.6
ERBB2 AND NOT-OPCML	Colon	0.75	1	0.6
ERBB2 AND NOT-GALR2	Colon	0.75	1	0.6
MUC1 AND NOT-JPH3	Colon	0.75	1	0.6
ERBB2 AND NOT-GPR78	Colon	0.75	1	0.6
ERBB2 AND NOT-MPL	Colon	0.75	1	0.6
ERBB2 AND NOT-ADCY8	Colon	0.75	1	0.6
ERBB2 AND NOT-CLDN16	Colon	0.75	1	0.6
MUC1 AND NOT-GRM1	Colon	0.75	1	0.6
MUC1 AND NOT-HTR5A	Colon	0.75	1	0.6
MUC1 AND NOT-SV2C	Colon	0.75	1	0.6
MUC1 AND NOT-KCNA10	Colon	0.75	1	0.6
MUC1 AND NOT-PTPRT	Colon	0.75	1	0.6
ERBB2 AND NOT-HTR1D	Colon	0.75	1	0.6
MUC1 AND NOT-LYPD1	Colon	0.75	1	0.6
ERBB2 AND NOT-OPRM1	Colon	0.75	1	0.6
MUC1 AND NOT-DCSTAMP	Colon	0.6	0.6	0.6
MUC1 AND NOT-NPHS1	Colon	0.6667	0.75	0.6
MUC1 AND NOT-ADAM30	Colon	0.6667	0.75	0.6
MUC1 AND NOT-NMBR	Colon	0.75	1	0.6
MUC1 AND NOT-SCN8A	Colon	0.75	1	0.6
MUC1 AND NOT-SLC30A8	Colon	0.75	1	0.6
MUC1 AND NOT-OPRM1	Colon	0.75	1	0.6
MUC1 AND NOT-CLRN1	Colon	0.75	1	0.6
ERBB2 AND NOT-CRB2	Colon	0.75	1	0.6
MUC1 AND NOT-GABRD	Colon	0.75	1	0.6
ERBB2 AND NOT-ABCG5	Colon	0.75	1	0.6
LYPD1 AND NOT-EPHA2	Glioma	0.8769	0.9344	0.8261
CDH11 AND NOT-EPHA2	Glioma	0.7287	0.7833	0.6812
CDH11 AND NOT-MUC1	Glioma	0.7361	0.7067	0.7681
CSMD2 AND NOT-MUC1	Glioma	0.6765	0.6866	0.6667
CDH11 AND NOT-ERBB2	Glioma	0.7143	0.7042	0.7246
OXTR AND NOT-MUC1	Glioma	0.6475	0.6429	0.6522
OXTR AND NOT-EPHA2	Glioma	0.6232	0.6232	0.6232
F2R AND NOT-EPHA2	Glioma	0.6196	0.4957	0.8261
OXTR AND NOT-ERBB2	Glioma	0.662	0.6438	0.6812
NPFFR1 AND NOT-EPHA2	Glioma	0.6923	0.6207	0.7826
NPFFR1 AND NOT-MUC1	Glioma	0.6512	0.5437	0.8116

FIG. 10 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
CDHR5 AND NOT-MUC1	Liver	0.92308	0.8571	1
CACFD1 AND NOT-MUC1	Liver	0.72727	0.8	0.6667
FZD5 AND NOT-MUC1	Liver	0.90909	1	0.8333
ABCC3 AND NOT-MUC1	Liver	0.76923	0.7143	0.8333
RHBG AND NOT-TNFRSF10A	Liver	0.66667	1	0.5
CLPTM1 AND NOT-MSLN	Liver	0.90909	1	0.8333
PMEPA1 AND NOT-GPC3	Colon	0.88889	1	0.8
SLC12A2 AND NOT-GPC3	Colon	0.75	1	0.6
SLC12A2 AND NOT-ERBB2	Colon	0.75	1	0.6
LY6G6D AND NOT-GPC3	Colon	0.75	1	0.6
MMP14 AND NOT-GPC3	Colon	0.75	1	0.6
CRB3 AND NOT-MUC1	Colon	0.66667	0.75	0.6
TNFSF15 AND NOT-GPC3	Colon	0.75	1	0.6
EPHB4 AND NOT-GPC3	Colon	0.8	0.8	0.8
SLC2A8 AND NOT-GPC3	Colon	0.75	1	0.6
LY6G6D AND NOT-ERBB2	Colon	0.75	1	0.6
KCNN4 AND NOT-GPC3	Colon	0.72727	0.6667	0.8
SIGMAR1 AND NOT-GPC3	Colon	0.8	0.8	0.8
MUC1 AND NOT-CLCA2	Colon	0.75	1	0.6
LY6G6D AND NOT-MUC1	Colon	0.75	1	0.6
MUC1 AND NOT-S1PR2	Colon	0.75	1	0.6
SLC5A6 AND NOT-GPC3	Colon	0.6	0.6	0.6
MUC1 AND NOT-CTLA4	Colon	0.75	1	0.6
ERBB2 AND NOT-PLA2R1	Colon	0.75	1	0.6
SLC39A4 AND NOT-ERBB2	Colon	0.6	0.6	0.6
ERBB2 AND NOT-P2RY1	Colon	0.75	1	0.6
MUC1 AND NOT-ITGAL	Colon	0.75	1	0.6
MUC1 AND NOT-CLECL1	Colon	0.75	1	0.6
ERBB2 AND NOT-GABRE	Colon	0.66667	0.75	0.6
MUC1 AND NOT-TNFRSF10C	Colon	0.75	1	0.6
VANGL2 AND NOT-ERBB2	Glioblastoma	0.61538	0.8889	0.4706
AQP4 AND NOT-ERBB2	Glioblastoma	0.60465	0.5	0.7647
JAM2 AND NOT-ERBB2	Glioblastoma	0.77419	0.8571	0.7059
HRH1 AND NOT-ERBB2	Glioblastoma	0.66667	0.9	0.5294
OR51B6 AND NOT-CD19	Glioblastoma	0.78049	0.6667	0.9412
PON2 AND NOT-ERBB2	Glioblastoma	0.68085	0.5333	0.9412
PTPRZ1 AND NOT-ERBB2	Glioma	0.9781	0.9853	0.971
PTPRZ1 AND NOT-MUC1	Glioma	0.98551	0.9855	0.9855
NRCAM AND NOT-EPHA2	Glioma	0.91729	0.9531	0.8841
PTPRZ1 AND NOT-EPHA2	Glioma	0.97059	0.9851	0.9565
PCDHB10 AND NOT-MUC1	Glioma	0.896	1	0.8116
PCDHB10 AND NOT-EPHA2	Glioma	0.90476	1	0.8261
PCDHB10 AND NOT-ERBB2	Glioma	0.896	1	0.8116
PON2 AND NOT-MUC1	Glioma	0.96241	1	0.9275
GOLM1 AND NOT-MUC1	Glioma	0.93333	0.9545	0.913
JAM2 AND NOT-EPHA2	Glioma	0.84034	1	0.7246
SRR AND NOT-ERBB2	Glioma	0.79195	0.7375	0.8551
JAM2 AND NOT-MUC1	Glioma	0.90476	1	0.8261
SGCB AND NOT-EPHA2	Glioma	0.85507	0.8551	0.8551
TREM2 AND NOT-MUC1	Glioma	0.90625	0.9831	0.8406
SRR AND NOT-MUC1	Glioma	0.77551	0.7308	0.8261
PON2 AND NOT-EPHA2	Glioma	0.84034	1	0.7246
JAM2 AND NOT-ERBB2	Glioma	0.92188	1	0.8551
SRR AND NOT-EPHA2	Glioma	0.76712	0.7273	0.8116
GOLM1 AND NOT-EPHA2	Glioma	0.8125	0.8814	0.7536
AQP4 AND NOT-MUC1	Glioma	0.79688	0.8644	0.7391
XPR1 AND NOT-ERBB2	Glioma	0.75641	0.6782	0.8551
XPR1 AND NOT-EPHA2	Glioma	0.75472	0.6667	0.8696
XPR1 AND NOT-MUC1	Glioma	0.75	0.6593	0.8696
PON2 AND NOT-ERBB2	Glioma	0.90476	1	0.8261
LDLRAD3 AND NOT-ERBB2	Glioma	0.74214	0.6556	0.8551
AQP4 AND NOT-EPHA2	Glioma	0.7874	0.8621	0.7246
LRP4 AND NOT-ERBB2	Glioma	0.73394	1	0.5797
ROM1 AND NOT-ERBB2	Glioma	0.73171	0.6316	0.8696
ATP2A2 AND NOT-EPHA2	Glioma	0.76106	0.9773	0.6232
LDLRAD3 AND NOT-EPHA2	Glioma	0.72848	0.6707	0.7971
FAM57A AND NOT-EPHA2	Glioma	0.816	0.9107	0.7391
PTPRG AND NOT-EPHA2	Glioma	0.74783	0.9348	0.6232
SLC16A2 AND NOT-EPHA2	Glioma	0.72593	0.7424	0.7101
ROM1 AND NOT-EPHA2	Glioma	0.72483	0.675	0.7826
PCDH17 AND NOT-EPHA2	Glioma	0.72393	0.6277	0.8551
FZD3 AND NOT-MUC1	Glioma	0.72258	0.6512	0.8116
PCDH17 AND NOT-MUC1	Glioma	0.72189	0.61	0.8841
TSPAN6 AND NOT-EPHA2	Glioma	0.78261	0.9783	0.6522
SLC16A2 AND NOT-MUC1	Glioma	0.78378	0.7342	0.8406
TREM2 AND NOT-EPHA2	Glioma	0.84298	0.9808	0.7391
FZD3 AND NOT-EPHA2	Glioma	0.71429	0.6471	0.7971

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
CD163 AND NOT-EPHA2	Glioma	0.6667	0.8444	0.5507
NPFFR1 AND NOT-ERBB2	Glioma	0.7186	0.6122	0.8696
CD163 AND NOT-MUC1	Glioma	0.6667	0.7593	0.5942
CD33 AND NOT-TGFBI	AML	0.8977	0.943	0.8566
CD33 AND NOT-S1PR1	AML	0.894	0.9348	0.8566
CD33 AND NOT-STEAP4	AML	0.9023	0.8851	0.9203
CD33 AND NOT-BTN3A1	AML	0.9398	0.9474	0.9323
CD33 AND NOT-BTN3A3	AML	0.9388	0.9623	0.9163
CD33 AND NOT-IFNAR2	AML	0.9053	0.9598	0.8566
CD33 AND NOT-LAT	AML	0.8986	0.8968	0.9004
CD33 AND NOT-PIK3IP1	AML	0.8718	0.8633	0.8805
CD33 AND NOT-SEMA4D	AML	0.8814	0.8357	0.9323
CD33 AND NOT-ATP7A	AML	0.8716	0.8517	0.8924
CD33 AND NOT-FLVCR1	AML	0.8305	0.7956	0.8685
CD33 AND NOT-MSMO1	AML	0.8187	0.8375	0.8008
CD33 AND NOT-KCNK6	AML	0.8139	0.7222	0.9323
CD33 AND NOT-SEMA4B	AML	0.8112	0.7227	0.9243
CD33 AND NOT-SYT11	AML	0.8512	0.8071	0.9004
CD33 AND NOT-FMNL1	AML	0.8427	0.7715	0.9283
CD33 AND NOT-CD74	AML	0.8286	0.7508	0.9243
CD33 AND NOT-SELP	AML	0.8101	0.7887	0.8327
CD33 AND NOT-SLC22A5	AML	0.8423	0.7655	0.9363
CD33 AND NOT-AGER	AML	0.802	0.7015	0.9363
CD33 AND NOT-PPAPDC1B	AML	0.8124	0.8419	0.7849
CD33 AND NOT-ATP8B1	AML	0.7932	0.6903	0.9323
CD33 AND NOT-AOC3	AML	0.7932	0.6903	0.9323
CD33 AND NOT-SLC31A1	AML	0.789	0.6928	0.9163
CD33 AND NOT-SLC38A1	AML	0.7978	0.7294	0.8805
CD33 AND NOT-CDH11	AML	0.786	0.6772	0.9363
CD33 AND NOT-SCARA5	AML	0.7833	0.6734	0.9363
CD33 AND NOT-MRGPRF	AML	0.782	0.6714	0.9363
CD33 AND NOT-LRIG3	AML	0.7807	0.6695	0.9363
CD33 AND NOT-SLC50A1	AML	0.7795	0.6826	0.9084
CD33 AND NOT-PLVAP	AML	0.7794	0.6676	0.9363
CD33 AND NOT-GHR	AML	0.7785	0.6725	0.9243
CD33 AND NOT-SLC6A6	AML	0.7781	0.6657	0.9363
CD33 AND NOT-CD163	AML	0.7824	0.9337	0.6733
CD33 AND NOT-IL17RA	AML	0.7736	0.6716	0.9124
CD33 AND NOT-PAQR8	AML	0.8674	0.9196	0.8207
CD33 AND NOT-FXYD6	AML	0.7679	0.6609	0.9163
CD33 AND NOT-TNFRSF10B	AML	0.7642	0.6456	0.9363
CD33 AND NOT-SLC9A1	AML	0.7642	0.6456	0.9363
CD33 AND NOT-ZACN	AML	0.763	0.6438	0.9363
CD33 AND NOT-ESAM	AML	0.763	0.6438	0.9363
CD33 AND NOT-CXCL16	AML	0.7585	0.81	0.7131
CD33 AND NOT-PROCR	AML	0.7556	0.6617	0.8805
CD33 AND NOT-IL15RA	AML	0.7402	0.645	0.8685
CD33 AND NOT-SLC16A5	AML	0.733	0.6522	0.8367
CD33 AND NOT-ZDHHC2	AML	0.7183	0.7154	0.7211
CD33 AND NOT-ANPEP	AML	0.7027	0.6414	0.7769
CD33 AND NOT-SORL1	AML	0.6992	0.6858	0.7131
CD33 AND NOT-BST2	AML	0.6974	0.7756	0.6335
CD33 AND NOT-THBD	AML	0.6966	0.7512	0.6494
CD33 AND NOT-SLC22A18	AML	0.6201	0.686	0.5657
CD33 AND NOT-IFI6	AML	0.6054	0.636	0.5777
CD19 AND NOT-CD93	B-Cell Diffuse	0.6875	0.8148	0.5946
CD19 AND NOT-CCR9	B-Cell Diffuse	0.6667	0.8077	0.5676
CD19 AND NOT-ACSL6	B-Cell Diffuse	0.6563	0.7778	0.5676
CD19 AND NOT-GPR22	B-Cell Diffuse	0.6471	0.7097	0.5946
CD19 AND NOT-ATP8B2	B-Cell Diffuse	0.6462	0.75	0.5676
CD19 AND NOT-CNTNAP2	B-Cell Diffuse	0.6452	0.8	0.5405
MS4A1 AND NOT-CCR9	B-Cell Diffuse	0.6389	0.6571	0.6216
CD19 AND NOT-DRD5	B-Cell Diffuse	0.6349	0.7692	0.5405
CD19 AND NOT-BTN3A1	B-Cell Diffuse	0.6207	0.8571	0.4865
CD19 AND NOT-ATP8A2	B-Cell Diffuse	0.6154	0.7143	0.5405
MS4A1 AND NOT-PAQR8	B-Cell Diffuse	0.6133	0.6053	0.6216
MS4A1 AND NOT-CNTNAP2	B-Cell Diffuse	0.6111	0.6286	0.5946
CD19 AND NOT-BTN3A3	B-Cell Diffuse	0.6102	0.8182	0.4865
CD19 AND NOT-SLC16A5	B-Cell Diffuse	0.6102	0.8182	0.4865
MS4A1 AND NOT-DSC1	B-Cell Diffuse	0.6061	0.6897	0.5405
CD19 AND NOT-CD1A	B-Cell Diffuse	0.6032	0.7308	0.5135
CD19 AND NOT-PPAPDC1B	B-Cell Diffuse	0.6	0.7826	0.4865
CD19 AND NOT-MPL	B-Cell Diffuse	0.6	0.7826	0.4865
CD19 AND NOT-SEMA4B	Mantle-Cell Lymphoma	0.9143	1	0.8421
CD19 AND NOT-KCNC2	Mantle-Cell Lymphoma	0.8986	1	0.8158
CD19 AND NOT-CD1A	Mantle-Cell Lymphoma	0.9143	1	0.8421
CD19 AND NOT-SYT6	Mantle-Cell Lymphoma	0.8986	1	0.8158

FIG. 10 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
MERTK AND NOT-EPHA2	Glioma	0.77586	0.9574	0.6522
TENM4 AND NOT-EPHA2	Glioma	0.79389	0.8387	0.7536
SLC16A2 AND NOT-ERBB2	Glioma	0.79195	0.7375	0.8551
TENM4 AND NOT-ERBB2	Glioma	0.80303	0.8413	0.7681
LRP4 AND NOT-MUC1	Glioma	0.71028	1	0.5507
MERTK AND NOT-MUC1	Glioma	0.79389	0.8387	0.7536
PTPRG AND NOT-MUC1	Glioma	0.704	0.7857	0.6377
TENM4 AND NOT-MUC1	Glioma	0.78462	0.8361	0.7391
ABHD12 AND NOT-ERBB2	Glioma	0.77193	0.9778	0.6377
ABHD12 AND NOT-EPHA2	Glioma	0.75	0.9767	0.6087
ABHD12 AND NOT-MUC1	Glioma	0.72727	0.9756	0.5797
GOLM1 AND NOT-ERBB2	Glioma	0.90226	0.9375	0.8696
TMEM231 AND NOT-ERBB2	Glioma	0.69737	0.6386	0.7681
LDLRAD3 AND NOT-MUC1	Glioma	0.69281	0.631	0.7681
MERTK AND NOT-ERBB2	Glioma	0.79389	0.8387	0.7536
TMEM231 AND NOT-MUC1	Glioma	0.68966	0.6579	0.7246
TSPAN6 AND NOT-MUC1	Glioma	0.83453	0.8286	0.8406
ADORA3 AND NOT-EPHA2	Glioma	0.73438	0.7966	0.6812
DISP1 AND NOT-EPHA2	Glioma	0.68657	0.7077	0.6667
SEMA5A AND NOT-ERBB2	Glioma	0.68571	1	0.5217
GPR34 AND NOT-EPHA2	Glioma	0.8	0.8929	0.7246
SLC6A9 AND NOT-ERBB2	Glioma	0.72368	0.6627	0.7971
GPR161 AND NOT-EPHA2	Glioma	0.68056	0.6533	0.7101
SLC6A9 AND NOT-EPHA2	Glioma	0.68027	0.641	0.7246
ROM1 AND NOT-MUC1	Glioma	0.67901	0.5914	0.7971
ATP2A2 AND NOT-ERBB2	Glioma	0.77165	0.8448	0.7101
PTPRG AND NOT-ERBB2	Glioma	0.736	0.8214	0.6667
TMEM100 AND NOT-EPHA2	Glioma	0.70866	0.7759	0.6522
TMEM231 AND NOT-EPHA2	Glioma	0.67606	0.6575	0.6957
SLC3A2 AND NOT-EPHA2	Glioma	0.73684	0.9333	0.6087
SEMA5A AND NOT-MUC1	Glioma	0.67308	1	0.5072
LRP4 AND NOT-EPHA2	Glioma	0.67308	1	0.5072
VANGL2 AND NOT-ERBB2	Glioma	0.67257	0.8636	0.5507
GPR161 AND NOT-ERBB2	Glioma	0.70833	0.68	0.7391
WDR19 AND NOT-EPHA2	Glioma	0.78571	0.7746	0.7971
TMEM100 AND NOT-MUC1	Glioma	0.71318	0.7667	0.6667
PODXL AND NOT-MUC1	Glioma	0.74074	0.7576	0.7246
CD33 AND NOT-IL4R	AML	0.88655	0.9378	0.8406
CD33 AND NOT-AMIGO2	AML	0.92585	0.9315	0.9203
CD33 AND NOT-NPC1	AML	0.90722	0.9402	0.8765
CD33 AND NOT-GPR171	AML	0.92901	0.9463	0.9124
CD33 AND NOT-ADTRP	AML	0.91977	0.9038	0.9363
CD33 AND NOT-STX7	AML	0.94715	0.9668	0.9283
CD33 AND NOT-ST3GAL5	AML	0.91383	0.9194	0.9084
CD33 AND NOT-IL7R	AML	0.92857	0.9249	0.9323
CD33 AND NOT-SLC41A1	AML	0.87398	0.8921	0.8566
CD33 AND NOT-DPP4	AML	0.92771	0.9352	0.9203
CD33 AND NOT-IFITM1	AML	0.87794	0.9491	0.8167
CD33 AND NOT-SLC44A2	AML	0.90873	0.9051	0.9124
CD33 AND NOT-CRTAM	AML	0.89613	0.9167	0.8765
CD33 AND NOT-C6orf89	AML	0.8632	0.8358	0.8924
CD33 AND NOT-BTN3A2	AML	0.93443	0.962	0.9084
CD33 AND NOT-KLRB1	AML	0.88085	0.9452	0.8247
CD33 AND NOT-GPR18	AML	0.88247	0.9145	0.8526
CD33 AND NOT-MAL	AML	0.89655	0.9132	0.8805
CD33 AND NOT-VIPR1	AML	0.86312	0.8255	0.9044
CD33 AND NOT-LTB	AML	0.88136	0.9412	0.8287
CD33 AND NOT-CD2	AML	0.924	0.9277	0.9203
CD33 AND NOT-RER1	AML	0.85657	0.8689	0.8446
CD33 AND NOT-EPHA4	AML	0.90909	0.8835	0.9363
CD33 AND NOT-IFNAR1	AML	0.90945	0.8988	0.9203
CD33 AND NOT-PLXDC1	AML	0.89524	0.8577	0.9363
CD33 AND NOT-PTGDR	AML	0.92857	0.9249	0.9323
CD33 AND NOT-KLRF1	AML	0.91235	0.9124	0.9124
CD33 AND NOT-HMOX2	AML	0.88168	0.8462	0.9203
CD33 AND NOT-TNFSF10	AML	0.88492	0.8814	0.8884
CD33 AND NOT-SLC46A2	AML	0.88588	0.8609	0.9124
CD33 AND NOT-IL2RB	AML	0.91235	0.9124	0.9124
CD33 AND NOT-CD274	AML	0.8785	0.8275	0.9363
CD33 AND NOT-MFAP3L	AML	0.90485	0.8826	0.9283
CD33 AND NOT-TRPC1	AML	0.85926	0.8028	0.9243
CD33 AND NOT-BMPR2	AML	0.8486	0.7993	0.9044
CD33 AND NOT-SLC16A6	AML	0.84889	0.9598	0.761
CD33 AND NOT-VAPA	AML	0.85775	0.9182	0.8048
CD33 AND NOT-SIRPG	AML	0.91406	0.8966	0.9323
CD33 AND NOT-CD28	AML	0.92615	0.928	0.9243
CD33 AND NOT-SLC17A5	AML	0.85246	0.7852	0.9323

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
CD19 AND NOT-GPR78	Mantle-Cell Lymphoma	0.8986	1	0.8158
CD19 AND NOT-CD82	Mantle-Cell Lymphoma	0.8986	1	0.8158
CD19 AND NOT-SLC32A1	Mantle-Cell Lymphoma	0.9143	1	0.8421
CD19 AND NOT-CACNG4	Mantle-Cell Lymphoma	0.9143	1	0.8421
CD19 AND NOT-SORL1	Mantle-Cell Lymphoma	0.8986	1	0.8158
MLANA AND NOT-NOX1	Melanoma	1	1	1
MLANA AND NOT-ATP13A5	Melanoma	1	1	1
MLANA AND NOT-NPFFR1	Melanoma	1	1	1
MLANA AND NOT-NMUR2	Melanoma	1	1	1
MLANA AND NOT-GHR	Melanoma	1	1	1
MLANA AND NOT-SCARA5	Melanoma	1	1	1
MLANA AND NOT-OR10J1	Melanoma	1	1	1
MLANA AND NOT-CDH20	Melanoma	1	1	1
MLANA AND NOT-HTR1E	Melanoma	1	1	1
MLANA AND NOT-TMEM235	Melanoma	1	1	1
MLANA AND NOT-CCKBR	Melanoma	1	1	1
MLANA AND NOT-NKAIN2	Melanoma	0.9524	1	0.9091
MLANA AND NOT-KCND2	Melanoma	0.9524	1	0.9091
MLANA AND NOT-BEST3	Melanoma	0.9524	1	0.9091
MLANA AND NOT-RGSL1	Melanoma	1	1	1
MLANA AND NOT-KCNF1	Melanoma	1	1	1
MLANA AND NOT-EPHA10	Melanoma	1	1	1
MLANA AND NOT-GPR26	Melanoma	1	1	1
MLANA AND NOT-KCNA10	Melanoma	1	1	1
MLANA AND NOT-CRB2	Melanoma	1	1	1
MLANA AND NOT-HTR6	Melanoma	1	1	1
MLANA AND NOT-GRIK4	Melanoma	1	1	1
MLANA AND NOT-GRIN1	Melanoma	1	1	1
MLANA AND NOT-GRIN2B	Melanoma	1	1	1
MLANA AND NOT-PCDHB1	Melanoma	1	1	1
MLANA AND NOT-SLC6A18	Melanoma	1	1	1
MLANA AND NOT-HTR5A	Melanoma	1	1	1
MLANA AND NOT-HTR1D	Melanoma	1	1	1
MLANA AND NOT-HCN4	Melanoma	1	1	1
MLANA AND NOT-GPR78	Melanoma	1	1	1
MLANA AND NOT-CACNG4	Melanoma	1	1	1
MLANA AND NOT-SEZ6	Melanoma	1	1	1
MLANA AND NOT-CNGA4	Melanoma	1	1	1
MLANA AND NOT-SLC32A1	Melanoma	1	1	1
MLANA AND NOT-CLDN19	Melanoma	1	1	1
MLANA AND NOT-CYP4A11	Melanoma	1	1	1
MLANA AND NOT-OR1Q1	Melanoma	1	1	1
MLANA AND NOT-SLC5A8	Melanoma	1	1	1
MLANA AND NOT-LRRN4	Melanoma	1	1	1
MLANA AND NOT-GPR156	Melanoma	1	1	1
MLANA AND NOT-FGF6	Melanoma	1	1	1
MLANA AND NOT-KCNH4	Melanoma	1	1	1
MLANA AND NOT-FSHR	Melanoma	1	1	1
MLANA AND NOT-CALHM1	Melanoma	1	1	1
MLANA AND NOT-OR1C1	Melanoma	1	1	1
MLANA AND NOT-OR2L2	Melanoma	1	1	1
MLANA AND NOT-TSPAN16	Melanoma	1	1	1
MLANA AND NOT-GHSR	Melanoma	1	1	1
MLANA AND NOT-AMHR2	Melanoma	1	1	1
MLANA AND NOT-KCNJ9	Melanoma	1	1	1
MLANA AND NOT-KCNH1	Melanoma	1	1	1
MLANA AND NOT-OR52D1	Melanoma	1	1	1
MLANA AND NOT-KCNQ2	Melanoma	1	1	1
MLANA AND NOT-CALN1	Melanoma	1	1	1
MLANA AND NOT-SLC1A6	Melanoma	1	1	1
MLANA AND NOT-SLC10A1	Melanoma	1	1	1
MLANA AND NOT-SLC10A2	Melanoma	1	1	1
MLANA AND NOT-SLC34A1	Melanoma	0.9524	1	0.9091
MLANA AND NOT-SLC22A5	Melanoma	1	1	1
MLANA AND NOT-UPK3A	Melanoma	1	1	1
MLANA AND NOT-CACNA1S	Melanoma	1	1	1
MLANA AND NOT-PCDH11Y	Melanoma	0.9524	1	0.9091
MLANA AND NOT-GJA10	Melanoma	1	1	1
MLANA AND NOT-GPR135	Melanoma	1	1	1
MLANA AND NOT-CASR	Melanoma	1	1	1
MLANA AND NOT-CCKAR	Melanoma	1	1	1
MLANA AND NOT-SLC28A1	Melanoma	1	1	1
MLANA AND NOT-GPR50	Melanoma	1	1	1
MLANA AND NOT-SLC22A6	Melanoma	1	1	1
MLANA AND NOT-OTOF	Melanoma	1	1	1
MLANA AND NOT-SLC22A13	Melanoma	1	1	1
MLANA AND NOT-IGDCC3	Melanoma	1	1	1

FIG. 10 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
CD33 AND NOT-P2RY10	AML	0.92247	0.9206	0.9243
CD33 AND NOT-PROS1	AML	0.8743	0.8262	0.9283
CD33 AND NOT-ABHD6	AML	0.87199	0.816	0.9363
CD33 AND NOT-CD247	AML	0.91235	0.9124	0.9124
CD33 AND NOT-LAMP3	AML	0.86876	0.8103	0.9363
CD33 AND NOT-FPR1	AML	0.84979	0.9209	0.7888
CD33 AND NOT-CD8A	AML	0.91235	0.9124	0.9124
CD33 AND NOT-ICOS	AML	0.904	0.9076	0.9004
CD33 AND NOT-TNFRSF1A	AML	0.84381	0.8595	0.8287
CD33 AND NOT-CD27	AML	0.91732	0.9066	0.9283
CD33 AND NOT-PMEPA1	AML	0.85714	0.7932	0.9323
CD33 AND NOT-AQP3	AML	0.86076	0.9148	0.8127
CD33 AND NOT-KLRD1	AML	0.92338	0.9109	0.9363
CD33 AND NOT-CD8B	AML	0.917	0.9098	0.9243
CD33 AND NOT-TSPAN18	AML	0.84229	0.7655	0.9363
CD33 AND NOT-SIT1	AML	0.92673	0.9213	0.9323
CD33 AND NOT-ATP10D	AML	0.87726	0.8862	0.8685
CD33 AND NOT-SLC22A3	AML	0.84381	0.768	0.9363
CD33 AND NOT-TIGIT	AML	0.89313	0.8571	0.9323
CD33 AND NOT-CCR6	AML	0.90173	0.8731	0.9323
CD33 AND NOT-BTLA	AML	0.91945	0.907	0.9323
CD33 AND NOT-LY9	AML	0.91262	0.8902	0.9363
CD33 AND NOT-NKG7	AML	0.88485	0.8975	0.8725
CD33 AND NOT-PAG1	AML	0.91383	0.9194	0.9084
CD33 AND NOT-PTCH1	AML	0.89922	0.8755	0.9243
CD33 AND NOT-GLG1	AML	0.83951	0.8681	0.8127
CD33 AND NOT-DRAM2	AML	0.84567	0.9009	0.7968
CD33 AND NOT-TSPAN14	AML	0.83817	0.8745	0.8048
CD33 AND NOT-SLC38A6	AML	0.85421	0.8814	0.8287
CD33 AND NOT-FCRL3	AML	0.91765	0.9035	0.9323
CD33 AND NOT-KCNJ2	AML	0.89024	0.9087	0.8725
CD33 AND NOT-IL6ST	AML	0.83594	0.8199	0.8526
CD33 AND NOT-TMX3	AML	0.88224	0.884	0.8805
CD33 AND NOT-PTPRCAP	AML	0.84556	0.8202	0.8725
CD33 AND NOT-CD40LG	AML	0.91228	0.8931	0.9323
CD33 AND NOT-VSIG1	AML	0.84079	0.763	0.9363
CD33 AND NOT-HVCN1	AML	0.89571	0.9202	0.8725
CD33 AND NOT-LGR6	AML	0.83696	0.7674	0.9203
CD33 AND NOT-CD3G	AML	0.92	0.9237	0.9163
CD33 AND NOT-CLEC2D	AML	0.92126	0.9105	0.9323
CD33 AND NOT-FCGR2B	AML	0.90683	0.944	0.8725
CD33 AND NOT-TGFBR2	AML	0.84229	0.7655	0.9363
CD33 AND NOT-SLC16A10	AML	0.85662	0.7952	0.9283
CD33 AND NOT-TMEM204	AML	0.83186	0.7484	0.9363
CD33 AND NOT-ITPR3	AML	0.83186	0.7484	0.9363
CD33 AND NOT-CD5	AML	0.90874	0.8864	0.9323
CD33 AND NOT-BBS4	AML	0.83107	0.8106	0.8526
CD33 AND NOT-FLT3LG	AML	0.89101	0.8566	0.9283
CD33 AND NOT-ABCG1	AML	0.85214	0.8327	0.8725
CD19 AND NOT-SCNN1D	B-Cell Diffuse	0.72131	0.9167	0.5946
CD19 AND NOT-CXCR2	B-Cell Diffuse	0.70968	0.88	0.5946
CD19 AND NOT-ABCA2	B-Cell Diffuse	0.70968	0.88	0.5946
CD19 AND NOT-CD244	B-Cell Diffuse	0.69841	0.8462	0.5946
CD19 AND NOT-SLC46A2	B-Cell Diffuse	0.69841	0.8462	0.5946
CD19 AND NOT-SLC24A1	B-Cell Diffuse	0.69841	0.8462	0.5946
CD19 AND NOT-S1PR5	B-Cell Diffuse	0.69841	0.8462	0.5946
CD19 AND NOT-STX3	B-Cell Diffuse	0.69841	0.8462	0.5946
CD19 AND NOT-FCER1A	B-Cell Diffuse	0.68852	0.875	0.5676
CD19 AND NOT-BST1	B-Cell Diffuse	0.6875	0.8148	0.5946
CD19 AND NOT-TLR5	B-Cell Diffuse	0.6875	0.8148	0.5946
CD19 AND NOT-NAALAD2	B-Cell Diffuse	0.6875	0.8148	0.5946
CD19 AND NOT-ITGA6	B-Cell Diffuse	0.6875	0.8148	0.5946
CD19 AND NOT-ASGR1	B-Cell Diffuse	0.67742	0.84	0.5676
CD19 AND NOT-BTN2A1	B-Cell Diffuse	0.67742	0.84	0.5676
CD19 AND NOT-ADAM23	B-Cell Diffuse	0.67742	0.84	0.5676
CD19 AND NOT-SCIMP	B-Cell Diffuse	0.67742	0.84	0.5676
CD19 AND NOT-TMIGD2	B-Cell Diffuse	0.67742	0.84	0.5676
CD19 AND NOT-ANO8	B-Cell Diffuse	0.67742	0.84	0.5676
CD19 AND NOT-MFAP3L	B-Cell Diffuse	0.67692	0.7857	0.5946
CD19 AND NOT-GP9	B-Cell Diffuse	0.67692	0.7857	0.5946
CD19 AND NOT-GPBAR1	B-Cell Diffuse	0.67692	0.7857	0.5946
MS4A1 AND NOT-TSPAN32	B-Cell Diffuse	0.67647	0.7419	0.6216
MS4A1 AND NOT-CD244	B-Cell Diffuse	0.66667	0.6857	0.6486
MS4A1 AND NOT-SLC46A2	B-Cell Diffuse	0.66667	0.6857	0.6486
MS4A1 AND NOT-ABCA2	B-Cell Diffuse	0.66667	0.6857	0.6486
CD19 AND NOT-CD300LB	B-Cell Diffuse	0.66667	0.7586	0.5946
CD19 AND NOT-CX3CR1	B-Cell Diffuse	0.66667	0.7586	0.5946

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
MLANA AND NOT-GABBR2	Melanoma	1	1	1
MLANA AND NOT-SLC1A2	Melanoma	0.9524	1	0.9091
MLANA AND NOT-ABCG8	Melanoma	1	1	1
MLANA AND NOT-CALHM3	Melanoma	1	1	1
MLANA AND NOT-RXFP3	Melanoma	1	1	1
MLANA AND NOT-MC2R	Melanoma	1	1	1
MLANA AND NOT-MIP	Melanoma	1	1	1
MLANA AND NOT-MPL	Melanoma	0.9524	1	0.9091
MLANA AND NOT-OR3A1	Melanoma	1	1	1
MLANA AND NOT-P2RX3	Melanoma	1	1	1
MLANA AND NOT-P2RY4	Melanoma	0.9524	1	0.9091
MLANA AND NOT-CALY	Melanoma	1	1	1
MLANA AND NOT-KCNK4	Melanoma	1	1	1
MLANA AND NOT-PCDH8	Melanoma	1	1	1
MLANA AND NOT-ATP8B1	Melanoma	1	1	1
MLANA AND NOT-CACNG6	Melanoma	1	1	1
MLANA AND NOT-BEST2	Melanoma	1	1	1
MLANA AND NOT-PCDHGC4	Melanoma	1	1	1
MLANA AND NOT-PCDHAC2	Melanoma	0.9524	1	0.9091
MLANA AND NOT-KCNK12	Melanoma	1	1	1
MLANA AND NOT-GJD2	Melanoma	1	1	1
MLANA AND NOT-LRFN2	Melanoma	1	1	1
MLANA AND NOT-SLC4A5	Melanoma	1	1	1
MLANA AND NOT-CACNG8	Melanoma	1	1	1
MLANA AND NOT-CACNG7	Melanoma	1	1	1
MLANA AND NOT-LRIG3	Melanoma	1	1	1
MLANA AND NOT-UMOD	Melanoma	1	1	1
GPR85 AND NOT-B4GALNT1	Neuroblastoma	0.7805	0.8	0.7619
CACNA1B AND NOT-B4GALNT1	Neuroblastoma	0.7624	0.7113	0.8214
SLC6A15 AND NOT-B4GALNT1	Neuroblastoma	0.6961	0.6495	0.75
KCNQ2 AND NOT-B4GALNT1	Neuroblastoma	0.6736	0.5963	0.7738
B4GALNT1 AND NOT-NKAIN2	Neuroblastoma	0.6383	0.7895	0.5357
B4GALNT1 AND NOT-KCNA1	Neuroblastoma	0.6667	0.6923	0.6429
B4GALNT1 AND NOT-GRM3	Neuroblastoma	0.6386	0.6463	0.631
B4GALNT1 AND NOT-KCNA2	Neuroblastoma	0.6232	0.7963	0.5119
B4GALNT1 AND NOT-CSMD3	Neuroblastoma	0.7101	0.9074	0.5833
B4GALNT1 AND NOT-GABRB2	Neuroblastoma	0.6369	0.6849	0.5952
B4GALNT1 AND NOT-SLCO1C1	Neuroblastoma	0.6933	0.7879	0.619
B4GALNT1 AND NOT-MLC1	Neuroblastoma	0.6391	0.6353	0.6429
B4GALNT1 AND NOT-DPP10	Neuroblastoma	0.6027	0.7097	0.5238
B4GALNT1 AND NOT-KCNJ10	Neuroblastoma	0.6026	0.6528	0.5595
EPHA10 AND NOT-ERBB2	Ovarian	0.6207	0.5294	0.75
FOLH1 AND NOT-NTSR2	Prostate	0.6667	1	0.5
FOLH1 AND NOT-GRIN2C	Prostate	0.8889	0.8	1
FOLH1 AND NOT-SLC6A13	Prostate	0.6667	1	0.5
FOLH1 AND NOT-SLC22A8	Prostate	0.6667	1	0.5
FOLH1 AND NOT-C8B	Prostate	0.8571	1	0.75
FOLH1 AND NOT-MEGF11	Prostate	0.6667	0.6	0.75
FOLH1 AND NOT-SLC13A5	Prostate	0.6667	0.6	0.75
FOLH1 AND NOT-PAQR8	Prostate	0.6154	0.4444	1
FOLH1 AND NOT-SLC4A8	Prostate	0.8889	0.8	1
FOLH1 AND NOT-SLC30A10	Prostate	0.8	0.6667	1
FOLH1 AND NOT-SLC1A2	Prostate	0.6667	0.6	0.75
FOLH1 AND NOT-ATP13A5	Prostate	0.8571	1	0.75
FOLH1 AND NOT-CD82	Prostate	0.8571	1	0.75
FOLH1 AND NOT-NRG3	Prostate	0.6667	0.6	0.75
FOLH1 AND NOT-CHRNA2	Prostate	0.6667	1	0.5
FOLH1 AND NOT-GRIN2A	Prostate	0.6667	1	0.5
FOLH1 AND NOT-CLDN17	Prostate	0.6667	0.6	0.75
FOLH1 AND NOT-SLCO1A2	Prostate	0.8571	1	0.75
FOLH1 AND NOT-KIRREL3	Prostate	0.8571	1	0.75
FOLH1 AND NOT-SLC6A12	Prostate	0.75	0.75	0.75
FOLH1 AND NOT-FMNL1	Prostate	0.75	0.75	0.75
FOLH1 AND NOT-OPALIN	Prostate	0.8571	1	0.75
FOLH1 AND NOT-SLC39A2	Prostate	0.8571	1	0.75
FOLH1 AND NOT-CNTNAP4	Prostate	0.8571	1	0.75
FOLH1 AND NOT-LCT	Prostate	0.6667	0.6	0.75
FOLH1 AND NOT-OR1Q1	Prostate	0.8571	1	0.75
FOLH1 AND NOT-GJA10	Prostate	0.6	0.5	0.75
FOLH1 AND NOT-KCNJ10	Prostate	0.6667	1	0.5
FOLH1 AND NOT-PIRT	Prostate	0.6	0.5	0.75
FOLH1 AND NOT-TTYH2	Prostate	0.6154	0.4444	1
FOLH1 AND NOT-KCNC1	Prostate	0.6	0.5	0.75
FOLH1 AND NOT-SLC12A5	Prostate	0.6667	0.6	0.75
FOLH1 AND NOT-VN1R2	Prostate	0.8571	1	0.75
FOLH1 AND NOT-MC2R	Prostate	0.75	0.75	0.75
FOLH1 AND NOT-MAG	Prostate	0.6667	0.6	0.75

FIG. 10 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
MS4A1 AND NOT-RNF144A	B-Cell Diffuse	0.66667	0.6857	0.6486
CD19 AND NOT-SLC18A2	B-Cell Diffuse	0.66667	0.7586	0.5946
MS4A1 AND NOT-BST1	B-Cell Diffuse	0.66667	0.6857	0.6486
MS4A1 AND NOT-CXCR2	B-Cell Diffuse	0.66667	0.6857	0.6486
CD19 AND NOT-TSPAN32	B-Cell Diffuse	0.66667	0.8077	0.5676
CD19 AND NOT-C19orf26	B-Cell Diffuse	0.66667	0.8077	0.5676
CD19 AND NOT-CCR3	B-Cell Diffuse	0.66667	0.8696	0.5405
MS4A1 AND NOT-MFAP3L	B-Cell Diffuse	0.65753	0.6667	0.6486
MS4A1 AND NOT-ITGA6	B-Cell Diffuse	0.65753	0.6667	0.6486
MS4A1 AND NOT-SIRPB1	B-Cell Diffuse	0.65753	0.6667	0.6486
CD19 AND NOT-CLEC1B	B-Cell Diffuse	0.65672	0.7333	0.5946
CD19 AND NOT-SIRPB1	B-Cell Diffuse	0.65672	0.7333	0.5946
CD19 AND NOT-SLC24A4	B-Cell Diffuse	0.65672	0.7333	0.5946
CD19 AND NOT-PTGDR2	B-Cell Diffuse	0.65574	0.8333	0.5405
CD19 AND NOT-MGAM	B-Cell Diffuse	0.65574	0.8333	0.5405
CD19 AND NOT-EPOR	B-Cell Diffuse	0.65574	0.8333	0.5405
CD19 AND NOT-IL1RAP	B-Cell Diffuse	0.65574	0.8333	0.5405
CD19 AND NOT-VIPR1	B-Cell Diffuse	0.65574	0.8333	0.5405
CD19 AND NOT-RNF144A	B-Cell Diffuse	0.65517	0.9048	0.5135
MS4A1 AND NOT-S1PR5	B-Cell Diffuse	0.64865	0.6486	0.6486
MS4A1 AND NOT-CX3CR1	B-Cell Diffuse	0.64865	0.6486	0.6486
MS4A1 AND NOT-ASGR1	B-Cell Diffuse	0.64789	0.6765	0.6216
MS4A1 AND NOT-CLEC1B	B-Cell Diffuse	0.64789	0.6765	0.6216
MS4A1 AND NOT-TMIGD2	B-Cell Diffuse	0.64789	0.6765	0.6216
MS4A1 AND NOT-FCER1A	B-Cell Diffuse	0.64789	0.6765	0.6216
CD19 AND NOT-VSIG2	B-Cell Diffuse	0.64706	0.7097	0.5946
CD19 AND NOT-LGR6	B-Cell Diffuse	0.64706	0.7097	0.5946
CD19 AND NOT-AQP8	Mantle-Cell Lymphoma	0.91429	1	0.8421
CD19 AND NOT-GPM6B	Mantle-Cell Lymphoma	0.91429	1	0.8421
CD19 AND NOT-TRPV2	Mantle-Cell Lymphoma	0.91429	1	0.8421
CD19 AND NOT-MS4A2	Mantle-Cell Lymphoma	0.91429	1	0.8421
CD19 AND NOT-FCER2	Mantle-Cell Lymphoma	0.91429	1	0.8421
CD19 AND NOT-SLC52A3	Mantle-Cell Lymphoma	0.91429	1	0.8421
CD19 AND NOT-LRRTM2	Mantle-Cell Lymphoma	0.91429	1	0.8421
CD19 AND NOT-CYP4F12	Mantle-Cell Lymphoma	0.91429	1	0.8421
MLANA AND NOT-RET	Melanoma	1	1	1
MLANA AND NOT-FAT4	Melanoma	1	1	1
MLANA AND NOT-FRRS1L	Melanoma	1	1	1
MLANA AND NOT-PCDH18	Melanoma	1	1	1
MLANA AND NOT-CADM2	Melanoma	1	1	1
MLANA AND NOT-ABCC9	Melanoma	1	1	1
MLANA AND NOT-FXYD4	Melanoma	1	1	1
MLANA AND NOT-KCNK3	Melanoma	1	1	1
MLANA AND NOT-SGMS1	Melanoma	1	1	1
MLANA AND NOT-XG	Melanoma	1	1	1
MLANA AND NOT-CFTR	Melanoma	1	1	1
MLANA AND NOT-KCNB1	Melanoma	1	1	1
MLANA AND NOT-ZP2	Melanoma	1	1	1
MLANA AND NOT-PCDH19	Melanoma	1	1	1
MLANA AND NOT-LY6G6D	Melanoma	1	1	1
MLANA AND NOT-GPRC5C	Melanoma	1	1	1
MLANA AND NOT-OR1J2	Melanoma	1	1	1
MLANA AND NOT-AQP10	Melanoma	1	1	1
MLANA AND NOT-GPR101	Melanoma	1	1	1
MLANA AND NOT-CHRM2	Melanoma	1	1	1
MLANA AND NOT-SGCG	Melanoma	1	1	1
MLANA AND NOT-NPR3	Melanoma	1	1	1
MLANA AND NOT-BMP2	Melanoma	1	1	1
MLANA AND NOT-TMEM100	Melanoma	1	1	1
MLANA AND NOT-NPY5R	Melanoma	1	1	1
MLANA AND NOT-AGTR1	Melanoma	1	1	1
MLANA AND NOT-IL31RA	Melanoma	1	1	1
MLANA AND NOT-HRH2	Melanoma	1	1	1
MLANA AND NOT-ADCYAP1R1	Melanoma	1	1	1
MLANA AND NOT-FAT3	Melanoma	1	1	1
MLANA AND NOT-NTRK3	Melanoma	1	1	1
MLANA AND NOT-FZD8	Melanoma	0.95238	1	0.9091
MLANA AND NOT-SYT8	Melanoma	0.95238	1	0.9091
MLANA AND NOT-GPRC5D	Melanoma	0.95238	1	0.9091
MLANA AND NOT-SGCA	Melanoma	0.95238	1	0.9091
MLANA AND NOT-OR51M1	Melanoma	0.95238	1	0.9091
MLANA AND NOT-SLC6A20	Melanoma	1	1	1
MLANA AND NOT-NPY1R	Melanoma	1	1	1
MLANA AND NOT-NTRK1	Melanoma	1	1	1
MLANA AND NOT-CCR4	Melanoma	1	1	1
MLANA AND NOT-ATP4A	Melanoma	1	1	1
MLANA AND NOT-GDPD2	Melanoma	1	1	1

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
FOLH1 AND NOT-GPR19	Prostate	0.75	0.75	0.75
FOLH1 AND NOT-SLCO1C1	Prostate	0.8571	1	0.75
FOLH1 AND NOT-TRPC3	Prostate	0.6667	1	0.5
FOLH1 AND NOT-MOG	Prostate	0.8571	1	0.75
FOLH1 AND NOT-NKAIN2	Prostate	0.6667	0.5	1
PTGIS AND NOT-B4GALNT1	Sarcoma	0.7143	0.9091	0.5882
NOT-CLDN10 AND MUC1	Colon	0.6667	0.75	0.6
NOT-CLDN10 AND ERBB2	Colon	0.6	0.6	0.6
NOT-PORCN AND ERBB2	Colon	0.6667	0.75	0.6
NOT-CNIH2 AND ERBB2	Colon	0.75	1	0.6
NOT-GPR158 AND ERBB2	Colon	0.75	1	0.6
NOT-AGTR2 AND ERBB2	Colon	0.75	1	0.6
NOT-KCNA7 AND MUC1	Colon	0.75	1	0.6
NOT-KCNJ10 AND MUC1	Colon	0.75	1	0.6
NOT-CNIH2 AND MUC1	Colon	0.75	1	0.6
NOT-KCNJ10 AND ERBB2	Colon	0.75	1	0.6
NOT-CACNG8 AND ERBB2	Colon	0.75	1	0.6
NOT-MEGF10 AND ERBB2	Colon	0.6667	0.5714	0.8
NOT-EPHA10 AND ERBB2	Colon	0.75	1	0.6
NOT-SCN2A AND ERBB2	Colon	0.6667	0.75	0.6
NOT-CACNG8 AND MUC1	Colon	0.75	1	0.6
NOT-NPHS1 AND ERBB2	Colon	0.6667	0.75	0.6
NOT-GRM7 AND MUC1	Colon	0.6667	0.75	0.6
NOT-KCNA7 AND ERBB2	Colon	0.75	1	0.6
NOT-ATP8A2 AND MUC1	Colon	0.75	1	0.6
NOT-ADAM30 AND ERBB2	Colon	0.6667	0.75	0.6
NOT-MEGF10 AND MUC1	Colon	0.6667	0.5714	0.8
NOT-KIRREL3 AND ERBB2	Colon	0.75	1	0.6
NOT-ATP8A2 AND ERBB2	Colon	0.75	1	0.6
NOT-SYT6 AND ERBB2	Colon	0.75	1	0.6
NOT-GALR1 AND ERBB2	Colon	0.6667	0.75	0.6
NOT-PORCN AND MUC1	Colon	0.6667	0.75	0.6
NOT-GABRA2 AND MUC1	Colon	0.8889	1	0.8
NOT-CD1A AND MUC1	Glioma	0.7191	0.5872	0.9275
NOT-CD1A AND ERBB2	Glioma	0.6802	0.5234	0.971
NOT-PHLDB2 AND CD33	AML	0.9391	0.9264	0.9522
NOT-GPR137B AND CD33	AML	0.8584	0.9302	0.7968
NOT-PHLDB2 AND IL3RA	AML	0.6649	0.9552	0.51
FAP AND NOT-ROR1	Breast	0.8515	0.9149	0.7963
FAP AND NOT-TYR	Breast	0.8381	0.8627	0.8148
EPCAM AND NOT-ALDH1A1	Breast	0.7209	0.9688	0.5741
EPCAM AND NOT-MUC13	Breast	0.7304	0.6885	0.7778
VTCN1 AND NOT-TYR	Breast	0.6905	0.9667	0.537
VTCN1 AND NOT-MET	Breast	0.6905	0.9667	0.537
EPCAM AND NOT-MUC4	Breast	0.6829	0.6087	0.7778
EPCAM AND NOT-SSTR1	Breast	0.6565	0.5584	0.7963
EPCAM AND NOT-GPA33	Breast	0.6829	0.6087	0.7778
EPCAM AND NOT-ENPP3	Breast	0.6496	0.6032	0.7037
EPCAM AND NOT-HHLA2	Breast	0.6412	0.5455	0.7778
EPCAM AND NOT-GUCY2C	Breast	0.6615	0.5658	0.7963
EPCAM AND NOT-SST	Breast	0.7167	0.6515	0.7963
TPBG AND NOT-TYR	Breast	0.62	0.6739	0.5741
EPCAM AND NOT-CLDN8	Breast	0.6116	0.5522	0.6852
EPCAM AND NOT-FOLR1	Breast	0.6187	0.5059	0.7963
FAP AND NOT-MET	Breast	0.6047	0.8125	0.4815
TPBG AND NOT-ROR1	Breast	0.6042	0.6905	0.537
EPCAM AND NOT-CLDN7	Breast	0.6107	0.5195	0.7407
EPCAM AND NOT-LGR5	Breast	0.6457	0.5616	0.7593
EPCAM AND NOT-CLDN2	Breast	0.6563	0.5676	0.7778
AXL AND NOT-ROR1	Breast	0.6207	0.5806	0.6667
EPCAM AND NOT-CLDN4	Breast	0.6087	0.5	0.7778
EPCAM AND NOT-TNFRSF13C	Breast	0.6056	0.4886	0.7963
GPNMB AND NOT-ROR1	Breast	0.6154	0.5263	0.7407
EPCAM AND NOT-EGFR	Breast	0.6067	0.7714	0.5
EPCAM AND NOT-PROM1	Breast	0.6538	0.68	0.6296
CLDN2 AND NOT-MUC1	Liver	0.625	0.5	0.8333
CLDN12 AND NOT-MUC1	Liver	1	1	1
DPEP1 AND NOT-GPC3	Colon	1	1	1
EPHB2 AND NOT-MUC1	Colon	0.75	1	0.6
RNF43 AND NOT-ERBB2	Colon	0.75	1	0.6
RNF43 AND NOT-GPC3	Colon	0.75	1	0.6
RNF43 AND NOT-MUC1	Colon	0.75	1	0.6
EPHB2 AND NOT-GPC3	Colon	0.7273	0.6667	0.8
EPHB2 AND NOT-ERBB2	Colon	0.8	0.8	0.8
CLDN2 AND NOT-GPC3	Colon	0.75	1	0.6
BIRC5 AND NOT-GPC3	Colon	0.625	0.4545	1
MET AND NOT-GPC3	Colon	0.75	1	0.6

FIG. 10 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
MLANA AND NOT-SLC51B	Melanoma	0.95238	1	0.9091
MLANA AND NOT-DCHS2	Melanoma	1	1	1
MLANA AND NOT-OPRD1	Melanoma	1	1	1
MLANA AND NOT-TAS2R8	Melanoma	1	1	1
MLANA AND NOT-PDGFRB	Melanoma	0.95238	1	0.9091
MLANA AND NOT-OPRL1	Melanoma	1	1	1
MLANA AND NOT-DLL4	Melanoma	1	1	1
MLANA AND NOT-TAS2R5	Melanoma	1	1	1
MLANA AND NOT-CDHR5	Melanoma	1	1	1
MLANA AND NOT-OR1F1	Melanoma	1	1	1
MLANA AND NOT-P2RX1	Melanoma	1	1	1
MLANA AND NOT-S1PR5	Melanoma	1	1	1
MLANA AND NOT-DUOX2	Melanoma	1	1	1
MLANA AND NOT-PKD1	Melanoma	1	1	1
MLANA AND NOT-TAS2R3	Melanoma	1	1	1
MLANA AND NOT-RHBDL2	Melanoma	0.95238	1	0.9091
MLANA AND NOT-TAS2R9	Melanoma	1	1	1
MLANA AND NOT-CLCNKB	Melanoma	1	1	1
MLANA AND NOT-NAALAD2	Melanoma	1	1	1
MLANA AND NOT-PCDHGA11	Melanoma	1	1	1
MLANA AND NOT-SLC38A4	Melanoma	1	1	1
MLANA AND NOT-ECSCR	Melanoma	0.95238	1	0.9091
MLANA AND NOT-SCN3A	Melanoma	0.95238	1	0.9091
MLANA AND NOT-SCN4A	Melanoma	1	1	1
MLANA AND NOT-SCN4B	Melanoma	0.95238	1	0.9091
MLANA AND NOT-SCNN1A	Melanoma	1	1	1
MLANA AND NOT-GPR182	Melanoma	1	1	1
MLANA AND NOT-SCTR	Melanoma	1	1	1
MLANA AND NOT-CDH23	Melanoma	1	1	1
MLANA AND NOT-SLC1A7	Melanoma	1	1	1
MLANA AND NOT-OPN1SW	Melanoma	1	1	1
MLANA AND NOT-SLC2A4	Melanoma	1	1	1
MLANA AND NOT-SLC5A5	Melanoma	1	1	1
MLANA AND NOT-SLC6A2	Melanoma	1	1	1
MLANA AND NOT-SLC6A3	Melanoma	1	1	1
MLANA AND NOT-GPR45	Melanoma	1	1	1
MLANA AND NOT-SLC6A9	Melanoma	1	1	1
MLANA AND NOT-SLC9A2	Melanoma	1	1	1
MLANA AND NOT-SCN2B	Melanoma	1	1	1
MLANA AND NOT-HRH4	Melanoma	1	1	1
MLANA AND NOT-ANO1	Melanoma	1	1	1
MLANA AND NOT-PCDHA10	Melanoma	1	1	1
MLANA AND NOT-AVPR1A	Melanoma	1	1	1
MLANA AND NOT-AVPR2	Melanoma	1	1	1
MLANA AND NOT-GABRQ	Melanoma	1	1	1
MLANA AND NOT-PCDHGC5	Melanoma	1	1	1
CELSR3 AND NOT-B4GALNT1	Neuroblastoma	0.75258	0.6636	0.869
SCN3A AND NOT-B4GALNT1	Neuroblastoma	0.71958	0.6476	0.8095
CPT1C AND NOT-B4GALNT1	Neuroblastoma	0.71006	0.7059	0.7143
FZD3 AND NOT-B4GALNT1	Neuroblastoma	0.67879	0.6914	0.6667
GABRB3 AND NOT-B4GALNT1	Neuroblastoma	0.65031	0.6709	0.631
B4GALNT1 AND NOT-SCN1B	Neuroblastoma	0.65734	0.7966	0.5595
KCNQ5 AND NOT-B4GALNT1	Neuroblastoma	0.64286	0.6429	0.6429
EPHA5 AND NOT-B4GALNT1	Neuroblastoma	0.62069	0.7377	0.5357
B4GALNT1 AND NOT-AQP4	Neuroblastoma	0.62162	0.7188	0.5476
B4GALNT1 AND NOT-ART3	Neuroblastoma	0.60645	0.662	0.5595
SCN3B AND NOT-B4GALNT1	Neuroblastoma	0.60377	0.5	0.7619
B4GALNT1 AND NOT-STYK1	Neuroblastoma	0.63226	0.6901	0.5833
B4GALNT1 AND NOT-SLC15A2	Neuroblastoma	0.67925	0.72	0.6429
B4GALNT1 AND NOT-SLC17A7	Neuroblastoma	0.64151	0.68	0.6071
B4GALNT1 AND NOT-GJB6	Neuroblastoma	0.67516	0.726	0.631
B4GALNT1 AND NOT-SLC4A4	Neuroblastoma	0.71533	0.9245	0.5833
B4GALNT1 AND NOT-SLC24A4	Neuroblastoma	0.60227	0.5761	0.631
B4GALNT1 AND NOT-ADRB1	Neuroblastoma	0.62069	0.6	0.6429
B4GALNT1 AND NOT-KCNS1	Neuroblastoma	0.60526	0.6765	0.5476
B4GALNT1 AND NOT-CACNA1A	Neuroblastoma	0.61905	0.619	0.619
KCNK15 AND NOT-ERBB2	Ovarian	0.8	0.6667	1
FOLH1 AND NOT-ACVR1C	Prostate	0.72727	0.5714	1
FOLH1 AND NOT-TMEM63B	Prostate	0.66667	1	0.5
FOLH1 AND NOT-PLB1	Prostate	0.66667	1	0.5
FOLH1 AND NOT-CYSLTR2	Prostate	0.66667	1	0.5
FOLH1 AND NOT-ATP1B1	Prostate	0.85714	1	0.75
FOLH1 AND NOT-UGT8	Prostate	0.75	0.75	0.75
FOLH1 AND NOT-APLNR	Prostate	0.66667	0.6	0.75
FOLH1 AND NOT-PCDHB12	Prostate	0.66667	1	0.5
FOLH1 AND NOT-CSF1	Prostate	0.66667	1	0.5
FOLH1 AND NOT-ITM2A	Prostate	0.66667	1	0.5

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
BIRC5 AND NOT-ERBB2	Colon	0.625	0.4545	1
MUC1 AND NOT-CLDN18	Colon	0.75	1	0.6
MUC1 AND NOT-GAGE1	Colon	0.75	1	0.6
ITGAV AND NOT-EPHA2	Glioma	0.8722	0.9063	0.8406
SLC39A6 AND NOT-EPHA2	Glioma	0.7656	0.8305	0.7101
SLC39A6 AND NOT-ERBB2	Glioma	0.7612	0.7846	0.7391
ITGAV AND NOT-ERBB2	Glioma	0.9209	0.9143	0.9275
ITGAV AND NOT-MUC1	Glioma	0.8489	0.8429	0.8551
EDNRB AND NOT-MUC1	Glioma	0.662	0.6438	0.6812
EDNRB AND NOT-ERBB2	Glioma	0.6963	0.7121	0.6812
EDNRB AND NOT-EPHA2	Glioma	0.6271	0.7551	0.5362
CLDN12 AND NOT-EPHA2	Glioma	0.6168	0.8684	0.4783
CLDN12 AND NOT-MUC1	Glioma	0.7193	0.9111	0.5942
SLC39A6 AND NOT-MUC1	Glioma	0.7647	0.7761	0.7536
CLDN12 AND NOT-ERBB2	Glioma	0.7304	0.913	0.6087
CD276 AND NOT-ERBB2	Glioma	0.6824	0.5743	0.8406
CD276 AND NOT-MUC1	Glioma	0.6237	0.4957	0.8406
CD33 AND NOT-CD79B	AML	0.8583	0.8856	0.8327
CD33 AND NOT-IL11RA	AML	0.8565	0.8957	0.8207
CD33 AND NOT-SLAMF7	AML	0.9206	0.917	0.9243
CD33 AND NOT-FCRL1	AML	0.903	0.8976	0.9084
CD33 AND NOT-MS4A1	AML	0.9134	0.9027	0.9243
CD33 AND NOT-FCRL5	AML	0.9116	0.8992	0.9243
CD33 AND NOT-CD22	AML	0.8944	0.863	0.9283
CD33 AND NOT-TNFRSF17	AML	0.9109	0.9055	0.9163
CD33 AND NOT-CD72	AML	0.8906	0.8593	0.9243
CD33 AND NOT-CLDN23	AML	0.8357	0.7573	0.9323
CD33 AND NOT-CD79A	AML	0.8657	0.814	0.9243
CD33 AND NOT-CD19	AML	0.8929	0.8893	0.8964
CD33 AND NOT-P2RX5	AML	0.7974	0.8578	0.745
CD33 AND NOT-STEAP2	AML	0.786	0.6772	0.9363
CD33 AND NOT-KDR	AML	0.782	0.6714	0.9363
CD33 AND NOT-ROR1	AML	0.7807	0.6695	0.9363
CD33 AND NOT-AXL	AML	0.78	0.6705	0.9323
CD33 AND NOT-STEAP1	AML	0.7798	0.6744	0.9243
CD33 AND NOT-CLDN5	AML	0.7769	0.6638	0.9363
CD33 AND NOT-CLDN11	AML	0.7756	0.662	0.9363
CD33 AND NOT-EDNRB	AML	0.7752	0.6696	0.9203
CD33 AND NOT-TNFRSF10A	AML	0.7655	0.6474	0.9363
CD33 AND NOT-ERBB2	AML	0.763	0.6438	0.9363
CD33 AND NOT-CD276	AML	0.7629	0.6534	0.9163
CD33 AND NOT-ENG	AML	0.7492	0.6519	0.8805
CD33 AND NOT-VCAM1	AML	0.7483	0.6596	0.8645
CD33 AND NOT-MUC1	AML	0.73	0.6275	0.8725
CD33 AND NOT-IL13RA1	AML	0.6559	0.6667	0.6454
CD33 AND NOT-CD52	AML	0.6166	0.8815	0.4741
SLC34A2 AND NOT-ERBB2	Lung Adenocarcinoma	0.625	0.5556	0.7143
CD19 AND NOT-CD33	B-Cell Diffuse	0.6667	0.8077	0.5676
MS4A1 AND NOT-ITGB3	B-Cell Diffuse	0.6575	0.6667	0.6486
CD19 AND NOT-NCAM1	B-Cell Diffuse	0.6452	0.8	0.5405
CD19 AND NOT-ITGB3	B-Cell Diffuse	0.6207	0.8571	0.4865
MS4A1 AND NOT-NCAM1	B-Cell Diffuse	0.6111	0.6286	0.5946
CD19 AND NOT-SSTR2	Mantle-Cell Lymphoma	0.9143	1	0.8421
CD19 AND NOT-TNFRSF17	Mantle-Cell Lymphoma	0.8986	1	0.8158
CD19 AND NOT-IL13RA1	Mantle-Cell Lymphoma	0.8986	1	0.8158
CD19 AND NOT-CLDN8	Mantle-Cell Lymphoma	0.9143	1	0.8421
CD19 AND NOT-TRPM4	Mantle-Cell Lymphoma	0.8986	1	0.8158
CD19 AND NOT-SSTR3	Mantle-Cell Lymphoma	0.9143	1	0.8421
CD19 AND NOT-MST1R	Mantle-Cell Lymphoma	0.8657	1	0.7632
CD19 AND NOT-SLAMF7	Mantle-Cell Lymphoma	0.8986	1	0.8158
CD19 AND NOT-CD180	Mantle-Cell Lymphoma	0.9143	1	0.8421
CD19 AND NOT-CLDN6	Mantle-Cell Lymphoma	0.8485	1	0.7368
CD19 AND NOT-CD72	Mantle-Cell Lymphoma	0.6897	1	0.5263
CD19 AND NOT-FCRL1	Mantle-Cell Lymphoma	0.6667	1	0.5
MLANA AND NOT-EPCAM	Melanoma	1	1	1
MLANA AND NOT-SSTR1	Melanoma	1	1	1
MLANA AND NOT-LGR5	Melanoma	1	1	1
MLANA AND NOT-CD34	Melanoma	0.9524	1	0.9091
MLANA AND NOT-GPA33	Melanoma	0.9524	1	0.9091
MLANA AND NOT-EGFR	Melanoma	1	1	1
MLANA AND NOT-ENPP3	Melanoma	0.9524	1	0.9091
MLANA AND NOT-SSTR3	Melanoma	1	1	1
MLANA AND NOT-SSTR5	Melanoma	1	1	1
MLANA AND NOT-CLDN5	Melanoma	0.9524	1	0.9091
MLANA AND NOT-VTCN1	Melanoma	1	1	1
MLANA AND NOT-ULBP1	Melanoma	1	1	1
MLANA AND NOT-ERBB2	Melanoma	1	1	1

FIG. 10 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
FOLH1 AND NOT-ATP10B	Prostate	0.85714	1	0.75
FOLH1 AND NOT-MAL	Prostate	0.66667	1	0.5
FOLH1 AND NOT-GRIK5	Prostate	0.85714	1	0.75
FOLH1 AND NOT-SLC34A3	Prostate	0.66667	1	0.5
FOLH1 AND NOT-SLC5A5	Prostate	0.66667	1	0.5
FOLH1 AND NOT-GAL3ST1	Prostate	0.75	0.75	0.75
FOLH1 AND NOT-CDH19	Prostate	0.85714	1	0.75
FOLH1 AND NOT-STX1B	Prostate	0.8	0.6667	1
FOLH1 AND NOT-GPRC5B	Prostate	0.66667	0.6	0.75
FOLH1 AND NOT-ATP6AP2	Prostate	0.66667	1	0.5
FOLH1 AND NOT-SLC4A1	Prostate	0.85714	1	0.75
FOLH1 AND NOT-OR10H3	Prostate	0.66667	1	0.5
FOLH1 AND NOT-OR51B2	Prostate	0.88889	0.8	1
FOLH1 AND NOT-OR7A10	Prostate	0.66667	0.6	0.75
FGFR1 AND NOT-B4GALNT1	Sarcoma	0.66667	0.9	0.5294
MMP14 AND NOT-B4GALNT1	Sarcoma	0.74074	1	0.5882
FLRT2 AND NOT-B4GALNT1	Sarcoma	0.73333	0.8462	0.6471
VASN AND NOT-B4GALNT1	Sarcoma	0.69231	1	0.5294
BFAR AND NOT-B4GALNT1	Sarcoma	0.73333	0.8462	0.6471
PCDHB14 AND NOT-B4GALNT1	Sarcoma	0.75862	0.9167	0.6471
NOT-KIAA0754 AND MSLN	Colon	0.66667	0.75	0.6
NOT-CLEC2D AND MSLN	Colon	0.66667	0.75	0.6
NOT-S1PR2 AND ERBB2	Colon	0.75	1	0.6
NOT-PLXNA4 AND ERBB2	Colon	0.75	1	0.6
NOT-CD200R1 AND ERBB2	Colon	0.75	1	0.6
NOT-CDH26 AND ERBB2	Colon	0.75	1	0.6
NOT-SLC6A2 AND ERBB2	Colon	0.75	1	0.6
NOT-ZP2 AND MSLN	Colon	0.66667	0.75	0.6
NOT-NAALAD2 AND MSLN	Colon	0.66667	0.75	0.6
NOT-TACR1 AND ERBB2	Colon	0.75	1	0.6
NOT-SCN4B AND ERBB2	Colon	0.75	1	0.6
NOT-PTPRS AND ERBB2	Colon	0.75	1	0.6
NOT-GRID1 AND MUC1	Colon	0.75	1	0.6
NOT-DIO3 AND ERBB2	Colon	0.75	1	0.6
NOT-KCNH8 AND ERBB2	Colon	0.75	1	0.6
NOT-PCDH9 AND MSLN	Colon	0.66667	0.75	0.6
NOT-GPR37 AND ERBB2	Colon	0.6	0.6	0.6
NOT-GPC3 AND EFNB1	Colon	0.6	0.6	0.6
NOT-GPC3 AND ANTXR1	Colon	0.75	1	0.6
NOT-C1orf210 AND MUC1	Glioma	0.82051	0.7356	0.9275
NOT-WLS AND CD33	AML	0.91585	0.9	0.9323
NOT-HEG1 AND CD33	AML	0.90083	0.9356	0.8685
NOT-TGFBR3 AND CD33	AML	0.89669	0.9313	0.8645
NOT-PTPRK AND CD33	AML	0.88645	0.8203	0.9641
NOT-ACVR2A AND CD33	AML	0.88469	0.8417	0.9323
NOT-PERP AND CD33	AML	0.86176	0.7843	0.9562
NOT-ATP9A AND CD33	AML	0.88258	0.8412	0.9283
NOT-PODXL AND CD33	AML	0.84283	0.7439	0.9721
NOT-BMPR1A AND CD33	AML	0.83871	0.7622	0.9323
NOT-CLCA4 AND MUC1	Lung Adenocarcinoma	0.60748	0.4514	0.9286
NOT-LRP4 AND B4GALNT1	Neuroblastoma	0.77019	0.8052	0.7381
NOT-SLC2A12 AND B4GALNT1	Neuroblastoma	0.73684	0.7241	0.75
NOT-SDC4 AND B4GALNT1	Neuroblastoma	0.66272	0.6588	0.6667
NOT-KCNK1 AND B4GALNT1	Neuroblastoma	0.62687	0.84	0.5
EPCAM AND NOT-CDH17	Breast	0.70588	0.75	0.6667
EPCAM AND NOT-PTPRR	Breast	0.71795	0.6667	0.7778
EPCAM AND NOT-SCARA5	Breast	0.62712	0.5781	0.6852
EPCAM AND NOT-SLC30A10	Breast	0.63636	0.5385	0.7778
EPCAM AND NOT-GRM7	Breast	0.62774	0.5181	0.7963
EPCAM AND NOT-FUT3	Breast	0.69091	0.6786	0.7037
ENPP1 AND NOT-TYR	Breast	0.65079	0.5694	0.7593
EPCAM AND NOT-ABHD3	Breast	0.64179	0.5375	0.7963
EPCAM AND NOT-DPP10	Breast	0.61069	0.5195	0.7407
LRRC8E AND NOT-TYR	Breast	0.68333	0.6212	0.7593
EPCAM AND NOT-MUC12	Breast	0.65152	0.5513	0.7963
EPCAM AND NOT-BEST2	Breast	0.64179	0.5375	0.7963
EPCAM AND NOT-NOX1	Breast	0.65152	0.5513	0.7963
EPCAM AND NOT-TMPRSS11E	Breast	0.61538	0.5263	0.7407

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
MLANA AND NOT-CLDN1	Melanoma	1	1	1
MLANA AND NOT-TNFRSF13C	Melanoma	1	1	1
MLANA AND NOT-SLC34A2	Melanoma	1	1	1
MLANA AND NOT-CLDN8	Melanoma	1	1	1
PMEL AND NOT-EPCAM	Melanoma	0.9524	1	0.9091
PMEL AND NOT-ENPP3	Melanoma	0.9524	1	0.9091
PMEL AND NOT-SSTR1	Melanoma	0.9524	1	0.9091
PMEL AND NOT-VTCN1	Melanoma	0.9524	1	0.9091
PMEL AND NOT-LGR5	Melanoma	0.9524	1	0.9091
PMEL AND NOT-CLDN1	Melanoma	0.9524	1	0.9091
PMEL AND NOT-ERBB2	Melanoma	0.9524	1	0.9091
PMEL AND NOT-CD34	Melanoma	0.9524	1	0.9091
PMEL AND NOT-CLDN7	Melanoma	0.9524	1	0.9091
MLANA AND NOT-SSTR4	Melanoma	1	1	1
MLANA AND NOT-ENG	Melanoma	1	1	1
MLANA AND NOT-RAET1E	Melanoma	0.9524	1	0.9091
MLANA AND NOT-MST1R	Melanoma	1	1	1
MLANA AND NOT-CLDN9	Melanoma	1	1	1
MLANA AND NOT-CLDN2	Melanoma	1	1	1
MLANA AND NOT-CLDN6	Melanoma	1	1	1
MLANA AND NOT-GUCY2C	Melanoma	0.9524	1	0.9091
MLANA AND NOT-FOLR1	Melanoma	1	1	1
MLANA AND NOT-CLDN7	Melanoma	1	1	1
MLANA AND NOT-IL20RA	Melanoma	0.9524	1	0.9091
PMEL AND NOT-SLC34A2	Melanoma	0.9524	1	0.9091
PMEL AND NOT-CLDN23	Melanoma	0.9524	1	0.9091
PMEL AND NOT-CLDN8	Melanoma	0.9524	1	0.9091
MLANA AND NOT-MUC13	Melanoma	1	1	1
MLANA AND NOT-STEAP2	Melanoma	0.9524	1	0.9091
MLANA AND NOT-CLDN11	Melanoma	0.9	1	0.8182
MLANA AND NOT-CLDN18	Melanoma	0.9	1	0.8182
MLANA AND NOT-AXL	Melanoma	0.9	1	0.8182
EDNRB AND NOT-KDR	Melanoma	0.8421	1	0.7273
ABCB5 AND NOT-B4GALNT1	Melanoma	0.7059	1	0.5455
ABCB5 AND NOT-KDR	Melanoma	0.625	1	0.4545
CLDN12 AND NOT-B4GALNT1	Melanoma	0.625	1	0.4545
RNF43 AND NOT-B4GALNT1	Melanoma	0.625	1	0.4545
L1CAM AND NOT-B4GALNT1	Neuroblastoma	0.6364	0.5147	0.8333
B4GALNT1 AND NOT-ERBB4	Neuroblastoma	0.6043	0.7636	0.5
MUC16 AND NOT-ERBB2	Ovarian	0.6957	0.7273	0.6667
FOLH1 AND NOT-FCRL5	Prostate	0.6667	1	0.5
FOLH1 AND NOT-TNFRSF13C	Prostate	0.6667	1	0.5
FOLH1 AND NOT-ERBB3	Prostate	0.6	0.5	0.75
FOLH1 AND NOT-LGR5	Prostate	0.8889	0.8	1
FOLH1 AND NOT-GPA33	Prostate	0.6	0.5	0.75
FOLH1 AND NOT-ALK	Prostate	0.6667	1	0.5
FOLH1 AND NOT-NCAM1	Prostate	0.6154	0.4444	1
NOT-ERBB4 AND MUC1	Colon	0.6	0.6	0.6
NOT-ITGB3 AND ERBB2	Colon	0.6667	0.75	0.6
NOT-ERBB4 AND ERBB2	Colon	0.6	0.6	0.6
NOT-GPC3 AND CD276	Colon	0.75	1	0.6
NOT-GPC3 AND IL20RA	Colon	0.6667	0.5	1
NOT-GAGE1 AND MSLN	Colon	0.6667	0.75	0.6
NOT-GPC3 AND PTK7	Colon	0.9091	0.8333	1
NOT-EPHA2 AND CD276	Glioma	0.6824	0.5743	0.8406
NOT-SLAMF7 AND MUC1	Glioma	0.6256	0.4648	0.9565
NOT-MUC1 AND PROM1	Glioma	0.6065	0.5465	0.6812
NOT-RAET1E AND MUC1	Glioma	0.6738	0.5339	0.913
NOT-GPA33 AND MUC1	Glioma	0.6979	0.5447	0.971
NOT-ITGAV AND CD33	AML	0.8119	0.7622	0.8685
NOT-GPNMB AND CD33	AML	0.7606	0.6627	0.8924
NOT-CLDN12 AND CD33	AML	0.7427	0.6281	0.9084
NOT-SLC34A2 AND KDR	Melanoma	1	1	1
NOT-VTCN1 AND KDR	Melanoma	0.9	1	0.8182
NOT-CLDN7 AND KDR	Melanoma	0.8696	0.8333	0.9091
NOT-TNFRSF13C AND KDR	Melanoma	0.8421	1	0.7273
NOT-CA9 AND KDR	Melanoma	0.7778	1	0.6364
NOT-FCRL5 AND KDR	Melanoma	0.625	1	0.4545

FIG. 11

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
PTPRZ1 AND ITGB8	Astrocytoma	0.904762	1	0.826087
PTPRZ1 AND ITM2B	Astrocytoma	0.876404	0.906977	0.847826
PTPRZ1 AND MPZL1	Astrocytoma	0.904762	1	0.826087
PTPRZ1 AND SLC44A2	Astrocytoma	0.857143	0.866667	0.847826
PTPRZ1 AND CSF1R	Astrocytoma	0.847826	0.847826	0.847826
PLP1 AND F2RL1	Astrocytoma	0.843373	0.945946	0.76087
PTPRZ1 AND ABCA1	Astrocytoma	0.835443	1	0.717391
PTPRZ1 AND NRCAM	Astrocytoma	0.833333	0.8	0.869565
PTPRZ1 AND ATP8B4	Astrocytoma	0.829268	0.944444	0.73913
PTPRZ1 AND CLCN5	Astrocytoma	0.828283	0.773585	0.891304
PTPRZ1 AND BMP2	Astrocytoma	0.827586	0.878049	0.782609
PTPRZ1 AND VANGL2	Astrocytoma	0.826087	0.826087	0.826087
PLP1 AND ABCA1	Astrocytoma	0.825	0.970588	0.717391
PTPRZ1 AND CD320	Astrocytoma	0.822222	0.840909	0.804348
PTPRZ1 AND CSF2RB	Astrocytoma	0.820513	1	0.695652
PTPRZ1 AND ADAM28	Astrocytoma	0.820513	1	0.695652
PTPRZ1 AND ANTXR1	Astrocytoma	0.819277	0.918919	0.73913
PTPRZ1 AND ADAM17	Astrocytoma	0.818182	0.857143	0.782609
PTPRZ1 AND DDR2	Astrocytoma	0.817204	0.808511	0.826087
PTPRZ1 AND LRRC8D	Astrocytoma	0.816327	0.769231	0.869565
PTPRZ1 AND DLL1	Astrocytoma	0.818182	0.857143	0.782609
PTPRZ1 AND AGTRAP	Astrocytoma	0.814815	0.942857	0.717391
PTPRZ1 AND CD68	Astrocytoma	0.814815	0.942857	0.717391
NRCAM AND TSPAN11	Astrocytoma	0.813953	0.875	0.76087
PTPRZ1 AND MACF1	Astrocytoma	0.813187	0.822222	0.804348
PTPRZ1 AND BMPR1A	Astrocytoma	0.811881	0.745455	0.891304
AQP4 AND VANGL2	Astrocytoma	0.808989	0.837209	0.782609
PTPRZ1 AND BVES	Astrocytoma	0.808989	0.837209	0.782609
PLP1 AND MPZL1	Astrocytoma	0.804878	0.916667	0.717391
NRCAM AND F2RL1	Astrocytoma	0.804348	0.804348	0.804348
PTPRZ1 AND PCDHB2	Astrocytoma	0.804124	0.764706	0.847826
SLC1A3 AND VANGL2	Astrocytoma	0.804124	0.764706	0.847826
FAIM2 AND VANGL2	Astrocytoma	0.8	0.818182	0.782609
PTPRZ1 AND SLCO2B1	Astrocytoma	0.8	0.941176	0.695652
NRCAM AND HAS2	Astrocytoma	0.8	0.871795	0.73913
FAIM2 AND F2RL1	Astrocytoma	0.8	0.871795	0.73913
AQP4 AND TSPAN11	Astrocytoma	0.8	0.871795	0.73913
PTPRZ1 AND TSPAN11	Astrocytoma	0.8	0.871795	0.73913
NRCAM AND VANGL2	Astrocytoma	0.8	0.871795	0.73913
PTPRZ1 AND CACNA1F	Astrocytoma	0.795918	0.75	0.847826
APLP1 AND VANGL2	Astrocytoma	0.795918	0.75	0.847826
PTPRZ1 AND MERTK	Astrocytoma	0.795699	0.787234	0.804348
NRCAM AND BMPR1A	Astrocytoma	0.795455	0.833333	0.76087
PTPRZ1 AND TSPAN12	Astrocytoma	0.795455	0.833333	0.76087
PTPRZ1 AND IL12RB1	Astrocytoma	0.795455	0.833333	0.76087
PTPRZ1 AND ATP13A4	Astrocytoma	0.794872	0.96875	0.673913
ADAM22 AND SLC3A2	Astrocytoma	0.791209	0.8	0.782609
PTPRZ1 AND CSF1	Astrocytoma	0.791209	0.8	0.782609
NRCAM AND TSPAN12	Astrocytoma	0.790698	0.85	0.73913
PTPRZ1 AND ADAM9	Astrocytoma	0.790698	0.85	0.73913
PTPRZ1 AND FCGRT	Astrocytoma	0.789474	1	0.652174
PTPRZ1 AND CLEC4A	Astrocytoma	0.789474	1	0.652174
GPM6B AND F2RL1	Astrocytoma	0.789474	1	0.652174
PTPRZ1 AND PANX1	Astrocytoma	0.788462	0.706897	0.891304
PTPRZ1 AND C5AR1	Astrocytoma	0.787234	0.770833	0.804348
PLP1 AND VANGL2	Astrocytoma	0.786517	0.813953	0.76087
NRCAM AND ABCC4	Astrocytoma	0.785714	0.868421	0.717391
CADM2 AND F2RL1	Astrocytoma	0.785714	0.868421	0.717391
GPM6B AND HAS2	Astrocytoma	0.78481	0.939394	0.673913
PTPRZ1 AND SLC1A3	Astrocytoma	0.784314	0.714286	0.869565
PTPRZ1 AND ATP13A1	Astrocytoma	0.784314	0.714286	0.869565
PTPRZ1 AND KCNJ8	Astrocytoma	0.784314	0.714286	0.869565
ADAM22 AND VANGL2	Astrocytoma	0.808989	0.837209	0.782609
PTPRZ1 AND ATP6AP2	Astrocytoma	0.783505	0.745098	0.826087
PCDH17 AND TSPAN11	Astrocytoma	0.782609	0.782609	0.782609
PTPRZ1 AND HTRA2	Astrocytoma	0.782609	0.782609	0.782609
PTPRZ1 AND CCR5	Astrocytoma	0.789474	1	0.652174
PLP1 AND TSPAN12	Astrocytoma	0.781609	0.829268	0.73913
PTPRZ1 AND P2RY1	Astrocytoma	0.780952	0.694915	0.891304
PTPRZ1 AND CLEC1A	Astrocytoma	0.780952	0.694915	0.891304
PTPRZ1 AND TPCN2	Astrocytoma	0.780952	0.694915	0.891304
PTPRZ1 AND TMEM119	Astrocytoma	0.779221	0.967742	0.652174
PTPRZ1 AND SRR	Astrocytoma	0.778947	0.755102	0.804348
ADAM22 AND TSPAN12	Astrocytoma	0.777778	0.795455	0.76087
PLP1 AND SNAP23	Astrocytoma	0.777778	0.795455	0.76087
ADAM12 AND PMEPA1	Breast	0.857143	0.882353	0.833333
ADAM12 AND JAG2	Breast	0.854167	0.97619	0.759259

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
STEAP2 AND STEAP4	Prostate	0.857143	1	0.75
STEAP1 AND CDH10	Prostate	0.857143	1	0.75
STEAP2 AND ADAM2	Prostate	0.857143	1	0.75
STEAP2 AND ATP8A2	Prostate	0.75	0.75	0.75
STEAP2 AND CSPG5	Prostate	0.857143	1	0.75
STEAP2 AND FLVCR1	Prostate	0.8	0.666667	1
STEAP1 AND GABRG3	Prostate	0.75	0.75	0.75
STEAP1 AND PCDH15	Prostate	0.75	0.75	0.75
STEAP1 AND TMEFF2	Prostate	0.75	0.75	0.75
STEAP1 AND PCDH8	Prostate	0.75	0.75	0.75
STEAP1 AND TTYH3	Prostate	0.75	0.75	0.75
STEAP1 AND NKAIN1	Prostate	0.857143	1	0.75
STEAP2 AND IL15RA	Prostate	0.727273	0.571429	1
STEAP2 AND UMODL1	Prostate	0.727273	0.571429	1
STEAP2 AND NMBR	Prostate	0.727273	0.571429	1
STEAP2 AND PCDHB1	Prostate	0.727273	0.571429	1
STEAP2 AND GRM6	Prostate	0.727273	0.571429	1
STEAP2 AND GRM5	Prostate	0.727273	0.571429	1
STEAP2 AND TAS1R1	Prostate	0.727273	0.571429	1
STEAP2 AND CHRNB4	Prostate	0.727273	0.571429	1
STEAP2 AND ATP6V0A4	Prostate	0.727273	0.571429	1
STEAP2 AND SLC13A2	Prostate	0.727273	0.571429	1
STEAP2 AND ZACN	Prostate	0.727273	0.571429	1
STEAP2 AND CHRNG	Prostate	0.727273	0.571429	1
STEAP2 AND KCNA7	Prostate	0.727273	0.571429	1
STEAP2 AND ATP13A5	Prostate	0.727273	0.571429	1
STEAP2 AND GLRA3	Prostate	0.727273	0.571429	1
STEAP2 AND AGTR2	Prostate	0.727273	0.571429	1
STEAP2 AND KCNQ2	Prostate	0.727273	0.571429	1
STEAP2 AND HTR5A	Prostate	0.727273	0.571429	1
STEAP2 AND DSC1	Prostate	0.727273	0.571429	1
STEAP2 AND HTR1F	Prostate	0.727273	0.571429	1
STEAP2 AND ACSL6	Prostate	0.727273	0.571429	1
STEAP2 AND TTYH3	Prostate	0.727273	0.571429	1
STEAP2 AND SLC9B1	Prostate	0.727273	0.571429	1
STEAP2 AND OR8B2	Prostate	0.727273	0.571429	1
STEAP2 AND ADAM21	Prostate	0.727273	0.571429	1
STEAP2 AND GPR78	Prostate	0.727273	0.571429	1
STEAP2 AND GHSR	Prostate	0.727273	0.571429	1
STEAP2 AND GHRHR	Prostate	0.727273	0.571429	1
STEAP2 AND PCDHA9	Prostate	0.727273	0.571429	1
STEAP2 AND CCR9	Prostate	0.727273	0.571429	1
STEAP2 AND KCNK16	Prostate	0.727273	0.571429	1
STEAP2 AND PRND	Prostate	0.727273	0.571429	1
STEAP2 AND KIRREL2	Prostate	0.727273	0.571429	1
STEAP2 AND CD74	Prostate	0.727273	0.571429	1
STEAP2 AND CD44	Prostate	0.727273	0.571429	1
STEAP2 AND MEGF11	Prostate	0.727273	0.571429	1
STEAP2 AND TMPRSS11D	Prostate	0.727273	0.571429	1
STEAP2 AND CNTNAP2	Prostate	0.727273	0.571429	1
STEAP2 AND GNRHR	Prostate	0.727273	0.571429	1
STEAP2 AND EPHA10	Prostate	0.727273	0.571429	1
FAP AND TGFBI	Sarcoma	0.785714	1	0.647059
FAP AND LAPTM5	Sarcoma	0.827586	1	0.705882
FAP AND CD163	Sarcoma	0.827586	1	0.705882
TGFBI AND ULBP2	Sarcoma	0.692308	1	0.529412
FAP AND SLC1A5	Sarcoma	0.692308	1	0.529412
FAP AND SLC31A1	Sarcoma	0.785714	1	0.647059
CDH17 AND SPON2	Stomach	0.896552	0.886364	0.906977
CDH17 AND THY1	Stomach	0.767677	0.678571	0.883721
CDH17 AND CLDN1	Stomach	0.921348	0.891304	0.953488
CLDN18 AND FAT1	Stomach	0.864198	0.921053	0.813953
CLDN18 AND CDH17	Stomach	0.897436	1	0.813953
CLDN18 AND SLC2A1	Stomach	0.847059	0.857143	0.837209
CLDN18 AND TTYH3	Stomach	0.843373	0.875	0.813953
CLDN18 AND PCDHA6	Stomach	0.747253	0.708333	0.790698
CLDN18 AND ATP4B	Stomach	0.769231	0.729167	0.813953
CDH17 AND PSCA	Stomach	0.739726	0.9	0.627907
CLDN18 AND ABCG5	Stomach	0.786517	0.76087	0.813953
CLDN18 AND VANGL1	Stomach	0.875	0.945946	0.813953
MUC13 AND BST2	Stomach	0.837838	1	0.72093
CLDN18 AND F2R	Stomach	0.86747	0.9	0.837209
CLDN18 AND RXFP3	Stomach	0.738095	0.756098	0.72093
CLDN18 AND FLVCR1	Stomach	0.809524	0.829268	0.790698
CLDN18 AND LRIG3	Stomach	0.860759	0.944444	0.790698
CLDN18 AND ABHD3	Stomach	0.864198	0.921053	0.813953
CLDN18 AND ATP8B1	Stomach	0.837838	1	0.72093

FIG. 11 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
ADAM12 AND CELSR2	Breast	0.851485	0.914894	0.796296
ADAM12 AND TSPAN17	Breast	0.851064	1	0.740741
ADAM12 AND CNNM4	Breast	0.851064	1	0.740741
ADAM12 AND SLC9A7	Breast	0.851485	0.914894	0.796296
ADAM12 AND CX3CL1	Breast	0.849057	0.865385	0.833333
ADAM12 AND CD9	Breast	0.848485	0.933333	0.777778
ADAM12 AND UNC5B	Breast	0.842105	0.97561	0.740741
ADAM12 AND CELSR1	Breast	0.841121	0.849057	0.833333
ADAM12 AND SLC12A7	Breast	0.83871	1	0.722222
ADAM12 AND KIAA1324	Breast	0.834951	0.877551	0.796296
ADAM12 AND GJB2	Breast	0.834951	0.877551	0.796296
ADAM12 AND CXADR	Breast	0.826923	0.86	0.796296
ADAM12 AND SLC27A1	Breast	0.824742	0.930233	0.740741
ADAM12 AND ACVR1B	Breast	0.823529	0.875	0.777778
ADAM12 AND ST14	Breast	0.821053	0.95122	0.722222
ADAM12 AND PROM2	Breast	0.821053	0.95122	0.722222
ADAM12 AND CCR5	Breast	0.818182	0.803571	0.833333
ADAM12 AND ABCA3	Breast	0.817204	0.974359	0.703704
ADAM12 AND LSR	Breast	0.817204	0.974359	0.703704
ADAM12 AND LAX1	Breast	0.807339	0.8	0.814815
ADAM12 AND ITGA11	Breast	0.804348	0.973684	0.685185
ADAM12 AND PLXDC1	Breast	0.804124	0.906977	0.722222
ADAM12 AND ITGAX	Breast	0.8	0.926829	0.703704
ADAM12 AND IL17RB	Breast	0.8	0.869565	0.740741
FGG AND SLC26A6	Liver	0.923077	0.857143	1
FGG AND LRRC8D	Liver	0.909091	1	0.833333
FGG AND SLC38A6	Liver	0.909091	1	0.833333
FGG AND HM13	Liver	1	1	1
FGG AND SLC39A1	Liver	0.909091	1	0.833333
FGG AND NCSTN	Liver	1	1	1
FGG AND C6orf89	Liver	1	1	1
FGG AND LRP11	Liver	1	1	1
FGG AND ATP2A2	Liver	0.909091	1	0.833333
FGG AND ADCY6	Liver	0.909091	1	0.833333
FGG AND FGFRL1	Liver	1	1	1
FGG AND CELSR3	Liver	0.857143	0.75	1
FGG AND ANO10	Liver	0.909091	1	0.833333
FGG AND NIPA1	Liver	0.909091	1	0.833333
FGG AND SLC17A5	Liver	0.909091	1	0.833333
HPN AND SLC26A6	Liver	1	1	1
TFR2 AND SLC26A6	Liver	0.833333	0.833333	0.833333
FGG AND TSPAN17	Liver	0.909091	1	0.833333
FGG AND CLN3	Liver	0.909091	1	0.833333
FGG AND NOX4	Liver	0.923077	0.857143	1
FGG AND SLC26A11	Liver	0.909091	1	0.833333
FGG AND NPBWR2	Liver	0.909091	1	0.833333
FGG AND ABCB8	Liver	0.857143	0.75	1
FGG AND PANX2	Liver	1	1	1
FGG AND PIGU	Liver	0.8	1	0.666667
FGG AND PHEX	Liver	1	1	1
FGG AND TMEM8B	Liver	0.8	1	0.666667
FGG AND PDCD1	Liver	0.909091	1	0.833333
FGG AND MRAP2	Liver	0.8	0.666667	1
FGG AND YIPF3	Liver	0.909091	1	0.833333
FGG AND SPPL3	Liver	0.8	1	0.666667
FGG AND CNNM3	Liver	0.8	0.666667	1
FGG AND OPRD1	Liver	1	1	1
FGG AND ADORA2B	Liver	0.909091	1	0.833333
FGG AND IGF2R	Liver	0.909091	1	0.833333
FGG AND ATP9B	Liver	0.8	1	0.666667
ASGR1 AND SMO	Liver	0.8	1	0.666667
FGG AND MMP14	Liver	1	1	1
FGG AND SLC5A5	Liver	0.909091	1	0.833333
FGG AND SLC2A5	Liver	0.909091	1	0.833333
FGG AND SMO	Liver	0.8	1	0.666667
FGG AND STIM1	Liver	0.909091	1	0.833333
FGG AND STX4	Liver	0.8	1	0.666667
FGG AND TMEM67	Liver	0.8	0.666667	1
SLC27A5 AND SLC26A6	Liver	0.923077	0.857143	1
FGG AND PLXNC1	Liver	0.8	1	0.666667
FGG AND MFSD5	Liver	0.8	1	0.666667
FGG AND DAGLB	Liver	0.8	1	0.666667
FGG AND TMPRSS5	Liver	0.8	0.666667	1
FGG AND KCNG3	Liver	0.8	0.666667	1
FGG AND SLC4A11	Liver	0.833333	0.833333	0.833333
FGG AND CRHR1	Liver	0.8	1	0.666667
SLC39A14 AND OLR1	Pancreas	0.736842	0.875	0.636364

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
CLDN18 AND KCNC2	Stomach	0.767442	0.767442	0.767442
CDH17 AND FAP	Stomach	0.837209	0.837209	0.837209
CLDN18 AND KCNK12	Stomach	0.759494	0.833333	0.697674
CLDN18 AND ENPP1	Stomach	0.755556	0.723404	0.790698
CLDN18 AND LHFPL5	Stomach	0.765957	0.705882	0.837209
MUC13 AND ESAM	Stomach	0.851064	0.784314	0.930233
MUC13 AND F2R	Stomach	0.857143	0.763636	0.976744
CLDN18 AND GJB4	Stomach	0.704545	0.688889	0.72093
CLDN18 AND PPAPDC1B	Stomach	0.847059	0.857143	0.837209
MST1R AND BST2	Stomach	0.837838	1	0.72093
CDH17 AND SLC7A5	Stomach	0.72549	0.627119	0.860465
CLDN18 AND GRIK4	Stomach	0.781609	0.772727	0.790698
CLDN18 AND CDH20	Stomach	0.765432	0.815789	0.72093
CLDN18 AND MLNR	Stomach	0.771084	0.8	0.744186
CLDN18 AND PCDHAC2	Stomach	0.795455	0.777778	0.813953
CLDN18 AND CACNA1B	Stomach	0.758621	0.75	0.767442
CLDN18 AND LRRC55	Stomach	0.782609	0.734694	0.837209
CLDN18 AND LCT	Stomach	0.765957	0.705882	0.837209
CLDN18 AND CACNG6	Stomach	0.765957	0.705882	0.837209
CLDN18 AND PAQR8	Stomach	0.837838	1	0.72093
CLDN18 AND LRFN2	Stomach	0.8	0.809524	0.790698
CLDN18 AND DIO1	Stomach	0.705882	0.714286	0.697674
CLDN18 AND GALR2	Stomach	0.790123	0.842105	0.744186
CLDN18 AND OPRM1	Stomach	0.777778	0.744681	0.813953
CLDN18 AND KCNH1	Stomach	0.755556	0.723404	0.790698
CLDN18 AND ABCA12	Stomach	0.777778	0.744681	0.813953
CLDN18 AND OR52D1	Stomach	0.764045	0.73913	0.790698
CLDN18 AND GPR135	Stomach	0.808989	0.782609	0.837209
CLDN18 AND CALHM1	Stomach	0.752688	0.7	0.813953
CLDN18 AND FSHR	Stomach	0.725275	0.6875	0.767442
CEACAM5 AND BST2	Stomach	0.771429	1	0.627907
CLDN18 AND GJA3	Stomach	0.771084	0.8	0.744186
CLDN18 AND GPR158	Stomach	0.755556	0.723404	0.790698
ATP8B1 AND THY1	Stomach	0.871795	0.971429	0.790698
CLDN18 AND SLC22A11	Stomach	0.712644	0.704545	0.72093
CLDN18 AND P2RY4	Stomach	0.765957	0.705882	0.837209
CLDN18 AND ABCG4	Stomach	0.853659	0.897436	0.813953
CLDN18 AND UMOD	Stomach	0.719101	0.695652	0.744186
CLDN18 AND CEACAM7	Stomach	0.813953	0.813953	0.813953
MUC17 AND CLDN1	Stomach	0.853333	1	0.744186
MUC13 AND IFI6	Stomach	0.729412	0.738095	0.72093
CLDN18 AND CACNG8	Stomach	0.818182	0.8	0.837209
CLDN18 AND CLRN1	Stomach	0.76087	0.714286	0.813953
CLDN18 AND CACNG2	Stomach	0.75	0.733333	0.767442
CLDN18 AND NPHS1	Stomach	0.791209	0.75	0.837209
CLDN18 AND GRIN2B	Stomach	0.755556	0.723404	0.790698
CLDN18 AND GABBR2	Stomach	0.711111	0.680851	0.744186
CLDN18 AND SLC17A1	Stomach	0.769231	0.729167	0.813953
CLDN18 AND CALY	Stomach	0.823529	0.833333	0.813953
CLDN18 AND SEZ6	Stomach	0.769231	0.729167	0.813953
PTPRZ1 AND CD33	Astrocytoma	0.829787	0.8125	0.847826
PTPRZ1 AND VCAM1	Astrocytoma	0.811881	0.745455	0.891304
PTPRZ1 AND CD38	Astrocytoma	0.764706	0.696429	0.847826
PTPRZ1 AND CD79A	Astrocytoma	0.761905	0.677966	0.869565
NCAM1 AND VANGL2	Astrocytoma	0.758621	0.804878	0.717391
PTPRZ1 AND EDNRB	Astrocytoma	0.758621	0.804878	0.717391
ADCYAP1R1 AND CD33	Astrocytoma	0.758621	0.804878	0.717391
NRCAM AND CD33	Astrocytoma	0.755102	0.711538	0.804348
PTPRZ1 AND MS4A1	Astrocytoma	0.75	0.672414	0.847826
PTPRZ1 AND NCAM1	Astrocytoma	0.745455	0.640625	0.891304
PTPRZ1 AND BMPR1B	Astrocytoma	0.733945	0.634921	0.869565
PTPRZ1 AND L1CAM	Astrocytoma	0.733945	0.634921	0.869565
PTPRZ1 AND P2RX5	Astrocytoma	0.728972	0.639344	0.847826
PTPRZ1 AND ITGB3	Astrocytoma	0.725275	0.733333	0.717391
PTPRZ1 AND TNFRSF10A	Astrocytoma	0.717391	0.717391	0.717391
PTPRZ1 AND GUCY2C	Astrocytoma	0.717391	0.717391	0.717391
NRCAM AND MUC1	Astrocytoma	0.716981	0.633333	0.826087
PTPRZ1 AND FCRL5	Astrocytoma	0.715789	0.693878	0.73913
PTPRZ1 AND FAP	Astrocytoma	0.712644	0.756098	0.673913
PTPRZ1 AND CLDN18	Astrocytoma	0.709677	0.702128	0.717391
PTPRZ1 AND FCRL2	Astrocytoma	0.709677	0.702128	0.717391
PTPRZ1 AND CLDN2	Astrocytoma	0.709677	0.702128	0.717391
NCAM1 AND ABCA1	Astrocytoma	0.708861	0.848485	0.608696
NCAM1 AND TSPAN11	Astrocytoma	0.708333	0.68	0.73913
PTPRZ1 AND TNFRSF13C	Astrocytoma	0.705882	0.769231	0.652174
PTPRZ1 AND CD276	Astrocytoma	0.704545	0.738095	0.673913
PTPRZ1 AND IL2RA	Astrocytoma	0.703297	0.711111	0.695652

FIG. 11 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
KCNE4 AND NTM	Pancreas	0.7	0.777778	0.636364
KCNE4 AND SYT13	Pancreas	0.695652	0.666667	0.727273
NOX4 AND TM4SF1	Pancreas	0.777778	1	0.636364
NOX4 AND ADAM28	Pancreas	0.705882	1	0.545455
NOX4 AND LITAF	Pancreas	0.705882	1	0.545455
TNFSF4 AND SYT13	Pancreas	0.736842	0.875	0.636364
KCNE4 AND MARCH1	Pancreas	0.666667	0.615385	0.727273
KCNE4 AND IGSF6	Pancreas	0.666667	0.615385	0.727273
KCNE4 AND FAM26F	Pancreas	0.666667	0.615385	0.727273
KCNE4 AND TACSTD2	Pancreas	0.666667	0.7	0.636364
KCNE4 AND TSPAN8	Pancreas	0.736842	0.875	0.636364
TREM2 AND TNFSF4	Pancreas	0.705882	1	0.545455
SYT13 AND ATP10D	Pancreas	0.727273	0.727273	0.727273
SYT13 AND LRRC32	Pancreas	0.72	0.642857	0.818182
KCNE4 AND SLC41A2	Pancreas	0.666667	0.857143	0.545455
KCNE4 AND GPRC5A	Pancreas	0.666667	0.857143	0.545455
NOX4 AND CXCR4	Pancreas	0.666667	0.857143	0.545455
KCNE4 AND SLC7A7	Pancreas	0.666667	0.615385	0.727273
KCNE4 AND CD48	Pancreas	0.666667	0.615385	0.727273
KCNE4 AND SLC44A3	Pancreas	0.695652	0.666667	0.727273
KCNE4 AND LSR	Pancreas	0.666667	0.857143	0.545455
TNFSF13B AND SYT13	Pancreas	0.842105	1	0.727273
TNFSF4 AND DSG2	Pancreas	0.736842	0.875	0.636364
VSIG1 AND GJA1	Pancreas	0.666667	0.857143	0.545455
NOX4 AND SLC38A5	Pancreas	0.695652	0.666667	0.727273
OSMR AND SYT13	Pancreas	0.666667	0.7	0.636364
KCNE4 AND ABCC8	Pancreas	0.736842	0.875	0.636364
GP2 AND TLR2	Pancreas	0.777778	1	0.636364
NTM AND DSG2	Pancreas	0.705882	1	0.545455
NOX4 AND GPRC5A	Pancreas	0.705882	1	0.545455
NOX4 AND PTPRH	Pancreas	0.736842	0.875	0.636364
TREM2 AND ADAM28	Pancreas	0.666667	0.857143	0.545455
KCNE4 AND ADAM12	Pancreas	0.666667	0.615385	0.727273
TM4SF1 AND SYT13	Pancreas	0.736842	0.875	0.636364
KCNE4 AND GABRP	Pancreas	0.695652	0.666667	0.727273
OLR1 AND ITGA11	Pancreas	0.736842	0.875	0.636364
SYT13 AND TSPAN4	Pancreas	0.761905	0.8	0.727273
TNFSF13B AND PPAP2C	Pancreas	0.705882	1	0.545455
SYT13 AND PKD2	Pancreas	0.64	0.571429	0.727273
KCNE4 AND TLR2	Pancreas	0.64	0.571429	0.727273
KCNE4 AND GPR65	Pancreas	0.64	0.571429	0.727273
GP2 AND OLR1	Pancreas	0.777778	1	0.636364
GP2 AND EDNRA	Pancreas	0.8	0.888889	0.727273
GP2 AND ITGB2	Pancreas	0.705882	1	0.545455
KCNE4 AND ITGA4	Pancreas	0.666667	0.615385	0.727273
GP2 AND NTM	Pancreas	0.777778	1	0.636364
SYT13 AND GLIPR1	Pancreas	0.952381	1	0.909091
KCNE4 AND SLC39A14	Pancreas	0.636364	0.636364	0.636364
NOX4 AND CD300LF	Pancreas	0.636364	0.636364	0.636364
KCNE4 AND ATP1A1	Pancreas	0.636364	0.636364	0.636364
KCNE4 AND PTPN2	Pancreas	0.666667	0.615385	0.727273
GP2 AND LRRC32	Pancreas	0.705882	1	0.545455
NOX4 AND VSIG1	Pancreas	0.7	0.777778	0.636364
KCNE4 AND TMEM30B	Pancreas	0.64	0.571429	0.727273
TNFSF13B AND GJB1	Pancreas	0.705882	1	0.545455
PDGFRB AND LITAF	Pancreas	0.705882	1	0.545455
SYT13 AND CXCR4	Pancreas	0.842105	1	0.727273
TM4SF1 AND ITGA11	Pancreas	0.631579	0.75	0.545455
NOX4 AND TREM1	Pancreas	0.64	0.571429	0.727273
NOX4 AND TNFRSF21	Pancreas	0.631579	0.75	0.545455
KCNE4 AND SLC1A1	Pancreas	0.727273	0.727273	0.727273
TNFSF4 AND GPRC5A	Pancreas	0.705882	1	0.545455
GP2 AND AMIGO2	Pancreas	0.842105	1	0.727273
GP2 AND ITGAM	Pancreas	0.842105	1	0.727273
TNFSF13B AND DUOX2	Pancreas	0.666667	0.857143	0.545455
SYT13 AND FXYD5	Pancreas	0.842105	1	0.727273
OLR1 AND DUOX2	Pancreas	0.705882	1	0.545455
GP2 AND MSR1	Pancreas	0.8	0.888889	0.727273
GJB1 AND CALHM2	Pancreas	0.705882	1	0.545455
GP2 AND CLEC5A	Pancreas	0.8	0.888889	0.727273
GP2 AND ADAM12	Pancreas	0.736842	0.875	0.636364
NOX4 AND SLCO2B1	Pancreas	0.7	0.777778	0.636364
NOX4 AND PTPN2	Pancreas	0.761905	0.8	0.727273
OLR1 AND SLC15A1	Pancreas	0.705882	1	0.545455
SYT13 AND CALCRL	Pancreas	0.64	0.571429	0.727273
TREM2 AND SLC28A3	Pancreas	0.666667	0.857143	0.545455
SYT13 AND ITGAM	Pancreas	0.846154	0.733333	1

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
PTPRZ1 AND CD79B	Astrocytoma	0.703297	0.711111	0.695652
PTPRZ1 AND PTK7	Astrocytoma	0.703297	0.711111	0.695652
PTPRZ1 AND TNFRSF17	Astrocytoma	0.702128	0.6875	0.717391
NCAM1 AND SLC3A2	Astrocytoma	0.702128	0.6875	0.717391
MMP16 AND ITGAV	Astrocytoma	0.701299	0.870968	0.586957
ADCYAP1R1 AND CLDN1	Astrocytoma	0.701031	0.666667	0.73913
NRCAM AND CD38	Astrocytoma	0.698113	0.616667	0.804348
PTPRZ1 AND SSTR1	Astrocytoma	0.698113	0.616667	0.804348
PTPRZ1 AND CD37	Astrocytoma	0.695652	0.695652	0.695652
PTPRZ1 AND CA9	Astrocytoma	0.694737	0.673469	0.717391
NCAM1 AND ATP13A4	Astrocytoma	0.692308	0.84375	0.586957
BMPR1B AND TLR4	Astrocytoma	0.691358	0.8	0.608696
MMP16 AND SLC39A6	Astrocytoma	0.690476	0.763158	0.630435
NRCAM AND FAP	Astrocytoma	0.690476	0.763158	0.630435
PTPRZ1 AND SLC7A5	Astrocytoma	0.688172	0.680851	0.695652
NCAM1 AND BMPR1A	Astrocytoma	0.688172	0.680851	0.695652
NCAM1 AND SLC16A2	Astrocytoma	0.688172	0.680851	0.695652
PTPRZ1 AND SSTR4	Astrocytoma	0.688172	0.680851	0.695652
PTPRZ1 AND ULBP1	Astrocytoma	0.6875	0.66	0.717391
PTPRZ1 AND FCRL1	Astrocytoma	0.6875	0.66	0.717391
PTPRZ1 AND CD19	Astrocytoma	0.685714	0.610169	0.782609
CALCRL AND BMPR1B	Astrocytoma	0.682927	0.777778	0.608696
MMP16 AND EDNRB	Astrocytoma	0.682927	0.777778	0.608696
NCAM1 AND SRR	Astrocytoma	0.680851	0.666667	0.695652
PTPRZ1 AND TNFSF11	Astrocytoma	0.680412	0.647059	0.717391
PTPRZ1 AND GPA33	Astrocytoma	0.680412	0.647059	0.717391
PTPRZ1 AND ABCB5	Astrocytoma	0.680412	0.647059	0.717391
PTPRZ1 AND MUC4	Astrocytoma	0.680412	0.647059	0.717391
PTPRZ1 AND CR2	Astrocytoma	0.68	0.62963	0.73913
NRCAM AND RAET1E	Astrocytoma	0.678899	0.587302	0.804348
KCNN3 AND CLDN1	Astrocytoma	0.690476	0.763158	0.630435
KCNN3 AND CD33	Astrocytoma	0.691358	0.8	0.608696
ADCYAP1R1 AND FAP	Astrocytoma	0.674699	0.756757	0.608696
PTPRZ1 AND CD180	Astrocytoma	0.674699	0.756757	0.608696
ADAM12 AND SDC1	Breast	0.875	1	0.777778
ADAM12 AND MUC1	Breast	0.857143	0.954545	0.777778
ADAM12 AND CLDN7	Breast	0.833333	0.952381	0.740741
FAP AND CD300LF	Breast	0.8	0.926829	0.703704
FAP AND CXCR4	Breast	0.826087	1	0.703704
FAP AND SLC16A3	Breast	0.821053	0.95122	0.722222
FAP AND IL17RB	Breast	0.821053	0.95122	0.722222
FAP AND SLC52A2	Breast	0.863158	1	0.759259
FAP AND CNNM4	Breast	0.813187	1	0.685185
FAP AND KIAA1324	Breast	0.786517	1	0.648148
FAP AND OR51B2	Breast	0.783505	0.883721	0.703704
FAP AND UNC5B	Breast	0.772727	1	0.62963
FAP AND MFSD10	Breast	0.758621	1	0.611111
ADAM12 AND ERBB3	Breast	0.758621	1	0.611111
SLC5A6 AND FAP	Breast	0.758621	1	0.611111
FAP AND CACFD1	Breast	0.886598	1	0.796296
FAP AND VMP1	Breast	0.795699	0.948718	0.685185
FAP AND LPAR2	Breast	0.758621	1	0.611111
FAP AND ADAM8	Breast	0.782609	0.947368	0.666667
FGG AND ERBB3	Liver	0.833333	0.833333	0.833333
FGG AND EPHB2	Liver	1	1	1
FGG AND CD34	Liver	0.8	1	0.666667
ASGR1 AND CD34	Liver	0.909091	1	0.833333
ASGR1 AND ERBB3	Liver	0.769231	0.714286	0.833333
FGG AND IL3RA	Liver	0.769231	0.714286	0.833333
FGG AND LGR5	Liver	0.833333	0.833333	0.833333
ASGR1 AND EPHB2	Liver	0.909091	1	0.833333
HPN AND ERBB3	Liver	0.833333	0.833333	0.833333
HPN AND CD34	Liver	0.909091	1	0.833333
FGG AND VTCN1	Liver	0.8	1	0.666667
ABCC6 AND CD34	Liver	0.8	1	0.666667
HPN AND EPHB2	Liver	1	1	1
TFR2 AND EPHB2	Liver	0.909091	1	0.833333
ASGR1 AND LGR5	Liver	0.769231	0.714286	0.833333
ABCB4 AND CD34	Liver	0.8	1	0.666667
SLC27A5 AND EPHB2	Liver	0.909091	1	0.833333
HPN AND IL3RA	Liver	0.833333	0.833333	0.833333
HPN AND LGR5	Liver	0.833333	0.833333	0.833333
SLC27A5 AND ERBB3	Liver	0.769231	0.714286	0.833333
SLC27A5 AND CD34	Liver	0.8	1	0.666667
FGG AND SSTR5	Liver	0.727273	0.8	0.666667
FGG AND SSTR3	Liver	0.769231	0.714286	0.833333
HPN AND ULBP1	Liver	0.769231	0.714286	0.833333

FIG. 11 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
NOX4 AND OLR1	Pancreas	0.625	1	0.454545
CLEC5A AND SYT13	Pancreas	0.625	1	0.454545
TREM2 AND DSG2	Pancreas	0.625	1	0.454545
TM4SF1 AND NALCN	Pancreas	0.625	1	0.454545
CLEC5A AND GJB1	Pancreas	0.625	1	0.454545
NOX4 AND TREM2	Pancreas	0.625	1	0.454545
TREM2 AND GJB2	Pancreas	0.625	1	0.454545
NOX4 AND SLC16A3	Pancreas	0.625	1	0.454545
SYT13 AND TYROBP	Pancreas	0.625	1	0.454545
KCNE4 AND GJB1	Pancreas	0.625	1	0.454545
NOX4 AND HLA-C	Pancreas	0.625	1	0.454545
KCNE4 AND TREM2	Pancreas	0.625	1	0.454545
GP2 AND CALHM2	Pancreas	0.666667	0.615385	0.727273
TNFSF13B AND RAMP1	Pancreas	0.777778	1	0.636364
CLEC5A AND TNFRSF21	Pancreas	0.625	1	0.454545
GP2 AND CD53	Pancreas	0.777778	1	0.636364
TNFSF13B AND DSG2	Pancreas	0.705882	1	0.545455
NOX4 AND SYT13	Pancreas	0.62069	0.5	0.818182
NOX4 AND SLAMF6	Pancreas	0.62069	0.5	0.818182
CDH6 AND ATRAID	Renal	0.857143	1	0.75
SLC3A1 AND SIRPA	Renal	0.857143	1	0.75
SLC3A1 AND TSPAN4	Renal	0.857143	1	0.75
CDH6 AND HLA-C	Renal	0.857143	1	0.75
CDH6 AND JTB	Renal	0.857143	1	0.75
CDH6 AND S100A10	Renal	0.857143	1	0.75
CDH6 AND HLA-B	Renal	0.857143	1	0.75
HEPH AND SLC39A10	Colon	1	1	1
MEP1A AND AMIGO2	Colon	0.909091	0.833333	1
ATP10B AND AMIGO2	Colon	0.909091	0.833333	1
GPR160 AND EVA1A	Colon	0.888889	1	0.8
GPR160 AND EDNRA	Colon	0.888889	1	0.8
GPR160 AND PCDH17	Colon	0.888889	1	0.8
GPR160 AND LDLRAD3	Colon	0.888889	1	0.8
GPR160 AND CDH3	Colon	0.888889	1	0.8
HEPH AND EVA1A	Colon	0.888889	1	0.8
GPR160 AND GJA4	Colon	0.888889	1	0.8
PMEPA1 AND LY75	Colon	0.888889	1	0.8
HEPH AND IL1RAP	Colon	0.888889	1	0.8
PMEPA1 AND GPR160	Colon	0.888889	1	0.8
PMEPA1 AND PON2	Colon	0.888889	1	0.8
GPR160 AND PODXL	Colon	0.888889	1	0.8
GPR160 AND TACSTD2	Colon	0.888889	1	0.8
MEP1A AND SLC39A10	Colon	1	1	1
HEPH AND TACSTD2	Colon	1	1	1
PMEPA1 AND HEPH	Colon	0.888889	1	0.8
PMEPA1 AND IFITM1	Colon	0.888889	1	0.8
GPR160 AND PDPN	Colon	0.888889	1	0.8
PMEPA1 AND SPINT1	Colon	0.888889	1	0.8
ATP10B AND SLC39A10	Colon	1	1	1
GPR160 AND EDAR	Colon	0.888889	1	0.8
HEPH AND EDAR	Colon	0.888889	1	0.8
PMEPA1 AND ATP1B1	Colon	0.888889	1	0.8
SLC5A1 AND CD81	Colon	0.888889	1	0.8
PMEPA1 AND ATP6AP2	Colon	0.888889	1	0.8
PMEPA1 AND GPRC5A	Colon	0.888889	1	0.8
PPAP2C AND IFITM1	Colon	0.888889	1	0.8
PMEPA1 AND IFNGR2	Colon	0.888889	1	0.8
AMIGO2 AND PSENEN	Colon	0.888889	1	0.8
IHH AND SLC39A10	Colon	0.888889	1	0.8
IYD AND SLC39A10	Colon	0.909091	0.833333	1
HEPH AND APCDD1	Colon	0.8	0.8	0.8
LSR AND AMIGO2	Colon	0.8	0.8	0.8
AMIGO2 AND IYD	Colon	0.8	0.8	0.8
AMIGO2 AND TM4SF5	Colon	0.8	0.8	0.8
GPM6B AND BMP2	Ependymoma	1	1	1
GPM6B AND WLS	Ependymoma	1	1	1
SLC52A2 AND TMEM123	Esophagus	0.782609	1	0.642857
PTPRZ1 AND OR1A1	Glioblastoma	0.909091	0.9375	0.882353
PTPRZ1 AND SIGLEC10	Glioblastoma	0.866667	1	0.764706
PTPRZ1 AND ADAM12	Glioblastoma	0.866667	1	0.764706
PTPRZ1 AND SLC4A2	Glioblastoma	0.9375	1	0.882353
PTPRZ1 AND SIRPG	Glioblastoma	0.9375	1	0.882353
NRCAM AND SLC4A2	Glioblastoma	0.875	0.933333	0.823529
PTPRZ1 AND CD84	Glioblastoma	0.866667	1	0.764706
PTPRZ1 AND ABCA1	Glioblastoma	0.875	0.933333	0.823529
NRCAM AND BRCA1	Glioblastoma	0.903226	1	0.823529
PTPRZ1 AND KCNK15	Glioblastoma	0.903226	1	0.823529

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
HPN AND VTCN1	Liver	0.8	1	0.666667
ABCB4 AND ERBB3	Liver	0.909091	1	0.833333
FGG AND CLDN9	Liver	0.857143	0.75	1
FGG AND ULBP1	Liver	0.714286	0.625	0.833333
SLC27A5 AND LGR5	Liver	0.769231	0.714286	0.833333
AQP9 AND ERBB3	Liver	0.769231	0.714286	0.833333
FGG AND ABCB5	Liver	0.714286	0.625	0.833333
SLC27A5 AND IL3RA	Liver	0.714286	0.625	0.833333
FGG AND ITGB6	Liver	0.8	0.666667	1
FGG AND GPA33	Liver	0.8	0.666667	1
FGG AND ULBP2	Liver	0.8	0.666667	1
ASGR1 AND VTCN1	Liver	0.8	1	0.666667
KCNE4 AND CLDN7	Pancreas	0.727273	0.727273	0.727273
KCNE4 AND STEAP1	Pancreas	0.736842	0.875	0.636364
FAP AND CXCR4	Pancreas	0.777778	1	0.636364
FAP AND DSG2	Pancreas	0.777778	1	0.636364
FAP AND SYT13	Pancreas	0.842105	1	0.727273
TNFSF4 AND EPCAM	Pancreas	0.777778	1	0.636364
STEAP1 AND CXCR4	Pancreas	0.777778	1	0.636364
FAP AND TNFRSF21	Pancreas	0.777778	1	0.636364
FAP AND TSPAN1	Pancreas	0.705882	1	0.545455
FAP AND ADAM28	Pancreas	0.705882	1	0.545455
TNFSF11 AND TM4SF1	Pancreas	0.666667	0.857143	0.545455
FAP AND CD53	Pancreas	0.777778	1	0.636364
GP2 AND GPNMB	Pancreas	0.777778	1	0.636364
FAP AND ILDR1	Pancreas	0.777778	1	0.636364
EPHA3 AND TM4SF1	Pancreas	0.705882	1	0.545455
FAP AND TM4SF1	Pancreas	0.842105	1	0.727273
NOX4 AND ITGB6	Pancreas	0.705882	1	0.545455
FAP AND FXYD3	Pancreas	0.705882	1	0.545455
FAP AND ATP1B1	Pancreas	0.777778	1	0.636364
FAP AND PTPN2	Pancreas	0.777778	1	0.636364
FAP AND TSPAN13	Pancreas	0.777778	1	0.636364
FAP AND UGT8	Pancreas	0.777778	1	0.636364
ADAM12 AND ITGB6	Pancreas	0.705882	1	0.545455
FAP AND CD300LF	Pancreas	0.777778	1	0.636364
FAP AND NPC1L1	Pancreas	0.842105	1	0.727273
FAP AND SLC44A3	Pancreas	0.842105	1	0.727273
CLDN2 AND OLR1	Pancreas	0.8	0.888889	0.727273
FAP AND LSR	Pancreas	0.777778	1	0.636364
FAP AND SLC6A14	Pancreas	0.705882	1	0.545455
KCNE4 AND STEAP2	Pancreas	0.636364	0.636364	0.636364
KCNE4 AND CLDN2	Pancreas	0.636364	0.636364	0.636364
FAP AND IL2RG	Pancreas	0.842105	1	0.727273
FAP AND GPRC5A	Pancreas	0.777778	1	0.636364
NOX4 AND FCRL5	Pancreas	0.692308	0.6	0.818182
FAP AND TNFSF13B	Pancreas	0.777778	1	0.636364
FAP AND F11R	Pancreas	0.777778	1	0.636364
FAP AND CD80	Pancreas	0.777778	1	0.636364
FAP AND OR1A2	Pancreas	0.842105	1	0.727273
FAP AND ST14	Pancreas	0.777778	1	0.636364
FAP AND CRB3	Pancreas	0.777778	1	0.636364
FAP AND TACSTD2	Pancreas	0.842105	1	0.727273
FAP AND GPR160	Pancreas	0.777778	1	0.636364
FAP AND CDH1	Pancreas	0.777778	1	0.636364
FAP AND DSC2	Pancreas	0.705882	1	0.545455
FAP AND C1orf210	Pancreas	0.777778	1	0.636364
TNFSF4 AND CLDN7	Pancreas	0.736842	0.875	0.636364
KCNE4 AND ITGB6	Pancreas	0.625	1	0.454545
FAP AND LITAF	Pancreas	0.625	1	0.454545
KCNE4 AND EPCAM	Pancreas	0.625	1	0.454545
FAP AND VNN2	Pancreas	0.842105	1	0.727273
FAP AND TAS2R16	Pancreas	0.777778	1	0.636364
FAP AND FPR1	Pancreas	0.842105	1	0.727273
FAP AND PCDHGA10	Pancreas	0.842105	1	0.727273
FAP AND SPINT1	Pancreas	0.842105	1	0.727273
FAP AND IL1R2	Pancreas	0.842105	1	0.727273
FAP AND TREM2	Pancreas	0.777778	1	0.636364
CDH6 AND TNFRSF13C	Renal	0.857143	1	0.75
CDH6 AND SLAMF7	Renal	0.857143	1	0.75
CDH6 AND ABCB5	Renal	0.857143	1	0.75
SLC3A1 AND CD276	Renal	0.818182	0.9	0.75
CDH6 AND CD70	Renal	0.857143	1	0.75
CDH6 AND CXCR5	Renal	0.857143	1	0.75
CDH6 AND ENPP3	Renal	0.8	1	0.666667
CDH6 AND CD52	Renal	0.857143	1	0.75
CDH6 AND ERBB3	Renal	0.8	1	0.666667

FIG. 11 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
NRCAM AND CYBA	Glioblastoma	0.9375	1	0.882353
PTPRZ1 AND SLC2A5	Glioblastoma	0.866667	1	0.764706
PTPRZ1 AND NRCAM	Glioblastoma	0.909091	0.9375	0.882353
PTPRZ1 AND ATP12A	Glioblastoma	0.866667	1	0.764706
NRCAM AND KCNK15	Glioblastoma	0.857143	0.833333	0.882353
PTPRZ1 AND GPR65	Glioblastoma	0.83871	0.928571	0.764706
PTPRZ1 AND CDH6	Glioblastoma	0.866667	1	0.764706
PTPRZ1 AND MR1	Glioblastoma	0.866667	1	0.764706
NRCAM AND SLC22A4	Glioblastoma	0.882353	0.882353	0.882353
PTPRZ1 AND MMP16	Glioblastoma	0.875	0.933333	0.823529
PTPRZ1 AND CLEC4A	Glioblastoma	0.9375	1	0.882353
PTPRZ1 AND CD99	Glioblastoma	0.9375	1	0.882353
PTPRZ1 AND FCGRT	Glioblastoma	0.866667	1	0.764706
NRCAM AND VANGL2	Glioblastoma	0.875	0.933333	0.823529
NRCAM AND PCDH18	Glioblastoma	0.866667	1	0.764706
PTPRZ1 AND GJC1	Glioblastoma	0.827586	1	0.705882
PTPRZ1 AND ATP13A1	Glioblastoma	0.827586	1	0.705882
PTPRZ1 AND DDR2	Glioblastoma	0.827586	1	0.705882
PTPRZ1 AND C5AR1	Glioblastoma	0.827586	1	0.705882
PTPRZ1 AND ABCC3	Glioblastoma	0.827586	1	0.705882
NRCAM AND CD99	Glioblastoma	0.827586	1	0.705882
NRCAM AND PSENEN	Glioblastoma	0.827586	1	0.705882
PTPRZ1 AND ITGB8	Glioblastoma	0.827586	1	0.705882
NRCAM AND AGTRAP	Glioblastoma	0.827586	1	0.705882
NRCAM AND EMP1	Glioblastoma	0.827586	1	0.705882
PTPRZ1 AND CD163L1	Glioblastoma	0.827586	1	0.705882
PTPRZ1 AND TMPRSS5	Glioblastoma	0.827586	1	0.705882
PTPRZ1 AND TLR7	Glioblastoma	0.827586	1	0.705882
PTPRZ1 AND F2RL2	Glioblastoma	0.827586	1	0.705882
PTPRZ1 AND CXCR4	Glioblastoma	0.827586	1	0.705882
NRCAM AND ADAM12	Glioblastoma	0.827586	1	0.705882
NRCAM AND CTNS	Glioblastoma	0.9375	1	0.882353
NRCAM AND SLC11A1	Glioblastoma	0.827586	1	0.705882
NRCAM AND BTN3A2	Glioblastoma	0.866667	1	0.764706
NRCAM AND BMPR1A	Glioblastoma	0.827586	1	0.705882
PTPRZ1 AND OR51B6	Glioblastoma	0.903226	1	0.823529
NLGN4X AND BRCA1	Glioblastoma	0.83871	0.928571	0.764706
PTPRZ1 AND F2RL1	Glioblastoma	0.866667	1	0.764706
PTPRZ1 AND SORT1	Glioblastoma	0.903226	1	0.823529
NRCAM AND LMAN2	Glioblastoma	0.83871	0.928571	0.764706
PTPRZ1 AND TLR3	Glioblastoma	0.827586	1	0.705882
PTPRZ1 AND SLC6A8	Glioblastoma	0.827586	1	0.705882
PTPRZ1 AND ITGB2	Glioblastoma	0.866667	1	0.764706
PTPRZ1 AND ENTPD1	Glioblastoma	0.903226	1	0.823529
PTPRZ1 AND EDNRA	Glioblastoma	0.827586	1	0.705882
PTPRZ1 AND SLC16A4	Glioblastoma	0.866667	1	0.764706
NRCAM AND OR51B6	Glioblastoma	0.903226	1	0.823529
PTPRZ1 AND TNFRSF19	Glioblastoma	0.866667	1	0.764706
NRCAM AND ANO6	Glioblastoma	0.827586	1	0.705882
NRCAM AND FAM57A	Glioblastoma	0.827586	1	0.705882
PTPRZ1 AND HRH1	Glioblastoma	0.827586	1	0.705882
GPM6A AND SLC7A7	Glioblastoma	0.882353	0.882353	0.882353
NRCAM AND SIGLEC10	Glioblastoma	0.83871	0.928571	0.764706
NRCAM AND SLC16A4	Glioblastoma	0.827586	1	0.705882
NRCAM AND CXCR4	Glioblastoma	0.827586	1	0.705882
GPM6B AND TSPAN6	Glioma	0.962406	1	0.927536
GPM6A AND TSPAN6	Glioma	0.970588	0.985075	0.956522
GPM6B AND PLGRKT	Glioma	0.970149	1	0.942029
GPM6B AND ABCA1	Glioma	0.94964	0.942857	0.956522
PTPRZ1 AND JAM2	Glioma	0.985294	1	0.971014
GPM6A AND SLC40A1	Glioma	0.970588	0.985075	0.956522
PTPRZ1 AND NRCAM	Glioma	0.946565	1	0.898551
GPM6A AND HLA-B	Glioma	0.94964	0.942857	0.956522
GPM6B AND ATP13A1	Glioma	0.954545	1	0.913043
GPM6A AND ABCA1	Glioma	0.963504	0.970588	0.956522
GPM6B AND ATRAID	Glioma	0.978102	0.985294	0.971014
GPM6A AND PLGRKT	Glioma	0.970149	1	0.942029
GPM6B AND JTB	Glioma	0.962406	1	0.927536
GPM6A AND JTB	Glioma	0.962406	1	0.927536
PTPRZ1 AND HLA-B	Glioma	0.992701	1	0.985507
PTPRZ1 AND BFAR	Glioma	0.992701	1	0.985507
PTPRZ1 AND XPR1	Glioma	0.970149	1	0.942029
GPM6A AND ATRAID	Glioma	0.985294	1	0.971014
PTPRZ1 AND ABCA1	Glioma	0.985294	1	0.971014
PTPRZ1 AND SLC40A1	Glioma	0.977778	1	0.956522
PTPRZ1 AND TSPAN6	Glioma	0.962406	1	0.927536
GPM6B AND XPR1	Glioma	0.970149	1	0.942029

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
CDH6 AND FOLR1	Renal	0.857143	1	0.75
SLC3A1 AND MET	Renal	0.8	1	0.666667
CDH6 AND ULBP1	Renal	0.857143	1	0.75
CDH6 AND CD37	Renal	0.857143	1	0.75
CDH6 AND CD19	Renal	0.857143	1	0.75
CDH6 AND VCAM1	Renal	0.8	1	0.666667
CDH6 AND EPCAM	Renal	0.8	1	0.666667
CDH6 AND SLC34A2	Renal	0.857143	1	0.75
CDH6 AND MET	Renal	0.857143	1	0.75
CDH6 AND GPNMB	Renal	0.857143	1	0.75
VCAM1 AND SYT13	Renal	0.857143	1	0.75
SLC3A1 AND VCAM1	Renal	0.761905	0.888889	0.666667
KCNJ16 AND VCAM1	Renal	0.761905	0.888889	0.666667
VCAM1 AND OR10H3	Renal	0.761905	0.888889	0.666667
DPEP1 AND CD81	Colon	1	1	1
DPEP1 AND IFITM1	Colon	1	1	1
DPEP1 AND AMIGO2	Colon	1	1	1
HEPH AND CLDN1	Colon	1	1	1
GUCY2C AND AMIGO2	Colon	0.909091	0.833333	1
MUC13 AND AMIGO2	Colon	0.909091	0.833333	1
DPEP1 AND TMEM123	Colon	1	1	1
CEACAM5 AND SLC39A10	Colon	1	1	1
CLDN2 AND AMIGO2	Colon	0.888889	1	0.8
CLDN2 AND ARL6IP5	Colon	0.888889	1	0.8
GPR160 AND FAP	Colon	0.888889	1	0.8
PMEPA1 AND CEACAM6	Colon	0.888889	1	0.8
DPEP1 AND SMAGP	Colon	0.888889	1	0.8
HEPH AND SLC7A5	Colon	0.888889	1	0.8
GPR160 AND THY1	Colon	0.888889	1	0.8
GPR160 AND SLC7A5	Colon	0.888889	1	0.8
GPR160 AND CLDN1	Colon	0.888889	1	0.8
DPEP1 AND SERINC5	Colon	0.888889	1	0.8
GUCY2C AND SLC39A10	Colon	1	1	1
GPA33 AND AMIGO2	Colon	0.909091	0.833333	1
MUC13 AND SLC39A10	Colon	1	1	1
HEPH AND THY1	Colon	1	1	1
PRNP AND ABCB5	Ependymoma	1	1	1
SSTR1 AND CD81	Ependymoma	1	1	1
GPM6B AND CD38	Ependymoma	1	1	1
PRNP AND ERBB2	Ependymoma	1	1	1
PRNP AND CLDN2	Ependymoma	1	1	1
PRNP AND VTCN1	Ependymoma	1	1	1
PRNP AND RNF43	Ependymoma	1	1	1
BCAP31 AND EPCAM	Esophagus	0.692308	0.75	0.642857
TM4SF5 AND MOK	Esophagus	0.727273	1	0.571429
GPRC5A AND MOK	Esophagus	0.666667	1	0.5
SLC52A2 AND ITGAV	Esophagus	0.782609	1	0.642857
CBX3 AND CDCP1	Esophagus	0.64	0.727273	0.571429
CBX3 AND TACSTD2	Esophagus	0.727273	1	0.571429
CBX3 AND PPAP2C	Esophagus	0.666667	0.8	0.571429
CBX3 AND CXADR	Esophagus	0.64	0.727273	0.571429
CBX3 AND SLC39A4	Esophagus	0.666667	0.8	0.571429
CBX3 AND IL22RA1	Esophagus	0.666667	0.8	0.571429
BCAP31 AND MUC13	Esophagus	0.64	0.727273	0.571429
EPCAM AND RHBDF2	Esophagus	0.666667	1	0.5
CBX3 AND EDNRA	Esophagus	0.666667	0.8	0.571429
CBX3 AND GPRC5A	Esophagus	0.64	0.727273	0.571429
SLC52A2 AND CLDN1	Esophagus	0.782609	1	0.642857
PTPRH AND MOK	Esophagus	0.636364	0.875	0.5
SLC52A2 AND ULBP2	Esophagus	0.6	1	0.428571
PTPRZ1 AND CD33	Glioblastoma	0.866667	1	0.764706
PTPRZ1 AND CD180	Glioblastoma	0.866667	1	0.764706
PTPRZ1 AND VCAM1	Glioblastoma	0.9375	1	0.882353
PTPRZ1 AND CD70	Glioblastoma	0.909091	0.9375	0.882353
PTPRZ1 AND SDC1	Glioblastoma	0.827586	1	0.705882
PTPRZ1 AND IL13RA1	Glioblastoma	0.903226	1	0.823529
PTPRZ1 AND ENG	Glioblastoma	0.785714	1	0.647059
PTPRZ1 AND NCAM1	Glioblastoma	0.909091	0.9375	0.882353
NRCAM AND VCAM1	Glioblastoma	0.909091	0.9375	0.882353
BMPR1B AND SIGLEC10	Glioblastoma	0.777778	0.736842	0.823529
PTPRZ1 AND SLC39A6	Glioblastoma	0.9375	1	0.882353
PTPRZ1 AND BMPR1B	Glioblastoma	0.909091	0.9375	0.882353
AQP4 AND EGFR	Glioblastoma	0.758621	0.916667	0.647059
NRCAM AND AXL	Glioblastoma	0.882353	0.882353	0.882353
PTPRZ1 AND KDR	Glioblastoma	0.866667	1	0.764706
NRCAM AND IL13RA1	Glioblastoma	0.789474	0.714286	0.882353
NRCAM AND CD180	Glioblastoma	0.756757	0.7	0.823529

FIG. 11 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
PTPRZ1 AND CD46	Glioma	0.962406	1	0.927536
PTPRZ1 AND PRCP	Glioma	0.977778	1	0.956522
PTPRZ1 AND SGCB	Glioma	0.985294	1	0.971014
NRCAM AND SLC3A2	Glioma	0.938462	1	0.884058
GPM6B AND JAM2	Glioma	0.964029	0.957143	0.971014
PTPRZ1 AND HLA-G	Glioma	0.992701	1	0.985507
PTPRZ1 AND ADAM9	Glioma	0.962406	1	0.927536
GPM6B AND DIABLO	Glioma	0.992701	1	0.985507
PTPRZ1 AND PLGRKT	Glioma	0.970149	1	0.942029
NRCAM AND HLA-G	Glioma	0.938462	1	0.884058
PTPRZ1 AND TNFRSF21	Glioma	0.977778	1	0.956522
GPM6B AND BFAR	Glioma	0.951049	0.918919	0.985507
PTPRZ1 AND CALCRL	Glioma	0.962406	1	0.927536
NRCAM AND IL10RB	Glioma	0.938462	1	0.884058
NRCAM AND HLA-C	Glioma	0.930233	1	0.869565
NRCAM AND CD81	Glioma	0.930233	1	0.869565
VMP1 AND SGCD	Leiomyosarcoma	0.727273	0.888889	0.615385
DDR2 AND BRCA1	Leiomyosarcoma	0.666667	0.875	0.538462
ADAM12 AND HTR2A	Liposarcoma	0.765432	0.688889	0.861111
ADAM12 AND CACNA1C	Liposarcoma	0.738095	0.645833	0.861111
ADAM12 AND STOML3	Liposarcoma	0.729412	0.632653	0.861111
ADAM12 AND SLC38A5	Liposarcoma	0.729412	0.632653	0.861111
ADAM12 AND EMP1	Liposarcoma	0.765432	0.688889	0.861111
ADAM12 AND GABRR1	Liposarcoma	0.731707	0.652174	0.833333
ADAM12 AND TNFRSF9	Liposarcoma	0.738095	0.645833	0.861111
CNTNAP1 AND SLC2A10	Liposarcoma	0.779221	0.731707	0.833333
ADAM12 AND OR2B2	Liposarcoma	0.72093	0.62	0.861111
ADAM12 AND GP5	Liposarcoma	0.740741	0.666667	0.833333
ADAM12 AND CORIN	Liposarcoma	0.72093	0.62	0.861111
ADAM12 AND SGCD	Liposarcoma	0.729412	0.632653	0.861111
ADAM12 AND GJB6	Liposarcoma	0.716049	0.644444	0.805556
ADAM12 AND MC3R	Liposarcoma	0.731707	0.652174	0.833333
ADAM12 AND PCDHB4	Liposarcoma	0.712644	0.607843	0.861111
ADAM12 AND S100A10	Liposarcoma	0.712644	0.607843	0.861111
ADAM12 AND PCDHGC3	Liposarcoma	0.712644	0.607843	0.861111
ADAM12 AND LAX1	Liposarcoma	0.712644	0.607843	0.861111
ADAM12 AND CLEC5A	Liposarcoma	0.712644	0.607843	0.861111
ADAM12 AND TRPA1	Liposarcoma	0.712644	0.607843	0.861111
ADAM12 AND MSR1	Liposarcoma	0.775	0.704545	0.861111
ADAM12 AND OR10A3	Liposarcoma	0.731707	0.652174	0.833333
ADAM12 AND UGT8	Liposarcoma	0.714286	0.625	0.833333
ADAM12 AND CHRM1	Liposarcoma	0.746988	0.659574	0.861111
ADAM12 AND PCDHGC5	Liposarcoma	0.746988	0.659574	0.861111
ADAM12 AND CRTAM	Liposarcoma	0.705882	0.612245	0.833333
ADAM12 AND SLC5A12	Liposarcoma	0.746988	0.659574	0.861111
ADAM12 AND PDCD1LG2	Liposarcoma	0.729412	0.632653	0.861111
ADAM12 AND TAS1R2	Liposarcoma	0.729412	0.632653	0.861111
ADAM12 AND FRRS1L	Liposarcoma	0.729412	0.632653	0.861111
ADAM12 AND SLC4A9	Liposarcoma	0.705882	0.612245	0.833333
ADAM12 AND EMCN	Liposarcoma	0.738095	0.645833	0.861111
ADAM12 AND HRH4	Liposarcoma	0.707317	0.630435	0.805556
ADAM12 AND TAAR1	Liposarcoma	0.729412	0.632653	0.861111
ADAM12 AND KCNG3	Liposarcoma	0.731707	0.652174	0.833333
ADAM12 AND TAS2R14	Liposarcoma	0.704545	0.596154	0.861111
ADAM12 AND KISS1R	Liposarcoma	0.704545	0.596154	0.861111
ADAM12 AND MTDH	Liposarcoma	0.704545	0.596154	0.861111
ADAM12 AND GPR84	Liposarcoma	0.705882	0.612245	0.833333
ADAM12 AND PILRA	Liposarcoma	0.712644	0.607843	0.861111
CNTNAP1 AND TYROBP	Liposarcoma	0.732394	0.742857	0.722222
ADAM12 AND VSIG1	Liposarcoma	0.704545	0.596154	0.861111
ADAM12 AND FPR3	Liposarcoma	0.75	0.681818	0.833333
ADAM12 AND CATSPER1	Liposarcoma	0.729412	0.632653	0.861111
ADAM12 AND PCDH7	Liposarcoma	0.739726	0.72973	0.75
ADAM12 AND PCDHB10	Liposarcoma	0.696629	0.584906	0.861111
ADAM12 AND C5AR1	Liposarcoma	0.696629	0.584906	0.861111
ADAM12 AND KCNH8	Liposarcoma	0.698795	0.617021	0.805556
ADAM12 AND TAS2R8	Liposarcoma	0.75	0.681818	0.833333
ADAM12 AND VMP1	Liposarcoma	0.692308	0.642857	0.75
ADAM12 AND OR5P2	Liposarcoma	0.691358	0.622222	0.777778
ADAM12 AND CHIC2	Liposarcoma	0.688889	0.574074	0.861111
ADAM12 AND GRIA2	Liposarcoma	0.75	0.681818	0.833333
ADAM12 AND IFITM2	Liposarcoma	0.704545	0.596154	0.861111
ADAM12 AND SLC23A1	Liposarcoma	0.729412	0.632653	0.861111
ADAM12 AND CNTNAP1	Liposarcoma	0.732394	0.742857	0.722222
CNTNAP1 AND IFNGR2	Liposarcoma	0.698413	0.814815	0.611111
ADAM12 AND GYPE	Liposarcoma	0.682353	0.591837	0.805556
ADAM12 AND SLC6A14	Liposarcoma	0.682353	0.591837	0.805556

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
PTPRZ1 AND AXL	Glioblastoma	0.875	0.933333	0.823529
NCAM1 AND SLC3A2	Glioblastoma	0.83871	0.928571	0.764706
PTPRZ1 AND SLC7A5	Glioblastoma	0.758621	0.916667	0.647059
NCAM1 AND TSPAN6	Glioblastoma	0.758621	0.916667	0.647059
NRCAM AND MUC1	Glioblastoma	0.740741	1	0.588235
NRCAM AND KDR	Glioblastoma	0.736842	0.666667	0.823529
NRCAM AND ENG	Glioblastoma	0.714286	0.909091	0.588235
PTPRZ1 AND P2RX5	Glioblastoma	0.714286	0.909091	0.588235
NCAM1 AND WNT5A	Glioblastoma	0.714286	0.909091	0.588235
NRCAM AND EDNRB	Glioblastoma	0.714286	0.6	0.882353
AQP4 AND CD33	Glioblastoma	0.733333	0.846154	0.647059
BMPR1B AND ADAM12	Glioblastoma	0.742857	0.722222	0.764706
FZD3 AND VCAM1	Glioblastoma	0.727273	0.75	0.705882
AQP4 AND CD180	Glioblastoma	0.709677	0.785714	0.647059
BMPR1B AND SLC3A2	Glioblastoma	0.777778	0.736842	0.823529
NRCAM AND ITGAV	Glioblastoma	0.866667	1	0.764706
PTPRZ1 AND CSPG4	Glioblastoma	0.692308	1	0.529412
PTPRZ1 AND MUC1	Glioblastoma	0.692308	1	0.529412
PTPRZ1 AND CA9	Glioblastoma	0.692308	1	0.529412
NCAM1 AND SRR	Glioblastoma	0.6875	0.733333	0.647059
BMPR1B AND ABCA1	Glioblastoma	0.684211	0.619048	0.764706
BMPR1B AND SLC11A1	Glioblastoma	0.785714	1	0.647059
NRCAM AND GPNMB	Glioblastoma	0.866667	1	0.764706
NCAM1 AND VANGL2	Glioblastoma	0.666667	0.9	0.529412
NLGN4X AND MUC1	Glioblastoma	0.666667	0.9	0.529412
NRCAM AND CSPG4	Glioblastoma	0.666667	0.769231	0.588235
GPM6B AND GPNMB	Glioblastoma	0.666667	0.631579	0.705882
NCAM1 AND FAM57A	Glioblastoma	0.666667	0.9	0.529412
SLC7A5 AND SLC11A1	Glioblastoma	0.666667	0.9	0.529412
CSPG4 AND PON2	Glioblastoma	0.692308	1	0.529412
NLGN4X AND ITGAV	Glioblastoma	0.666667	0.769231	0.588235
NCAM1 AND IGDCC4	Glioblastoma	0.733333	0.846154	0.647059
PTPRZ1 AND ITGAV	Glioblastoma	0.866667	1	0.764706
BMPR1B AND HAVCR2	Glioblastoma	0.764706	0.764706	0.764706
FZD3 AND AXL	Glioblastoma	0.666667	0.6875	0.647059
NRCAM AND CD33	Glioblastoma	0.651163	0.538462	0.823529
BMPR1B AND IGDCC4	Glioblastoma	0.651163	0.538462	0.823529
NRCAM AND SLC39A6	Glioblastoma	0.651163	0.538462	0.823529
BMPR1B AND CD86	Glioblastoma	0.709677	0.785714	0.647059
GPM6B AND VCAM1	Glioblastoma	0.65	0.565217	0.764706
AQP4 AND PROM1	Glioblastoma	0.648649	0.6	0.705882
AQP4 AND SLC7A5	Glioblastoma	0.648649	0.6	0.705882
NCAM1 AND JAM2	Glioblastoma	0.758621	0.916667	0.647059
BMPR1B AND MERTK	Glioblastoma	0.65	0.565217	0.764706
FZD3 AND CD33	Glioblastoma	0.645161	0.714286	0.588235
GRIA3 AND MUC1	Glioblastoma	0.642857	0.818182	0.529412
BMPR1B AND CD300A	Glioblastoma	0.666667	0.631579	0.705882
PTPRZ1 AND EPHA3	Glioblastoma	0.64	1	0.470588
PTPRZ1 AND EGFR	Glioblastoma	0.64	1	0.470588
PTPRZ1 AND STEAP1	Glioblastoma	0.64	1	0.470588
GPM6A AND ITGAV	Glioma	0.94964	0.942857	0.956522
GPM6B AND ITGAV	Glioma	0.93617	0.916667	0.956522
PTPRZ1 AND ITGAV	Glioma	0.985294	1	0.971014
NRCAM AND CD276	Glioma	0.931298	0.983871	0.884058
NLGN1 AND ITGAV	Glioma	0.930233	1	0.869565
NRCAM AND ITGAV	Glioma	0.921875	1	0.855072
PTPRZ1 AND SLC39A6	Glioma	0.970149	1	0.942029
NRCAM AND ABCB5	Glioma	0.917293	0.953125	0.884058
NRCAM AND SLC34A2	Glioma	0.917293	0.953125	0.884058
NRCAM AND VTCN1	Glioma	0.917293	0.953125	0.884058
NRCAM AND GUCY2C	Glioma	0.917293	0.953125	0.884058
NRCAM AND FCRL5	Glioma	0.917293	0.953125	0.884058
PTPRZ1 AND NCAM1	Glioma	0.985507	0.985507	0.985507
NRCAM AND SLAMF7	Glioma	0.917293	0.953125	0.884058
NRCAM AND SSTR4	Glioma	0.917293	0.953125	0.884058
NRCAM AND CLDN7	Glioma	0.917293	0.953125	0.884058
NRCAM AND CD180	Glioma	0.917293	0.953125	0.884058
NRCAM AND SSTR3	Glioma	0.917293	0.953125	0.884058
NRCAM AND ENG	Glioma	0.917293	0.953125	0.884058
NRCAM AND MUC4	Glioma	0.917293	0.953125	0.884058
NRCAM AND MUC13	Glioma	0.917293	0.953125	0.884058
PTPRZ1 AND CD276	Glioma	0.985294	1	0.971014
NRCAM AND GPA33	Glioma	0.917293	0.953125	0.884058
NRCAM AND CLDN23	Glioma	0.917293	0.953125	0.884058
APLP1 AND ITGAV	Glioma	0.920863	0.914286	0.927536
NLGN4X AND ITGAV	Glioma	0.923077	0.983607	0.869565
NRCAM AND RAET1E	Glioma	0.917293	0.953125	0.884058

FIG. 11 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
ADAM12 AND BTN3A2	Liposarcoma	0.681818	0.576923	0.833333
PON2 AND TACSTD2	Lung Adenocarcinoma	0.637168	0.837209	0.514286
OSMR AND TLCD1	Lung Adenocarcinoma	0.618182	0.85	0.485714
PON2 AND CDH3	Lung Adenocarcinoma	0.654545	0.9	0.514286
VMP1 AND NTM	Lung Adenocarcinoma	0.616667	0.74	0.528571
OSMR AND GOLM1	Lung Adenocarcinoma	0.615385	0.765957	0.514286
OSMR AND XPR1	Lung Adenocarcinoma	0.612613	0.829268	0.485714
VMP1 AND WNT3	Lung Adenocarcinoma	0.633333	0.76	0.542857
PSENEN AND GJA1	Lung Adenocarcinoma	0.608696	0.777778	0.5
VMP1 AND FGG	Lung Adenocarcinoma	0.607143	0.809524	0.485714
OSMR AND ATP1B1	Lung Adenocarcinoma	0.626087	0.8	0.514286
TREM1 AND PON2	Lung Adenocarcinoma	0.62963	0.894737	0.485714
OSMR AND C1orf210	Lung Adenocarcinoma	0.6	0.72	0.514286
OSMR AND BCAP31	Lung Adenocarcinoma	0.6	0.72	0.514286
OSMR AND SLC39A11	Lung Adenocarcinoma	0.610169	0.75	0.514286
PON2 AND AGTRAP	Lung Adenocarcinoma	0.612613	0.829268	0.485714
OLR1 AND PON2	Lung Adenocarcinoma	0.608696	0.777778	0.5
PON2 AND LAMP3	Lung Adenocarcinoma	0.630631	0.853659	0.5
TREM1 AND KCNK5	Lung Adenocarcinoma	0.6	0.825	0.471429
PON2 AND KCNJ15	Lung Adenocarcinoma	0.605505	0.846154	0.471429
PTPRCAP AND SLC2A5	B-Cell Diffuse	0.782609	0.84375	0.72973
CD48 AND SLC2A5	B-Cell Diffuse	0.757576	0.862069	0.675676
CXCL9 AND GPR18	B-Cell Diffuse	0.732394	0.764706	0.702703
PTPRCAP AND KCNMA1	B-Cell Diffuse	0.732394	0.764706	0.702703
CXCL9 AND ITGAE	B-Cell Diffuse	0.732394	0.764706	0.702703
IL2RB AND CDH5	Anaplastic Lymphoma	0.740741	0.833333	0.666667
IL2RB AND GJA1	Anaplastic Lymphoma	0.692308	0.818182	0.6
IL2RB AND OSMR	Anaplastic Lymphoma	0.692308	0.818182	0.6
IL2RB AND RAMP3	Anaplastic Lymphoma	0.64	0.8	0.533333
IL2RB AND ECSCR	Anaplastic Lymphoma	0.615385	0.727273	0.533333
CXCL9 AND IL2RB	Anaplastic Lymphoma	0.615385	0.727273	0.533333
IL2RB AND GJA4	Anaplastic Lymphoma	0.740741	0.833333	0.666667
CD2 AND CDH5	Anaplastic Lymphoma	0.615385	0.727273	0.533333
CXCL9 AND IL2RB	T-Cell, Peripheral	0.8	0.814815	0.785714
CXCL9 AND SIRPG	T-Cell, Peripheral	0.816327	0.952381	0.714286
CXCL9 AND CD3G	T-Cell, Peripheral	0.763636	0.777778	0.75
CXCL9 AND TRPV5	T-Cell, Peripheral	0.75	0.9	0.642857
CXCL9 AND KIAA0754	T-Cell, Peripheral	0.769231	0.833333	0.714286
CXCL9 AND PTPRC	T-Cell, Peripheral	0.75	0.75	0.75
PTPRM AND C11orf24	Melanoma	0.9	1	0.818182
C11orf24 AND BCAP31	Melanoma	0.9	1	0.818182
C11orf24 AND PRCP	Melanoma	0.9	1	0.818182
C11orf24 AND NPC1	Melanoma	0.9	1	0.818182
TNFSF9 AND VMP1	Melanoma	0.952381	1	0.909091
MC1R AND VMP1	Melanoma	0.9	1	0.818182
MC1R AND STOM	Melanoma	0.9	1	0.818182
TNFSF9 AND STOM	Melanoma	0.952381	1	0.909091
PTPRZ1 AND GPR52	Oligodendroglioma	0.888889	1	0.8
GRIA2 AND CALCRL	Oligodendroglioma	0.888889	1	0.8
PCDHGC3 AND GPM6B	Oligodendroglioma	0.888889	1	0.8
GPM6A AND CALCRL	Oligodendroglioma	0.888889	1	0.8
GRIA2 AND VANGL2	Oligodendroglioma	0.888889	1	0.8
PTPRZ1 AND ADAM22	Oligodendroglioma	0.888889	1	0.8
NRCAM AND TSPAN11	Oligodendroglioma	0.928571	1	0.866667
REEP2 AND CALCRL	Oligodendroglioma	0.888889	1	0.8
GPM6B AND BMP2	Oligodendroglioma	0.857143	0.923077	0.8
PTPRZ1 AND CLEC1A	Oligodendroglioma	0.888889	1	0.8
PTPRZ1 AND ANTXR1	Oligodendroglioma	0.888889	1	0.8
GPRC5B AND TSPAN11	Oligodendroglioma	0.848485	0.777778	0.933333
SCAMP5 AND VANGL2	Oligodendroglioma	0.888889	1	0.8
PCDHGC3 AND PTPRZ1	Oligodendroglioma	0.846154	1	0.733333
PTPRZ1 AND TMPRSS5	Oligodendroglioma	0.846154	1	0.733333
PTPRZ1 AND OR51B2	Oligodendroglioma	0.846154	1	0.733333
PTPRZ1 AND TREM1	Oligodendroglioma	0.846154	1	0.733333
PTPRZ1 AND VIPR2	Oligodendroglioma	0.846154	1	0.733333
NLGN1 AND VANGL2	Oligodendroglioma	0.846154	1	0.733333
PCDHGC3 AND PON2	Oligodendroglioma	0.846154	1	0.733333
PTPRZ1 AND FCRL3	Oligodendroglioma	0.846154	1	0.733333
PTPRZ1 AND TNFSF18	Oligodendroglioma	0.846154	1	0.733333
PTPRZ1 AND DLL1	Oligodendroglioma	0.846154	1	0.733333
PTPRZ1 AND CALCRL	Oligodendroglioma	0.846154	1	0.733333
PTPRZ1 AND CHRNA5	Oligodendroglioma	0.846154	1	0.733333
PTPRZ1 AND CD300LF	Oligodendroglioma	0.846154	1	0.733333
REEP2 AND VANGL2	Oligodendroglioma	0.846154	1	0.733333
GRIA2 AND ANTXR1	Oligodendroglioma	0.928571	1	0.866667
PCDHGC3 AND TSPAN7	Oligodendroglioma	0.888889	1	0.8
KCNK15 AND PCDH19	Ovarian	0.869565	0.909091	0.833333

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
NRCAM AND CD52	Glioma	0.917293	0.953125	0.884058
NRCAM AND TNFRSF17	Glioma	0.917293	0.953125	0.884058
NRCAM AND ULBP2	Glioma	0.917293	0.953125	0.884058
NRCAM AND TNFRSF13C	Glioma	0.917293	0.953125	0.884058
NRCAM AND CD34	Glioma	0.917293	0.953125	0.884058
NRCAM AND MS4A1	Glioma	0.917293	0.953125	0.884058
NRCAM AND ROR1	Glioma	0.917293	0.953125	0.884058
NRCAM AND PTK7	Glioma	0.909091	0.952381	0.869565
NRCAM AND CA9	Glioma	0.909091	0.952381	0.869565
NRCAM AND TNFSF11	Glioma	0.909091	0.952381	0.869565
NRCAM AND FCRL1	Glioma	0.909091	0.952381	0.869565
NRCAM AND IL2RA	Glioma	0.909091	0.952381	0.869565
NRCAM AND CD19	Glioma	0.909091	0.952381	0.869565
TNFSF4 AND GPNMB	Leiomyosarcoma	0.631579	1	0.461538
FGFR1 AND ULBP2	Leiomyosarcoma	0.727273	0.888889	0.615385
FLT3 AND CD37	AML	0.935818	0.974138	0.900398
ATP8B4 AND CD70	AML	0.884532	0.975962	0.808765
ADAM12 AND FOLH1	Liposarcoma	0.765432	0.688889	0.861111
ADAM12 AND IL2RA	Liposarcoma	0.756098	0.673913	0.861111
ADAM12 AND ULBP2	Liposarcoma	0.681319	0.563636	0.861111
OSMR AND EPCAM	Lung Adenocarcinoma	0.666667	0.947368	0.514286
EPCAM AND SLC39A10	Lung Adenocarcinoma	0.629921	0.701754	0.571429
VMP1 AND FAP	Lung Adenocarcinoma	0.628099	0.745098	0.542857
CXCR4 AND SDC1	Lung Adenocarcinoma	0.615385	0.765957	0.514286
VMP1 AND SLC34A2	Lung Adenocarcinoma	0.661017	0.8125	0.557143
SLC34A2 AND NTM	Lung Adenocarcinoma	0.611111	0.594595	0.628571
OLR1 AND SDC1	Lung Adenocarcinoma	0.610169	0.75	0.514286
CLEC7A AND EPCAM	Lung Adenocarcinoma	0.608696	0.777778	0.5
TSPAN4 AND CLDN7	Lung Adenocarcinoma	0.645161	0.740741	0.571429
PON2 AND FOLR1	Lung Adenocarcinoma	0.614035	0.795455	0.5
SLC34A2 AND SLC2A5	Lung Adenocarcinoma	0.608	0.690909	0.542857
CLEC5A AND EPCAM	Lung Adenocarcinoma	0.605505	0.846154	0.471429
VMP1 AND SDC1	Lung Adenocarcinoma	0.654545	0.9	0.514286
SLC34A2 AND PON2	Lung Adenocarcinoma	0.603774	0.888889	0.457143
TREM1 AND SDC1	Lung Adenocarcinoma	0.603448	0.76087	0.5
SLC34A2 AND SLC41A2	Lung Adenocarcinoma	0.603175	0.678571	0.542857
EPCAM AND VAMP5	Lung Adenocarcinoma	0.603175	0.678571	0.542857
FAP AND GPRC5A	Lung Adenocarcinoma	0.6	0.825	0.471429
PON2 AND PTK7	Lung Adenocarcinoma	0.62069	0.782609	0.514286
SLC34A2 AND WNT3	Lung Adenocarcinoma	0.650794	0.732143	0.585714
EPCAM AND PLXND1	Lung Adenocarcinoma	0.606061	0.645161	0.571429
SLC34A2 AND CDH5	Lung Adenocarcinoma	0.606061	0.526316	0.714286
FAP AND PON2	Lung Adenocarcinoma	0.616822	0.891892	0.471429
MAGEA11 AND SLC7A13	Lung Carcinoma	0.606061	0.857143	0.46875
CXCL9 AND CD79B	B-Cell Diffuse	0.793651	0.961538	0.675676
CXCL9 AND FCRL5	B-Cell Diffuse	0.761905	0.923077	0.648649
CXCL9 AND CXCR5	B-Cell Diffuse	0.754098	0.958333	0.621622
CXCL9 AND P2RX5	B-Cell Diffuse	0.753623	0.8125	0.702703
MCOLN2 AND CD180	B-Cell Diffuse	0.764706	0.83871	0.702703
FCRL3 AND GPNMB	B-Cell Diffuse	0.736842	0.717949	0.756757
CXCL9 AND CR2	B-Cell Diffuse	0.754098	0.958333	0.621622
CD79B AND RAMP3	B-Cell Diffuse	0.746269	0.833333	0.675676
PTPRCAP AND GPNMB	B-Cell Diffuse	0.735294	0.806452	0.675676
CXCL9 AND CD72	B-Cell Diffuse	0.733333	0.956522	0.594595
CD79B AND PPAP2B	B-Cell Diffuse	0.727273	0.827586	0.648649
CD79B AND ENPP2	B-Cell Diffuse	0.727273	0.827586	0.648649
CD79B AND TNFRSF9	B-Cell Diffuse	0.727273	0.827586	0.648649
CD79B AND SLCO2B1	B-Cell Diffuse	0.727273	0.827586	0.648649
MCOLN2 AND GPNMB	B-Cell Diffuse	0.764706	0.83871	0.702703
CD79B AND CDH5	B-Cell Diffuse	0.727273	0.827586	0.648649
CD79A AND BRCA1	B-Cell Diffuse	0.727273	0.827586	0.648649
CD79B AND TMEM119	B-Cell Diffuse	0.727273	0.827586	0.648649
CD79B AND ECSCR	B-Cell Diffuse	0.716418	0.8	0.648649
CD40 AND GPNMB	B-Cell Diffuse	0.716418	0.8	0.648649
TNFRSF9 AND GPNMB	B-Cell Diffuse	0.714286	0.757576	0.675676
CD40 AND VCAM1	B-Cell Diffuse	0.712329	0.722222	0.702703
CLECL1 AND GPNMB	B-Cell Diffuse	0.711864	0.954545	0.567568
CD79B AND MRGPRX4	B-Cell Diffuse	0.724638	0.78125	0.675676
CD79A AND SLC2A5	B-Cell Diffuse	0.738462	0.857143	0.648649
MCOLN2 AND P2RX5	B-Cell Diffuse	0.716418	0.8	0.648649
CD79B AND OSMR	B-Cell Diffuse	0.707692	0.821429	0.621622
CD79B AND DDR2	B-Cell Diffuse	0.707692	0.821429	0.621622
CD79B AND GJA1	B-Cell Diffuse	0.707692	0.821429	0.621622
CD79B AND AQP1	B-Cell Diffuse	0.707692	0.821429	0.621622
CD79B AND MCAM	B-Cell Diffuse	0.705882	0.774194	0.648649
CD79A AND SLC1A3	B-Cell Diffuse	0.705882	0.774194	0.648649
P2RX5 AND ENPP2	B-Cell Diffuse	0.704225	0.735294	0.675676

FIG. 11 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
KCNK15 AND SLC26A7	Ovarian	0.857143	1	0.75
KCNK15 AND TACR2	Ovarian	0.909091	1	0.833333
ADAM12 AND CD99	Sarcoma	0.785714	1	0.647059
DDR2 AND SLC20A1	Sarcoma	0.8	0.923077	0.705882
ADAM12 AND VMP1	Sarcoma	0.866667	1	0.764706
OSMR AND TNFSF4	Sarcoma	0.827586	1	0.705882
ADAM12 AND HLA-B	Sarcoma	0.969697	1	0.941176
OSMR AND ITGA11	Sarcoma	0.827586	1	0.705882
OSMR AND PDGFRB	Sarcoma	0.875	0.933333	0.823529
TSPAN8 AND FXYD5	Stomach	0.860759	0.944444	0.790698
GOLM1 AND FXYD5	Stomach	0.8	0.765957	0.837209
LSR AND FXYD5	Stomach	0.915663	0.95	0.883721
VSIG1 AND PTPRK	Stomach	0.735294	1	0.581395
VSIG1 AND GPRC5A	Stomach	0.735294	1	0.581395
VSIG1 AND PMEPA1	Stomach	0.724638	0.961538	0.581395
VSIG1 AND LRP11	Stomach	0.724638	0.961538	0.581395
VSIG1 AND CD151	Stomach	0.724638	0.961538	0.581395
GJB1 AND FXYD5	Stomach	0.95122	1	0.906977
PTPRH AND FXYD5	Stomach	0.879121	0.833333	0.930233
TSPAN8 AND MCAM	Stomach	0.785714	0.804878	0.767442
TM4SF5 AND FXYD5	Stomach	0.941176	0.952381	0.930233
B3GNT3 AND FXYD5	Stomach	0.97619	1	0.953488
VSIG1 AND PON2	Stomach	0.716418	1	0.55814
VSIG1 AND PDPN	Stomach	0.716418	1	0.55814
VSIG1 AND MYOF	Stomach	0.716418	1	0.55814
VSIG1 AND EMP1	Stomach	0.714286	0.925926	0.581395
SYT13 AND FXYD5	Stomach	0.953488	0.953488	0.953488
VSIG1 AND TSPAN3	Stomach	0.735294	1	0.581395
VSIG1 AND PERP	Stomach	0.714286	0.925926	0.581395
VSIG1 AND ITGA2	Stomach	0.716418	1	0.55814
VSIG1 AND LTBR	Stomach	0.735294	1	0.581395
VSIG1 AND ATP1A1	Stomach	0.716418	1	0.55814
VSIG1 AND GJB2	Stomach	0.724638	0.961538	0.581395
VSIG1 AND ST14	Stomach	0.705882	0.96	0.55814
VSIG1 AND CLSTN1	Stomach	0.705882	0.96	0.55814
VSIG1 AND PLXNA1	Stomach	0.704225	0.892857	0.581395
VSIG1 AND PPAP2C	Stomach	0.704225	0.892857	0.581395
VSIG1 AND LGR4	Stomach	0.724638	0.961538	0.581395
VSIG1 AND EMP2	Stomach	0.714286	0.925926	0.581395
TM4SF5 AND CDH3	Stomach	0.795181	0.825	0.767442
VSIG1 AND LSR	Stomach	0.704225	0.892857	0.581395
IYD AND FXYD5	Stomach	0.860759	0.944444	0.790698
TSPAN8 AND EDNRA	Stomach	0.842105	0.969697	0.744186
VSIG1 AND FZD6	Stomach	0.716418	1	0.55814
VSIG1 AND WNT5A	Stomach	0.69697	1	0.534884
VSIG1 AND TM4SF1	Stomach	0.69697	1	0.534884
VSIG1 AND GJA1	Stomach	0.69697	1	0.534884
VSIG1 AND DSG2	Stomach	0.704225	0.892857	0.581395
VSIG1 AND LDLR	Stomach	0.714286	0.925926	0.581395
VSIG1 AND PTPRF	Stomach	0.704225	0.892857	0.581395
VSIG1 AND SLCO2B1	Stomach	0.695652	0.923077	0.55814
VSIG1 AND CD9	Stomach	0.695652	0.923077	0.55814
VSIG1 AND DAG1	Stomach	0.695652	0.923077	0.55814
VSIG1 AND OSMR	Stomach	0.694444	0.862069	0.581395
VSIG1 AND TNFRSF21	Stomach	0.694444	0.862069	0.581395
VSIG1 AND EPHB4	Stomach	0.694444	0.862069	0.581395
PTPRH AND TACSTD2	Stomach	0.831461	0.804348	0.860465
VSIG1 AND ABHD12	Stomach	0.716418	1	0.55814
VSIG1 AND AKAP1	Stomach	0.694444	0.862069	0.581395
VSIG1 AND PRRG1	Stomach	0.694444	0.862069	0.581395
KCNE3 AND DIO2	Stomach	0.787234	0.72549	0.860465
TSPAN8 AND PDGFRB	Stomach	0.842105	0.969697	0.744186
LSR AND TNFSF15	Stomach	0.840909	0.822222	0.860465
VSIG1 AND SPINT1	Stomach	0.735294	1	0.581395
VSIG1 AND SDF4	Stomach	0.695652	0.923077	0.55814
VSIG1 AND TNFRSF11A	Stomach	0.735294	1	0.581395
VSIG1 AND S100A10	Stomach	0.714286	0.925926	0.581395
VSIG1 AND ATP2A2	Stomach	0.714286	0.925926	0.581395
TSPAN1 AND FXYD5	Stomach	0.934783	0.877551	1
TM4SF5 AND PDPN	Stomach	0.808511	0.745098	0.883721
VSIG1 AND SGCB	Stomach	0.716418	1	0.55814
TSPAN8 AND PDPN	Stomach	0.780488	0.820513	0.744186
VSIG1 AND B3GNT3	Stomach	0.694444	0.862069	0.581395
VSIG1 AND FAM57A	Stomach	0.704225	0.892857	0.581395
VSIG1 AND NRP1	Stomach	0.686567	0.958333	0.534884
VSIG1 AND CAV2	Stomach	0.685714	0.888889	0.55814
VSIG1 AND AQP1	Stomach	0.685714	0.888889	0.55814

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
CD180 AND SLC2A5	B-Cell Diffuse	0.75	0.771429	0.72973
MS4A1 AND CELSR1	Mantle-Cell Lymphoma	0.962025	0.926829	1
MS4A1 AND ST14	Mantle-Cell Lymphoma	0.944444	1	0.894737
MS4A1 AND CLECL1	Mantle-Cell Lymphoma	0.894118	0.808511	1
P2RX5 AND PKD2L1	Mantle-Cell Lymphoma	0.933333	0.945946	0.921053
P2RX5 AND STRA6	Mantle-Cell Lymphoma	0.935065	0.923077	0.947368
P2RX5 AND CELSR1	Mantle-Cell Lymphoma	0.909091	0.897436	0.921053
MS4A1 AND KCNN4	Mantle-Cell Lymphoma	0.974359	0.95	1
P2RX5 AND MRGPRX4	Mantle-Cell Lymphoma	0.894737	0.894737	0.894737
P2RX5 AND SLC13A3	Mantle-Cell Lymphoma	0.909091	0.897436	0.921053
P2RX5 AND TSPAN1	Mantle-Cell Lymphoma	0.876712	0.914286	0.842105
CD52 AND KCNN4	Mantle-Cell Lymphoma	0.888889	0.837209	0.947368
P2RX5 AND OR2S2	Mantle-Cell Lymphoma	0.911392	0.878049	0.947368
P2RX5 AND IL17RC	Mantle-Cell Lymphoma	0.891892	0.916667	0.868421
P2RX5 AND ADAM15	Mantle-Cell Lymphoma	0.868421	0.868421	0.868421
CD52 AND ST14	Mantle-Cell Lymphoma	0.882353	1	0.789474
P2RX5 AND ST14	Mantle-Cell Lymphoma	0.865672	1	0.763158
P2RX5 AND LRRC32	Mantle-Cell Lymphoma	0.9	0.857143	0.947368
P2RX5 AND P2RX2	Mantle-Cell Lymphoma	0.923077	0.9	0.947368
P2RX5 AND SLC22A24	Mantle-Cell Lymphoma	0.935065	0.923077	0.947368
P2RX5 AND PTH1R	Mantle-Cell Lymphoma	0.911392	0.878049	0.947368
P2RX5 AND TIE1	Mantle-Cell Lymphoma	0.911392	0.878049	0.947368
P2RX5 AND KCNE2	Mantle-Cell Lymphoma	0.894737	0.894737	0.894737
P2RX5 AND SLC34A3	Mantle-Cell Lymphoma	0.933333	0.945946	0.921053
P2RX5 AND KCNK1	Mantle-Cell Lymphoma	0.911392	0.878049	0.947368
P2RX5 AND EDNRA	Mantle-Cell Lymphoma	0.86747	0.8	0.947368
P2RX5 AND HEPH	Mantle-Cell Lymphoma	0.861111	0.911765	0.815789
P2RX5 AND UPK2	Mantle-Cell Lymphoma	0.897436	0.875	0.921053
P2RX5 AND OR51B4	Mantle-Cell Lymphoma	0.86747	0.8	0.947368
MS4A1 AND LRRC32	Mantle-Cell Lymphoma	0.853933	0.745098	1
P2RX5 AND FUT1	Mantle-Cell Lymphoma	0.853333	0.864865	0.842105
P2RX5 AND FXYD3	Mantle-Cell Lymphoma	0.911392	0.878049	0.947368
P2RX5 AND CLSTN2	Mantle-Cell Lymphoma	0.886076	0.853659	0.921053
P2RX5 AND TMEM150B	Mantle-Cell Lymphoma	0.911392	0.878049	0.947368
P2RX5 AND ABCC8	Mantle-Cell Lymphoma	0.911392	0.878049	0.947368
P2RX5 AND PROKR2	Mantle-Cell Lymphoma	0.935065	0.923077	0.947368
P2RX5 AND DCHS2	Mantle-Cell Lymphoma	0.888889	0.837209	0.947368
P2RX5 AND AGTR1	Mantle-Cell Lymphoma	0.878049	0.818182	0.947368
P2RX5 AND OR51M1	Mantle-Cell Lymphoma	0.9	0.857143	0.947368
P2RX5 AND PAQR5	Mantle-Cell Lymphoma	0.891892	0.916667	0.868421
P2RX5 AND PCDHB12	Mantle-Cell Lymphoma	0.923077	0.9	0.947368
P2RX5 AND FCRL4	Mantle-Cell Lymphoma	0.878049	0.818182	0.947368
P2RX5 AND FAIM2	Mantle-Cell Lymphoma	0.90411	0.942857	0.868421
P2RX5 AND HRH1	Mantle-Cell Lymphoma	0.853333	0.864865	0.842105
CXCL9 AND CR2	T-Cell, Peripheral	0.784314	0.869565	0.714286
CXCL9 AND MS4A1	T-Cell, Peripheral	0.740741	0.769231	0.714286
CXCL9 AND P2RX5	T-Cell, Peripheral	0.740741	0.769231	0.714286
CXCL9 AND CD22	T-Cell, Peripheral	0.692308	0.75	0.642857
CXCL9 AND ULBP2	T-Cell, Peripheral	0.72	0.818182	0.642857
CXCL9 AND IL2RA	T-Cell, Peripheral	0.679245	0.72	0.642857
CXCL9 AND CD38	T-Cell, Peripheral	0.641509	0.68	0.607143
CXCL9 AND CD79A	T-Cell, Peripheral	0.62963	0.653846	0.607143
TNFSF11 AND IL2RB	T-Cell, Peripheral	0.619048	0.928571	0.464286
GPR174 AND GPNMB	T-Cell, Peripheral	0.638889	0.522727	0.821429
BTLA AND TPBG	T-Cell, Peripheral	0.615385	0.666667	0.571429
TNFSF11 AND CLEC2D	T-Cell, Peripheral	0.622222	0.823529	0.5
TNFRSF9 AND TPBG	T-Cell, Peripheral	0.625	0.75	0.535714
TNFSF11 AND GPR174	T-Cell, Peripheral	0.622222	0.823529	0.5
ITGAL AND GPNMB	T-Cell, Peripheral	0.631579	0.5	0.857143
TNFSF11 AND CD28	T-Cell, Peripheral	0.608696	0.777778	0.5
EDNRB AND IL6R	Melanoma	0.9	1	0.818182
EDNRB AND IL1RAP	Melanoma	0.9	1	0.818182
EDNRB AND ABCC5	Melanoma	0.9	1	0.818182
EDNRB AND JAG2	Melanoma	0.9	1	0.818182
EDNRB AND CLCN5	Melanoma	0.9	1	0.818182
EDNRB AND JTB	Melanoma	0.9	1	0.818182
EDNRB AND GPR37	Melanoma	0.9	1	0.818182
EDNRB AND DLL1	Melanoma	0.9	1	0.818182
EDNRB AND SLC2A8	Melanoma	0.9	1	0.818182
SLC10A4 AND CD276	Neuroblastoma	0.953488	0.931818	0.97619
SLC10A4 AND PTK7	Neuroblastoma	0.947977	0.921348	0.97619
SLC10A4 AND L1CAM	Neuroblastoma	0.943182	0.902174	0.988095
SLC10A4 AND CLDN12	Neuroblastoma	0.935673	0.91954	0.952381
CHRNA3 AND THY1	Neuroblastoma	0.909091	0.925926	0.892857
SLC10A4 AND CBX3	Neuroblastoma	0.907975	0.936709	0.880952
GRIA2 AND EGFR	Oligodendroglioma	0.903226	0.875	0.933333
GPRC5B AND EGFR	Oligodendroglioma	0.896552	0.928571	0.866667

FIG. 11 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
VSIG1 AND CLPTM1	Stomach	0.685714	0.888889	0.55814
VSIG1 AND MFSD10	Stomach	0.685714	0.888889	0.55814
VSIG1 AND SLC4A2	Stomach	0.714286	0.925926	0.581395
VSIG1 AND SLC39A10	Stomach	0.684932	0.833333	0.581395
VSIG1 AND HEPH	Stomach	0.684932	0.833333	0.581395
VSIG1 AND ANO1	Stomach	0.684932	0.833333	0.581395
VSIG1 AND SLC12A7	Stomach	0.714286	0.925926	0.581395
VSIG1 AND KCNK1	Stomach	0.684932	0.833333	0.581395
VSIG1 AND CALCRL	Stomach	0.684932	0.833333	0.581395
VSIG1 AND APOLD1	Stomach	0.684932	0.833333	0.581395
VSIG1 AND JAG1	Stomach	0.684932	0.833333	0.581395
VSIG1 AND BMPR1A	Stomach	0.694444	0.862069	0.581395
LSR AND PDPN	Stomach	0.86747	0.9	0.837209
TSPAN8 AND DIO2	Stomach	0.717949	0.8	0.651163
VSIG1 AND BCAP31	Stomach	0.714286	0.925926	0.581395
VSIG1 AND VIMP	Stomach	0.735294	1	0.581395
VSIG1 AND CDCP1	Stomach	0.694444	0.862069	0.581395
FXYD6 AND BEST3	Astrocytoma	0.831461	0.860465	0.804348
CRB1 AND F2R	Astrocytoma	0.795699	0.787234	0.804348
KCNJ10 AND F2R	Astrocytoma	0.774194	0.765957	0.782609
NKAIN4 AND GPR137B	Astrocytoma	0.772727	0.809524	0.73913
NKAIN4 AND F2R	Astrocytoma	0.764045	0.790698	0.73913
MLC1 AND F2R	Astrocytoma	0.76087	0.76087	0.76087
SLCO1C1 AND F2R	Astrocytoma	0.758621	0.804878	0.717391
NKAIN4 AND LAPTM5	Astrocytoma	0.756098	0.861111	0.673913
GABBR2 AND F2R	Astrocytoma	0.782609	0.782609	0.782609
CRB1 AND BEST3	Astrocytoma	0.752941	0.820513	0.695652
NKAIN4 AND CD44	Astrocytoma	0.752941	0.820513	0.695652
MLC1 AND BEST3	Astrocytoma	0.741573	0.767442	0.717391
GABBR2 AND BEST3	Astrocytoma	0.738095	0.815789	0.673913
NKAIN4 AND SORL1	Astrocytoma	0.736842	0.933333	0.608696
TTYH2 AND BEST3	Astrocytoma	0.735632	0.780488	0.695652
CRB1 AND SEMA4B	Astrocytoma	0.734694	0.692308	0.782609
MLC1 AND SEMA4B	Astrocytoma	0.729167	0.7	0.76087
KCNJ10 AND BEST3	Astrocytoma	0.722892	0.810811	0.652174
SLC1A2 AND F2R	Astrocytoma	0.716981	0.633333	0.826087
SLCO1C1 AND BEST3	Astrocytoma	0.708861	0.848485	0.608696
NKAIN4 AND CXCL16	Astrocytoma	0.708861	0.848485	0.608696
SLCO1C1 AND LAPTM5	Astrocytoma	0.707317	0.805556	0.630435
SLC39A12 AND F2R	Astrocytoma	0.705882	0.769231	0.652174
MOG AND F2R	Astrocytoma	0.704545	0.738095	0.673913
PPAPDC1A AND VANGL1	Breast	0.744681	0.875	0.648148
PPAPDC1A AND LRRC8E	Breast	0.76087	0.921053	0.648148
PPAPDC1A AND ATP8B1	Breast	0.77551	0.863636	0.703704
PPAPDC1A AND VAMP8	Breast	0.767677	0.844444	0.703704
PPAPDC1A AND TGFBI	Breast	0.765957	0.9	0.666667
PPAPDC1A AND TNFRSF12A	Breast	0.715789	0.829268	0.62963
PPAPDC1A AND AOC3	Breast	0.714286	0.795455	0.648148
BST2 AND VANGL1	Breast	0.712644	0.939394	0.574074
BST2 AND LRRC8E	Breast	0.727273	0.941176	0.592593
PPAPDC1A AND SLC22A18	Breast	0.709677	0.846154	0.611111
PPAPDC1A AND F2R	Breast	0.709677	0.846154	0.611111
PPAPDC1A AND PPAPDC1B	Breast	0.762887	0.860465	0.685185
PPAPDC1A AND PHLDB2	Breast	0.74	0.804348	0.685185
PPAPDC1A AND BST2	Breast	0.772727	1	0.62963
PPAPDC1A AND ANTXR2	Breast	0.76	0.826087	0.703704
PPAPDC1A AND BTN3A1	Breast	0.696629	0.885714	0.574074
ENPP1 AND LRRC8E	Breast	0.72381	0.745098	0.703704
PPAPDC1A AND PTGER4	Breast	0.711538	0.74	0.685185
PPAPDC1A AND CXCL16	Breast	0.707071	0.777778	0.648148
PPAPDC1A AND CD163	Breast	0.770833	0.880952	0.685185
PPAPDC1A AND ATP7A	Breast	0.678899	0.672727	0.685185
PPAPDC1A AND SLC31A1	Breast	0.778947	0.902439	0.685185
PPAPDC1A AND CD58	Breast	0.723404	0.85	0.62963
PPAPDC1A AND TNFRSF10B	Breast	0.666667	0.708333	0.62963
CDH11 AND VANGL1	Breast	0.697674	0.9375	0.555556
PPAPDC1A AND BTN3A3	Breast	0.658824	0.903226	0.518519
PPAPDC1A AND PROCR	Breast	0.695652	0.842105	0.592593
PPAPDC1A AND SLC38A1	Breast	0.652174	0.789474	0.555556
PPAPDC1A AND IFNAR2	Breast	0.650602	0.931034	0.5
TNFRSF12A AND VANGL1	Breast	0.679612	0.714286	0.648148
PPAPDC1A AND CDH11	Breast	0.65	1	0.481481
CDH11 AND LRRC8E	Breast	0.65	1	0.481481
TNFRSF12A AND LRRC8E	Breast	0.747253	0.918919	0.62963
ENPP1 AND LAPTM5	Breast	0.704545	0.911765	0.574074
CDH11 AND FLVCR1	Breast	0.714286	1	0.555556
ENPP1 AND SLC38A1	Breast	0.647059	0.6875	0.611111

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
PTPRZ1 AND EGFR	Oligodendroglioma	0.846154	1	0.733333
PTPRZ1 AND CD38	Oligodendroglioma	0.888889	1	0.8
GPM6A AND EGFR	Oligodendroglioma	0.933333	0.933333	0.933333
NRXN2 AND EGFR	Oligodendroglioma	0.827586	0.857143	0.8
NRCAM AND EGFR	Oligodendroglioma	0.888889	1	0.8
LSAMP AND EGFR	Oligodendroglioma	0.827586	0.857143	0.8
PTPRZ1 AND CD180	Oligodendroglioma	0.846154	1	0.733333
AQP4 AND EGFR	Oligodendroglioma	0.814815	0.916667	0.733333
APLP1 AND EGFR	Oligodendroglioma	0.814815	0.916667	0.733333
NCAM2 AND EGFR	Oligodendroglioma	0.888889	1	0.8
SLC1A3 AND EGFR	Oligodendroglioma	0.814815	0.916667	0.733333
PTPRZ1 AND CA9	Oligodendroglioma	0.8	1	0.666667
PTPRZ1 AND CD33	Oligodendroglioma	0.8	1	0.666667
SLC6A1 AND EGFR	Oligodendroglioma	0.814815	0.916667	0.733333
NLGN1 AND EGFR	Oligodendroglioma	0.8	1	0.666667
SLC22A17 AND EGFR	Oligodendroglioma	0.928571	1	0.866667
PTPRZ1 AND CLDN1	Oligodendroglioma	0.857143	0.923077	0.8
ADAM22 AND EGFR	Oligodendroglioma	0.8	0.8	0.8
PTPRZ1 AND CR2	Oligodendroglioma	0.8	1	0.666667
PTPRZ1 AND CSPG4	Oligodendroglioma	0.8	1	0.666667
MUC16 AND SLC1A3	Ovarian	0.736842	1	0.583333
KCNK15 AND VTCN1	Ovarian	0.736842	1	0.583333
MUC16 AND SLC16A10	Ovarian	0.761905	0.888889	0.666667
MUC16 AND ADORA3	Ovarian	0.727273	0.8	0.666667
MUC16 AND NRCAM	Ovarian	0.736842	1	0.583333
MUC16 AND PCDH19	Ovarian	0.666667	1	0.5
KCNK15 AND SDC1	Ovarian	0.666667	0.6	0.75
MUC16 AND CDH2	Ovarian	0.7	0.875	0.583333
MUC16 AND SUSD3	Ovarian	0.761905	0.888889	0.666667
OR51E2 AND EPCAM	Prostate	0.857143	1	0.75
ATP2C1 AND EPCAM	Prostate	0.857143	1	0.75
ANO7 AND EPCAM	Prostate	0.857143	1	0.75
FAP AND JTB	Sarcoma	0.785714	1	0.647059
MUC13 AND FXYD5	Stomach	0.965517	0.954545	0.976744
CLDN18 AND ST14	Stomach	0.886076	0.972222	0.813953
MUC13 AND MCAM	Stomach	0.901099	0.854167	0.953488
TSPAN8 AND THY1	Stomach	0.815789	0.939394	0.72093
CLDN18 AND GJB2	Stomach	0.9	0.972973	0.837209
TSPAN1 AND THY1	Stomach	0.8	0.701754	0.930233
MUC13 AND CDH5	Stomach	0.91954	0.909091	0.930233
CLDN18 AND BRCA1	Stomach	0.871795	0.971429	0.790698
CLDN18 AND B3GNT3	Stomach	0.847059	0.857143	0.837209
CLDN18 AND F2RL2	Stomach	0.85	0.918919	0.790698
MUC13 AND PDPN	Stomach	0.898876	0.869565	0.930233
CLDN18 AND LGR4	Stomach	0.878049	0.923077	0.837209
CLDN18 AND TM4SF5	Stomach	0.875	0.945946	0.813953
CEACAM5 AND FXYD5	Stomach	0.883117	1	0.790698
CLDN18 AND PPAP2C	Stomach	0.888889	0.947368	0.837209
MUC13 AND LDLRAD3	Stomach	0.866667	0.829787	0.906977
MUC13 AND EDNRA	Stomach	0.963855	1	0.930233
MUC13 AND PDGFRB	Stomach	0.941176	0.952381	0.930233
CLDN18 AND TNFRSF11A	Stomach	0.911392	1	0.837209
CLDN18 AND SLC16A1	Stomach	0.837209	0.837209	0.837209
CLDN18 AND F2RL1	Stomach	0.911392	1	0.837209
MUC13 AND APOLD1	Stomach	0.875	0.792453	0.976744
CLDN18 AND GOLM1	Stomach	0.843373	0.875	0.813953
CLDN18 AND TSPAN8	Stomach	0.835443	0.916667	0.767442
MUC13 AND CALCRL	Stomach	0.880952	0.902439	0.860465
CLDN18 AND SEMA4G	Stomach	0.808989	0.782609	0.837209
CLDN18 AND ABCC3	Stomach	0.86747	0.9	0.837209
CLDN18 AND SLC7A1	Stomach	0.888889	0.947368	0.837209
CLDN18 AND ATP10B	Stomach	0.772727	0.755556	0.790698
VSIG1 AND CLDN1	Stomach	0.735294	1	0.581395
CLDN18 AND SYT13	Stomach	0.839506	0.894737	0.790698
CLDN18 AND SLC39A14	Stomach	0.837209	0.837209	0.837209
B3GNT3 AND THY1	Stomach	0.952381	0.97561	0.930233
CLDN18 AND SLC12A2	Stomach	0.839506	0.894737	0.790698
PTPRH AND THY1	Stomach	0.83871	0.78	0.906977
CLDN18 AND DIO2	Stomach	0.775	0.837838	0.72093
CLDN18 AND CDH3	Stomach	0.853333	1	0.744186
EPCAM AND DIO2	Stomach	0.747475	0.660714	0.860465
MUC13 AND IFITM2	Stomach	0.911111	0.87234	0.953488
CEACAM5 AND MCAM	Stomach	0.777778	0.744681	0.813953
VSIG1 AND ERBB2	Stomach	0.724638	0.961538	0.581395
NCAM1 AND SLC39A6	Astrocytoma	0.606742	0.627907	0.586957
NCAM1 AND EDNRB	Astrocytoma	0.606061	0.566038	0.652174
FAP AND CLDN4	Breast	0.791209	0.972973	0.666667

FIG. 11 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
SLC2A2 AND ALCAM	Liver	1	1	1
SLC2A2 AND DTNA	Liver	0.909091	1	0.833333
SLC2A2 AND GPR158	Liver	0.909091	1	0.833333
SLC2A2 AND FGF6	Liver	0.833333	0.833333	0.833333
SLC13A5 AND PLVAP	Liver	0.8	1	0.666667
SLCO1B1 AND FGF6	Liver	0.833333	0.833333	0.833333
SLCO1B1 AND PLVAP	Liver	0.8	1	0.666667
SLC2A2 AND NPFFR1	Liver	0.8	1	0.666667
SLC2A2 AND GABRG3	Liver	0.8	1	0.666667
SLC2A2 AND PCDHB1	Liver	0.857143	0.75	1
SLCO1B1 AND SMPD2	Liver	1	1	1
SLC2A2 AND SLC22A11	Liver	0.909091	1	0.833333
SLC2A2 AND ZACN	Liver	0.909091	1	0.833333
SLC2A2 AND SEZ6	Liver	0.857143	0.75	1
SLCO1B1 AND ZACN	Liver	0.909091	1	0.833333
SLCO1B1 AND SLC22A12	Liver	0.769231	0.714286	0.833333
SLC2A2 AND MRGPRX2	Liver	0.769231	0.714286	0.833333
SLC2A2 AND KCNJ6	Liver	0.909091	1	0.833333
SLC2A2 AND SLC22A12	Liver	0.769231	0.714286	0.833333
SLC2A2 AND CNTNAP4	Liver	0.769231	0.714286	0.833333
SLCO1B1 AND NPFFR1	Liver	0.8	1	0.666667
SLC2A2 AND SLC12A9	Liver	0.8	1	0.666667
SLC2A2 AND SLC30A8	Liver	0.769231	0.714286	0.833333
ABCG5 AND ALCAM	Liver	1	1	1
SLC17A2 AND SLC22A12	Liver	0.769231	0.714286	0.833333
SLC2A2 AND GRM3	Liver	0.769231	0.714286	0.833333
SLCO1B1 AND ALCAM	Liver	1	1	1
SLCO1B1 AND KCNJ6	Liver	0.909091	1	0.833333
SLC17A2 AND SMPD2	Liver	0.909091	1	0.833333
SLCO1B1 AND DTNA	Liver	0.909091	1	0.833333
SLC2A2 AND SMPD2	Liver	1	1	1
SLC2A2 AND HTR1E	Liver	0.769231	0.714286	0.833333
SLC2A2 AND SLC8A3	Liver	0.8	1	0.666667
SLC2A2 AND MAG	Liver	0.833333	0.833333	0.833333
SLC2A2 AND UPK3A	Liver	0.8	1	0.666667
SLC2A2 AND GRM1	Liver	0.857143	0.75	1
SLCO1B1 AND GABRG3	Liver	0.8	1	0.666667
SLC2A2 AND KCNK16	Liver	0.833333	0.833333	0.833333
SLC43A1 AND ALCAM	Liver	0.909091	1	0.833333
SLCO1B1 AND SLC22A5	Liver	0.833333	0.833333	0.833333
SLC2A2 AND ROS1	Liver	0.857143	0.75	1
SLC2A2 AND CACNA1E	Liver	0.769231	0.714286	0.833333
SLC2A2 AND GABRA5	Liver	0.857143	0.75	1
SLC2A2 AND PVRL1	Liver	0.857143	0.75	1
SLC2A2 AND SLC43A1	Liver	0.909091	1	0.833333
SLC2A2 AND CLSTN3	Liver	0.909091	1	0.833333
SLCO1B1 AND GPR158	Liver	0.75	0.6	1
SLCO1B1 AND SEZ6	Liver	0.75	0.6	1
SLCO1B1 AND PCDHB1	Liver	0.75	0.6	1
SLCO1B1 AND MUC12	Liver	0.833333	0.833333	0.833333
SLCO1B1 AND SLC30A8	Liver	0.75	0.6	1
SLCO1B1 AND KCNK16	Liver	0.833333	0.833333	0.833333
ABCG5 AND DTNA	Liver	0.909091	1	0.833333
SLC17A2 AND ZACN	Liver	0.8	1	0.666667
SLCO1B3 AND ALCAM	Liver	0.8	1	0.666667
SLCO1B1 AND PVRL1	Liver	0.857143	0.75	1
SLC2A2 AND OPCML	Liver	0.857143	0.75	1
SLCO1B1 AND SLC12A9	Liver	0.857143	0.75	1
SLCO1B1 AND RGSL1	Liver	0.769231	0.714286	0.833333
SLCO1B1 AND SLC6A11	Liver	0.8	1	0.666667
SLCO1B1 AND MMP24	Liver	0.833333	0.833333	0.833333
SLC2A2 AND TSHR	Liver	0.769231	0.714286	0.833333
ABCG8 AND ALCAM	Liver	0.8	1	0.666667
SLC17A2 AND NPFFR1	Liver	0.8	1	0.666667
SLCO1B1 AND SLC22A11	Liver	0.833333	0.833333	0.833333
SLC2A2 AND NKAIN4	Liver	0.8	1	0.666667
SLCO1B1 AND CALN1	Liver	0.8	0.666667	1
SLCO1B1 AND DRD5	Liver	0.75	0.6	1
SLCO1B1 AND OPCML	Liver	0.75	0.6	1
SLC2A2 AND ACSL6	Liver	1	1	1
SLCO1B1 AND NKAIN4	Liver	0.8	1	0.666667
SLC17A2 AND ALCAM	Liver	0.909091	1	0.833333
SLC17A2 AND DTNA	Liver	0.909091	1	0.833333
SLCO1B1 AND MAG	Liver	0.769231	0.714286	0.833333
C8B AND SMPD2	Liver	0.8	1	0.666667
SLCO1B1 AND TRPM3	Liver	0.833333	0.833333	0.833333
SLCO1B1 AND SCN2A	Liver	0.769231	0.714286	0.833333

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
FAP AND SDC1	Breast	0.875	1	0.777778
FAP AND WT1	Breast	0.808081	0.888889	0.740741
FAP AND MS4A1	Breast	0.744681	0.875	0.648148
FAP AND SLC7A5	Breast	0.868687	0.955556	0.796296
FAP AND FOLH1	Breast	0.828283	0.911111	0.759259
FAP AND CLDN7	Breast	0.698795	1	0.537037
FAP AND BIRC5	Breast	0.82	0.891304	0.759259
FAP AND SSTR4	Breast	0.744681	0.875	0.648148
FAP AND RNF43	Breast	0.857143	0.954545	0.777778
FAP AND CD22	Breast	0.757895	0.878049	0.666667
TPBG AND FAP	Breast	0.705882	0.967742	0.555556
FAP AND CXCR5	Breast	0.755102	0.840909	0.685185
VTCN1 AND VCAM1	Breast	0.690476	0.966667	0.537037
FAP AND BCAN	Breast	0.723404	0.85	0.62963
VTCN1 AND KDR	Breast	0.690476	0.966667	0.537037
FAP AND MUC13	Breast	0.795918	0.886364	0.722222
VTCN1 AND GPNMB	Breast	0.690476	0.966667	0.537037
VTCN1 AND THY1	Breast	0.682927	1	0.518519
VTCN1 AND SLC7A5	Breast	0.682927	1	0.518519
FAP AND CD70	Breast	0.816327	0.909091	0.740741
FAP AND MUC4	Breast	0.681319	0.837838	0.574074
FAP AND SLAMF7	Breast	0.782609	0.947368	0.666667
VTCN1 AND CD34	Breast	0.674699	0.965517	0.518519
FAP AND DNAJB8	Breast	0.674157	0.857143	0.555556
TPBG AND CLDN4	Breast	0.674157	0.857143	0.555556
CLDN2 AND CD276	Liver	0.909091	1	0.833333
CLDN2 AND CD70	Liver	0.769231	0.714286	0.833333
CLDN2 AND SDC1	Liver	0.8	1	0.666667
CLDN2 AND CLDN1	Liver	0.8	1	0.666667
CLDN2 AND IL11RA	Liver	0.615385	0.571429	0.666667
FAP AND EPCAM	Pancreas	0.777778	1	0.636364
FAP AND CLDN7	Pancreas	0.842105	1	0.727273
FAP AND ITGB6	Pancreas	0.777778	1	0.636364
FAP AND PROM1	Pancreas	0.777778	1	0.636364
FAP AND MUC13	Pancreas	0.705882	1	0.545455
FAP AND ERBB3	Pancreas	0.842105	1	0.727273
FAP AND CLDN18	Pancreas	0.625	1	0.454545
FAP AND CD79A	Pancreas	0.705882	1	0.545455
FAP AND CD52	Pancreas	0.777778	1	0.636364
FAP AND FCRL5	Pancreas	0.842105	1	0.727273
FAP AND CD37	Pancreas	0.705882	1	0.545455
FAP AND MST1R	Pancreas	0.705882	1	0.545455
FAP AND VCAM1	Pancreas	0.777778	1	0.636364
FAP AND CLDN23	Pancreas	0.777778	1	0.636364
FAP AND SLAMF7	Pancreas	0.705882	1	0.545455
FAP AND STEAP1	Pancreas	0.777778	1	0.636364
FAP AND TNFSF11	Pancreas	0.842105	1	0.727273
FAP AND TNFRSF17	Pancreas	0.842105	1	0.727273
FAP AND SDC1	Pancreas	0.705882	1	0.545455
FAP AND IGF1R	Pancreas	0.842105	1	0.727273
FAP AND TNFRSF10A	Pancreas	0.777778	1	0.636364
FAP AND FOLH1	Pancreas	0.842105	1	0.727273
CLDN2 AND GPNMB	Pancreas	0.842105	1	0.727273
FAP AND SLC7A5	Pancreas	0.842105	1	0.727273
FAP AND MUC4	Pancreas	0.777778	1	0.636364
EPHA3 AND MUC13	Pancreas	0.631579	0.75	0.545455
FAP AND CD72	Pancreas	0.777778	1	0.636364
FAP AND CD38	Pancreas	0.777778	1	0.636364
FAP AND FOLR1	Pancreas	0.777778	1	0.636364
FAP AND ULBP1	Pancreas	0.777778	1	0.636364
FAP AND STEAP2	Pancreas	0.777778	1	0.636364
FAP AND CLDN2	Pancreas	0.842105	1	0.727273
FAP AND IL2RA	Pancreas	0.705882	1	0.545455
FAP AND BMPR1B	Pancreas	0.777778	1	0.636364
FAP AND EPHB2	Pancreas	0.842105	1	0.727273
EPHA3 AND CLDN1	Pancreas	0.625	1	0.454545
EPHA3 AND CLDN2	Pancreas	0.705882	1	0.545455
FAP AND RNF43	Pancreas	0.777778	1	0.636364
EPHA3 AND CD70	Pancreas	0.631579	0.75	0.545455
EPHA3 AND PROM1	Pancreas	0.625	1	0.454545
FAP AND VTCN1	Pancreas	0.777778	1	0.636364
EPHA3 AND CD52	Pancreas	0.736842	0.875	0.636364
TNFSF11 AND CLDN18	Pancreas	0.625	1	0.454545
EPHA3 AND IL13RA1	Pancreas	0.705882	1	0.545455
EPHA3 AND TNFSF11	Pancreas	0.666667	0.857143	0.545455
FAP AND IL13RA1	Pancreas	0.842105	1	0.727273
FAP AND TRPM4	Pancreas	0.705882	1	0.545455

FIG. 11 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
SLC2A2 AND PCDHAC2	Liver	0.833333	0.833333	0.833333
SLC2A2 AND CACNG1	Liver	0.769231	0.714286	0.833333
SLC17A2 AND KCNJ6	Liver	0.8	1	0.666667
SLCO1B1 AND GRM1	Liver	0.8	0.666667	1
SLC2A2 AND LRRN4	Liver	0.8	1	0.666667
SLC13A5 AND ZACN	Liver	0.833333	0.833333	0.833333
SLC2A2 AND CRB2	Liver	0.769231	0.714286	0.833333
SLC2A2 AND OR7C1	Liver	0.769231	0.714286	0.833333
SLC2A2 AND GABRA3	Liver	0.909091	1	0.833333
SLC2A2 AND MMP24	Liver	0.833333	0.833333	0.833333
SLCO1B1 AND UPK3A	Liver	0.727273	0.8	0.666667
SLCO1B1 AND MLANA	Liver	0.727273	0.8	0.666667
SLC2A2 AND KCNA7	Liver	0.727273	0.8	0.666667
SLC17A2 AND FGF6	Liver	0.727273	0.8	0.666667
SLC2A2 AND HCN1	Liver	0.727273	0.8	0.666667
SLCO1B1 AND GABRA5	Liver	0.857143	0.75	1
SLC2A2 AND OR8D1	Liver	0.769231	0.714286	0.833333
SLCO1B1 AND LRRN4	Liver	0.8	1	0.666667
SLCO1B1 AND GPR61	Liver	0.909091	1	0.833333
SLC2A2 AND DPP10	Liver	0.727273	0.8	0.666667
SLC2A2 AND MLANA	Liver	0.727273	0.8	0.666667
SLC30A8 AND HLA-DRB1	Pancreas	0.777778	1	0.636364
CLDN10 AND BST2	Pancreas	0.625	1	0.454545
SLC30A8 AND BST2	Pancreas	0.777778	1	0.636364
PPAPDC1A AND CD58	Pancreas	0.705882	1	0.545455
SLC30A8 AND LAPTM5	Pancreas	0.777778	1	0.636364
SLC22A2 AND HLA-DRB1	Renal	0.7	0.875	0.583333
SLC17A1 AND IFNAR2	Renal	0.666667	1	0.5
SLC6A13 AND HLA-DRB1	Renal	0.631579	0.857143	0.5
SLC22A2 AND VAMP8	Renal	0.631579	0.857143	0.5
SLC13A1 AND EMP3	Renal	0.6	0.75	0.5
SLC17A1 AND HLA-DRB1	Renal	0.631579	0.857143	0.5
SLC17A1 AND CXCL16	Renal	0.666667	1	0.5
NOX1 AND TGFBI	Colon	0.909091	0.833333	1
FUT3 AND TGFBI	Colon	0.909091	0.833333	1
IFI6 AND NOX1	Colon	1	1	1
IFI6 AND FUT3	Colon	1	1	1
CDH17 AND TGFBI	Colon	0.909091	0.833333	1
IFI6 AND FAT1	Colon	0.888889	1	0.8
IFI6 AND CDH17	Colon	0.888889	1	0.8
IFI6 AND CEACAM7	Colon	1	1	1
IFI6 AND MUC12	Colon	1	1	1
MUC12 AND TGFBI	Colon	0.909091	0.833333	1
IFI6 AND EBP	Colon	0.909091	0.833333	1
IFI6 AND SLC22A18	Colon	0.75	1	0.6
IFI6 AND SLC50A1	Colon	0.727273	0.666667	0.8
IFI6 AND LRIG3	Colon	0.8	0.8	0.8
IFI6 AND VANGL1	Colon	0.909091	0.833333	1
CDH17 AND PPAPDC1A	Colon	0.75	1	0.6
CDH17 AND BST2	Colon	0.75	1	0.6
IFI6 AND ATP8B1	Colon	0.888889	1	0.8
CDH17 AND CDH11	Colon	0.75	1	0.6
SLC22A18 AND TGFBI	Colon	0.666667	0.75	0.6
CLDN15 AND TGFBI	Colon	0.666667	0.75	0.6
IFI6 AND CLDN15	Colon	0.75	1	0.6
IFI6 AND PAQR8	Colon	0.8	0.8	0.8
IFI6 AND EFNB2	Colon	0.888889	1	0.8
IFI6 AND PPAPDC1A	Colon	0.888889	1	0.8
IFI6 AND SLC1A5	Colon	0.888889	1	0.8
SLC22A18 AND CDH11	Colon	0.75	1	0.6
CEACAM7 AND TGFBI	Colon	0.909091	0.833333	1
IFI6 AND FLVCR1	Colon	0.833333	0.714286	1
GRIN2A AND TGFBI	Ependymoma	1	1	1
CXCL16 AND CDH17	Esophagus	0.666667	0.692308	0.642857
CXCL16 AND FUT3	Esophagus	0.782609	1	0.642857
SLCO1B3 AND VANGL1	Esophagus	0.636364	0.875	0.5
CXCL16 AND MUC17	Esophagus	0.782609	1	0.642857
SLCO1B3 AND SLC6A6	Esophagus	0.636364	0.875	0.5
SLCO1B3 AND LRIG3	Esophagus	0.636364	0.875	0.5
GPR158 AND F2R	Glioblastoma	0.83871	0.928571	0.764706
LYPD1 AND LAPTM5	Glioblastoma	0.903226	1	0.823529
LYPD1 AND ADAM29	Glioblastoma	0.777778	0.736842	0.823529
LYPD1 AND HLA-DRB1	Glioblastoma	0.777778	0.736842	0.823529
CRB1 AND CD58	Glioblastoma	0.758621	0.916667	0.647059
LYPD1 AND SPAM1	Glioblastoma	0.742857	0.722222	0.764706
LYPD1 AND F2R	Glioblastoma	0.740741	1	0.588235
MLC1 AND BTN3A3	Glioblastoma	0.736842	0.666667	0.823529

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
FAP AND MS4A1	Pancreas	0.705882	1	0.545455
FAP AND EGFR	Pancreas	0.842105	1	0.727273
FAP AND MET	Pancreas	0.705882	1	0.545455
CLDN2 AND FCRL5	Pancreas	0.666667	0.5625	0.818182
FAP AND GUCY2C	Pancreas	0.842105	1	0.727273
EPHA3 AND STEAP1	Pancreas	0.666667	0.857143	0.545455
FAP AND FCRL2	Pancreas	0.842105	1	0.727273
EPHA3 AND STEAP2	Pancreas	0.6	0.666667	0.545455
FAP AND FCRL1	Pancreas	0.842105	1	0.727273
FAP AND CLDN12	Pancreas	0.777778	1	0.636364
EPHA3 AND EPHB2	Pancreas	0.7	0.777778	0.636364
FAP AND SSTR5	Pancreas	0.705882	1	0.545455
FAP AND SLC34A2	Pancreas	0.777778	1	0.636364
FAP AND CLDN1	Pancreas	0.777778	1	0.636364
FAP AND MUC16	Pancreas	0.842105	1	0.727273
FAP AND LGR5	Pancreas	0.842105	1	0.727273
EPHA3 AND CD72	Pancreas	0.666667	0.7	0.636364
EPHA3 AND IL2RA	Pancreas	0.608696	0.583333	0.636364
FAP AND GPA33	Pancreas	0.842105	1	0.727273
CLDN2 AND VCAM1	Renal	0.761905	0.888889	0.666667
VCAM1 AND ERBB3	Renal	0.736842	1	0.583333
CLDN2 AND GPNMB	Renal	0.695652	0.727273	0.666667
CLDN2 AND CD276	Renal	0.695652	0.727273	0.666667
CLDN2 AND AXL	Renal	0.761905	0.888889	0.666667
VCAM1 AND EPCAM	Renal	0.666667	1	0.5
VCAM1 AND FOLR1	Renal	0.857143	1	0.75
VCAM1 AND ERBB2	Renal	0.666667	1	0.5
VCAM1 AND MUC1	Renal	0.666667	1	0.5
VCAM1 AND CLDN12	Renal	0.8	1	0.666667
VCAM1 AND ABCB5	Renal	0.666667	0.6	0.75
VCAM1 AND SDC1	Renal	0.8	1	0.666667
VCAM1 AND CLDN1	Renal	0.857143	1	0.75
VCAM1 AND EGFR	Renal	0.736842	1	0.583333
DPEP1 AND ITGAV	Colon	0.909091	0.833333	1
DPEP1 AND CEACAM6	Colon	1	1	1
GPA33 AND CLDN1	Colon	1	1	1
CEACAM5 AND THY1	Colon	0.888889	1	0.8
DPEP1 AND CBX3	Colon	0.888889	1	0.8
DPEP1 AND BIRC5	Colon	1	1	1
GUCY2C AND CLDN1	Colon	1	1	1
MUC13 AND THY1	Colon	1	1	1
MUC13 AND CLDN1	Colon	1	1	1
GUCY2C AND THY1	Colon	1	1	1
GPA33 AND THY1	Colon	1	1	1
EPHB2 AND ITGAV	Colon	0.888889	1	0.8
DPEP1 AND SLC7A5	Colon	0.909091	0.833333	1
CLDN3 AND THY1	Colon	1	1	1
MUC13 AND SLC7A5	Colon	1	1	1
GUCY2C AND SLC7A5	Colon	1	1	1
GPA33 AND SLC7A5	Colon	1	1	1
CEACAM6 AND THY1	Colon	0.888889	1	0.8
DPEP1 AND STEAP1	Colon	0.888889	1	0.8
CLDN7 AND THY1	Colon	1	1	1
CLDN3 AND SLC7A5	Colon	1	1	1
CLDN3 AND ITGAV	Colon	0.909091	0.833333	1
CLDN2 AND ITGAV	Colon	0.8	0.8	0.8
CEACAM5 AND FAP	Colon	0.888889	1	0.8
CLDN2 AND CEACAM6	Colon	0.888889	1	0.8
GUCY2C AND ITGAV	Colon	1	1	1
EPHB2 AND THY1	Colon	0.888889	1	0.8
MUC13 AND FAP	Colon	0.888889	1	0.8
CEACAM6 AND FAP	Colon	0.888889	1	0.8
CLDN2 AND BIRC5	Colon	0.888889	1	0.8
GUCY2C AND FAP	Colon	0.888889	1	0.8
EPHB2 AND CLDN1	Colon	0.888889	1	0.8
DPEP1 AND EPHB2	Colon	0.888889	1	0.8
GPA33 AND FAP	Colon	0.888889	1	0.8
EPHB2 AND FAP	Colon	0.888889	1	0.8
CLDN2 AND LGR5	Colon	0.888889	1	0.8
RNF43 AND CEACAM5	Colon	0.75	1	0.6
RNF43 AND CEACAM6	Colon	0.75	1	0.6
DPEP1 AND IL2RA	Colon	0.888889	1	0.8
CLDN7 AND ITGAV	Colon	1	1	1
GPA33 AND ITGAV	Colon	1	1	1
MUC13 AND ITGAV	Colon	1	1	1
DPEP1 AND LGR5	Colon	0.888889	1	0.8
CLDN7 AND FAP	Colon	0.888889	1	0.8

FIG. 11 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
SLCO1C1 AND LAPTM5	Glioblastoma	0.823529	0.823529	0.823529
SLCO1C1 AND TNFRSF12A	Glioblastoma	0.733333	0.846154	0.647059
SLCO1C1 AND ANTXR2	Glioblastoma	0.727273	0.75	0.705882
LYPD1 AND ANTXR2	Glioblastoma	0.740741	1	0.588235
CDH10 AND STAB1	Glioblastoma	0.758621	0.916667	0.647059
LYPD1 AND CD44	Glioblastoma	0.758621	0.916667	0.647059
LYPD1 AND CD163	Glioblastoma	0.903226	1	0.823529
CDH10 AND LAPTM5	Glioblastoma	0.810811	0.75	0.882353
MLC1 AND BST2	Glioblastoma	0.736842	0.666667	0.823529
LYPD1 AND CD93	Glioblastoma	0.785714	1	0.647059
LYPD1 AND TGFBI	Glioblastoma	0.8	0.923077	0.705882
LYPD1 AND TNFRSF12A	Glioblastoma	0.714286	0.909091	0.588235
CDH10 AND ANTXR2	Glioblastoma	0.785714	1	0.647059
SLCO1C1 AND CD93	Glioblastoma	0.727273	0.75	0.705882
ASTN1 AND EMP3	Glioblastoma	0.709677	0.785714	0.647059
MLC1 AND TNFRSF12A	Glioblastoma	0.709677	0.785714	0.647059
CRB1 AND LAPTM5	Glioblastoma	0.709677	0.785714	0.647059
LYPD1 AND CXCL16	Glioblastoma	0.827586	1	0.705882
SLCO1C1 AND CD58	Glioblastoma	0.787879	0.8125	0.764706
LYPD1 AND CDH11	Glioblastoma	0.83871	0.928571	0.764706
LYPD1 AND VAMP8	Glioblastoma	0.777778	0.736842	0.823529
SYT11 AND TNFRSF12A	Glioblastoma	0.705882	0.705882	0.705882
SLCO1C1 AND CDH11	Glioblastoma	0.714286	0.6	0.882353
LYPD1 AND OR1C1	Glioblastoma	0.702703	0.65	0.764706
LYPD1 AND SLCO1B3	Glioblastoma	0.736842	0.666667	0.823529
GPR158 AND SLC39A8	Glioblastoma	0.733333	0.846154	0.647059
PCDH8 AND LAPTM5	Glioblastoma	0.8125	0.866667	0.764706
LYPD1 AND BST2	Glioblastoma	0.875	0.933333	0.823529
LYPD1 AND DTNA	Glioblastoma	0.7	0.608696	0.823529
LYPD1 AND KCNJ10	Glioblastoma	0.7	0.608696	0.823529
ASTN1 AND CD58	Glioblastoma	0.733333	0.846154	0.647059
LYPD1 AND SYT11	Glioblastoma	0.7	0.608696	0.823529
CDH10 AND EMP3	Glioblastoma	0.785714	1	0.647059
LYPD1 AND CD58	Glioblastoma	0.692308	1	0.529412
LYPD1 AND EMP3	Glioblastoma	0.692308	1	0.529412
GPR158 AND CD163	Glioblastoma	0.764706	0.764706	0.764706
CRB1 AND BTN3A1	Glioblastoma	0.689655	0.833333	0.588235
CDH10 AND TNFRSF12A	Glioblastoma	0.689655	0.833333	0.588235
LYPD1 AND STAB1	Glioblastoma	0.689655	0.833333	0.588235
SLC1A2 AND BST2	Glioblastoma	0.689655	0.833333	0.588235
OMG AND F2R	Glioblastoma	0.6875	0.733333	0.647059
SLCO1C1 AND STAB1	Glioblastoma	0.6875	0.733333	0.647059
ASTN1 AND ANTXR2	Glioblastoma	0.6875	0.733333	0.647059
LYPD1 AND BTN3A3	Glioblastoma	0.689655	0.833333	0.588235
GPR158 AND CD93	Glioblastoma	0.684211	0.619048	0.764706
LYPD1 AND SLCO1C1	Glioblastoma	0.682927	0.583333	0.823529
LYPD1 AND CDH10	Glioblastoma	0.682927	0.583333	0.823529
SLCO1C1 AND SLC39A8	Glioblastoma	0.733333	0.846154	0.647059
SLCO1C1 AND CD163	Glioblastoma	0.774194	0.857143	0.705882
CDH10 AND VAMP8	Glioblastoma	0.714286	0.6	0.882353
ASTN1 AND CD93	Glioblastoma	0.666667	0.6875	0.647059
CRB1 AND BST2	Glioblastoma	0.666667	0.6875	0.647059
SLCO1C1 AND VAMP8	Glioblastoma	0.666667	0.535714	0.882353
GABBR2 AND LAPTM5	Glioblastoma	0.666667	0.590909	0.764706
SLC1A2 AND LAPTM5	Glioblastoma	0.666667	0.769231	0.588235
CRB1 AND SLC31A1	Glioblastoma	0.666667	0.9	0.529412
CRB1 AND CXCL16	Glioblastoma	0.666667	0.769231	0.588235
CDH10 AND CDH11	Glioblastoma	0.848485	0.875	0.823529
MLC1 AND SLC31A1	Glioblastoma	0.666667	0.631579	0.705882
SLCO1C1 AND TGFBI	Glioblastoma	0.666667	0.590909	0.764706
CRB1 AND F2R	Glioblastoma	0.666667	0.9	0.529412
LYPD1 AND TTYH2	Glioblastoma	0.666667	0.590909	0.764706
MLC1 AND F2R	Glioblastoma	0.666667	0.631579	0.705882
LYPD1 AND KCNA2	Glioblastoma	0.666667	0.590909	0.764706
SLC1A2 AND CD163	Glioblastoma	0.666667	0.769231	0.588235
CRB1 AND CDH11	Glioblastoma	0.666667	0.6875	0.647059
LYPD1 AND DSCAM	Glioblastoma	0.666667	0.56	0.823529
SYT11 AND F2R	Glioma	0.992701	1	0.985507
GPR158 AND F2R	Glioma	0.977778	1	0.956522
SYT11 AND BTN3A3	Glioma	0.933333	0.954545	0.913043
SYT11 AND HLA-DRB1	Glioma	0.934307	0.941176	0.927536
SYT11 AND LAPTM5	Glioma	0.971429	0.957746	0.985507
SYT11 AND VAMP8	Glioma	0.948905	0.955882	0.942029
SYT11 AND STAB1	Glioma	0.964539	0.944444	0.985507
SYT11 AND CD74	Glioma	0.957143	0.943662	0.971014
OMG AND F2R	Glioma	0.9	0.887324	0.913043
SYT11 AND CLDN15	Glioma	0.978417	0.971429	0.985507

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
CLDN2 AND FAP	Colon	0.888889	1	0.8
CLDN3 AND FAP	Colon	0.888889	1	0.8
CLDN2 AND IL20RA	Colon	0.888889	1	0.8
CLDN2 AND EPHB2	Colon	0.888889	1	0.8
EPCAM AND SLC7A5	Esophagus	0.8125	0.722222	0.928571
EPCAM AND ITGAV	Esophagus	0.692308	0.75	0.642857
CBX3 AND EPCAM	Esophagus	0.666667	0.8	0.571429
EPCAM AND SDC1	Esophagus	0.88	1	0.785714
CBX3 AND MUC1	Esophagus	0.608696	0.777778	0.5
CBX3 AND ERBB3	Esophagus	0.666667	0.8	0.571429
CBX3 AND CLDN7	Esophagus	0.666667	0.8	0.571429
CBX3 AND RNF43	Esophagus	0.608696	0.777778	0.5
CBX3 AND CLDN1	Esophagus	0.666667	1	0.5
MUC13 AND SLC7A5	Esophagus	0.923077	1	0.857143
CBX3 AND EGFR	Esophagus	0.64	0.727273	0.571429
CBX3 AND ITGB6	Esophagus	0.666667	0.8	0.571429
MUC13 AND THY1	Esophagus	0.857143	0.857143	0.857143
CBX3 AND CLDN4	Esophagus	0.666667	0.8	0.571429
CBX3 AND CLDN12	Esophagus	0.64	0.727273	0.571429
CBX3 AND KDR	Esophagus	0.615385	0.666667	0.571429
CBX3 AND IL20RA	Esophagus	0.666667	0.8	0.571429
EPHB2 AND CLDN1	Esophagus	0.923077	1	0.857143
CBX3 AND MUC13	Esophagus	0.608696	0.777778	0.5
BIRC5 AND EPCAM	Esophagus	0.782609	1	0.642857
EPHB2 AND SLC7A5	Esophagus	0.8	0.75	0.857143
ITGAV AND CLDN3	Esophagus	0.608696	0.777778	0.5
CEACAM5 AND THY1	Esophagus	0.896552	0.866667	0.928571
CBX3 AND CLDN3	Esophagus	0.666667	0.8	0.571429
BMPR1B AND ITGAV	Glioblastoma	0.702703	0.65	0.764706
NCAM1 AND SLC39A6	Glioma	0.829268	0.944444	0.73913
NCAM1 AND ITGAV	Glioma	0.784314	0.714286	0.869565
ITGAV AND BMPR1B	Glioma	0.859259	0.878788	0.84058
ITGAV AND SLC7A5	Glioma	0.925373	0.953846	0.898551
NCAM1 AND IL13RA1	Glioma	0.753623	0.753623	0.753623
ITGAV AND ULBP1	Glioma	0.859259	0.878788	0.84058
ITGAV AND SLC39A6	Glioma	0.868852	1	0.768116
NCAM1 AND EDNRB	Glioma	0.716418	0.738462	0.695652
ITGAV AND ERBB4	Glioma	0.817518	0.823529	0.811594
NCAM1 AND CD276	Glioma	0.71345	0.598039	0.884058
ITGAV AND EDNRB	Glioma	0.8	0.941176	0.695652
NCAM1 AND PROM1	Glioma	0.717557	0.758065	0.681159
NCAM1 AND CLDN12	Glioma	0.702703	0.928571	0.565217
ITGAV AND CD19	Glioma	0.834532	0.828571	0.84058
ITGAV AND GUCY2C	Glioma	0.884058	0.884058	0.884058
ITGAV AND CLDN12	Glioma	0.809917	0.942308	0.710145
ITGAV AND CR2	Glioma	0.884058	0.884058	0.884058
ITGAV AND MSLN	Glioma	0.916667	0.88	0.956522
ITGAV AND SSTR4	Glioma	0.884058	0.884058	0.884058
ITGAV AND L1CAM	Glioma	0.907801	0.888889	0.927536
ITGAV AND FCRL5	Glioma	0.833333	0.873016	0.797101
ITGAV AND FCRL1	Glioma	0.9	0.887324	0.913043
CD37 AND SLC7A5	AML	0.818505	0.73955	0.916335
CD38 AND SLC7A5	AML	0.776291	0.746324	0.808765
CD37 AND MUC1	AML	0.725173	0.862637	0.625498
P2RX5 AND SLC7A5	AML	0.683465	0.565104	0.864542
CD38 AND CD70	AML	0.666667	0.664032	0.669323
P2RX5 AND CD70	AML	0.649266	0.549724	0.792829
SLC7A5 AND CD70	AML	0.648464	0.567164	0.756972
SLC7A5 AND CD52	AML	0.755556	0.627968	0.948207
CD38 AND PROM1	AML	0.64466	0.628788	0.661355
CD276 AND EPHB2	Liposarcoma	0.779221	0.731707	0.833333
EPHB2 AND GPNMB	Liposarcoma	0.716049	0.644444	0.805556
CD276 AND FOLH1	Liposarcoma	0.712329	0.702703	0.722222
EPHB2 AND ENG	Liposarcoma	0.704545	0.596154	0.861111
EPHB2 AND PTK7	Liposarcoma	0.736842	0.7	0.777778
CD276 AND B4GALNT1	Liposarcoma	0.622951	0.76	0.527778
EPHB2 AND FOLH1	Liposarcoma	0.638298	0.517241	0.833333
CD276 AND MS4A1	Liposarcoma	0.652174	0.535714	0.833333
SLC34A2 AND SLC7A5	Lung Adenocarcinoma	0.666667	0.636364	0.7
SLC34A2 AND FCRL5	Lung Adenocarcinoma	0.639344	0.75	0.557143
EPCAM AND SLC7A5	Lung Adenocarcinoma	0.632479	0.787234	0.528571
FAP AND EPCAM	Lung Adenocarcinoma	0.625	0.833333	0.5
FAP AND CEACAM6	Lung Adenocarcinoma	0.634615	0.970588	0.471429
SLC34A2 AND FOLR2	Lung Adenocarcinoma	0.621118	0.549451	0.714286
SLC34A2 AND BIRC5	Lung Adenocarcinoma	0.616667	0.74	0.528571
FAP AND CLDN7	Lung Adenocarcinoma	0.614035	0.795455	0.5
SLC34A2 AND VCAM1	Lung Adenocarcinoma	0.606061	0.526316	0.714286

FIG. 11 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
SYT11 AND GPR50	Glioma	0.956522	0.956522	0.956522
SYT11 AND BTN3A1	Glioma	0.933333	0.954545	0.913043
SYT11 AND CD163	Glioma	0.924242	0.968254	0.884058
SYT11 AND TGFBI	Glioma	0.94964	0.942857	0.956522
SYT11 AND ADAM7	Glioma	0.957143	0.943662	0.971014
SYT11 AND FSHR	Glioma	0.94964	0.942857	0.956522
SYT11 AND CD36	Glioma	0.942029	0.942029	0.942029
SYT11 AND ATP6V0A4	Glioma	0.948905	0.955882	0.942029
SYT11 AND CD93	Glioma	0.957143	0.943662	0.971014
SYT11 AND SV2C	Glioma	0.94964	0.942857	0.956522
SYT11 AND MIP	Glioma	0.964029	0.957143	0.971014
SYT11 AND SLC22A13	Glioma	0.971429	0.957746	0.985507
SYT11 AND ICAM1	Glioma	0.934307	0.941176	0.927536
SYT11 AND GYPC	Glioma	0.964539	0.944444	0.985507
SYT11 AND FGF6	Glioma	0.971429	0.957746	0.985507
SYT11 AND GABRG3	Glioma	0.971429	0.957746	0.985507
SYT11 AND BEST2	Glioma	0.971429	0.957746	0.985507
SYT11 AND SLC12A9	Glioma	0.971429	0.957746	0.985507
SYT11 AND OR2L2	Glioma	0.964029	0.957143	0.971014
SYT11 AND TSHR	Glioma	0.957143	0.943662	0.971014
SYT11 AND ADAM20	Glioma	0.94964	0.942857	0.956522
SYT11 AND P2RX4	Glioma	0.970588	0.985075	0.956522
SYT11 AND OR4N4	Glioma	0.957143	0.943662	0.971014
SYT11 AND LRRN4	Glioma	0.971429	0.957746	0.985507
SYT11 AND SELP	Glioma	0.957143	0.943662	0.971014
SYT11 AND ZP4	Glioma	0.964539	0.944444	0.985507
SYT11 AND GHSR	Glioma	0.957143	0.943662	0.971014
SYT11 AND KCNA10	Glioma	0.964029	0.957143	0.971014
SYT11 AND SLC22A12	Glioma	0.948905	0.955882	0.942029
SYT11 AND GJA3	Glioma	0.964539	0.944444	0.985507
SYT11 AND MLANA	Glioma	0.942029	0.942029	0.942029
GPR19 AND F2R	Glioma	0.921875	1	0.855072
SYT11 AND CCR9	Glioma	0.957143	0.943662	0.971014
SYT11 AND TRPC7	Glioma	0.94964	0.942857	0.956522
SYT11 AND PROCR	Glioma	0.956522	0.956522	0.956522
SYT11 AND ESAM	Glioma	0.964029	0.957143	0.971014
SYT11 AND ROS1	Glioma	0.964029	0.957143	0.971014
SYT11 AND KCNU1	Glioma	0.964539	0.944444	0.985507
SYT11 AND CATSPERD	Glioma	0.957143	0.943662	0.971014
SYT11 AND TNFRSF10B	Glioma	0.956522	0.956522	0.956522
SYT11 AND MRGPRX2	Glioma	0.942029	0.942029	0.942029
SYT11 AND GALR1	Glioma	0.934307	0.941176	0.927536
SYT11 AND CHRNA9	Glioma	0.942029	0.942029	0.942029
SYT11 AND CDH11	Glioma	0.924242	0.968254	0.884058
SYT11 AND STEAP4	Glioma	0.957143	0.943662	0.971014
SLC22A16 AND EMP3	AML	0.909449	0.898833	0.920319
SLC22A16 AND CD44	AML	0.929006	0.946281	0.912351
SLC22A16 AND FMNL1	AML	0.895706	0.920168	0.87251
SLC22A16 AND SLC39A8	AML	0.882227	0.953704	0.820717
SLC22A16 AND MLC1	AML	0.865424	0.900862	0.832669
SLC22A16 AND SLC43A1	AML	0.834061	0.922705	0.760956
SLC22A16 AND SEMA4D	AML	0.828452	0.872247	0.788845
SLC22A16 AND TTYH2	AML	0.811839	0.864865	0.76494
SLC22A16 AND STAB1	AML	0.778523	0.887755	0.693227
TTYH3 AND PTGIS	Liposarcoma	0.636364	0.7	0.583333
TTYH3 AND SLC6A7	Liposarcoma	0.626866	0.677419	0.583333
TTYH3 AND ATP8B2	Liposarcoma	0.621622	0.605263	0.638889
CD163 AND SLC6A7	Liposarcoma	0.658228	0.604651	0.722222
TTYH3 AND PHLDB2	Liposarcoma	0.605263	0.575	0.638889
TTYH3 AND CDH11	Liposarcoma	0.605263	0.575	0.638889
TTYH3 AND TNFRSF12A	Liposarcoma	0.630137	0.621622	0.638889
STAB1 AND SLC6A7	Liposarcoma	0.666667	0.641026	0.694444
STAB1 AND EMP3	Liposarcoma	0.648649	0.631579	0.666667
STAB1 AND TGFBI	Liposarcoma	0.648649	0.631579	0.666667
STAB1 AND IFNAR2	Liposarcoma	0.6	0.545455	0.666667
STAB1 AND LRIG3	Liposarcoma	0.657895	0.625	0.694444
EMP3 AND VANGL1	Liposarcoma	0.666667	0.568627	0.805556
STAB1 AND FAT1	Liposarcoma	0.724638	0.757576	0.694444
STAB1 AND CD93	Liposarcoma	0.675676	0.657895	0.694444
CD163 AND VANGL1	Liposarcoma	0.717949	0.666667	0.777778
CD163 AND FAT1	Liposarcoma	0.701299	0.658537	0.75
STAB1 AND GPR137B	Liposarcoma	0.621622	0.605263	0.638889
STAB1 AND GPR158	Liposarcoma	0.623377	0.585366	0.666667
STAB1 AND ABHD3	Liposarcoma	0.617284	0.555556	0.694444
STAB1 AND CDH20	Liposarcoma	0.625	0.568182	0.694444
KCNJ10 AND HLA-DRB1	B-Cell Diffuse	0.75	0.771429	0.72973
KCNJ10 AND LAPTM5	B-Cell Diffuse	0.724638	0.78125	0.675676

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
SLC34A2 AND SDC1	Lung Adenocarcinoma	0.603448	0.76087	0.5
SLC34A2 AND FAP	Lung Adenocarcinoma	0.6	0.72	0.514286
CEACAM6 AND SLC7A5	Lung Adenocarcinoma	0.603774	0.888889	0.457143
SLC34A2 AND HLA-DOB	Lung Adenocarcinoma	0.632258	0.576471	0.7
EPCAM AND PTK7	Lung Adenocarcinoma	0.605505	0.846154	0.471429
MAGEA11 AND ULBP2	Lung Carcinoma	0.606061	0.857143	0.46875
CXCR5 AND GPNMB	B-Cell Diffuse	0.776119	0.866667	0.702703
CD180 AND GPNMB	B-Cell Diffuse	0.746269	0.833333	0.675676
CD79B AND GPNMB	B-Cell Diffuse	0.738462	0.857143	0.648649
CD79A AND GPNMB	B-Cell Diffuse	0.733333	0.956522	0.594595
CD79B AND IL2RA	B-Cell Diffuse	0.714286	0.757576	0.675676
CD72 AND GPNMB	B-Cell Diffuse	0.711864	0.954545	0.567568
P2RX5 AND GPNMB	B-Cell Diffuse	0.711864	0.954545	0.567568
CD79B AND VCAM1	B-Cell Diffuse	0.707692	0.821429	0.621622
FCRL2 AND GPNMB	B-Cell Diffuse	0.689655	0.952381	0.540541
P2RX5 AND VCAM1	B-Cell Diffuse	0.676056	0.705882	0.648649
CD72 AND VCAM1	B-Cell Diffuse	0.666667	0.71875	0.621622
CD79B AND TPBG	B-Cell Diffuse	0.685714	0.727273	0.648649
CD180 AND VCAM1	B-Cell Diffuse	0.658537	0.6	0.72973
FCRL5 AND VCAM1	B-Cell Diffuse	0.738462	0.857143	0.648649
CD79B AND CD276	B-Cell Diffuse	0.6875	0.814815	0.594595
FCRL5 AND GPNMB	B-Cell Diffuse	0.745763	1	0.594595
CXCR5 AND VCAM1	B-Cell Diffuse	0.642857	0.574468	0.72973
CR2 AND GPNMB	B-Cell Diffuse	0.642857	0.947368	0.486486
CD79A AND VCAM1	B-Cell Diffuse	0.64	0.631579	0.648649
CD72 AND EDNRB	B-Cell Diffuse	0.637681	0.6875	0.594595
CD37 AND GPNMB	B-Cell Diffuse	0.705882	0.774194	0.648649
IL2RA AND GPNMB	B-Cell Diffuse	0.632911	0.595238	0.675676
FCRL1 AND GPNMB	B-Cell Diffuse	0.631579	0.9	0.486486
CD72 AND TPBG	B-Cell Diffuse	0.630137	0.638889	0.621622
CD79A AND EDNRB	B-Cell Diffuse	0.628571	0.666667	0.594595
CD79B AND EDNRB	B-Cell Diffuse	0.623377	0.6	0.648649
CD180 AND EDNRB	B-Cell Diffuse	0.613333	0.605263	0.621622
P2RX5 AND EDNRB	B-Cell Diffuse	0.608696	0.65625	0.567568
FCRL5 AND IL2RA	B-Cell Diffuse	0.698413	0.846154	0.594595
CD72 AND IL2RA	B-Cell Diffuse	0.605263	0.589744	0.621622
FCRL5 AND TPBG	B-Cell Diffuse	0.676923	0.785714	0.594595
P2RX5 AND IL2RA	B-Cell Diffuse	0.60274	0.611111	0.594595
FCRL5 AND CD34	B-Cell Diffuse	0.644068	0.863636	0.513514
CD37 AND VCAM1	B-Cell Diffuse	0.611765	0.541667	0.702703
CD79A AND SLC7A5	B-Cell Diffuse	0.666667	0.807692	0.567568
CD79A AND IL2RA	B-Cell Diffuse	0.625	0.740741	0.540541
CD79B AND KDR	B-Cell Diffuse	0.6875	0.814815	0.594595
P2RX5 AND SLC7A5	B-Cell Diffuse	0.612903	0.76	0.513514
IL2RA AND SLC7A5	Anaplastic Lymphoma	0.642857	0.692308	0.6
IL2RA AND VCAM1	Anaplastic Lymphoma	0.62069	0.642857	0.6
IL2RA AND TPBG	Anaplastic Lymphoma	0.692308	0.818182	0.6
IL2RA AND CLDN5	Anaplastic Lymphoma	0.692308	0.818182	0.6
CD79B AND GUCY2C	Mantle-Cell Lymphoma	0.835821	0.965517	0.736842
P2RX5 AND GUCY2C	Mantle-Cell Lymphoma	0.828571	0.90625	0.763158
P2RX5 AND FOLR1	Mantle-Cell Lymphoma	0.857143	0.846154	0.868421
CD79B AND CD34	Mantle-Cell Lymphoma	0.823529	0.933333	0.736842
P2RX5 AND CA9	Mantle-Cell Lymphoma	0.80597	0.931034	0.710526
P2RX5 AND FOLH1	Mantle-Cell Lymphoma	0.8	0.875	0.736842
MS4A1 AND PMEL	Mantle-Cell Lymphoma	0.791667	0.655172	1
P2RX5 AND CD70	Mantle-Cell Lymphoma	0.914286	1	0.842105
MS4A1 AND ERBB3	Mantle-Cell Lymphoma	0.783505	0.644068	1
P2RX5 AND ROR1	Mantle-Cell Lymphoma	0.782609	0.870968	0.710526
MS4A1 AND CD70	Mantle-Cell Lymphoma	0.898551	1	0.815789
CD79B AND FAP	Mantle-Cell Lymphoma	0.78125	0.961538	0.657895
MS4A1 AND ROR1	Mantle-Cell Lymphoma	0.779221	0.769231	0.789474
MS4A1 AND ERBB2	Mantle-Cell Lymphoma	0.778947	0.649123	0.973684
MS4A1 AND CD52	Mantle-Cell Lymphoma	0.778947	0.649123	0.973684
P2RX5 AND CSPG4	Mantle-Cell Lymphoma	0.776119	0.896552	0.684211
MS4A1 AND TNFRSF10A	Mantle-Cell Lymphoma	0.77551	0.633333	1
P2RX5 AND ERBB3	Mantle-Cell Lymphoma	0.774194	0.654545	0.947368
MS4A1 AND LGR5	Mantle-Cell Lymphoma	0.774194	0.654545	0.947368
MS4A1 AND ERBB4	Mantle-Cell Lymphoma	0.774194	0.654545	0.947368
P2RX5 AND SLC34A2	Mantle-Cell Lymphoma	0.769231	0.925926	0.657895
MS4A1 AND MUC16	Mantle-Cell Lymphoma	0.767677	0.622951	1
CD79B AND MUC16	Mantle-Cell Lymphoma	0.767123	0.8	0.736842
MS4A1 AND CLDN2	Mantle-Cell Lymphoma	0.765957	0.642857	0.947368
CD72 AND CD70	Mantle-Cell Lymphoma	0.818182	0.964286	0.710526
MS4A1 AND TNFRSF13C	Mantle-Cell Lymphoma	0.894118	0.808511	1
CD79B AND CLDN2	Mantle-Cell Lymphoma	0.760563	0.818182	0.710526
MS4A1 AND SDC1	Mantle-Cell Lymphoma	0.76	0.612903	1
CD70 AND CD52	Mantle-Cell Lymphoma	0.882353	1	0.789474

FIG. 11 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall	Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
KCNJ10 AND VAMP8	B-Cell Diffuse	0.622951	0.791667	0.513514	MS4A1 AND SLC7A5	Mantle-Cell Lymphoma	0.756098	0.704545	0.815789
GPR19 AND IL15RA	Anaplastic Lymphoma	0.606061	0.555556	0.666667	EDNRB AND IL11RA	Melanoma	0.9	1	0.818182
GPR137B AND SLC6A15	Melanoma	0.9	1	0.818182	EDNRB AND CSPG4	Melanoma	0.9	1	0.818182
GPR19 AND TGFBI	Melanoma	0.9	1	0.818182	EDNRB AND VCAM1	Melanoma	0.9	1	0.818182
GPR137B AND GPR19	Melanoma	1	1	1	EDNRB AND GPNMB	Melanoma	0.9	1	0.818182
GPR137B AND GPR158	Melanoma	0.842105	1	0.727273	EDNRB AND L1CAM	Melanoma	0.9	1	0.818182
FAT1 AND GPR19	Melanoma	0.842105	1	0.727273	EDNRB AND AXL	Melanoma	0.842105	1	0.727273
PAQR7 AND GPR19	Melanoma	0.9	1	0.818182	EDNRB AND TRPM4	Melanoma	0.842105	1	0.727273
SLC22A5 AND GPR19	Melanoma	0.842105	1	0.727273	EDNRB AND STEAP1	Melanoma	0.842105	1	0.727273
GPR19 AND SLC50A1	Melanoma	0.9	1	0.818182	EDNRB AND CD34	Melanoma	0.842105	1	0.727273
GPR137B AND IFI6	Melanoma	0.818182	0.818182	0.818182	EDNRB AND SLC7A5	Melanoma	0.842105	1	0.727273
GPR19 AND ATP7A	Melanoma	0.857143	0.9	0.818182	EDNRB AND CLDN12	Melanoma	0.842105	1	0.727273
GPR19 AND THBD	Melanoma	0.9	1	0.818182	EDNRB AND FAP	Melanoma	0.842105	1	0.727273
GPR137B AND TTYH2	Melanoma	0.785714	0.647059	1	EDNRB AND IGF1R	Melanoma	0.842105	1	0.727273
GPR137B AND GPR85	Melanoma	0.857143	0.9	0.818182	EDNRB AND ERBB2	Melanoma	0.842105	1	0.727273
FAT1 AND IFI6	Melanoma	0.777778	1	0.636364	EDNRB AND CD70	Melanoma	0.842105	1	0.727273
GPR137B AND SLC23A2	Melanoma	0.777778	1	0.636364	EDNRB AND CLDN23	Melanoma	0.777778	1	0.636364
GPR19 AND ATP7A	Neuroblastoma	0.895349	0.875	0.916667	CLDN12 AND GPNMB	Melanoma	0.777778	1	0.636364
CACNA1B AND ATP7A	Neuroblastoma	0.882353	0.872093	0.892857	ALK AND GPNMB	Neuroblastoma	0.825806	0.901408	0.761905
TRPM8 AND SLCO1A2	Prostate	0.727273	0.571429	1	ALK AND FOLR1	Neuroblastoma	0.8125	0.855263	0.77381
UPK3A AND ATP8B1	Prostate	0.666667	0.6	0.75	L1CAM AND CBX3	Neuroblastoma	0.8	1	0.666667
GGTLC1 AND CDH10	Prostate	0.8	0.666667	1	ALK AND GPC3	Neuroblastoma	0.797468	0.851351	0.75
ADAM2 AND GHR	Prostate	0.666667	1	0.5	ALK AND CD52	Neuroblastoma	0.794872	0.861111	0.738095
UPK3A AND SLC43A1	Prostate	0.857143	1	0.75	ALK AND ERBB3	Neuroblastoma	0.792453	0.84	0.75
UPK3A AND ADAM2	Prostate	0.666667	1	0.5	L1CAM AND GPNMB	Neuroblastoma	0.790698	0.772727	0.809524
UPK3A AND CNTNAP2	Prostate	0.666667	0.6	0.75	ALK AND OAS1	Neuroblastoma	0.782051	0.847222	0.72619
UPK3A AND SLC2A2	Prostate	0.666667	0.6	0.75	ALK AND CD34	Neuroblastoma	0.781457	0.880597	0.702381
UPK3A AND GRIA4	Prostate	0.666667	0.6	0.75	ALK AND MST1R	Neuroblastoma	0.77707	0.835616	0.72619
UPK3A AND SELP	Prostate	0.666667	0.6	0.75	L1CAM AND CD276	Neuroblastoma	0.814815	0.846154	0.785714
UPK3A AND STEAP4	Prostate	0.666667	1	0.5	ALK AND CLDN3	Neuroblastoma	0.771242	0.855072	0.702381
UPK3A AND GABRG3	Prostate	0.666667	0.6	0.75	ALK AND MSLN	Neuroblastoma	0.763158	0.852941	0.690476
ADAM2 AND ATP8B1	Prostate	0.857143	1	0.75	ALK AND ALDH1A1	Neuroblastoma	0.76129	0.830986	0.702381
UPK3A AND GPR158	Prostate	0.666667	0.6	0.75	ALK AND MUC4	Neuroblastoma	0.76129	0.830986	0.702381
STAB1 AND SLC6A7	Sarcoma	0.689655	0.833333	0.588235	ALK AND CD72	Neuroblastoma	0.754967	0.850746	0.678571
STAB1 AND EMP3	Sarcoma	0.727273	0.75	0.705882	ALK AND FOLH1	Neuroblastoma	0.753247	0.828571	0.690476
STAB1 AND TGFBI	Sarcoma	0.645161	0.714286	0.588235	ALK AND FCRL2	Neuroblastoma	0.75	0.838235	0.678571
CDH17 AND BST2	Stomach	0.829268	0.871795	0.790698	ALK AND IL3RA	Neuroblastoma	0.728477	0.820896	0.654762
CDH17 AND SLC6A6	Stomach	0.816327	0.727273	0.930233	ALK AND EDNRB	Neuroblastoma	0.724832	0.830769	0.642857
ATP8B1 AND CDH11	Stomach	0.871795	0.971429	0.790698	ALK AND ULBP1	Neuroblastoma	0.721088	0.84127	0.630952
ATP8B1 AND CD93	Stomach	0.871795	0.971429	0.790698	ALK AND CXCR5	Neuroblastoma	0.72	0.818182	0.642857
ATP8B1 AND F2R	Stomach	0.91358	0.973684	0.860465	ALK AND ABCB5	Neuroblastoma	0.72	0.818182	0.642857
MUC17 AND ESAM	Stomach	0.826667	0.96875	0.72093	L1CAM AND BMPR1B	Neuroblastoma	0.712329	0.83871	0.619048
MUC17 AND IFI6	Stomach	0.753623	1	0.604651	ALK AND TNFRSF17	Neuroblastoma	0.711409	0.815385	0.630952
CDH17 AND IFI6	Stomach	0.621359	0.533333	0.744186	NCAM1 AND EGFR	Oligodendroglioma	0.666667	0.541667	0.866667
MUC17 AND THBD	Stomach	0.826667	0.96875	0.72093	SLC7A5 AND MUC13	Oligodendroglioma	0.72	0.9	0.6
ATP8B1 AND PLVAP	Stomach	0.860465	0.860465	0.860465	SLC7A5 AND MSLN	Oligodendroglioma	0.692308	0.818182	0.6
MUC17 AND CD163	Stomach	0.815789	0.939394	0.72093	SLC7A5 AND CLDN9	Oligodendroglioma	0.689655	0.714286	0.666667
CDH17 AND CDH11	Stomach	0.810127	0.888889	0.744186	ABCB5 AND BMPR1B	Oligodendroglioma	0.714286	0.769231	0.666667
MUC17 AND SYT11	Stomach	0.842105	0.969697	0.744186	ABCB5 AND SLC7A5	Oligodendroglioma	0.647059	0.578947	0.733333
MUC17 AND EMP3	Stomach	0.815789	0.939394	0.72093	SLC7A5 AND GPA33	Oligodendroglioma	0.6	0.6	0.6
ATP8B1 AND TGFBI	Stomach	0.839506	0.894737	0.790698	NCAM1 AND EDNRB	Oligodendroglioma	0.615385	0.727273	0.533333
MUC17 AND F2R	Stomach	0.853333	1	0.744186	SSTR2 AND BMPR1B	Oligodendroglioma	0.695652	1	0.533333
MUC17 AND STEAP4	Stomach	0.842105	0.969697	0.744186	SSTR3 AND SLC7A5	Oligodendroglioma	0.642857	0.692308	0.6
MUC17 AND CD44	Stomach	0.842105	0.969697	0.744186	SLC7A5 AND CLDN18	Oligodendroglioma	0.625	0.588235	0.666667
MUC17 AND SLC1A5	Stomach	0.831169	0.941176	0.744186	SLC7A5 AND ITGB3	Oligodendroglioma	0.606061	0.555556	0.666667
MUC17 AND EPHA2	Stomach	0.831169	0.941176	0.744186	CLDN11 AND GPA33	Oligodendroglioma	0.714286	0.769231	0.666667
MUC17 AND SEMA4D	Stomach	0.842105	0.969697	0.744186	SSTR2 AND EDNRB	Oligodendroglioma	0.615385	0.727273	0.533333
MUC17 AND GYPC	Stomach	0.815789	0.939394	0.72093	ABCB5 AND SLC39A6	Oligodendroglioma	0.666667	0.611111	0.733333
MUC17 AND BST2	Stomach	0.771429	1	0.627907	MUC16 AND NCAM1	Ovarian	0.666667	0.777778	0.583333
MUC17 AND PROCR	Stomach	0.826667	0.96875	0.72093	STEAP2 AND EPCAM	Prostate	0.857143	1	0.75
MUC17 AND PLVAP	Stomach	0.831169	0.941176	0.744186	STEAP1 AND EPCAM	Prostate	0.857143	1	0.75
MUC17 AND S1PR1	Stomach	0.826667	0.96875	0.72093	STEAP2 AND CLDN8	Prostate	1	1	1
MUC17 AND CLSTN3	Stomach	0.842105	0.969697	0.744186	STEAP2 AND EPHA3	Prostate	0.8	0.666667	1
MUC17 AND VANGL1	Stomach	0.831169	0.941176	0.744186	STEAP2 AND MUC1	Prostate	1	1	1
MUC17 AND PHLDB2	Stomach	0.826667	0.96875	0.72093	STEAP2 AND VCAM1	Prostate	1	1	1
MUC17 AND STAB1	Stomach	0.831169	0.941176	0.744186	STEAP2 AND CBX3	Prostate	0.857143	1	0.75
MUC17 AND NRG3	Stomach	0.842105	0.969697	0.744186	STEAP1 AND P2RX5	Prostate	0.75	0.75	0.75
MUC17 AND ATP8B2	Stomach	0.8	0.9375	0.697674	STEAP2 AND IL13RA1	Prostate	0.888889	0.8	1
ATP8B1 AND STAB1	Stomach	0.847059	0.857143	0.837209	STEAP1 AND MUC1	Prostate	0.857143	1	0.75
MUC17 AND TTYH2	Stomach	0.842105	0.969697	0.744186	STEAP2 AND IL20RA	Prostate	0.857143	1	0.75
ATP8B1 AND ESAM	Stomach	0.888889	0.947368	0.837209	STEAP2 AND IL2RA	Prostate	0.727273	0.571429	1
ATP8B1 AND BST2	Stomach	0.837838	1	0.72093	STEAP2 AND BIRC5	Prostate	0.727273	0.571429	1
ATP8B1 AND MRGPRF	Stomach	0.819277	0.85	0.790698	STEAP2 AND ABCB5	Prostate	0.727273	0.571429	1
NKAIN4 AND ITGAV	Astrocytoma	0.746988	0.837838	0.673913	STEAP2 AND STEAP1	Prostate	0.727273	0.571429	1
BEST3 AND SLC39A6	Astrocytoma	0.708861	0.848485	0.608696	STEAP2 AND HLA-DOB	Prostate	0.727273	0.571429	1
CRB1 AND SLC39A6	Astrocytoma	0.708333	0.68	0.73913	STEAP2 AND FAP	Prostate	0.727273	0.571429	1
CRB1 AND EDNRB	Astrocytoma	0.704762	0.627119	0.804348	STEAP2 AND LGR5	Prostate	0.727273	0.571429	1

FIG. 11 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall	Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
NCAM1 AND BEST3	Astrocytoma	0.694737	0.673469	0.717391	STEAP2 AND TNFRSF10A	Prostate	0.727273	0.571429	1
BEST3 AND EDNRB	Astrocytoma	0.692308	0.84375	0.586957	STEAP2 AND CLDN3	Prostate	0.727273	0.571429	1
NCAM1 AND LAPTM5	Astrocytoma	0.690476	0.763158	0.630435	STEAP2 AND CLDN2	Prostate	0.727273	0.571429	1
SLCO1C1 AND EDNRB	Astrocytoma	0.680412	0.647059	0.717391	STEAP2 AND MS4A1	Prostate	0.727273	0.571429	1
NCAM1 AND F2R	Astrocytoma	0.680412	0.647059	0.717391	STEAP2 AND MUC16	Prostate	0.727273	0.571429	1
BEST3 AND BMPR1B	Astrocytoma	0.666667	0.8125	0.565217	STEAP2 AND CD33	Prostate	0.727273	0.571429	1
CRB1 AND NCAM1	Astrocytoma	0.659091	0.690476	0.630435	STEAP2 AND CLDN23	Prostate	0.727273	0.571429	1
BEST3 AND MUC1	Astrocytoma	0.658824	0.717949	0.608696	STEAP2 AND CD37	Prostate	0.727273	0.571429	1
BEST3 AND FOLH1	Astrocytoma	0.651163	0.7	0.608696	STEAP2 AND SST	Prostate	0.727273	0.571429	1
KCNJ10 AND CD276	Astrocytoma	0.650602	0.72973	0.586957	STEAP2 AND MOK	Prostate	0.727273	0.571429	1
NKAIN4 AND AXL	Astrocytoma	0.649351	0.806452	0.543478	STEAP2 AND SSTR1	Prostate	0.727273	0.571429	1
BEST3 AND VCAM1	Astrocytoma	0.649351	0.806452	0.543478	STEAP2 AND P2RX5	Prostate	0.727273	0.571429	1
NCAM1 AND SEMA4B	Astrocytoma	0.647619	0.576271	0.73913	STEAP2 AND WT1	Prostate	0.727273	0.571429	1
BMPR1B AND F2R	Astrocytoma	0.647619	0.576271	0.73913	STEAP2 AND CD160	Prostate	0.727273	0.571429	1
MLC1 AND CD276	Astrocytoma	0.641975	0.742857	0.565217	STEAP1 AND EPHA3	Prostate	0.857143	1	0.75
SLC1A2 AND EDNRB	Astrocytoma	0.637931	0.528571	0.804348	STEAP1 AND CD72	Prostate	0.75	0.75	0.75
ASTN1 AND EDNRB	Astrocytoma	0.635294	0.692308	0.586957	STEAP2 AND TNFRSF13C	Prostate	0.727273	0.571429	1
FXYD6 AND EDNRB	Astrocytoma	0.630631	0.538462	0.76087	STEAP1 AND IGF1R	Prostate	0.75	0.75	0.75
CSPG5 AND EDNRB	Astrocytoma	0.626506	0.702703	0.565217	STEAP1 AND SLC39A6	Prostate	0.75	0.75	0.75
BEST3 AND L1CAM	Astrocytoma	0.626506	0.702703	0.565217	STEAP1 AND CLDN4	Prostate	0.75	0.75	0.75
GABBR2 AND SLC39A6	Astrocytoma	0.622642	0.55	0.717391	STEAP1 AND ERBB2	Prostate	0.75	0.75	0.75
BEST3 AND ERBB4	Astrocytoma	0.621622	0.821429	0.5	STEAP1 AND IL20RA	Prostate	0.857143	1	0.75
CRB1 AND SLC7A5	Astrocytoma	0.62069	0.658537	0.586957	STEAP1 AND BMPR1B	Prostate	0.75	0.75	0.75
SLCO1C1 AND SLC39A6	Astrocytoma	0.619048	0.684211	0.565217	STEAP1 AND CLDN8	Prostate	0.857143	1	0.75
BEST3 AND CD79A	Astrocytoma	0.617284	0.714286	0.543478	STEAP1 AND CLDN3	Prostate	0.75	0.75	0.75
CRB1 AND SDC1	Astrocytoma	0.617021	0.604167	0.630435	STEAP2 AND MSLN	Prostate	0.727273	0.571429	1
SYT6 AND SLC39A6	Astrocytoma	0.613636	0.642857	0.586957	STEAP2 AND DNAJB8	Prostate	0.727273	0.571429	1
CRB1 AND CLDN9	Astrocytoma	0.612245	0.576923	0.652174	STEAP1 AND PTK7	Prostate	0.75	0.75	0.75
CRB1 AND PTK7	Astrocytoma	0.609756	0.694444	0.543478	STEAP1 AND SSTR1	Prostate	0.75	0.75	0.75
CRB1 AND TNFRSF10A	Astrocytoma	0.607843	0.553571	0.673913	STEAP1 AND ENPP3	Prostate	0.666667	1	0.5
SLC1A2 AND CD180	Astrocytoma	0.606742	0.627907	0.586957	STEAP2 AND ENPP3	Prostate	0.666667	1	0.5
SLC1A2 AND PTK7	Astrocytoma	0.606742	0.627907	0.586957	STEAP1 AND PROM1	Prostate	0.666667	1	0.5
KCNJ10 AND CD180	Astrocytoma	0.60241	0.675676	0.543478	STEAP1 AND CLDN7	Prostate	0.75	0.75	0.75
CRB1 AND IL3RA	Astrocytoma	0.601942	0.54386	0.673913	ABCA5 AND EPCAM	Prostate	0.666667	1	0.5
ADCY8 AND EDNRB	Astrocytoma	0.613861	0.563636	0.673913	STEAP1 AND ROR1	Prostate	0.666667	1	0.5
FAP AND PPAPDC1A	Breast	0.845361	0.953488	0.759259	STEAP1 AND CLDN18	Prostate	0.666667	0.6	0.75
FAP AND SORL1	Breast	0.75	0.970588	0.611111	STEAP1 AND NCAM1	Prostate	0.666667	0.6	0.75
PPAPDC1A AND CLDN4	Breast	0.73913	0.894737	0.62963	STEAP1 AND CLDN11	Prostate	0.666667	0.6	0.75
PPAPDC1A AND CLDN7	Breast	0.73913	0.894737	0.62963	STEAP1 AND ULBP1	Prostate	0.666667	0.6	0.75
PPAPDC1A AND SPON2	Breast	0.741573	0.942857	0.611111	STEAP1 AND ULBP2	Prostate	0.666667	0.6	0.75
PPAPDC1A AND MUC1	Breast	0.765957	0.9	0.666667	STEAP1 AND PSCA	Prostate	0.666667	0.6	0.75
PPAPDC1A AND SDC1	Breast	0.744681	0.875	0.648148	STEAP1 AND ULBP3	Prostate	0.666667	0.6	0.75
FAP AND FLVCR1	Breast	0.875	1	0.777778	STEAP1 AND SSTR2	Prostate	0.666667	0.6	0.75
FAP AND TNFRSF12A	Breast	0.795918	0.886364	0.722222	STEAP1 AND SST	Prostate	0.666667	0.6	0.75
FAP AND PPAPDC1B	Breast	0.829787	0.975	0.722222	STEAP1 AND FCRL2	Prostate	0.666667	0.6	0.75
FAP AND VANGL1	Breast	0.808511	0.95	0.703704	STEAP1 AND CR2	Prostate	0.666667	0.6	0.75
FAP AND BST2	Breast	0.83871	1	0.722222	STEAP1 AND TNFRSF10A	Prostate	0.666667	0.6	0.75
FAP AND LRRC8E	Breast	0.795699	0.948718	0.685185	STEAP1 AND MST1R	Prostate	0.666667	0.6	0.75
PPAPDC1A AND GPNMB	Breast	0.747253	0.918919	0.62963	STEAP1 AND IL3RA	Prostate	0.666667	0.6	0.75
PPAPDC1A AND ITGB6	Breast	0.742268	0.837209	0.666667	STEAP1 AND DLL3	Prostate	0.666667	0.6	0.75
FAP AND SLCO1B1	Breast	0.762887	0.860465	0.685185	FAP AND ULBP2	Sarcoma	0.740741	1	0.588235
FAP AND CXCL16	Breast	0.857143	0.954545	0.777778	FAP AND SLC7A5	Sarcoma	0.692308	1	0.529412
FAP AND IFNAR2	Breast	0.741573	0.942857	0.611111	FAP AND FOLH1	Sarcoma	0.740741	1	0.588235
TPBG AND BST2	Breast	0.712644	0.939394	0.574074	FAP AND CD276	Sarcoma	0.827586	1	0.705882
FAP AND CACNG6	Breast	0.803922	0.854167	0.759259	FAP AND IL13RA1	Sarcoma	0.692308	1	0.529412
FAP AND SLC10A1	Breast	0.838095	0.862745	0.814815	FAP AND EGFR	Sarcoma	0.827586	1	0.705882
FAP AND SEMA4D	Breast	0.833333	0.952381	0.740741	FAP AND TPBG	Sarcoma	0.740741	1	0.588235
FAP AND OXTR	Breast	0.703297	0.864865	0.592593	FAP AND ABCB5	Sarcoma	0.827586	1	0.705882
FAP AND ADAM29	Breast	0.838095	0.862745	0.814815	FAP AND ULBP1	Sarcoma	0.866667	1	0.764706
FAP AND CALHM1	Breast	0.75	0.857143	0.666667	FAP AND FCRL5	Sarcoma	0.827586	1	0.705882
PPAPDC1A AND IL13RA1	Breast	0.731183	0.871795	0.62963	MUC13 AND THY1	Stomach	0.939759	0.975	0.906977
FAP AND GABRG2	Breast	0.717391	0.868421	0.611111	EPCAM AND THY1	Stomach	0.816327	0.727273	0.930233
PPAPDC1A AND AXL	Breast	0.704545	0.911765	0.574074	CEACAM5 AND THY1	Stomach	0.857143	0.970588	0.767442
FAP AND KCND2	Breast	0.871287	0.93617	0.814815	CLDN18 AND MUC13	Stomach	0.878049	0.923077	0.837209
PPAPDC1A AND STEAP1	Breast	0.770833	0.880952	0.685185	MUC13 AND SPON2	Stomach	0.963855	1	0.930233
TPBG AND SLC38A1	Breast	0.698795	1	0.537037	CLDN18 AND BIRC5	Stomach	0.777778	0.744681	0.813953
FAP AND GUCY2D	Breast	0.77551	0.863636	0.703704	CLDN18 AND STEAP1	Stomach	0.897436	1	0.813953
PPAPDC1A AND ANXA1	Breast	0.712871	0.765957	0.666667	CLDN18 AND EPHB2	Stomach	0.857143	0.970588	0.767442
TPBG AND ENPP1	Breast	0.696629	0.885714	0.574074	CLDN18 AND THY1	Stomach	0.871795	0.971429	0.790698
FAP AND LPPR3	Breast	0.715789	0.829268	0.62963	MUC13 AND FAP	Stomach	0.860465	0.860465	0.860465
SLC2A2 AND CD34	Liver	0.8	1	0.666667	CLDN18 AND EPCAM	Stomach	0.888889	0.947368	0.837209
SLCO1B1 AND CD34	Liver	0.8	1	0.666667	CLDN18 AND MST1R	Stomach	0.810127	0.888889	0.744186
SLC43A1 AND ERBB3	Liver	0.909091	1	0.833333	CLDN18 AND CEACAM5	Stomach	0.805556	1	0.674419
SLCO1B1 AND EPHB2	Liver	1	1	1	EPCAM AND SLC7A5	Stomach	0.917647	0.928571	0.906977
SLC2A2 AND ABCB5	Liver	0.769231	0.714286	0.833333	CLDN18 AND PROM1	Stomach	0.746988	0.775	0.72093
SLC2A2 AND ERBB3	Liver	0.769231	0.714286	0.833333	MUC13 AND SLC7A5	Stomach	0.938272	1	0.883721
SLC2A2 AND IL3RA	Liver	0.769231	0.714286	0.833333	CLDN18 AND SST	Stomach	0.764045	0.73913	0.790698

FIG. 11 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
SLCO1B1 AND LGR5	Liver	0.769231	0.714286	0.833333
SLCO1B1 AND ERBB3	Liver	0.769231	0.714286	0.833333
SLC2A2 AND LGR5	Liver	0.769231	0.714286	0.833333
SLC2A2 AND ULBP1	Liver	0.769231	0.714286	0.833333
SLC13A5 AND ERBB3	Liver	0.769231	0.714286	0.833333
SLC17A2 AND EPHB2	Liver	0.909091	1	0.833333
SLC17A2 AND LGR5	Liver	0.769231	0.714286	0.833333
SLC2A2 AND SSTR5	Liver	0.8	1	0.666667
SLCO1B1 AND ITGB6	Liver	0.75	0.6	1
SLCO1B1 AND VTCN1	Liver	0.8	1	0.666667
SLC2A2 AND EPHB2	Liver	1	1	1
SLCO1B1 AND SSTR5	Liver	0.727273	0.8	0.666667
SLC2A2 AND ULBP2	Liver	0.857143	0.75	1
SLC13A5 AND CD34	Liver	0.727273	0.8	0.666667
SLC6A12 AND ERBB3	Liver	0.769231	0.714286	0.833333
SLC13A5 AND EPHB2	Liver	0.769231	0.714286	0.833333
SLCO1B1 AND IL3RA	Liver	0.714286	0.625	0.833333
SLC2A2 AND VTCN1	Liver	0.8	1	0.666667
SLCO1B1 AND ULBP2	Liver	0.75	0.6	1
SLCO1B1 AND SSTR3	Liver	0.714286	0.625	0.833333
SLC17A2 AND VTCN1	Liver	0.8	1	0.666667
SLC2A2 AND CD22	Liver	0.8	1	0.666667
SLC17A2 AND SSTR5	Liver	0.727273	0.8	0.666667
SLC10A1 AND EPHB2	Liver	0.8	1	0.666667
SLC2A2 AND L1CAM	Liver	0.857143	0.75	1
SLCO1B1 AND CLDN9	Liver	0.8	0.666667	1
SLCO1B1 AND GPA33	Liver	0.75	0.6	1
SLCO1B1 AND GUCY2C	Liver	0.75	0.6	1
SLC2A2 AND CLDN9	Liver	0.8	1	0.666667
SLC2A2 AND ROR1	Liver	0.8	1	0.666667
SLC10A1 AND IL3RA	Liver	0.714286	0.625	0.833333
SLC17A2 AND SSTR3	Liver	0.714286	0.625	0.833333
SLCO1B1 AND L1CAM	Liver	0.75	0.6	1
SLCO1B1 AND MUC4	Liver	0.75	0.6	1
SLC2A2 AND ITGB6	Liver	0.857143	0.75	1
SLCO1B1 AND ULBP1	Liver	0.666667	0.555556	0.833333
SLC17A2 AND ULBP2	Liver	0.666667	0.555556	0.833333
SLC2A2 AND TNFRSF8	Liver	0.8	1	0.666667
SLCO1B1 AND ABCB5	Liver	0.666667	0.555556	0.833333
SLC17A2 AND ITGB6	Liver	0.666667	0.555556	0.833333
SLCO1B1 AND TNFRSF8	Liver	0.8	1	0.666667
SLC17A2 AND CLDN9	Liver	0.714286	0.625	0.833333
SLC17A2 AND CD34	Liver	0.666667	1	0.5
SLC30A10 AND CLDN1	Liver	0.666667	1	0.5
SLCO1B1 AND CD22	Liver	0.8	1	0.666667
SLCO1B1 AND MST1R	Liver	0.666667	1	0.5
SLCO1B1 AND PTK7	Liver	0.666667	1	0.5
SLC2A2 AND IL2RA	Liver	0.666667	0.666667	0.666667
SLC2A2 AND PTK7	Liver	0.666667	1	0.5
SLC2A2 AND CSPG4	Liver	0.666667	0.666667	0.666667
ABCG8 AND RAET1E	Liver	0.666667	1	0.5
ALCAM AND SDC1	Liver	0.909091	1	0.833333
SLC17A2 AND ERBB3	Liver	0.666667	0.666667	0.666667
SLC2A2 AND CLDN6	Liver	0.833333	0.833333	0.833333
C8B AND EPHB2	Liver	0.666667	1	0.5
ABCG5 AND CD34	Liver	0.666667	1	0.5
ABCG8 AND CD70	Liver	0.8	1	0.666667
SLC10A1 AND LGR5	Liver	0.666667	0.666667	0.666667
SLC2A2 AND MUC4	Liver	0.857143	0.75	1
SLC17A2 AND GPA33	Liver	0.666667	0.555556	0.833333
SLC2A2 AND CR2	Liver	0.666667	0.666667	0.666667
SLC10A1 AND CD34	Liver	0.666667	1	0.5
SLC2A2 AND SSTR4	Liver	0.666667	0.666667	0.666667
SLC17A2 AND GUCY2C	Liver	0.666667	0.555556	0.833333
FAP AND CD82	Pancreas	0.842105	1	0.727273
CLDN2 AND LAMP5	Pancreas	0.736842	0.875	0.636364
FAP AND HLA-DRB1	Pancreas	0.777778	1	0.636364
FAP AND LAPTM5	Pancreas	0.705882	1	0.545455
FAP AND CD74	Pancreas	0.777778	1	0.636364
FAP AND LRRC52	Pancreas	0.777778	1	0.636364
FAP AND P2RX4	Pancreas	0.777778	1	0.636364
FAP AND TNFRSF10B	Pancreas	0.842105	1	0.727273
FAP AND CCR9	Pancreas	0.842105	1	0.727273
FAP AND SORL1	Pancreas	0.842105	1	0.727273
FAP AND CXCL16	Pancreas	0.842105	1	0.727273
FAP AND SEMA4D	Pancreas	0.842105	1	0.727273
FAP AND UMODL1	Pancreas	0.842105	1	0.727273

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
CLDN18 AND GUCY2C	Stomach	0.73913	0.693878	0.790698
EPCAM AND FAP	Stomach	0.817204	0.76	0.883721
CLDN18 AND SPON2	Stomach	0.795455	0.777778	0.813953
CLDN18 AND PSCA	Stomach	0.682927	0.717949	0.651163
CLDN18 AND ERBB3	Stomach	0.888889	0.947368	0.837209
CLDN18 AND CBX3	Stomach	0.853333	1	0.744186
CLDN18 AND SSTR4	Stomach	0.725275	0.6875	0.767442
CLDN18 AND MAGEA1	Stomach	0.808989	0.782609	0.837209
CLDN18 AND FCRL5	Stomach	0.8	0.765957	0.837209
CLDN18 AND AFP	Stomach	0.765957	0.705882	0.837209
MUC13 AND CLDN1	Stomach	0.965517	0.954545	0.976744
CA9 AND EPCAM	Stomach	0.712329	0.866667	0.604651
CEACAM5 AND SPON2	Stomach	0.839506	0.894737	0.790698
CLDN18 AND GAGE1	Stomach	0.774194	0.72	0.837209
CLDN18 AND SSTR2	Stomach	0.765957	0.705882	0.837209
CLDN18 AND DLL3	Stomach	0.795455	0.777778	0.813953
CEACAM5 AND FAP	Stomach	0.837838	1	0.72093
CLDN18 AND ULBP1	Stomach	0.765957	0.705882	0.837209
CLDN18 AND TNFSF11	Stomach	0.786517	0.76087	0.813953
CLDN18 AND NCAM1	Stomach	0.75	0.733333	0.767442
MUC13 AND CD34	Stomach	0.845361	0.759259	0.953488
CLDN18 AND MAGEA11	Stomach	0.747253	0.708333	0.790698
CLDN18 AND ABCB5	Stomach	0.765957	0.705882	0.837209
CLDN18 AND MUC16	Stomach	0.73913	0.693878	0.790698
CLDN18 AND DNAJB8	Stomach	0.808989	0.782609	0.837209
CLDN18 AND MUC4	Stomach	0.774194	0.72	0.837209
CLDN18 AND SSX1	Stomach	0.75	0.733333	0.767442
CLDN18 AND CA9	Stomach	0.674699	0.7	0.651163
CLDN18 AND SSTR5	Stomach	0.808989	0.782609	0.837209
CLDN18 AND RNF43	Stomach	0.837209	0.837209	0.837209
CLDN18 AND L1CAM	Stomach	0.777778	0.744681	0.813953
EPCAM AND SPON2	Stomach	0.931818	0.911111	0.953488
CLDN18 AND CD19	Stomach	0.765957	0.705882	0.837209
CLDN7 AND THY1	Stomach	0.926829	0.974359	0.883721
CLDN18 AND IL2RA	Stomach	0.781609	0.772727	0.790698
CLDN18 AND TRPM4	Stomach	0.888889	0.947368	0.837209
CLDN18 AND CLDN1	Stomach	0.9	0.972973	0.837209
CLDN18 AND WT1	Stomach	0.765957	0.705882	0.837209
CLDN18 AND SSTR3	Stomach	0.791209	0.75	0.837209
CLDN18 AND BCAN	Stomach	0.808989	0.782609	0.837209
CLDN18 AND ERBB2	Stomach	0.888889	0.947368	0.837209
CLDN18 AND PMEL	Stomach	0.765957	0.705882	0.837209
CLDN18 AND OAS1	Stomach	0.735294	1	0.581395
CLDN18 AND SLC39A6	Stomach	0.86747	0.9	0.837209
PTPRZ1 AND F2R	Astrocytoma	0.866667	0.886364	0.847826
PTPRZ1 AND BEST3	Astrocytoma	0.857143	0.866667	0.847826
FAIM2 AND F2R	Astrocytoma	0.840909	0.880952	0.804348
PLP1 AND F2R	Astrocytoma	0.831461	0.860465	0.804348
CRB1 AND ABCA1	Astrocytoma	0.822222	0.840909	0.804348
NRCAM AND F2R	Astrocytoma	0.813953	0.875	0.76087
NKAIN4 AND PTPRZ1	Astrocytoma	0.809524	0.894737	0.73913
MLC1 AND HAS2	Astrocytoma	0.804598	0.853659	0.76087
PTPRZ1 AND CD82	Astrocytoma	0.804348	0.804348	0.804348
SLC1A2 AND VANGL2	Astrocytoma	0.804124	0.764706	0.847826
KCNJ10 AND TSPAN11	Astrocytoma	0.8	0.818182	0.782609
KCNJ10 AND VANGL2	Astrocytoma	0.8	0.818182	0.782609
PTPRZ1 AND FXYD6	Astrocytoma	0.8	0.740741	0.869565
GABBR2 AND VANGL2	Astrocytoma	0.795699	0.787234	0.804348
PTPRZ1 AND GALR1	Astrocytoma	0.795699	0.787234	0.804348
PTPRZ1 AND STAB1	Astrocytoma	0.791667	0.76	0.826087
PTPRZ1 AND SYT6	Astrocytoma	0.791667	0.76	0.826087
CRB1 AND ITGB8	Astrocytoma	0.791209	0.8	0.782609
SLCO1C1 AND ABCA1	Astrocytoma	0.790698	0.85	0.73913
GPM6B AND BEST3	Astrocytoma	0.790123	0.914286	0.695652
ADAM22 AND F2R	Astrocytoma	0.786517	0.813953	0.76087
NRCAM AND SEMA4B	Astrocytoma	0.786517	0.813953	0.76087
MLC1 AND ABCA1	Astrocytoma	0.785714	0.868421	0.717391
PLP1 AND BEST3	Astrocytoma	0.78481	0.939394	0.673913
NKAIN4 AND ABCA1	Astrocytoma	0.781609	0.829268	0.73913
PTPRZ1 AND SLC10A1	Astrocytoma	0.78	0.722222	0.847826
CRB1 AND VANGL2	Astrocytoma	0.778947	0.755102	0.804348
MLC1 AND BMPR1A	Astrocytoma	0.777778	0.795455	0.76087
CRB1 AND DLL1	Astrocytoma	0.777778	0.795455	0.76087
CRB1 AND CALCRL	Astrocytoma	0.776471	0.846154	0.717391
NKAIN4 AND SNAP23	Astrocytoma	0.776471	0.846154	0.717391
CRB1 AND ITM2B	Astrocytoma	0.776471	0.846154	0.717391
TTYH2 AND VANGL2	Astrocytoma	0.774194	0.765957	0.782609

FIG. 11 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
EPHA3 AND CLDN10	Pancreas	0.625	1	0.454545
FAP AND GRM4	Pancreas	0.625	1	0.454545
FAP AND NMBR	Pancreas	0.842105	1	0.727273
FAP AND CRB1	Pancreas	0.842105	1	0.727273
FAP AND EFNB2	Pancreas	0.777778	1	0.636364
FAP AND ICAM1	Pancreas	0.842105	1	0.727273
FAP AND CCKAR	Pancreas	0.705882	1	0.545455
EPHA3 AND CD58	Pancreas	0.736842	0.875	0.636364
FAP AND KCNU1	Pancreas	0.842105	1	0.727273
FAP AND SLC6A18	Pancreas	0.777778	1	0.636364
FAP AND PPAPDC1A	Pancreas	0.777778	1	0.636364
FAP AND OTOF	Pancreas	0.842105	1	0.727273
FAP AND KCNV2	Pancreas	0.842105	1	0.727273
FAP AND SLC2A2	Pancreas	0.842105	1	0.727273
FAP AND MEGF11	Pancreas	0.842105	1	0.727273
FAP AND DPP10	Pancreas	0.705882	1	0.545455
SLC30A8 AND CD52	Pancreas	0.777778	1	0.636364
CLDN2 AND KCND2	Pancreas	0.8	0.888889	0.727273
EPHA3 AND LAMP5	Pancreas	0.777778	1	0.636364
EPHA3 AND PTGIR	Pancreas	0.736842	0.875	0.636364
FAP AND KCNK12	Pancreas	0.705882	1	0.545455
FAP AND CDH18	Pancreas	0.842105	1	0.727273
FAP AND CD163	Pancreas	0.777778	1	0.636364
FAP AND CATSPERD	Pancreas	0.842105	1	0.727273
FAP AND SLC17A1	Pancreas	0.842105	1	0.727273
FAP AND GGTLC1	Pancreas	0.625	1	0.454545
FAP AND OPRK1	Pancreas	0.705882	1	0.545455
FAP AND IFNAR2	Pancreas	0.705882	1	0.545455
FAP AND PPAPDC1B	Pancreas	0.705882	1	0.545455
FAP AND PCDHA6	Pancreas	0.777778	1	0.636364
FAP AND EBP	Pancreas	0.625	1	0.454545
FAP AND CD58	Pancreas	0.842105	1	0.727273
FAP AND SLC17A3	Pancreas	0.842105	1	0.727273
FAP AND KCNJ6	Pancreas	0.842105	1	0.727273
FAP AND SLC26A8	Pancreas	0.842105	1	0.727273
FAP AND NPFFR1	Pancreas	0.777778	1	0.636364
CLDN18 AND KCND2	Pancreas	0.631579	0.75	0.545455
SLC30A8 AND GPNMB	Pancreas	0.777778	1	0.636364
FAP AND GPR22	Pancreas	0.777778	1	0.636364
FAP AND SPAM1	Pancreas	0.705882	1	0.545455
FAP AND FCAMR	Pancreas	0.842105	1	0.727273
FAP AND OR52D1	Pancreas	0.777778	1	0.636364
FAP AND DIO1	Pancreas	0.842105	1	0.727273
EPHA3 AND PTGER4	Pancreas	0.777778	1	0.636364
FAP AND OR2L2	Pancreas	0.842105	1	0.727273
FAP AND AJAP1	Pancreas	0.705882	1	0.545455
EPHA3 AND HLA-DRB1	Pancreas	0.777778	1	0.636364
FAP AND CLDN10	Pancreas	0.705882	1	0.545455
FAP AND USH2A	Pancreas	0.777778	1	0.636364
FAP AND ZP4	Pancreas	0.842105	1	0.727273
FAP AND OPRM1	Pancreas	0.842105	1	0.727273
FAP AND SLC12A5	Pancreas	0.842105	1	0.727273
FAP AND TNFRSF12A	Pancreas	0.842105	1	0.727273
FAP AND GPR83	Pancreas	0.842105	1	0.727273
FAP AND CHRNA1	Pancreas	0.842105	1	0.727273
VCAM1 AND GGTLC1	Renal	0.8	1	0.666667
VCAM1 AND FAT1	Renal	0.8	1	0.666667
VCAM1 AND SLC22A5	Renal	0.736842	1	0.583333
VCAM1 AND SLC22A18	Renal	0.8	1	0.666667
CLDN2 AND GYPC	Renal	0.727273	0.8	0.666667
CLDN2 AND EMP3	Renal	0.727273	0.8	0.666667
VCAM1 AND CLDN10	Renal	0.8	1	0.666667
CLDN2 AND SLC38A1	Renal	0.736842	1	0.583333
SLC22A2 AND VCAM1	Renal	0.7	0.875	0.583333
CLDN2 AND LAPTM5	Renal	0.8	1	0.666667
SLC17A1 AND VCAM1	Renal	0.666667	1	0.5
VCAM1 AND SLC6A12	Renal	0.666667	1	0.5
VCAM1 AND SLC22A11	Renal	0.666667	1	0.5
CLDN2 AND P2RX4	Renal	0.727273	0.8	0.666667
CLDN2 AND GPR137B	Renal	0.64	0.615385	0.666667
SLC13A1 AND VCAM1	Renal	0.631579	0.857143	0.5
SLC6A13 AND VCAM1	Renal	0.631579	0.857143	0.5
VCAM1 AND TNFRSF12A	Renal	0.666667	1	0.5
CLDN2 AND VANGL1	Renal	0.636364	0.7	0.583333
VCAM1 AND ABCA12	Renal	0.666667	0.6	0.75
VCAM1 AND TRPM3	Renal	0.631579	0.857143	0.5
CLDN2 AND CD58	Renal	0.695652	0.727273	0.666667

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
PTPRZ1 AND CD44	Astrocytoma	0.773333	1	0.630435
NRCAM AND BEST3	Astrocytoma	0.772727	0.809524	0.73913
SLCO1C1 AND CALCRL	Astrocytoma	0.771084	0.864865	0.695652
PCDH10 AND BEST3	Astrocytoma	0.771084	0.864865	0.695652
CRB1 AND ABCC4	Astrocytoma	0.771084	0.864865	0.695652
SLCO1C1 AND F2RL1	Astrocytoma	0.771084	0.864865	0.695652
NKAIN4 AND WLS	Astrocytoma	0.771084	0.864865	0.695652
MLC1 AND VANGL2	Astrocytoma	0.769231	0.777778	0.76087
GPM6A AND F2R	Astrocytoma	0.769231	0.777778	0.76087
ADCYAP1R1 AND BEST3	Astrocytoma	0.767442	0.825	0.717391
PTPRZ1 AND CLDN17	Astrocytoma	0.766355	0.672131	0.891304
NKAIN4 AND SLC25A3	Astrocytoma	0.765432	0.885714	0.673913
ADAM12 AND PPAPDC1A	Breast	0.830189	0.846154	0.814815
ADAM12 AND SORL1	Breast	0.815534	0.857143	0.777778
ADAM12 AND LAMP5	Breast	0.807339	0.8	0.814815
ADAM12 AND LRRC8E	Breast	0.787879	0.866667	0.722222
ADAM12 AND VANGL1	Breast	0.783505	0.883721	0.703704
BST2 AND PPAP2C	Breast	0.777778	0.972222	0.648148
ADAM12 AND MEGF10	Breast	0.77551	0.863636	0.703704
ADAM12 AND UNC5A	Breast	0.770833	0.880952	0.685185
SLC5A6 AND KCNA4	Breast	0.777778	0.972222	0.648148
PRLR AND FLVCR1	Breast	0.76087	0.921053	0.648148
ADAM12 AND NKAIN1	Breast	0.757895	0.878049	0.666667
SLC5A6 AND ADAM29	Breast	0.777778	0.972222	0.648148
SLC5A6 AND SLC10A1	Breast	0.777778	0.972222	0.648148
SLC5A6 AND SLC22A14	Breast	0.777778	0.972222	0.648148
SLC5A6 AND GLRA2	Breast	0.777778	0.972222	0.648148
PPAPDC1A AND MFSD10	Breast	0.744186	1	0.592593
BST2 AND PROM2	Breast	0.782609	0.947368	0.666667
PPAPDC1A AND FXYD3	Breast	0.787234	0.925	0.685185
PPAPDC1A AND F11R	Breast	0.744681	0.875	0.648148
PPAPDC1A AND PROM2	Breast	0.755102	0.840909	0.685185
PPAPDC1A AND SLC52A2	Breast	0.786517	1	0.648148
CELSR2 AND BST2	Breast	0.736842	0.853659	0.648148
SLC5A6 AND SCN10A	Breast	0.777778	0.972222	0.648148
SLC5A6 AND MIP	Breast	0.735632	0.969697	0.592593
ADAM12 AND CNIH2	Breast	0.734694	0.818182	0.666667
PPAPDC1A AND LSR	Breast	0.733333	0.916667	0.611111
PRLR AND CXCL16	Breast	0.73913	0.894737	0.62963
SLC5A6 AND SLC10A2	Breast	0.764045	0.971429	0.62963
SLC5A6 AND KCND2	Breast	0.777778	0.972222	0.648148
PPAPDC1A AND PLP2	Breast	0.744186	1	0.592593
PPAPDC1A AND CDH1	Breast	0.744681	0.875	0.648148
PPAPDC1A AND CALHM2	Breast	0.741573	0.942857	0.611111
PPAPDC1A AND EDNRA	Breast	0.782609	0.947368	0.666667
PRLR AND IL17RA	Breast	0.731183	0.871795	0.62963
PPAPDC1A AND TCIRG1	Breast	0.755556	0.944444	0.62963
SLC5A6 AND OR1C1	Breast	0.735632	0.969697	0.592593
SLC5A6 AND PPAPDC1A	Breast	0.729412	1	0.574074
SLC5A6 AND LRRC8E	Breast	0.729412	1	0.574074
PPAPDC1A AND TNFSF10	Breast	0.730769	0.76	0.703704
SLC5A6 AND CLDN19	Breast	0.735632	0.969697	0.592593
SLC5A6 AND GPR19	Breast	0.755556	0.944444	0.62963
PPAPDC1A AND SECTM1	Breast	0.747253	0.918919	0.62963
PPAPDC1A AND NOX4	Breast	0.76	0.826087	0.703704
PPAPDC1A AND PLXND1	Breast	0.744186	1	0.592593
MFSD10 AND CDH11	Breast	0.735632	0.969697	0.592593
SLC2A2 AND FGFRL1	Liver	1	1	1
FGG AND FGF6	Liver	0.909091	1	0.833333
SLCO1B1 AND SLC26A6	Liver	0.923077	0.857143	1
SLC2A2 AND CELSR3	Liver	0.857143	0.75	1
SLC2A2 AND CNNM3	Liver	0.857143	0.75	1
FGG AND SLC22A5	Liver	0.833333	0.833333	0.833333
SLC2A2 AND TREML2	Liver	0.909091	1	0.833333
SLC2A2 AND SLC38A6	Liver	0.909091	1	0.833333
SLC2A2 AND PANX2	Liver	1	1	1
SLC2A2 AND SLC26A6	Liver	1	1	1
SLC17A2 AND SLC26A6	Liver	0.833333	0.833333	0.833333
SLC2A2 AND ANO8	Liver	0.833333	0.833333	0.833333
SLC2A2 AND ABCB8	Liver	0.909091	1	0.833333
FGG AND ALCAM	Liver	1	1	1
FGG AND KCNJ6	Liver	0.909091	1	0.833333
ABCG5 AND FGFRL1	Liver	1	1	1
SLCO1B1 AND HM13	Liver	1	1	1
SLC2A2 AND BCAM	Liver	0.909091	1	0.833333
SLC2A2 AND TP53I13	Liver	1	1	1
SLC2A2 AND NPBWR2	Liver	0.909091	1	0.833333

FIG. 11 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
VCAM1 AND SLC38A1	Renal	0.615385	0.571429	0.666667
CLDN2 AND ABCG4	Renal	0.615385	0.571429	0.666667
CLDN2 AND CD44	Renal	0.666667	0.666667	0.666667
CLDN2 AND HLA-DRB1	Renal	0.761905	0.888889	0.666667
VCAM1 AND EPHA8	Renal	0.615385	0.571429	0.666667
CLDN2 AND ICAM1	Renal	0.666667	1	0.5
VCAM1 AND CALHM3	Renal	0.666667	0.6	0.75
VCAM1 AND LRRN4	Renal	0.666667	0.6	0.75
SLC13A1 AND CD276	Renal	0.6	0.75	0.5
SLC13A1 AND AXL	Renal	0.6	0.75	0.5
VCAM1 AND SLC2A1	Renal	0.636364	0.7	0.583333
VCAM1 AND TNFRSF10B	Renal	0.615385	0.571429	0.666667
DPEP1 AND TGFBI	Colon	1	1	1
IFI6 AND CEACAM5	Colon	1	1	1
CDH17 AND CLDN1	Colon	1	1	1
IFI6 AND GUCY2C	Colon	1	1	1
GPA33 AND TGFBI	Colon	0.909091	0.833333	1
CLDN7 AND TGFBI	Colon	0.909091	0.833333	1
IFI6 AND GPA33	Colon	1	1	1
CDH17 AND THY1	Colon	1	1	1
MUC13 AND TGFBI	Colon	0.909091	0.833333	1
IFI6 AND MUC13	Colon	1	1	1
CEACAM5 AND TGFBI	Colon	0.909091	0.833333	1
GUCY2C AND TGFBI	Colon	0.909091	0.833333	1
DPEP1 AND IFI6	Colon	1	1	1
CLDN3 AND TGFBI	Colon	0.909091	0.833333	1
CDH17 AND SLC7A5	Colon	1	1	1
IFI6 AND EPHB2	Colon	0.909091	0.833333	1
CLDN2 AND TGFBI	Colon	0.888889	1	0.8
NOX1 AND CLDN1	Colon	1	1	1
IFI6 AND STEAP1	Colon	1	1	1
DPEP1 AND NOX1	Colon	1	1	1
NOX1 AND THY1	Colon	1	1	1
IFI6 AND CLDN3	Colon	1	1	1
CDH17 AND FAP	Colon	0.888889	1	0.8
CXCL16 AND MUC13	Esophagus	0.782609	1	0.642857
CDH17 AND CLDN1	Esophagus	0.928571	0.928571	0.928571
SLCO1B3 AND EPCAM	Esophagus	0.636364	0.875	0.5
CXCL16 AND CEACAM5	Esophagus	0.782609	1	0.642857
CDH17 AND MOK	Esophagus	0.666667	1	0.5
MUC17 AND CLDN1	Esophagus	0.88	1	0.785714
SLCO1B3 AND SLC7A5	Esophagus	0.6	1	0.428571
MUC13 AND BST2	Esophagus	0.6	1	0.428571
SLCO1B3 AND BIRC5	Esophagus	0.636364	0.875	0.5
CBX3 AND SLC2A1	Esophagus	0.695652	0.888889	0.571429
CDH17 AND SPON2	Esophagus	0.685714	0.571429	0.857143
SLCO1B3 AND MUC1	Esophagus	0.636364	0.875	0.5
SLCO1B3 AND IL20RA	Esophagus	0.636364	0.875	0.5
CXCL16 AND GUCY2C	Esophagus	0.727273	1	0.571429
CBX3 AND EFNB2	Esophagus	0.695652	0.888889	0.571429
CXCL16 AND EPCAM	Esophagus	0.782609	1	0.642857
LYPD1 AND CD33	Glioblastoma	0.764706	0.764706	0.764706
MLC1 AND IL13RA1	Glioblastoma	0.717949	0.636364	0.823529
LYPD1 AND CD180	Glioblastoma	0.702703	0.65	0.764706
LYPD1 AND AXL	Glioblastoma	0.848485	0.875	0.823529
LYPD1 AND SLC7A5	Glioblastoma	0.7	0.608696	0.823529
LYPD1 AND CD70	Glioblastoma	0.7	0.608696	0.823529
LYPD1 AND CLDN5	Glioblastoma	0.7	0.608696	0.823529
MLC1 AND VCAM1	Glioblastoma	0.742857	0.722222	0.764706
CRB1 AND ITGAV	Glioblastoma	0.692308	1	0.529412
LYPD1 AND CSPG4	Glioblastoma	0.692308	1	0.529412
LYPD1 AND IL13RA1	Glioblastoma	0.823529	0.823529	0.823529
LYPD1 AND SLC39A6	Glioblastoma	0.684211	0.619048	0.764706
MLC1 AND GPNMB	Glioblastoma	0.709677	0.785714	0.647059
LYPD1 AND BMPR1B	Glioblastoma	0.717949	0.636364	0.823529
LYPD1 AND NCAM1	Glioblastoma	0.666667	0.590909	0.764706
LYPD1 AND MUC1	Glioblastoma	0.666667	0.9	0.529412
SYT11 AND GPNMB	Glioblastoma	0.666667	0.56	0.823529
SYT11 AND ITGAV	Glioblastoma	0.717949	0.636364	0.823529
SLC1A2 AND IL13RA1	Glioblastoma	0.692308	1	0.529412
MLC1 AND EGFR	Glioblastoma	0.647059	0.647059	0.647059
SYT11 AND MUC1	Glioblastoma	0.642857	0.818182	0.529412
CDH10 AND MUC1	Glioblastoma	0.642857	0.818182	0.529412
MLC1 AND ITGAV	Glioblastoma	0.666667	0.769231	0.588235
LYPD1 AND ITGAV	Glioblastoma	0.740741	1	0.588235
LYPD1 AND PROM1	Glioblastoma	0.631579	0.571429	0.705882
GPR158 AND ITGAV	Glioblastoma	0.666667	0.56	0.823529

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
SLC2A2 AND TMEM67	Liver	0.857143	0.75	1
SLCO1B1 AND ANO10	Liver	0.909091	1	0.833333
ABCG5 AND SLC38A6	Liver	0.909091	1	0.833333
HPN AND PLVAP	Liver	0.8	1	0.666667
FGG AND MLANA	Liver	0.8	1	0.666667
SLCO1B1 AND CELSR3	Liver	0.8	0.666667	1
SLC2A2 AND SLC5A5	Liver	0.909091	1	0.833333
FGG AND GPR158	Liver	0.8	0.666667	1
ABCG5 AND SMO	Liver	0.8	1	0.666667
ASGR1 AND PLVAP	Liver	0.8	1	0.666667
SLCO1B1 AND C6orf89	Liver	1	1	1
SLC2A2 AND DAGLB	Liver	0.8	1	0.666667
SLC2A2 AND SMO	Liver	0.8	1	0.666667
SLC2A2 AND HM13	Liver	1	1	1
FGG AND SLC50A1	Liver	1	1	1
SLC2A2 AND SLC4A11	Liver	0.8	1	0.666667
SLCO1B1 AND SMO	Liver	0.8	1	0.666667
FGG AND PLVAP	Liver	0.8	1	0.666667
FGG AND NPFFR1	Liver	0.8	1	0.666667
SLC2A2 AND PTCHD1	Liver	0.833333	0.833333	0.833333
SLC2A2 AND PLXNC1	Liver	0.8	1	0.666667
SLCO1B1 AND SLC4A11	Liver	0.833333	0.833333	0.833333
SLCO1B1 AND CD200	Liver	0.923077	0.857143	1
SLC2A2 AND KCNN3	Liver	0.909091	1	0.833333
SLC2A2 AND CRHR1	Liver	0.8	1	0.666667
SLC2A2 AND ATP2A2	Liver	0.909091	1	0.833333
SLC2A2 AND NOTCH4	Liver	1	1	1
FGG AND PCDHB1	Liver	0.8	0.666667	1
SLCO1B1 AND DAGLB	Liver	0.8	1	0.666667
SLC2A2 AND SLC12A7	Liver	0.909091	1	0.833333
SLCO1B1 AND TLCD1	Liver	1	1	1
SLC17A2 AND HM13	Liver	0.909091	1	0.833333
SLCO1B1 AND FGFRL1	Liver	1	1	1
ABCB4 AND PLVAP	Liver	0.8	1	0.666667
SLC2A2 AND IGSF8	Liver	1	1	1
SLC2A2 AND SLC39A1	Liver	0.909091	1	0.833333
FGG AND DTNA	Liver	0.909091	1	0.833333
SLC2A2 AND TSPAN5	Liver	0.833333	0.833333	0.833333
SLC2A2 AND QSOX2	Liver	0.909091	1	0.833333
FGG AND SLC22A12	Liver	0.833333	0.833333	0.833333
SLC13A5 AND SMO	Liver	0.8	1	0.666667
SLC2A2 AND C6orf89	Liver	1	1	1
SLC2A2 AND SLC26A11	Liver	0.909091	1	0.833333
SLCO1B1 AND SLC7A11	Liver	1	1	1
SLCO1B1 AND NIPA1	Liver	0.909091	1	0.833333
SLC2A2 AND MMP14	Liver	0.923077	0.857143	1
SLC2A2 AND TIE1	Liver	0.857143	0.75	1
SLCO1B1 AND SLC38A6	Liver	0.909091	1	0.833333
SLCO1B1 AND SLC39A1	Liver	0.909091	1	0.833333
SLCO1B1 AND NPBWR2	Liver	0.909091	1	0.833333
ABCG5 AND CELSR3	Liver	0.923077	0.857143	1
ABCC6 AND PLVAP	Liver	0.8	1	0.666667
SLCO1B1 AND NCSTN	Liver	1	1	1
SLC2A2 AND SLC16A11	Liver	0.833333	0.833333	0.833333
SLCO1B1 AND TP53I13	Liver	0.923077	0.857143	1
ASGR1 AND FGF6	Liver	0.833333	0.833333	0.833333
SLCO1B1 AND TREML2	Liver	0.909091	1	0.833333
SLC27A5 AND PLVAP	Liver	0.8	1	0.666667
SLCO1B1 AND PANX2	Liver	1	1	1
SLCO1B1 AND LRRC8D	Liver	0.909091	1	0.833333
SLCO1B1 AND ABCB8	Liver	0.857143	0.75	1
SLC17A2 AND ANO10	Liver	0.8	1	0.666667
TNFRSF12A AND ADAM28	Pancreas	0.777778	1	0.636364
KCNE4 AND IFNAR2	Pancreas	0.666667	0.615385	0.727273
SYT13 AND EMP3	Pancreas	0.8	0.888889	0.727273
NOX4 AND TNFRSF10B	Pancreas	0.666667	0.7	0.636364
PTGIS AND GJB1	Pancreas	0.666667	0.615385	0.727273
NOX4 AND TNFRSF12A	Pancreas	0.636364	0.636364	0.636364
KCNE4 AND TNFRSF12A	Pancreas	0.636364	0.636364	0.636364
TNFRSF12A AND TNFSF4	Pancreas	0.705882	1	0.545455
NOX4 AND UMODL1	Pancreas	0.72	0.642857	0.818182
SYT13 AND BST2	Pancreas	0.842105	1	0.727273
SYT13 AND PTGIS	Pancreas	0.628571	0.458333	1
NOX4 AND LAPTM5	Pancreas	0.625	1	0.454545
NOX4 AND CD58	Pancreas	0.705882	1	0.545455
CLDN10 AND CXCR4	Pancreas	0.705882	1	0.545455
GP2 AND PTGIS	Pancreas	0.8	0.888889	0.727273

FIG. 11 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
NCAM1 AND PPAPDC1B	Glioblastoma	0.645161	0.714286	0.588235
SLCO1C1 AND IL13RA1	Glioblastoma	0.625	0.483871	0.882353
BEST3 AND BMPR1B	Glioblastoma	0.625	0.666667	0.588235
SLCO1C1 AND MUC1	Glioblastoma	0.62069	0.75	0.529412
CRB1 AND VCAM1	Glioblastoma	0.611111	0.578947	0.647059
SCN1A AND ITGAV	Glioblastoma	0.666667	0.769231	0.588235
GPR19 AND ITGAV	Glioblastoma	0.62069	0.75	0.529412
SLCO1C1 AND GPNMB	Glioblastoma	0.684211	0.619048	0.764706
GPR158 AND GPNMB	Glioblastoma	0.603774	0.444444	0.941176
LYPD1 AND VCAM1	Glioblastoma	0.827586	1	0.705882
SLCO1C1 AND ITGAV	Glioblastoma	0.742857	0.722222	0.764706
ASTN1 AND ITGAV	Glioblastoma	0.666667	0.9	0.529412
BMPR1B AND CD163	Glioblastoma	0.827586	1	0.705882
NCAM1 AND IFNAR2	Glioblastoma	0.666667	0.6875	0.647059
SLCO1C1 AND CLDN12	Glioblastoma	0.615385	0.457143	0.941176
PPAPDC1A AND ITGAV	Glioblastoma	0.631579	0.571429	0.705882
NCAM1 AND SLC31A1	Glioblastoma	0.689655	0.833333	0.588235
SYT11 AND ITGAV	Glioma	0.977778	1	0.956522
GPR158 AND ITGAV	Glioma	0.948905	0.955882	0.942029
KCNJ10 AND ITGAV	Glioma	0.906475	0.9	0.913043
SYT11 AND CLDN8	Glioma	0.957143	0.943662	0.971014
GPR19 AND ITGAV	Glioma	0.913386	1	0.84058
SYT11 AND FCRL5	Glioma	0.971429	0.957746	0.985507
SYT11 AND CD276	Glioma	0.964029	0.957143	0.971014
SYT11 AND ENG	Glioma	0.964539	0.944444	0.985507
SYT11 AND GUCY2C	Glioma	0.957143	0.943662	0.971014
SYT11 AND CLDN23	Glioma	0.964029	0.957143	0.971014
SYT11 AND EDNRB	Glioma	0.942029	0.942029	0.942029
SYT11 AND FCRL1	Glioma	0.934307	0.941176	0.927536
SYT11 AND CD37	Glioma	0.964539	0.944444	0.985507
SYT11 AND ABCB5	Glioma	0.957143	0.943662	0.971014
SYT11 AND ITGB6	Glioma	0.957143	0.943662	0.971014
SYT11 AND TNFRSF17	Glioma	0.957143	0.943662	0.971014
SYT11 AND CD180	Glioma	0.964539	0.944444	0.985507
SYT11 AND TNFSF11	Glioma	0.956522	0.956522	0.956522
SYT11 AND PTK7	Glioma	0.94964	0.942857	0.956522
SYT11 AND SDC1	Glioma	0.942029	0.942029	0.942029
SYT11 AND ULBP1	Glioma	0.948905	0.955882	0.942029
SYT11 AND VTCN1	Glioma	0.94964	0.942857	0.956522
SYT11 AND SSTR4	Glioma	0.94964	0.942857	0.956522
ASTN1 AND ITGAV	Glioma	0.888889	0.982456	0.811594
SYT11 AND IL2RA	Glioma	0.964539	0.944444	0.985507
SYT11 AND CD72	Glioma	0.957143	0.943662	0.971014
SYT11 AND CLDN2	Glioma	0.94964	0.942857	0.956522
SYT11 AND SLC34A2	Glioma	0.942029	0.942029	0.942029
SYT11 AND MUC4	Glioma	0.964539	0.944444	0.985507
SYT11 AND SSTR3	Glioma	0.964539	0.944444	0.985507
SYT11 AND CD52	Glioma	0.964029	0.957143	0.971014
CDH10 AND ITGAV	Glioma	0.878049	1	0.782609
SYT11 AND SLC39A6	Glioma	0.956522	0.956522	0.956522
SYT11 AND KDR	Glioma	0.964539	0.944444	0.985507
SYT11 AND CR2	Glioma	0.94964	0.942857	0.956522
SYT11 AND MS4A1	Glioma	0.957143	0.943662	0.971014
SYT11 AND MSLN	Glioma	0.964539	0.944444	0.985507
SYT11 AND CD33	Glioma	0.964539	0.944444	0.985507
SYT11 AND CLDN18	Glioma	0.918519	0.939394	0.898551
SYT11 AND AXL	Glioma	0.957143	0.943662	0.971014
SYT11 AND IL3RA	Glioma	0.957143	0.943662	0.971014
SYT11 AND CD34	Glioma	0.964539	0.944444	0.985507
SYT11 AND ROR1	Glioma	0.964539	0.944444	0.985507
SYT11 AND FCRL2	Glioma	0.957143	0.943662	0.971014
SYT11 AND MUC16	Glioma	0.964539	0.944444	0.985507
SYT11 AND RNF43	Glioma	0.957143	0.943662	0.971014
SYT11 AND GPA33	Glioma	0.957143	0.943662	0.971014
SYT11 AND RAET1E	Glioma	0.964539	0.944444	0.985507
SYT11 AND CA9	Glioma	0.957143	0.943662	0.971014
SYT11 AND TNFRSF10A	Glioma	0.94964	0.942857	0.956522
SYT11 AND MST1R	Glioma	0.948905	0.955882	0.942029
SYT11 AND SSTR5	Glioma	0.94964	0.942857	0.956522
SYT11 AND PMEL	Glioma	0.964539	0.944444	0.985507
SYT11 AND SLAMF7	Glioma	0.964539	0.944444	0.985507
SYT11 AND CXCR5	Glioma	0.957143	0.943662	0.971014
OMG AND ITGAV	Glioma	0.867133	0.837838	0.898551
SYT11 AND TNFRSF13C	Glioma	0.964539	0.944444	0.985507
SYT11 AND CD38	Glioma	0.964539	0.944444	0.985507
SYT11 AND IL20RA	Glioma	0.94964	0.942857	0.956522
SYT11 AND CLDN9	Glioma	0.957143	0.943662	0.971014

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
GP2 AND CDH11	Pancreas	0.777778	1	0.636364
SLC30A8 AND VMP1	Pancreas	0.705882	1	0.545455
GP2 AND BST2	Pancreas	0.705882	1	0.545455
NOX4 AND PCDHA6	Pancreas	0.64	0.571429	0.727273
GP2 AND F2R	Pancreas	0.842105	1	0.727273
GP2 AND LAMP5	Pancreas	0.705882	1	0.545455
NOX4 AND CD82	Pancreas	0.608696	0.583333	0.636364
KCNE4 AND LAPTM5	Pancreas	0.608696	0.583333	0.636364
NOX4 AND NMBR	Pancreas	0.72	0.642857	0.818182
GP2 AND CD163	Pancreas	0.842105	1	0.727273
KCNE4 AND HLA-DRB1	Pancreas	0.608696	0.583333	0.636364
KCNE4 AND CD74	Pancreas	0.608696	0.583333	0.636364
SLC30A8 AND TM4SF1	Pancreas	0.705882	1	0.545455
GJB1 AND EMP3	Pancreas	0.666667	0.857143	0.545455
TM4SF1 AND KCND2	Pancreas	0.666667	0.857143	0.545455
NOX4 AND KCNU1	Pancreas	0.6	0.473684	0.818182
NOX4 AND OTOF	Pancreas	0.6	0.473684	0.818182
GP2 AND CD93	Pancreas	0.777778	1	0.636364
TM4SF1 AND CACNG4	Pancreas	0.631579	0.75	0.545455
GP2 AND LAPTM5	Pancreas	0.705882	1	0.545455
OLR1 AND CEACAM7	Pancreas	0.705882	1	0.545455
GP2 AND PPAPDC1A	Pancreas	0.615385	0.533333	0.727273
SLC30A8 AND PTPRK	Pancreas	0.625	1	0.454545
SLC30A8 AND CD9	Pancreas	0.777778	1	0.636364
GP2 AND HLA-DRB1	Pancreas	0.625	1	0.454545
GP2 AND SLC6A6	Pancreas	0.636364	0.636364	0.636364
NOX4 AND PTGIR	Pancreas	0.736842	0.875	0.636364
GP2 AND PTGIR	Pancreas	0.666667	0.857143	0.545455
PTGIS AND TNFRSF21	Pancreas	0.666667	0.7	0.636364
NOX4 AND LAMP5	Pancreas	0.8	0.888889	0.727273
CLDN10 AND TNFSF13B	Pancreas	0.625	1	0.454545
CLDN10 AND SLAMF6	Pancreas	0.6	0.666667	0.545455
GP2 AND PROCR	Pancreas	0.625	1	0.454545
TREM2 AND LAMP5	Pancreas	0.631579	0.75	0.545455
TNFRSF12A AND VSIG1	Pancreas	0.631579	0.75	0.545455
SLC30A8 AND S100A10	Pancreas	0.777778	1	0.636364
CDH6 AND GYPC	Renal	0.857143	1	0.75
CDH6 AND NOX1	Renal	0.857143	1	0.75
CDH6 AND P2RX4	Renal	0.857143	1	0.75
CDH6 AND IL13	Renal	0.857143	1	0.75
CDH6 AND CLDN15	Renal	0.857143	1	0.75
CDH6 AND HLA-DRB1	Renal	0.857143	1	0.75
CDH6 AND ABCA12	Renal	0.857143	1	0.75
CDH6 AND C8B	Renal	0.857143	1	0.75
CDH6 AND SLC22A14	Renal	0.857143	1	0.75
JPH1 AND TGFBI	Colon	1	1	1
IFI6 AND HEPH	Colon	1	1	1
CDH17 AND SLC39A10	Colon	1	1	1
NOX1 AND AMIGO2	Colon	0.909091	0.833333	1
IFI6 AND SLC39A4	Colon	1	1	1
IFI6 AND GPR160	Colon	0.909091	0.833333	1
IFI6 AND MEP1A	Colon	1	1	1
IFI6 AND IHH	Colon	1	1	1
CDH17 AND AMIGO2	Colon	0.909091	0.833333	1
ATP10B AND TGFBI	Colon	0.909091	0.833333	1
MEP1A AND TGFBI	Colon	0.909091	0.833333	1
IFI6 AND CFTR	Colon	1	1	1
IFI6 AND SLC26A3	Colon	1	1	1
HEPH AND TGFBI	Colon	0.909091	0.833333	1
IFI6 AND ATP10B	Colon	1	1	1
IFI6 AND SLC5A1	Colon	1	1	1
IFI6 AND CEACAM1	Colon	0.909091	0.833333	1
IYD AND TGFBI	Colon	0.909091	0.833333	1
TM4SF5 AND TGFBI	Colon	0.909091	0.833333	1
NOX1 AND SLC39A10	Colon	1	1	1
IFI6 AND ATP1A1	Colon	0.888889	1	0.8
IFI6 AND SLC35G1	Colon	1	1	1
IFI6 AND IYD	Colon	0.909091	0.833333	1
SLC26A3 AND TGFBI	Colon	0.909091	0.833333	1
IFI6 AND TM4SF5	Colon	1	1	1
NOX1 AND TACSTD2	Colon	1	1	1
IFI6 AND JPH1	Colon	1	1	1
FUT3 AND SLC39A10	Colon	1	1	1
IFI6 AND XK	Colon	0.909091	0.833333	1
IFI6 AND TSPAN8	Colon	1	1	1
IFI6 AND ST14	Colon	1	1	1
IFI6 AND PTPRH	Colon	0.833333	0.714286	1

FIG. 11 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
SYT11 AND CD79A	Glioma	0.964539	0.944444	0.985507
SCN1A AND ITGAV	Glioma	0.90625	0.983051	0.84058
SYT11 AND LGR5	Glioma	0.942029	0.942029	0.942029
ITGAV AND SLC4A8	Glioma	0.882353	0.895522	0.869565
SYT11 AND VCAM1	Glioma	0.957143	0.943662	0.971014
SYT11 AND FOLR1	Glioma	0.964539	0.944444	0.985507
SYT11 AND MUC13	Glioma	0.964539	0.944444	0.985507
SYT11 AND CLDN1	Glioma	0.964539	0.944444	0.985507
SYT11 AND CD70	Glioma	0.956522	0.956522	0.956522
SYT11 AND EGFR	Glioma	0.875	0.949153	0.811594
SYT11 AND CD79B	Glioma	0.918519	0.939394	0.898551
ITGAV AND SORL1	Glioma	0.916031	0.967742	0.869565
SYT11 AND ENPP3	Glioma	0.926471	0.940299	0.913043
GPR158 AND SLC39A6	Glioma	0.852941	0.865672	0.84058
GPR19 AND EDNRB	Glioma	0.878788	0.920635	0.84058
SYT11 AND CSPG4	Glioma	0.850394	0.931034	0.782609
LYPD1 AND ITGAV	Glioma	0.904762	1	0.826087
NCAM1 AND LAPTM5	Glioma	0.873016	0.964912	0.797101
SCN2A AND ITGAV	Glioma	0.883721	0.95	0.826087
ITGAV AND CNIH2	Glioma	0.880597	0.907692	0.855072
MLC1 AND ITGAV	Glioma	0.83871	0.945455	0.753623
GPR19 AND SLC39A6	Glioma	0.836066	0.962264	0.73913
FXYD6 AND ITGAV	Glioma	0.840336	1	0.724638
SLC22A16 AND CD37	AML	0.924949	0.942149	0.908367
SLC22A16 AND SLC7A5	AML	0.901734	0.873134	0.932271
SLC22A16 AND P2RX5	AML	0.874751	0.873016	0.876494
SLC22A16 AND ENG	AML	0.859504	0.892704	0.828685
SLC22A16 AND IL11RA	AML	0.824268	0.867841	0.784861
CD37 AND SLC43A1	AML	0.772182	0.96988	0.641434
TTYH3 AND CD276	Liposarcoma	0.638889	0.638889	0.638889
TTYH3 AND GPNMB	Liposarcoma	0.637681	0.666667	0.611111
TTYH3 AND PTK7	Liposarcoma	0.608696	0.636364	0.583333
ROR1 AND ESAM	Liposarcoma	0.6	0.75	0.5
ROR1 AND PLVAP	Liposarcoma	0.6	0.75	0.5
EPHB2 AND GYPC	Liposarcoma	0.746988	0.659574	0.861111
EPHB2 AND CD93	Liposarcoma	0.697674	0.6	0.833333
EMP3 AND FOLH1	Liposarcoma	0.722222	0.722222	0.722222
STAB1 AND GPNMB	Liposarcoma	0.705882	0.75	0.666667
EMP3 AND EPHB2	Liposarcoma	0.72	0.692308	0.75
CD163 AND EPHB2	Liposarcoma	0.694444	0.694444	0.694444
ATP8B2 AND EPHB2	Liposarcoma	0.676923	0.758621	0.611111
EPHB2 AND PHLDB2	Liposarcoma	0.641975	0.577778	0.722222
CD163 AND CD276	Liposarcoma	0.717949	0.666667	0.777778
STAB1 AND TPBG	Liposarcoma	0.603175	0.703704	0.527778
SLC6A7 AND EPHB2	Liposarcoma	0.675325	0.634146	0.722222
EPHB2 AND TGFBI	Liposarcoma	0.657143	0.676471	0.638889
SLC34A2 AND SLC50A1	Lung Adenocarcinoma	0.612613	0.829268	0.485714
EPCAM AND CD93	Lung Adenocarcinoma	0.601504	0.634921	0.571429
SLC34A2 AND S1PR1	Lung Adenocarcinoma	0.614458	0.53125	0.728571
EPCAM AND CD163	Lung Adenocarcinoma	0.608	0.690909	0.542857
SLC34A2 AND ESAM	Lung Adenocarcinoma	0.608696	0.538462	0.7
CD163 AND CLDN7	Lung Adenocarcinoma	0.62069	0.782609	0.514286
SLC34A2 AND DYSF	Lung Adenocarcinoma	0.604938	0.532609	0.7
KCNJ10 AND GPNMB	B-Cell Diffuse	0.739726	0.75	0.72973
CD79B AND KCNJ10	B-Cell Diffuse	0.727273	0.827586	0.648649
CD180 AND KCNJ10	B-Cell Diffuse	0.722222	0.742857	0.702703
CD79B AND SLC22A8	B-Cell Diffuse	0.716418	0.8	0.648649
CD79B AND CDH11	B-Cell Diffuse	0.707692	0.821429	0.621622
CD79B AND GJD2	B-Cell Diffuse	0.707692	0.821429	0.621622
KCNJ10 AND CD52	B-Cell Diffuse	0.702703	0.702703	0.702703
CD79B AND SLC22A14	B-Cell Diffuse	0.695652	0.75	0.648649
KCNJ10 AND CD37	B-Cell Diffuse	0.684932	0.694444	0.675676
CD79A AND KCNJ10	B-Cell Diffuse	0.685714	0.727273	0.648649
CD79B AND FAT1	B-Cell Diffuse	0.676471	0.741935	0.621622
CD72 AND KCNJ10	B-Cell Diffuse	0.676471	0.741935	0.621622
CD79B AND CSPG5	B-Cell Diffuse	0.666667	0.71875	0.621622
KCNJ10 AND CD38	B-Cell Diffuse	0.666667	0.657895	0.675676
CD79A AND CDH11	B-Cell Diffuse	0.657534	0.666667	0.648649
CD79B AND AOC3	B-Cell Diffuse	0.6875	0.814815	0.594595
FCRL5 AND CSPG5	B-Cell Diffuse	0.686567	0.766667	0.621622
KCNJ10 AND TNFRSF17	B-Cell Diffuse	0.658228	0.619048	0.702703
CD72 AND CDH11	B-Cell Diffuse	0.637681	0.6875	0.594595
CD180 AND CDH11	B-Cell Diffuse	0.675	0.627907	0.72973
FCRL5 AND SLC22A8	B-Cell Diffuse	0.688525	0.875	0.567568
CD72 AND AOC3	B-Cell Diffuse	0.626866	0.7	0.567568
CD79B AND PLVAP	B-Cell Diffuse	0.666667	0.807692	0.567568
CD79B AND PROCR	B-Cell Diffuse	0.622951	0.791667	0.513514

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
CDH17 AND GJA4	Colon	0.888889	1	0.8
CDH17 AND PODXL	Colon	0.888889	1	0.8
CDH17 AND TMED1	Colon	0.888889	1	0.8
CDH17 AND IL1RAP	Colon	0.888889	1	0.8
CDH17 AND LDLRAD3	Colon	0.888889	1	0.8
IFI6 AND DUOX2	Colon	0.909091	0.833333	1
CDH17 AND CDH3	Colon	1	1	1
IFI6 AND TNFRSF11A	Colon	1	1	1
IFI6 AND GPR35	Colon	1	1	1
IFI6 AND SI	Colon	0.888889	1	0.8
CXCL16 AND TM4SF5	Esophagus	0.782609	1	0.642857
CXCL16 AND SYT13	Esophagus	0.727273	1	0.571429
CXCL16 AND B3GNT3	Esophagus	0.727273	1	0.571429
CXCL16 AND PTPRH	Esophagus	0.6	0.5625	0.642857
PTPRZ1 AND SPAM1	Glioblastoma	0.903226	1	0.823529
GPM6B AND F2R	Glioblastoma	0.866667	1	0.764706
PTPRZ1 AND ADAM29	Glioblastoma	0.903226	1	0.823529
PTPRZ1 AND STAB1	Glioblastoma	0.903226	1	0.823529
PTPRZ1 AND F2R	Glioblastoma	0.866667	1	0.764706
NLGN4X AND LAPTM5	Glioblastoma	0.903226	1	0.823529
PTPRZ1 AND ANTXR2	Glioblastoma	0.903226	1	0.823529
PTPRZ1 AND LAPTM5	Glioblastoma	0.9375	1	0.882353
NRCAM AND TGFBI	Glioblastoma	0.83871	0.928571	0.764706
PTPRZ1 AND CXCL16	Glioblastoma	0.866667	1	0.764706
GPM6A AND F2R	Glioblastoma	0.827586	1	0.705882
GPM6A AND EMP3	Glioblastoma	0.827586	1	0.705882
NRCAM AND F2R	Glioblastoma	0.827586	1	0.705882
PTPRZ1 AND BEST3	Glioblastoma	0.827586	1	0.705882
NRCAM AND TNFRSF12A	Glioblastoma	0.827586	1	0.705882
PTPRZ1 AND TNFRSF12A	Glioblastoma	0.827586	1	0.705882
PTPRZ1 AND PIRT	Glioblastoma	0.866667	1	0.764706
PTPRZ1 AND IL17RA	Glioblastoma	0.827586	1	0.705882
MLC1 AND OR51B6	Glioblastoma	0.909091	0.9375	0.882353
PTPRZ1 AND LYPD1	Glioblastoma	0.875	0.933333	0.823529
NRCAM AND LAPTM5	Glioblastoma	0.9375	1	0.882353
AQP4 AND SPAM1	Glioblastoma	0.827586	1	0.705882
NLGN1 AND LAPTM5	Glioblastoma	0.827586	1	0.705882
MLC1 AND FAM57A	Glioblastoma	0.827586	1	0.705882
MLC1 AND BMPR1A	Glioblastoma	0.8125	0.866667	0.764706
PTPRZ1 AND TGFBI	Glioblastoma	0.827586	1	0.705882
NRCAM AND HLA-DRB1	Glioblastoma	0.833333	0.789474	0.882353
SYT11 AND ADAM12	Glioblastoma	0.823529	0.823529	0.823529
MLC1 AND SLC2A10	Glioblastoma	0.866667	1	0.764706
NRCAM AND CDH11	Glioblastoma	0.903226	1	0.823529
PTPRZ1 AND HLA-DRB1	Glioblastoma	0.903226	1	0.823529
PTPRZ1 AND CD93	Glioblastoma	0.827586	1	0.705882
NRCAM AND CXCL16	Glioblastoma	0.903226	1	0.823529
GPM6A AND CD93	Glioblastoma	0.8	0.923077	0.705882
GPM6A AND ANTXR2	Glioblastoma	0.8	0.923077	0.705882
SYT11 AND PCDH12	Glioblastoma	0.8	0.923077	0.705882
AQP4 AND ADAM29	Glioblastoma	0.8	0.923077	0.705882
MLC1 AND TNFRSF19	Glioblastoma	0.8	0.923077	0.705882
GPR158 AND ABCA1	Glioblastoma	0.8	0.777778	0.823529
PTPRZ1 AND LRIG3	Glioblastoma	0.827586	1	0.705882
NRCAM AND CD93	Glioblastoma	0.827586	1	0.705882
NRCAM AND BTN3A3	Glioblastoma	0.833333	0.789474	0.882353
PTPRZ1 AND NKAIN4	Glioblastoma	0.8	0.923077	0.705882
NRCAM AND VAMP8	Glioblastoma	0.882353	0.882353	0.882353
PTPRZ1 AND SLC5A10	Glioblastoma	0.866667	1	0.764706
GPM6B AND F2R	Glioma	0.992701	1	0.985507
GPM6A AND F2R	Glioma	0.992701	1	0.985507
PTPRZ1 AND F2R	Glioma	0.985294	1	0.971014
SYT11 AND HLA-B	Glioma	0.956522	0.956522	0.956522
SYT11 AND SLC40A1	Glioma	0.956522	0.956522	0.956522
NRCAM AND F2R	Glioma	0.938462	1	0.884058
SYT11 AND TSPAN6	Glioma	0.970149	1	0.942029
SYT11 AND ABCA1	Glioma	0.970588	0.985075	0.956522
SYT11 AND PLGRKT	Glioma	0.970149	1	0.942029
SYT11 AND JAM2	Glioma	0.985294	1	0.971014
PTPRZ1 AND LAPTM5	Glioma	0.985507	0.985507	0.985507
SYT11 AND JTB	Glioma	0.962406	1	0.927536
SYT11 AND ATRAID	Glioma	0.985294	1	0.971014
SYT11 AND HLA-C	Glioma	0.957143	0.943662	0.971014
SYT11 AND HLA-G	Glioma	0.964539	0.944444	0.985507
PTPRZ1 AND SYT11	Glioma	0.985294	1	0.971014
GPM6A AND BTN3A3	Glioma	0.970149	1	0.942029
PTPRZ1 AND BTN3A3	Glioma	0.962406	1	0.927536

FIG. 11 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
KCNJ10 AND SLAMF7	B-Cell Diffuse	0.622222	0.528302	0.756757
CD79B AND HTR1E	B-Cell Diffuse	0.666667	0.807692	0.567568
FCRL5 AND SLC22A14	B-Cell Diffuse	0.625	0.740741	0.540541
CD72 AND FAT1	B-Cell Diffuse	0.619718	0.647059	0.594595
P2RX5 AND CDH11	B-Cell Diffuse	0.637681	0.6875	0.594595
P2RX5 AND KCNJ10	B-Cell Diffuse	0.666667	0.71875	0.621622
FCRL5 AND KCNJ10	B-Cell Diffuse	0.709677	0.88	0.594595
FCRL5 AND SLCO1B3	B-Cell Diffuse	0.619048	0.553191	0.702703
FCRL5 AND GJD2	B-Cell Diffuse	0.62069	0.857143	0.486486
FCRL5 AND GLRA2	B-Cell Diffuse	0.644068	0.863636	0.513514
P2RX5 AND SLC22A2	Mantle-Cell Lymphoma	0.897436	0.875	0.921053
P2RX5 AND HCN2	Mantle-Cell Lymphoma	0.911392	0.878049	0.947368
P2RX5 AND ADCY10	Mantle-Cell Lymphoma	0.888889	0.837209	0.947368
P2RX5 AND PVRL1	Mantle-Cell Lymphoma	0.891892	0.916667	0.868421
P2RX5 AND NTSR2	Mantle-Cell Lymphoma	0.878049	0.818182	0.947368
P2RX5 AND GHSR	Mantle-Cell Lymphoma	0.906667	0.918919	0.894737
P2RX5 AND SLC30A8	Mantle-Cell Lymphoma	0.911392	0.878049	0.947368
P2RX5 AND PCDHGC4	Mantle-Cell Lymphoma	0.861111	0.911765	0.815789
P2RX5 AND NKAIN4	Mantle-Cell Lymphoma	0.88	0.891892	0.868421
P2RX5 AND CDH22	Mantle-Cell Lymphoma	0.9	0.857143	0.947368
P2RX5 AND GRIN2B	Mantle-Cell Lymphoma	0.853659	0.795455	0.921053
P2RX5 AND MC2R	Mantle-Cell Lymphoma	0.86747	0.8	0.947368
P2RX5 AND CACNG1	Mantle-Cell Lymphoma	0.875	0.833333	0.921053
P2RX5 AND ADAM7	Mantle-Cell Lymphoma	0.85	0.809524	0.894737
CD79B AND HTR5A	Mantle-Cell Lymphoma	0.848485	1	0.736842
CD79B AND HCN2	Mantle-Cell Lymphoma	0.848485	1	0.736842
P2RX5 AND GPR6	Mantle-Cell Lymphoma	0.876712	0.914286	0.842105
P2RX5 AND CYP4A11	Mantle-Cell Lymphoma	0.842105	0.842105	0.842105
P2RX5 AND GJD2	Mantle-Cell Lymphoma	0.933333	0.945946	0.921053
P2RX5 AND HTR5A	Mantle-Cell Lymphoma	0.897436	0.875	0.921053
P2RX5 AND DISP2	Mantle-Cell Lymphoma	0.86747	0.8	0.947368
CD79B AND ADCY10	Mantle-Cell Lymphoma	0.835821	0.965517	0.736842
CD79B AND SLC30A8	Mantle-Cell Lymphoma	0.835821	0.965517	0.736842
CD79B AND AMHR2	Mantle-Cell Lymphoma	0.835821	0.965517	0.736842
CD79B AND EPHA8	Mantle-Cell Lymphoma	0.835821	0.965517	0.736842
P2RX5 AND HTR1D	Mantle-Cell Lymphoma	0.835443	0.804878	0.868421
MS4A1 AND NTSR2	Mantle-Cell Lymphoma	0.835165	0.716981	1
MS4A1 AND SLC22A9	Mantle-Cell Lymphoma	0.835165	0.716981	1
P2RX5 AND AMHR2	Mantle-Cell Lymphoma	0.9	0.857143	0.947368
P2RX5 AND PIRT	Mantle-Cell Lymphoma	0.833333	0.882353	0.789474
P2RX5 AND EPHA8	Mantle-Cell Lymphoma	0.923077	0.9	0.947368
P2RX5 AND SLC22A14	Mantle-Cell Lymphoma	0.911392	0.878049	0.947368
CD79B AND GJD2	Mantle-Cell Lymphoma	0.830769	1	0.710526
CD79B AND GHSR	Mantle-Cell Lymphoma	0.830769	1	0.710526
CD79B AND FSHR	Mantle-Cell Lymphoma	0.830769	1	0.710526
P2RX5 AND CNTNAP4	Mantle-Cell Lymphoma	0.829268	0.772727	0.894737
P2RX5 AND ROS1	Mantle-Cell Lymphoma	0.828571	0.90625	0.763158
P2RX5 AND OR2L2	Mantle-Cell Lymphoma	0.853659	0.795455	0.921053
P2RX5 AND GPR12	Mantle-Cell Lymphoma	0.876712	0.914286	0.842105
P2RX5 AND JPH3	Mantle-Cell Lymphoma	0.827586	0.734694	0.947368
MS4A1 AND HTR5A	Mantle-Cell Lymphoma	0.826087	0.703704	1
MS4A1 AND ADCY10	Mantle-Cell Lymphoma	0.826087	0.703704	1
P2RX5 AND GABRB2	Mantle-Cell Lymphoma	0.825	0.785714	0.868421
P2RX5 AND SCN1A	Mantle-Cell Lymphoma	0.825	0.785714	0.868421
P2RX5 AND SLC2A1	Mantle-Cell Lymphoma	0.846154	0.825	0.868421
P2RX5 AND OR2C3	Mantle-Cell Lymphoma	0.829268	0.772727	0.894737
CD79B AND OR10J1	Mantle-Cell Lymphoma	0.823529	0.933333	0.736842
CD79B AND MC2R	Mantle-Cell Lymphoma	0.823529	0.933333	0.736842
CD79B AND CLDN20	Mantle-Cell Lymphoma	0.823529	0.933333	0.736842
CD79B AND SLC22A14	Mantle-Cell Lymphoma	0.823529	0.933333	0.736842
P2RX5 AND LCT	Mantle-Cell Lymphoma	0.823529	0.744681	0.921053
P2RX5 AND CHRNB2	Mantle-Cell Lymphoma	0.878049	0.818182	0.947368
P2RX5 AND GPRC6A	Mantle-Cell Lymphoma	0.820513	0.8	0.842105
P2RX5 AND KCNA7	Mantle-Cell Lymphoma	0.857143	0.846154	0.868421
P2RX5 AND GRM5	Mantle-Cell Lymphoma	0.842105	0.842105	0.842105
P2RX5 AND OR10J1	Mantle-Cell Lymphoma	0.819277	0.755556	0.894737
P2RX5 AND TRPM3	Mantle-Cell Lymphoma	0.819277	0.755556	0.894737
P2RX5 AND CEACAM7	Mantle-Cell Lymphoma	0.819277	0.755556	0.894737
P2RX5 AND SLC1A6	Mantle-Cell Lymphoma	0.819277	0.755556	0.894737
P2RX5 AND LRFN2	Mantle-Cell Lymphoma	0.84507	0.909091	0.789474
P2RX5 AND SLC22A9	Mantle-Cell Lymphoma	0.878049	0.818182	0.947368
CD79B AND NKAIN4	Mantle-Cell Lymphoma	0.818182	0.964286	0.710526
CD79B AND CDH22	Mantle-Cell Lymphoma	0.818182	0.964286	0.710526
CD79B AND SLC1A6	Mantle-Cell Lymphoma	0.818182	0.964286	0.710526
P2RX5 AND TMEM235	Mantle-Cell Lymphoma	0.818182	0.72	0.947368
MS4A1 AND CLDN20	Mantle-Cell Lymphoma	0.817204	0.690909	1
P2RX5 AND FSHR	Mantle-Cell Lymphoma	0.906667	0.918919	0.894737

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
DDR2 AND ATP8B2	Leiomyosarcoma	0.761905	1	0.615385
FLT3 AND EMP3	AML	0.944559	0.974576	0.916335
SLC22A16 AND HCST	AML	0.939759	0.947368	0.932271
SLC22A16 AND P2RX1	AML	0.937374	0.95082	0.924303
SLC22A16 AND P2RY8	AML	0.928571	0.924901	0.932271
CNTNAP1 AND CD163	Liposarcoma	0.8	0.823529	0.777778
ADAM12 AND KCNU1	Liposarcoma	0.740741	0.666667	0.833333
ADAM12 AND EMP3	Liposarcoma	0.727273	0.682927	0.777778
ADAM12 AND SLC6A7	Liposarcoma	0.725	0.659091	0.805556
ADAM12 AND MDGA2	Liposarcoma	0.75	0.681818	0.833333
ADAM12 AND OR2C3	Liposarcoma	0.746988	0.659574	0.861111
ADAM12 AND OR52D1	Liposarcoma	0.72093	0.62	0.861111
ADAM12 AND CSMD2	Liposarcoma	0.746988	0.659574	0.861111
ADAM12 AND SV2C	Liposarcoma	0.738095	0.645833	0.861111
ADAM12 AND HLA-DRB1	Liposarcoma	0.707317	0.630435	0.805556
ADAM12 AND DRD1	Liposarcoma	0.765432	0.688889	0.861111
ADAM12 AND LCT	Liposarcoma	0.729412	0.632653	0.861111
ADAM12 AND TRHR	Liposarcoma	0.738095	0.645833	0.861111
ADAM12 AND KCNV2	Liposarcoma	0.756098	0.673913	0.861111
ADAM12 AND ADAM7	Liposarcoma	0.759494	0.697674	0.833333
ADAM12 AND GABRA3	Liposarcoma	0.746988	0.659574	0.861111
ADAM12 AND OR8B2	Liposarcoma	0.705882	0.612245	0.833333
ADAM12 AND OR3A1	Liposarcoma	0.714286	0.625	0.833333
ADAM12 AND GRIN2B	Liposarcoma	0.698795	0.617021	0.805556
ADAM12 AND GPR85	Liposarcoma	0.696629	0.584906	0.861111
ADAM12 AND CACNG4	Liposarcoma	0.696629	0.584906	0.861111
ADAM12 AND NMBR	Liposarcoma	0.734177	0.674419	0.805556
ADAM12 AND TRPM8	Liposarcoma	0.738095	0.645833	0.861111
ADAM12 AND PCDHA9	Liposarcoma	0.738095	0.645833	0.861111
ADAM12 AND GLRA3	Liposarcoma	0.717949	0.666667	0.777778
ADAM12 AND GRM4	Liposarcoma	0.691358	0.622222	0.777778
ADAM12 AND GYPC	Liposarcoma	0.688889	0.574074	0.861111
ADAM12 AND LRRTM4	Liposarcoma	0.684211	0.65	0.722222
ADAM12 AND FCAMR	Liposarcoma	0.756098	0.673913	0.861111
ADAM12 AND P2RY4	Liposarcoma	0.681818	0.576923	0.833333
ADAM12 AND MLANA	Liposarcoma	0.722892	0.638298	0.833333
ADAM12 AND SLCO6A1	Liposarcoma	0.729412	0.632653	0.861111
ADAM12 AND ACSL6	Liposarcoma	0.674699	0.595745	0.777778
ADAM12 AND ATP8B2	Liposarcoma	0.722222	0.722222	0.722222
ADAM12 AND GPR83	Liposarcoma	0.740741	0.666667	0.833333
TTYH3 AND LRP1	Liposarcoma	0.686567	0.741935	0.638889
GJC1 AND LAPTM5	Liposarcoma	0.666667	0.6	0.75
ADAM12 AND GABRG2	Liposarcoma	0.659341	0.545455	0.833333
ADAM12 AND BEST3	Liposarcoma	0.753247	0.707317	0.805556
ADAM12 AND ADAM21	Liposarcoma	0.658537	0.586957	0.75
ADAM12 AND CD74	Liposarcoma	0.658537	0.586957	0.75
TTYH3 AND PKD2	Liposarcoma	0.657143	0.676471	0.638889
CNTNAP1 AND LAPTM5	Liposarcoma	0.777778	0.777778	0.777778
TTYH3 AND COLEC12	Liposarcoma	0.647887	0.657143	0.638889
TTYH3 AND PDGFRB	Liposarcoma	0.647059	0.6875	0.611111
ADAM12 AND CATSPERD	Liposarcoma	0.64	0.615385	0.666667
TTYH3 AND DCHS1	Liposarcoma	0.638889	0.638889	0.638889
TTYH3 AND FAT4	Liposarcoma	0.637681	0.666667	0.611111
CD163 AND PCDH7	Liposarcoma	0.658228	0.604651	0.722222
ADAM12 AND LAPTM5	Liposarcoma	0.634146	0.565217	0.722222
EMP3 AND SLC2A10	Liposarcoma	0.75	0.75	0.75
GPR34 AND SLC6A7	Liposarcoma	0.630137	0.621622	0.638889
TTYH3 AND PCDH12	Liposarcoma	0.630137	0.621622	0.638889
TTYH3 AND MMP14	Liposarcoma	0.647887	0.657143	0.638889
ADAM12 AND MUC12	Liposarcoma	0.717949	0.666667	0.777778
TTYH3 AND SGCD	Liposarcoma	0.625	0.714286	0.555556
ADAM12 AND GPR22	Liposarcoma	0.623656	0.508772	0.805556
ATP8B2 AND SLC2A10	Liposarcoma	0.717949	0.666667	0.777778
TTYH3 AND PLXDC1	Liposarcoma	0.622951	0.76	0.527778
TTYH3 AND EMP1	Liposarcoma	0.621622	0.605263	0.638889
TTYH3 AND CNTNAP1	Liposarcoma	0.621622	0.605263	0.638889
TTYH3 AND VASN	Liposarcoma	0.621622	0.605263	0.638889
ADAM12 AND SLC17A1	Liposarcoma	0.621622	0.605263	0.638889
TTYH3 AND PLXNA1	Liposarcoma	0.621622	0.605263	0.638889
TTYH3 AND SLC2A10	Liposarcoma	0.619718	0.628571	0.611111
OLR1 AND SLC50A1	Lung Adenocarcinoma	0.655172	0.826087	0.542857
OSMR AND SLC50A1	Lung Adenocarcinoma	0.625	0.833333	0.5
BST2 AND TLCD1	Lung Adenocarcinoma	0.619469	0.813953	0.5
PON2 AND CD44	Lung Adenocarcinoma	0.62069	0.782609	0.514286
PON2 AND IFNAR2	Lung Adenocarcinoma	0.641509	0.944444	0.485714
SLC50A1 AND FZD6	Lung Adenocarcinoma	0.62069	0.782609	0.514286
KCNJ10 AND CD53	B-Cell Diffuse	0.759494	0.714286	0.810811

FIG. 11 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
CD79B AND GPR6	Mantle-Cell Lymphoma	0.8125	1	0.684211
CD79B AND ADAM7	Mantle-Cell Lymphoma	0.8125	1	0.684211
CD79B AND LRFN2	Mantle-Cell Lymphoma	0.8125	1	0.684211
CD79B AND SLCO1A2	Mantle-Cell Lymphoma	0.8125	1	0.684211
CD79B AND GPR12	Mantle-Cell Lymphoma	0.8125	1	0.684211
CD79B AND CLDN15	Mantle-Cell Lymphoma	0.811594	0.903226	0.736842
CD79B AND GPR50	Mantle-Cell Lymphoma	0.811594	0.903226	0.736842
CD79B AND HRH3	Mantle-Cell Lymphoma	0.811594	0.903226	0.736842
EDNRB AND GPR19	Melanoma	0.9	1	0.818182
EDNRB AND LRIG3	Melanoma	0.9	1	0.818182
EDNRB AND ZACN	Melanoma	0.9	1	0.818182
EDNRB AND FAT1	Melanoma	0.9	1	0.818182
EDNRB AND EFNB2	Melanoma	0.9	1	0.818182
EDNRB AND GHR	Melanoma	0.9	1	0.818182
EDNRB AND SLC22A18	Melanoma	0.9	1	0.818182
NKAIN1 AND CBX3	Neuroblastoma	0.866242	0.931507	0.809524
NKAIN1 AND ALK	Neuroblastoma	0.867925	0.92	0.821429
GPR19 AND PTK7	Neuroblastoma	0.854054	0.782178	0.940476
GPR19 AND GPNMB	Neuroblastoma	0.842105	0.827586	0.857143
ALK AND CD163	Neuroblastoma	0.840764	0.90411	0.785714
KCNQ2 AND CBX3	Neuroblastoma	0.839506	0.871795	0.809524
CACNA1B AND CD276	Neuroblastoma	0.836957	0.77	0.916667
SYT11 AND ALK	Neuroblastoma	0.840764	0.90411	0.785714
SLC1A2 AND EGFR	Oligodendroglioma	0.933333	0.933333	0.933333
CSPG5 AND EGFR	Oligodendroglioma	0.888889	1	0.8
OMG AND EGFR	Oligodendroglioma	0.857143	0.923077	0.8
KCNJ10 AND EGFR	Oligodendroglioma	0.933333	0.933333	0.933333
MLC1 AND EGFR	Oligodendroglioma	0.896552	0.928571	0.866667
DTNA AND EGFR	Oligodendroglioma	0.888889	1	0.8
ASTN1 AND EGFR	Oligodendroglioma	0.888889	1	0.8
CDH10 AND EGFR	Oligodendroglioma	0.857143	0.923077	0.8
GPR19 AND EGFR	Oligodendroglioma	0.903226	0.875	0.933333
UNC5A AND EGFR	Oligodendroglioma	0.846154	1	0.733333
GPR158 AND EGFR	Oligodendroglioma	0.8	0.8	0.8
OPCML AND EGFR	Oligodendroglioma	0.8	0.8	0.8
SEZ6 AND EGFR	Oligodendroglioma	0.928571	1	0.866667
FXYD6 AND EGFR	Oligodendroglioma	0.928571	1	0.866667
CSPG5 AND EDNRB	Oligodendroglioma	0.8	0.8	0.8
ASTN1 AND CSPG4	Oligodendroglioma	0.8	0.7	0.933333
BEST3 AND BMPR1B	Oligodendroglioma	0.866667	0.866667	0.866667
CSPG5 AND MUC13	Oligodendroglioma	0.888889	1	0.8
BEST3 AND SLC7A5	Oligodendroglioma	0.777778	0.666667	0.933333
MLC1 AND CD38	Oligodendroglioma	0.774194	0.75	0.8
KCNC1 AND EGFR	Oligodendroglioma	0.774194	0.75	0.8
CSPG5 AND FOLR1	Oligodendroglioma	0.774194	0.75	0.8
MLC1 AND MUC1	Oligodendroglioma	0.785714	0.846154	0.733333
CSPG5 AND CLDN18	Oligodendroglioma	0.827586	0.857143	0.8
CSPG5 AND VTCN1	Oligodendroglioma	0.846154	1	0.733333
LRRTM4 AND EGFR	Oligodendroglioma	0.764706	0.684211	0.866667
GABBR2 AND EGFR	Oligodendroglioma	0.764706	0.684211	0.866667
CSPG5 AND MUC1	Oligodendroglioma	0.764706	0.684211	0.866667
DTNA AND MUC13	Oligodendroglioma	0.764706	0.684211	0.866667
BEST3 AND SLC39A6	Oligodendroglioma	0.823529	0.736842	0.933333
MLC1 AND VTCN1	Oligodendroglioma	0.8	1	0.666667
SLC1A2 AND VTCN1	Oligodendroglioma	0.814815	0.916667	0.733333
SLC1A2 AND PTK7	Oligodendroglioma	0.8	1	0.666667
CSPG5 AND GUCY2C	Oligodendroglioma	0.758621	0.785714	0.733333
DTNA AND MUC1	Oligodendroglioma	0.759231	0.909091	0.666667
KCNJ10 AND CD276	Oligodendroglioma	0.8125	0.764706	0.866667
SLCO1C1 AND EGFR	Oligodendroglioma	0.827586	0.857143	0.8
CSPG5 AND CA9	Oligodendroglioma	0.756757	0.636364	0.933333
MUC16 AND SYT11	Ovarian	0.736842	1	0.583333
EPHA10 AND VTCN1	Ovarian	0.666667	1	0.5
MUC16 AND CSPG5	Ovarian	0.666667	1	0.5
MUC16 AND PPAPDC1A	Ovarian	0.666667	0.777778	0.583333
MUC16 AND LAMP5	Ovarian	0.631579	0.857143	0.5
MUC16 AND GPRC6A	Ovarian	0.631579	0.857143	0.5
MUC16 AND KCNU1	Ovarian	0.608696	0.636364	0.583333
TRPM8 AND EPCAM	Prostate	0.857143	1	0.75
STEAP2 AND SLC43A1	Prostate	1	1	1

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
PTPRCAP AND KCNJ10	B-Cell Diffuse	0.722222	0.742857	0.702703
CD48 AND KCNJ10	B-Cell Diffuse	0.724638	0.78125	0.675676
IL2RB AND FAT1	Anaplastic Lymphoma	0.64	0.8	0.533333
CXCL9 AND GPR19	T-Cell, Peripheral	0.75	0.75	0.75
CXCL9 AND KCNJ10	T-Cell, Peripheral	0.769231	0.833333	0.714286
CXCL9 AND SEMA4D	T-Cell, Peripheral	0.784314	0.869565	0.714286
CXCL9 AND TTYH2	T-Cell, Peripheral	0.714286	0.714286	0.714286
CXCL9 AND SLC38A1	T-Cell, Peripheral	0.652174	0.833333	0.535714
BCAP31 AND TTYH2	Melanoma	0.952381	1	0.909091
GPR137B AND KCNN2	Melanoma	0.9	1	0.818182
PLP2 AND SLC6A15	Melanoma	0.9	1	0.818182
BCAP31 AND EMP3	Melanoma	0.952381	1	0.909091
TNFSF9 AND GPR137B	Melanoma	0.952381	1	0.909091
C11orf24 AND GPR137B	Melanoma	0.9	1	0.818182
NKAIN4 AND PTPRZ1	Oligodendroglioma	0.888889	1	0.8
KCNK15 AND CSPG5	Ovarian	0.8	1	0.666667
KCNK15 AND EPHA10	Ovarian	0.785714	0.6875	0.916667
KCNK15 AND SLC2A1	Ovarian	0.758621	0.647059	0.916667
KCNK15 AND NMUR2	Ovarian	0.75	0.75	0.75
ADAM12 AND EMP3	Sarcoma	1	1	1
CD99 AND PCDHA9	Sarcoma	0.740741	1	0.588235
ADAM12 AND SLC6A7	Sarcoma	0.8125	0.866667	0.764706
ADAM12 AND KCNU1	Sarcoma	0.75	0.652174	0.882353
CDH17 AND FXYD5	Stomach	0.911111	0.87234	0.953488
CDH17 AND CDH3	Stomach	0.8	0.765957	0.837209
CDH17 AND TACSTD2	Stomach	0.911392	1	0.837209
CDH17 AND EDNRA	Stomach	0.876404	0.847826	0.906977
CDH17 AND PDPN	Stomach	0.83871	0.78	0.906977
CDH17 AND MCAM	Stomach	0.727273	0.597015	0.930233
CDH17 AND CDH5	Stomach	0.829787	0.764706	0.906977
CDH17 AND IFITM2	Stomach	0.824742	0.740741	0.930233
VSIG1 AND TNFRSF12A	Stomach	0.714286	0.925926	0.581395
VSIG1 AND VANGL1	Stomach	0.714286	0.925926	0.581395
CDH17 AND APOLD1	Stomach	0.780952	0.66129	0.953488
VSIG1 AND FAT1	Stomach	0.705882	0.96	0.55814
VSIG1 AND PROCR	Stomach	0.704225	0.892857	0.581395
CDH17 AND OSMR	Stomach	0.866667	0.829787	0.906977
CDH17 AND C5AR1	Stomach	0.822222	0.787234	0.860465
VSIG1 AND CDH11	Stomach	0.695652	0.923077	0.55814
CDH17 AND PDGFRB	Stomach	0.853933	0.826087	0.883721
CDH17 AND LDLRAD3	Stomach	0.690265	0.557143	0.906977
VSIG1 AND EFNB2	Stomach	0.684932	0.833333	0.581395
CDH17 AND ANO1	Stomach	0.894118	0.904762	0.883721
CDH17 AND GJA4	Stomach	0.716981	0.603175	0.883721
CDH17 AND CALCRL	Stomach	0.823529	0.833333	0.813953
CDH17 AND PODXL	Stomach	0.847059	0.857143	0.837209
CDH17 AND SLC2A3	Stomach	0.690909	0.567164	0.883721
VSIG1 AND EPHA2	Stomach	0.676923	1	0.511628
CDH17 AND FCGR3B	Stomach	0.771084	0.8	0.744186
CDH17 AND CD81	Stomach	0.718447	0.616667	0.860465
VSIG1 AND LRIG3	Stomach	0.675676	0.806452	0.581395
CDH17 AND CXCL9	Stomach	0.707317	0.74359	0.674419
CDH17 AND SLC41A1	Stomach	0.727273	0.597015	0.930233
TSPAN8 AND BST2	Stomach	0.724638	0.961538	0.581395
ATP8B1 AND FXYD5	Stomach	0.91358	0.973684	0.860465
VSIG1 AND F2R	Stomach	0.714286	0.925926	0.581395
VSIG1 AND CDH17	Stomach	0.735294	1	0.581395
CDH17 AND ABCC1	Stomach	0.880952	0.902439	0.860465
CDH17 AND PLXDC1	Stomach	0.813187	0.770833	0.860465
CDH17 AND SLC29A1	Stomach	0.772277	0.672414	0.906977
CDH17 AND RHBDF2	Stomach	0.752688	0.7	0.813953
VSIG1 AND TTYH3	Stomach	0.666667	0.827586	0.55814
VSIG1 AND SLC1A5	Stomach	0.675676	0.806452	0.581395
CDH17 AND DIO2	Stomach	0.66	0.578947	0.767442
CDH17 AND F2RL2	Stomach	0.9	0.972973	0.837209
MUC17 AND CDH3	Stomach	0.788732	1	0.651163
ATP8B1 AND CXCR4	Stomach	0.826667	0.96875	0.72093
VSIG1 AND CD82	Stomach	0.657895	0.757576	0.581395
VSIG1 AND BST2	Stomach	0.675676	0.806452	0.581395

FIG. 12

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
VANGL2 AND NOT-TACSTD2	Astrocytoma	0.8387	0.8298	0.8478
VANGL2 AND NOT-AQP3	Astrocytoma	0.8298	0.8125	0.8478
VANGL2 AND NOT-FXYD3	Astrocytoma	0.8298	0.8125	0.8478
VANGL2 AND NOT-PERP	Astrocytoma	0.8298	0.8125	0.8478
VANGL2 AND NOT-F11R	Astrocytoma	0.8261	0.8261	0.8261
VANGL2 AND NOT-KRT5	Astrocytoma	0.8211	0.7959	0.8478
VANGL2 AND NOT-DUOX1	Astrocytoma	0.8211	0.7959	0.8478
VANGL2 AND NOT-SMAGP	Astrocytoma	0.8125	0.78	0.8478
PTPRZ1 AND NOT-SLC2A12	Astrocytoma	0.8119	0.7455	0.8913
NRCAM AND NOT-TENM2	Astrocytoma	0.8046	0.8537	0.7609
PTPRZ1 AND NOT-MTUS1	Astrocytoma	0.8041	0.7647	0.8478
PTPRZ1 AND NOT-ABCC5	Astrocytoma	0.8039	0.7321	0.8913
PTPRZ1 AND NOT-LRP11	Astrocytoma	0.8039	0.7321	0.8913
VANGL2 AND NOT-CDH1	Astrocytoma	0.8	0.8182	0.7826
PTPRZ1 AND NOT-FZD7	Astrocytoma	0.7921	0.7273	0.8696
PTPRZ1 AND NOT-CNST	Astrocytoma	0.7921	0.7273	0.8696
PTPRZ1 AND NOT-SACM1L	Astrocytoma	0.7885	0.7069	0.8913
VANGL2 AND NOT-TMEM30B	Astrocytoma	0.7879	0.7358	0.8478
VANGL2 AND NOT-DSG2	Astrocytoma	0.7872	0.7708	0.8043
VANGL2 AND NOT-ST14	Astrocytoma	0.7865	0.814	0.7609
VANGL2 AND NOT-PLP2	Astrocytoma	0.7835	0.7451	0.8261
PTPRZ1 AND NOT-FZD10	Astrocytoma	0.781	0.6949	0.8913
PTPRZ1 AND NOT-FAM73B	Astrocytoma	0.781	0.6949	0.8913
PTPRZ1 AND NOT-PLXNA2	Astrocytoma	0.7789	0.7551	0.8043
PTPRZ1 AND NOT-ANO3	Astrocytoma	0.7789	0.7551	0.8043
ADAM12 AND NOT-GPBAR1	Breast	0.8654	0.9	0.8333
ADAM12 AND NOT-TRPC6	Breast	0.8627	0.9167	0.8148
ADAM12 AND NOT-F10	Breast	0.8462	0.88	0.8148
ADAM12 AND NOT-GLDN	Breast	0.8462	0.88	0.8148
ADAM12 AND NOT-TNFRSF10D	Breast	0.8462	0.88	0.8148
ADAM12 AND NOT-AVPR1A	Breast	0.8462	0.88	0.8148
ADAM12 AND NOT-JAM3	Breast	0.8411	0.8491	0.8333
ADAM12 AND NOT-NAALAD2	Breast	0.8333	0.8333	0.8333
ADAM12 AND NOT-SLC16A7	Breast	0.8302	0.8462	0.8148
ADAM12 AND NOT-CXCR2	Breast	0.8283	0.9111	0.7593
ADAM12 AND NOT-F3	Breast	0.8257	0.8182	0.8333
ADAM12 AND NOT-FCGR3B	Breast	0.8247	0.9302	0.7407
ADAM12 AND NOT-GPR87	Breast	0.8235	0.875	0.7778
ADAM12 AND NOT-SLC22A3	Breast	0.8224	0.8302	0.8148
ADAM12 AND NOT-KLB	Breast	0.8182	0.8036	0.8333
ADAM12 AND NOT-OLR1	Breast	0.8182	0.8036	0.8333
ADAM12 AND NOT-C14orf180	Breast	0.8182	0.8036	0.8333
ADAM12 AND NOT-PLXNA4	Breast	0.8182	0.8036	0.8333
ADAM12 AND NOT-PPAP2A	Breast	0.8182	0.8036	0.8333
ADAM12 AND NOT-SLC4A4	Breast	0.8182	0.8036	0.8333
ADAM12 AND NOT-APOB	Breast	0.8182	0.8036	0.8333
ADAM12 AND NOT-SCN3A	Breast	0.8182	0.8036	0.8333
ADAM12 AND NOT-ENPEP	Breast	0.8182	0.8036	0.8333
ADAM12 AND NOT-SGCB	Breast	0.8182	0.8036	0.8333
ADAM12 AND NOT-DPP4	Breast	0.8182	0.8036	0.8333
ADAM12 AND NOT-SCN9A	Breast	0.8182	0.8036	0.8333
ADAM12 AND NOT-ATP8B4	Breast	0.8182	0.8036	0.8333
ADAM12 AND NOT-ABCC9	Breast	0.8182	0.8036	0.8333
ADAM12 AND NOT-SIGLEC6	Breast	0.8182	0.8036	0.8333
ADAM12 AND NOT-CD300LG	Breast	0.8182	0.8036	0.8333
ADAM12 AND NOT-ERVFRD-1	Breast	0.8182	0.8036	0.8333
ADAM12 AND NOT-SLC52A1	Breast	0.8182	0.8036	0.8333
ADAM12 AND NOT-SMAGP	Breast	0.8182	0.8036	0.8333
ADAM12 AND NOT-CDH6	Breast	0.8182	0.8036	0.8333
ADAM12 AND NOT-CDH5	Breast	0.8182	0.8036	0.8333
ADAM12 AND NOT-CAV2	Breast	0.8182	0.8036	0.8333
ADAM12 AND NOT-MCAM	Breast	0.8182	0.8036	0.8333
ADAM12 AND NOT-ITM2A	Breast	0.8155	0.8571	0.7778
ADAM12 AND NOT-MME	Breast	0.8155	0.8571	0.7778
ADAM12 AND NOT-FCGR2B	Breast	0.8148	0.8148	0.8148
ADAM12 AND NOT-C15orf27	Breast	0.8148	0.8148	0.8148
ADAM12 AND NOT-PAG1	Breast	0.8148	0.8148	0.8148
ADAM12 AND NOT-ACPP	Breast	0.8148	0.8148	0.8148
ADAM12 AND NOT-KL	Breast	0.8148	0.8148	0.8148
ADAM12 AND NOT-ADCYAP1R1	Breast	0.8148	0.8148	0.8148
ADAM12 AND NOT-PVRL3	Breast	0.8148	0.8148	0.8148
ADAM12 AND NOT-SLC13A4	Breast	0.8148	0.8148	0.8148
ADAM12 AND NOT-KIRREL	Breast	0.8119	0.8723	0.7593
ADAM12 AND NOT-IL1R1	Breast	0.8119	0.8723	0.7593
ADAM12 AND NOT-PCDH18	Breast	0.8113	0.8269	0.7963
ADAM12 AND NOT-P2RY1	Breast	0.8113	0.8269	0.7963
ADAM12 AND NOT-PTH1R	Breast	0.8113	0.8269	0.7963

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
FGFR4 AND NOT-IL3RA	Liver	0.8	1	0.6667
LRP5 AND NOT-TPBG	Liver	0.8333	0.8333	0.8333
FGFR4 AND NOT-CD52	Liver	0.8	1	0.6667
FGFR4 AND NOT-IL20RA	Liver	0.8	1	0.6667
FGFR4 AND NOT-FOLR1	Liver	0.8	1	0.6667
FGFR4 AND NOT-CD33	Liver	0.8	1	0.6667
FGFR4 AND NOT-CD37	Liver	0.8	1	0.6667
SLC4A2 AND NOT-PTK7	Liver	0.8	1	0.6667
FGFR4 AND NOT-PMEL	Liver	0.8	1	0.6667
FGFR4 AND NOT-TNFRSF17	Liver	0.8	1	0.6667
FGFR4 AND NOT-MS4A1	Liver	0.8	1	0.6667
FGFR4 AND NOT-CD22	Liver	0.8	1	0.6667
FGFR4 AND NOT-ERBB2	Liver	0.8	1	0.6667
FGFR4 AND NOT-MUC16	Liver	0.8	1	0.6667
FGFR4 AND NOT-SSTR2	Liver	0.8	1	0.6667
FGFR4 AND NOT-GUCY2C	Liver	0.8	1	0.6667
FGFR4 AND NOT-EPHB2	Liver	0.8	1	0.6667
FGFR4 AND NOT-CD70	Liver	0.8	1	0.6667
FGFR4 AND NOT-CD79A	Liver	0.8	1	0.6667
FGFR4 AND NOT-FCRL5	Liver	0.8	1	0.6667
FGFR4 AND NOT-ERBB4	Liver	0.8	1	0.6667
FGFR4 AND NOT-CR2	Liver	0.8	1	0.6667
FGFR4 AND NOT-FCRL2	Liver	0.8	1	0.6667
FGFR4 AND NOT-CD79B	Liver	0.8	1	0.6667
FGFR4 AND NOT-CLDN8	Liver	0.8	1	0.6667
FGFR4 AND NOT-L1CAM	Liver	0.8	1	0.6667
FGFR4 AND NOT-PTK7	Liver	0.8	1	0.6667
FGFR4 AND NOT-PROM1	Liver	0.8	1	0.6667
FGFR4 AND NOT-MUC13	Liver	0.8	1	0.6667
FGFR4 AND NOT-SLC7A5	Liver	0.8	1	0.6667
FGFR4 AND NOT-TRPM4	Liver	0.8	1	0.6667
FGFR4 AND NOT-CD180	Liver	0.8	1	0.6667
FGFR4 AND NOT-CLDN23	Liver	0.8	1	0.6667
FGFR4 AND NOT-MUC4	Liver	0.8	1	0.6667
FGFR4 AND NOT-MST1R	Liver	0.8	1	0.6667
FGFR4 AND NOT-CD19	Liver	0.8	1	0.6667
SLC4A2 AND NOT-CLDN11	Liver	0.8	1	0.6667
SLC4A2 AND NOT-IGF1R	Liver	0.8	1	0.6667
APOB AND NOT-PROM1	Liver	0.7273	0.8	0.6667
SLC4A2 AND NOT-ITGB6	Liver	0.8	1	0.6667
ITPR2 AND NOT-CLDN8	Liver	0.8333	0.8333	0.8333
SLC4A2 AND NOT-CSPG4	Liver	0.8	1	0.6667
SLC4A2 AND NOT-PROM1	Liver	0.8	1	0.6667
FGFR4 AND NOT-TNFSF11	Liver	0.8	1	0.6667
FGFR4 AND NOT-FAP	Liver	0.8	1	0.6667
FAP AND NOT-PTPRS	Pancreas	0.8421	1	0.7273
FAP AND NOT-SDC2	Pancreas	0.7059	1	0.5455
FAP AND NOT-KCNMA1	Pancreas	0.7778	1	0.6364
FAP AND NOT-TGFBR3	Pancreas	0.7778	1	0.6364
FAP AND NOT-LIFR	Pancreas	0.7778	1	0.6364
FAP AND NOT-GJC1	Pancreas	0.8421	1	0.7273
FAP AND NOT-PCDH19	Pancreas	0.8421	1	0.7273
FAP AND NOT-CALHM2	Pancreas	0.7778	1	0.6364
FAP AND NOT-PTGER3	Pancreas	0.7778	1	0.6364
FAP AND NOT-ESYT1	Pancreas	0.7778	1	0.6364
FAP AND NOT-ITGA7	Pancreas	0.7778	1	0.6364
FAP AND NOT-SGMS1	Pancreas	0.7059	1	0.5455
FAP AND NOT-CAV1	Pancreas	0.7778	1	0.6364
FAP AND NOT-SLC47A1	Pancreas	0.8421	1	0.7273
FAP AND NOT-NRXN3	Pancreas	0.7778	1	0.6364
FAP AND NOT-BVES	Pancreas	0.7778	1	0.6364
FAP AND NOT-REEP2	Pancreas	0.7059	1	0.5455
FAP AND NOT-SLC2A12	Pancreas	0.7778	1	0.6364
FAP AND NOT-PTH1R	Pancreas	0.7778	1	0.6364
FAP AND NOT-ADCY4	Pancreas	0.8421	1	0.7273
FAP AND NOT-ECSCR	Pancreas	0.8421	1	0.7273
FAP AND NOT-RECK	Pancreas	0.7059	1	0.5455
FAP AND NOT-PRLR	Pancreas	0.8421	1	0.7273
FAP AND NOT-EMCN	Pancreas	0.8421	1	0.7273
FAP AND NOT-ADRA2C	Pancreas	0.8421	1	0.7273
FAP AND NOT-PEAR1	Pancreas	0.7778	1	0.6364
FAP AND NOT-JAM2	Pancreas	0.7778	1	0.6364
FAP AND NOT-AVPR1A	Pancreas	0.7778	1	0.6364
FAP AND NOT-ABCG2	Pancreas	0.7059	1	0.5455
FAP AND NOT-SCN4B	Pancreas	0.7059	1	0.5455
FAP AND NOT-PTGFR	Pancreas	0.8421	1	0.7273
FAP AND NOT-ST6GALNAC6	Pancreas	0.7778	1	0.6364

FIG. 12 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
ADAM12 AND NOT-TRHDE	Breast	0.8077	0.84	0.7778
ADAM12 AND NOT-GPR146	Breast	0.8073	0.8	0.8148
ADAM12 AND NOT-LYVE1	Breast	0.8073	0.8	0.8148
ADAM12 AND NOT-TUSC3	Breast	0.8073	0.8	0.8148
ADAM12 AND NOT-ADRB1	Breast	0.8073	0.8	0.8148
ADAM12 AND NOT-GJA5	Breast	0.8073	0.8	0.8148
ADAM12 AND NOT-CAV1	Breast	0.8073	0.8	0.8148
ADAM12 AND NOT-THSD7A	Breast	0.8073	0.8	0.8148
GPRC5A AND NOT-BMP2	Breast	0.8235	0.875	0.7778
ADAM12 AND NOT-NT5E	Breast	0.8037	0.8113	0.7963
ADAM12 AND NOT-ENPP2	Breast	0.8037	0.8113	0.7963
ADAM12 AND NOT-IL1RAP	Breast	0.8037	0.8113	0.7963
ADAM12 AND NOT-SLC31A2	Breast	0.8	0.8235	0.7778
ADAM12 AND NOT-BMP2	Breast	0.8	0.8696	0.7407
ADAM12 AND NOT-HTR2B	Breast	0.7963	0.7963	0.7963
ADAM12 AND NOT-C10orf54	Breast	0.7963	0.7963	0.7963
ADAM12 AND NOT-CHL1	Breast	0.7963	0.7963	0.7963
ADAM12 AND NOT-TLR4	Breast	0.7963	0.7963	0.7963
ADAM12 AND NOT-SGCG	Breast	0.7963	0.7963	0.7963
ADAM12 AND NOT-SLC7A6	Breast	0.7963	0.7963	0.7963
ADAM12 AND NOT-SLC28A2	Breast	0.7963	0.7963	0.7963
ADAM12 AND NOT-LEPR	Breast	0.7961	0.8367	0.7593
SLC27A5 AND NOT-CLEC1B	Liver	0.9091	1	0.8333
FGG AND NOT-CLEC1B	Liver	0.9091	1	0.8333
HPN AND NOT-CLEC1B	Liver	0.9091	1	0.8333
FGG AND NOT-MARCO	Liver	0.8571	0.75	1
HPN AND NOT-MARCO	Liver	0.8571	0.75	1
ABCC6 AND NOT-CLEC1B	Liver	0.8333	0.8333	0.8333
APOB AND NOT-CDHR2	Liver	0.8333	0.8333	0.8333
HPN AND NOT-KCNN2	Liver	1	1	1
ABCB4 AND NOT-CLEC1B	Liver	0.9091	1	0.8333
SLC27A5 AND NOT-GPR182	Liver	0.8	1	0.6667
ABCC6 AND NOT-CLEC4G	Liver	0.8	1	0.6667
SCARB1 AND NOT-LYVE1	Liver	0.8	1	0.6667
SLC4A2 AND NOT-ATP1B3	Liver	0.8	1	0.6667
SLC4A2 AND NOT-PMP22	Liver	0.8	1	0.6667
GJB1 AND NOT-CLEC1B	Liver	0.8	1	0.6667
HPN AND NOT-CLEC4G	Liver	0.8	1	0.6667
SLC27A5 AND NOT-CLEC4G	Liver	0.8	1	0.6667
SLC4A2 AND NOT-KCNN4	Liver	0.8	1	0.6667
SLC4A2 AND NOT-JAM2	Liver	0.8	1	0.6667
FGG AND NOT-CLEC4G	Liver	0.8	1	0.6667
FGG AND NOT-CLEC4M	Liver	0.8	1	0.6667
SLC4A2 AND NOT-SLC24A3	Liver	0.8	1	0.6667
ABCC6 AND NOT-CLEC4M	Liver	0.8	1	0.6667
ABCC6 AND NOT-VIPR1	Liver	0.8	1	0.6667
SLC4A2 AND NOT-JAM3	Liver	0.8	1	0.6667
HPN AND NOT-CLEC4M	Liver	0.8	1	0.6667
ABCC6 AND NOT-MARCO	Liver	0.8	0.6667	1
SLC27A5 AND NOT-PTH1R	Liver	0.8	1	0.6667
SLC27A5 AND NOT-CLEC4M	Liver	0.8	1	0.6667
SLC4A2 AND NOT-SLC44A2	Liver	0.8	1	0.6667
SLC4A2 AND NOT-CLSTN1	Liver	0.8	1	0.6667
SLC4A2 AND NOT-MARVELD1	Liver	0.8	1	0.6667
GJB1 AND NOT-CLEC4G	Liver	0.8	1	0.6667
SMO AND NOT-GPM6B	Liver	0.8	1	0.6667
SLC27A5 AND NOT-MARCO	Liver	0.8	0.6667	1
ASGR1 AND NOT-CLEC1B	Liver	0.8	1	0.6667
SLC4A2 AND NOT-CD320	Liver	0.8	1	0.6667
FGG AND NOT-GPR182	Liver	0.8	1	0.6667
SLC4A2 AND NOT-PDPN	Liver	0.8	1	0.6667
HPN AND NOT-GPR182	Liver	0.8	1	0.6667
GJB1 AND NOT-CLEC4M	Liver	0.8	1	0.6667
GJB1 AND NOT-MARCO	Liver	0.8	1	0.6667
HPN AND NOT-PTH1R	Liver	0.8	1	0.6667
SLC4A2 AND NOT-TGFB3	Liver	0.8	1	0.6667
SMO AND NOT-PLA2R1	Liver	0.8	1	0.6667
SLC4A2 AND NOT-FZD7	Liver	0.8	1	0.6667
TFR2 AND NOT-CLEC1B	Liver	0.8	1	0.6667
HPN AND NOT-STAB2	Liver	0.8	1	0.6667
KCNE4 AND NOT-TSPAN18	Pancreas	0.64	0.5714	0.7273
NOX4 AND NOT-PTH1R	Pancreas	0.6316	0.75	0.5455
TREM2 AND NOT-FXYD1	Pancreas	0.625	1	0.4545
CDH6 AND NOT-CNTN1	Renal	0.8571	1	0.75
CDH6 AND NOT-COL25A1	Renal	0.8571	1	0.75
CDH6 AND NOT-FAM26E	Renal	0.8571	1	0.75
CDH6 AND NOT-TRPM7	Renal	0.8571	1	0.75

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
FAP AND NOT-TSPAN7	Pancreas	0.7778	1	0.6364
FAP AND NOT-SLC16A1	Pancreas	0.7059	1	0.5455
FAP AND NOT-GPR161	Pancreas	0.625	1	0.4545
FAP AND NOT-FGFR1	Pancreas	0.625	1	0.4545
FAP AND NOT-FZD7	Pancreas	0.625	1	0.4545
FAP AND NOT-NDRG4	Pancreas	0.8421	1	0.7273
FAP AND NOT-KCNB1	Pancreas	0.8421	1	0.7273
FAP AND NOT-PRIMA1	Pancreas	0.7778	1	0.6364
FAP AND NOT-TRPC1	Pancreas	0.7059	1	0.5455
FAP AND NOT-FZD4	Pancreas	0.7778	1	0.6364
FAP AND NOT-TMEM25	Pancreas	0.625	1	0.4545
FAP AND NOT-TMEM231	Pancreas	0.7059	1	0.5455
CDH6 AND NOT-ERBB4	Renal	0.8571	1	0.75
CDH6 AND NOT-BMPR1B	Renal	0.8571	1	0.75
CDH6 AND NOT-CLDN11	Renal	0.8571	1	0.75
CDH6 AND NOT-IL11RA	Renal	0.8571	1	0.75
CDH6 AND NOT-FAP	Renal	0.8571	1	0.75
CDH6 AND NOT-MUC16	Renal	0.8571	1	0.75
CDH6 AND NOT-IL13RA1	Renal	0.8571	1	0.75
CDH6 AND NOT-CLDN8	Renal	0.8571	1	0.75
CDH6 AND NOT-SLC7A5	Renal	0.8571	1	0.75
CDH6 AND NOT-LGR5	Renal	0.8571	1	0.75
CDH6 AND NOT-CLDN6	Renal	0.8571	1	0.75
CDH6 AND NOT-SSTR2	Renal	0.8571	1	0.75
CDH6 AND NOT-CSPG4	Renal	0.8571	1	0.75
CDH6 AND NOT-FCRL1	Renal	0.8571	1	0.75
CDH6 AND NOT-CD276	Renal	0.8571	1	0.75
CDH6 AND NOT-IL2RA	Renal	0.8571	1	0.75
CDH6 AND NOT-CR2	Renal	0.8571	1	0.75
CDH6 AND NOT-ULBP2	Renal	0.8571	1	0.75
CDH6 AND NOT-P2RX5	Renal	0.8571	1	0.75
CDH6 AND NOT-PMEL	Renal	0.8571	1	0.75
CDH6 AND NOT-SSTR1	Renal	0.8571	1	0.75
CDH6 AND NOT-CLDN12	Renal	0.8571	1	0.75
CDH6 AND NOT-MUC1	Renal	0.8571	1	0.75
CDH6 AND NOT-FOLH1	Renal	0.8571	1	0.75
CDH6 AND NOT-FCRL2	Renal	0.8571	1	0.75
CDH6 AND NOT-RNF43	Renal	0.8571	1	0.75
CDH6 AND NOT-CD72	Renal	0.8571	1	0.75
CDH6 AND NOT-MUC13	Renal	0.8	1	0.6667
KCNJ16 AND NOT-ERBB4	Renal	0.8	1	0.6667
CDH6 AND NOT-AXL	Renal	0.8	1	0.6667
CDH6 AND NOT-TNFRSF17	Renal	0.8	1	0.6667
CDH6 AND NOT-TRPM4	Renal	0.8571	1	0.75
CDH6 AND NOT-PTK7	Renal	0.8571	1	0.75
CDH6 AND NOT-GPA33	Renal	0.8571	1	0.75
CDH6 AND NOT-IL20RA	Renal	0.8571	1	0.75
CDH6 AND NOT-CLDN18	Renal	0.8571	1	0.75
CDH6 AND NOT-ENG	Renal	0.8571	1	0.75
CDH6 AND NOT-ROR1	Renal	0.8	1	0.6667
CDH6 AND NOT-EPHB2	Renal	0.8	1	0.6667
CDH6 AND NOT-ERBB2	Renal	0.8571	1	0.75
CDH6 AND NOT-EGFR	Renal	0.8571	1	0.75
CDH6 AND NOT-SLC39A6	Renal	0.8	1	0.6667
CDH6 AND NOT-STEAP2	Renal	0.8571	1	0.75
CDH6 AND NOT-STEAP1	Renal	0.8571	1	0.75
CDH6 AND NOT-GUCY2C	Renal	0.8571	1	0.75
CDH6 AND NOT-IGF1R	Renal	0.8571	1	0.75
CDH6 AND NOT-CLDN23	Renal	0.8	1	0.6667
CDH6 AND NOT-SSTR3	Renal	0.8571	1	0.75
CDH6 AND NOT-ITGB6	Renal	0.8	1	0.6667
CDH6 AND NOT-CD180	Renal	0.8571	1	0.75
CDH6 AND NOT-RAET1E	Renal	0.8571	1	0.75
CDH6 AND NOT-MUC4	Renal	0.8571	1	0.75
CDH6 AND NOT-EPHA3	Renal	0.8571	1	0.75
CDH6 AND NOT-IL3RA	Renal	0.8571	1	0.75
CDH6 AND NOT-SSTR4	Renal	0.8571	1	0.75
CDH6 AND NOT-CD22	Renal	0.8571	1	0.75
CDH6 AND NOT-CLDN9	Renal	0.8	1	0.6667
CDH6 AND NOT-TNFRSF8	Renal	0.8571	1	0.75
CDH6 AND NOT-CLDN1	Renal	0.8571	1	0.75
CDH6 AND NOT-TNFRSF10A	Renal	0.8	1	0.6667
CDH6 AND NOT-TNFSF11	Renal	0.8	1	0.6667
CDH6 AND NOT-FCRL5	Renal	0.8571	1	0.75
CDH6 AND NOT-CD38	Renal	0.8	1	0.6667
CDH6 AND NOT-MS4A1	Renal	0.8	1	0.6667
CDH6 AND NOT-CD79A	Renal	0.8	1	0.6667

FIG. 12 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
CDH6 AND NOT-SCN7A	Renal	0.8571	1	0.75
CDH16 AND NOT-SCNN1B	Renal	0.8571	1	0.75
CDH6 AND NOT-DST	Renal	0.8571	1	0.75
CDH6 AND NOT-ANTXR1	Renal	0.8571	1	0.75
CDH6 AND NOT-USP48	Renal	0.8571	1	0.75
CDH6 AND NOT-KCNE1	Renal	0.8571	1	0.75
CDH6 AND NOT-KIRREL	Renal	0.8571	1	0.75
CDH6 AND NOT-BMPR2	Renal	0.8571	1	0.75
CDH6 AND NOT-MRAP2	Renal	0.8571	1	0.75
CDH6 AND NOT-ZDHHC20	Renal	0.8571	1	0.75
CDH6 AND NOT-BMPR1A	Renal	0.8571	1	0.75
CDH6 AND NOT-SLC20A2	Renal	0.8571	1	0.75
CDH6 AND NOT-BVES	Renal	0.8571	1	0.75
CDH6 AND NOT-ADAM23	Renal	0.8571	1	0.75
CDH6 AND NOT-SLC2A12	Renal	0.8571	1	0.75
CDH6 AND NOT-GPR39	Renal	0.8571	1	0.75
CDH6 AND NOT-NRP2	Renal	0.8571	1	0.75
CDH6 AND NOT-SLC12A2	Renal	0.8571	1	0.75
CDH6 AND NOT-ST3GAL5	Renal	0.8571	1	0.75
CDH6 AND NOT-ATP11C	Renal	0.8571	1	0.75
CDH6 AND NOT-PLA2R1	Renal	0.8571	1	0.75
CDH6 AND NOT-GPR173	Renal	0.8571	1	0.75
CDH6 AND NOT-SLC8A1	Renal	0.8571	1	0.75
CDH6 AND NOT-PDGFRA	Renal	0.8571	1	0.75
CDH6 AND NOT-FAT3	Renal	0.8571	1	0.75
CDH6 AND NOT-PCDH9	Renal	0.8571	1	0.75
CDH6 AND NOT-BACE1	Renal	0.8571	1	0.75
CDH6 AND NOT-OXGR1	Renal	0.8571	1	0.75
CDH6 AND NOT-CRIM1	Renal	0.8571	1	0.75
CDH6 AND NOT-FLRT2	Renal	0.8571	1	0.75
CDH6 AND NOT-TRPV1	Renal	0.8571	1	0.75
CDH6 AND NOT-TGFBR1	Renal	0.8571	1	0.75
CDH6 AND NOT-RELL1	Renal	0.8571	1	0.75
CDH6 AND NOT-CLCN3	Renal	0.8571	1	0.75
CDH6 AND NOT-ATP11B	Renal	0.8571	1	0.75
CDH6 AND NOT-SLMAP	Renal	0.8571	1	0.75
CDH6 AND NOT-ATP13A3	Renal	0.8571	1	0.75
CDH6 AND NOT-GPM6B	Renal	0.8571	1	0.75
CDH6 AND NOT-CYBRD1	Renal	0.8571	1	0.75
CDH6 AND NOT-WLS	Renal	0.8571	1	0.75
CDH6 AND NOT-SYT1	Renal	0.8571	1	0.75
CDH6 AND NOT-ADAM22	Renal	0.8571	1	0.75
CDH6 AND NOT-COLEC12	Renal	0.8571	1	0.75
CDH6 AND NOT-GPR63	Renal	0.8571	1	0.75
CDH6 AND NOT-TPM1	Renal	0.8571	1	0.75
CDH6 AND NOT-TMEM65	Renal	0.8571	1	0.75
CDH6 AND NOT-IL6ST	Renal	0.8571	1	0.75
CDH6 AND NOT-SLC12A6	Renal	0.8571	1	0.75
CDH6 AND NOT-ACVR2A	Renal	0.8571	1	0.75
CDH6 AND NOT-LRPAP1	Renal	0.8571	1	0.75
CDH6 AND NOT-SLC9A6	Renal	0.8571	1	0.75
CDH6 AND NOT-ZDHHC21	Renal	0.8571	1	0.75
CDH6 AND NOT-PTPRA	Renal	0.8571	1	0.75
CDH6 AND NOT-LSAMP	Renal	0.8571	1	0.75
SLC3A1 AND NOT-TMEM30B	Renal	0.8571	1	0.75
CDH6 AND NOT-DIO2	Renal	0.8571	1	0.75
CDH6 AND NOT-PTPRJ	Renal	0.8571	1	0.75
CDH6 AND NOT-CD47	Renal	0.8571	1	0.75
CDH6 AND NOT-DPP6	Renal	0.8571	1	0.75
CDH6 AND NOT-LRRTM2	Renal	0.8571	1	0.75
CDH6 AND NOT-RET	Renal	0.8571	1	0.75
CDH6 AND NOT-TSPAN2	Renal	0.8571	1	0.75
CDH6 AND NOT-HHIP	Renal	0.8571	1	0.75
CDH6 AND NOT-SLC4A3	Renal	0.8571	1	0.75
CDH6 AND NOT-PRPH2	Renal	0.8571	1	0.75
CDH6 AND NOT-ROM1	Renal	0.8571	1	0.75
CDH6 AND NOT-SLC36A1	Renal	0.8571	1	0.75
CDH6 AND NOT-SLC7A10	Renal	0.8571	1	0.75
CDH6 AND NOT-SLC7A1	Renal	0.8571	1	0.75
CDH6 AND NOT-SLC6A9	Renal	0.8571	1	0.75
CDH6 AND NOT-GRIK2	Renal	0.8571	1	0.75
CDH6 AND NOT-SLC7A6	Renal	0.8571	1	0.75
CDH6 AND NOT-SLC25A4	Renal	0.8571	1	0.75
CDH6 AND NOT-M6PR	Renal	0.8571	1	0.75
CDH6 AND NOT-SLC5A3	Renal	0.8571	1	0.75
CDH6 AND NOT-PLXNA4	Renal	0.8571	1	0.75
CDH6 AND NOT-CORIN	Renal	0.8571	1	0.75
CDH6 AND NOT-CD79B	Renal	0.8571	1	0.75
CDH6 AND NOT-CLDN5	Renal	0.8571	1	0.75
CDH6 AND NOT-VTCN1	Renal	0.8571	1	0.75
CDH6 AND NOT-SSTR5	Renal	0.8	1	0.6667
CDH6 AND NOT-TPBG	Renal	0.8	1	0.6667
CDH6 AND NOT-SDC1	Renal	0.8	1	0.6667
CDH6 AND NOT-MST1R	Renal	0.8571	1	0.75
SLC3A1 AND NOT-LGR5	Renal	0.7826	0.8182	0.75
CDH16 AND NOT-LGR5	Renal	0.7826	0.8182	0.75
CDH16 AND NOT-CLDN8	Renal	0.7619	0.8889	0.6667
DPEP1 AND NOT-SLC16A7	Colon	1	1	1
DPEP1 AND NOT-GP2	Colon	1	1	1
DPEP1 AND NOT-ZP2	Colon	1	1	1
DPEP1 AND NOT-SLC22A7	Colon	1	1	1
DPEP1 AND NOT-P2RX6	Colon	1	1	1
MUC13 AND NOT-FLVCR2	Colon	0.9091	0.8333	1
GUCY2C AND NOT-FLVCR2	Colon	0.9091	0.8333	1
DPEP1 AND NOT-KCNJ13	Colon	1	1	1
DPEP1 AND NOT-RTP1	Colon	1	1	1
DPEP1 AND NOT-OR2C1	Colon	1	1	1
DPEP1 AND NOT-SLC5A12	Colon	1	1	1
DPEP1 AND NOT-OR10H1	Colon	1	1	1
DPEP1 AND NOT-GRM2	Colon	0.9091	0.8333	1
GPA33 AND NOT-FLVCR2	Colon	0.9091	0.8333	1
CLDN7 AND NOT-SLC22A23	Colon	0.9091	0.8333	1
DPEP1 AND NOT-NTRK3	Colon	1	1	1
EPHB2 AND NOT-VSIG2	Colon	0.8889	1	0.8
EPHB2 AND NOT-CNNM4	Colon	0.8889	1	0.8
CLDN2 AND NOT-SLC16A7	Colon	0.8889	1	0.8
CLDN2 AND NOT-SLC19A3	Colon	0.8889	1	0.8
GPA33 AND NOT-SEMA4G	Colon	0.8889	1	0.8
EPHB2 AND NOT-FLVCR2	Colon	0.8889	1	0.8
CLDN2 AND NOT-P2RX6	Colon	0.8889	1	0.8
EPHB2 AND NOT-NEO1	Colon	0.8889	1	0.8
EPHB2 AND NOT-ATP2A3	Colon	0.8889	1	0.8
CLDN2 AND NOT-ABCC2	Colon	0.8889	1	0.8
EPHB2 AND NOT-SLC4A4	Colon	0.8889	1	0.8
EPHB2 AND NOT-F2RL1	Colon	0.8889	1	0.8
DPEP1 AND NOT-OR10A5	Colon	0.8889	1	0.8
CLDN2 AND NOT-ADRA1B	Colon	0.8889	1	0.8
DPEP1 AND NOT-CYSLTR2	Colon	0.8889	1	0.8
DPEP1 AND NOT-CACNA1I	Colon	0.8889	1	0.8
GUCY2C AND NOT-SEMA4G	Colon	0.8889	1	0.8
DPEP1 AND NOT-CYP4F2	Colon	0.8889	1	0.8
PMEPA1 AND NOT-IL11RA	Colon	0.8889	1	0.8
EPHB2 AND NOT-ADCY6	Colon	0.8889	1	0.8
DPEP1 AND NOT-SPN	Colon	0.8889	1	0.8
MUC13 AND NOT-SEMA4G	Colon	0.8889	1	0.8
EPHB2 AND NOT-AMN	Colon	0.8889	1	0.8
DPEP1 AND NOT-C1orf101	Colon	0.8889	1	0.8
EPHB2 AND NOT-SEMA4G	Colon	0.8889	1	0.8
EPHB2 AND NOT-SCNN1B	Colon	0.8889	1	0.8
EPHB2 AND NOT-UGT8	Colon	0.8889	1	0.8
EPHB2 AND NOT-PSEN1	Colon	0.8889	1	0.8
EPHB2 AND NOT-PTPRD	Colon	0.8889	1	0.8
EPHB2 AND NOT-TMPRSS2	Colon	0.8889	1	0.8
EPHB2 AND NOT-SLC22A23	Colon	0.8889	1	0.8
EPHB2 AND NOT-CA12	Colon	0.8889	1	0.8
EPHB2 AND NOT-SEMA6A	Colon	0.8889	1	0.8
EPHB2 AND NOT-PTGDR	Colon	0.8889	1	0.8
GUCY2C AND NOT-SLC22A23	Colon	0.9091	0.8333	1
DPEP1 AND NOT-KCNG3	Colon	1	1	1
DPEP1 AND NOT-CNGB3	Colon	1	1	1
DPEP1 AND NOT-GRPR	Colon	1	1	1
DPEP1 AND NOT-ESYT3	Colon	1	1	1
DPEP1 AND NOT-ATP1B2	Colon	1	1	1
DPEP1 AND NOT-LPAR3	Colon	0.9091	0.8333	1
DPEP1 AND NOT-KCNJ1	Colon	1	1	1
DPEP1 AND NOT-UNC5D	Colon	1	1	1
DPEP1 AND NOT-SLC4A4	Colon	1	1	1
DPEP1 AND NOT-GDPD2	Colon	0.9091	0.8333	1
DPEP1 AND NOT-KCNJ5	Colon	1	1	1
FZD3 AND NOT-B4GALNT1	Ependymoma	1	1	1
SLC1A3 AND NOT-CD33	Ependymoma	1	1	1
FZD3 AND NOT-P2RX5	Ependymoma	1	1	1
SLC1A3 AND NOT-CD37	Ependymoma	1	1	1
SLC1A3 AND NOT-IL13RA1	Ependymoma	1	1	1

FIG. 12 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
CDH6 AND NOT-SORT1	Renal	0.8571	1	0.75
CDH6 AND NOT-ACVR1B	Renal	0.8571	1	0.75
CDH6 AND NOT-CNST	Renal	0.8571	1	0.75
CDH6 AND NOT-TMEM150C	Renal	0.8571	1	0.75
CDH6 AND NOT-HEPH	Renal	0.8571	1	0.75
CDH6 AND NOT-REEP2	Renal	0.8571	1	0.75
CDH6 AND NOT-OPN1SW	Renal	0.8571	1	0.75
CDH6 AND NOT-ROBO2	Renal	0.8571	1	0.75
CDH6 AND NOT-GRIA1	Renal	0.8571	1	0.75
CDH6 AND NOT-LRP8	Renal	0.8571	1	0.75
CDH6 AND NOT-APLP1	Renal	0.8571	1	0.75
CDH6 AND NOT-ATP13A2	Renal	0.8571	1	0.75
ATP10B AND NOT-FLVCR2	Colon	0.9091	0.8333	1
HEPH AND NOT-FLVCR2	Colon	0.9091	0.8333	1
MEP1A AND NOT-FLVCR2	Colon	0.9091	0.8333	1
PMEPA1 AND NOT-GABRE	Colon	0.8889	1	0.8
FGFR4 AND NOT-LIFR	Colon	0.8889	1	0.8
PMEPA1 AND NOT-FZD4	Colon	0.8889	1	0.8
PMEPA1 AND NOT-ANO3	Colon	0.8889	1	0.8
PMEPA1 AND NOT-LIFR	Colon	0.8889	1	0.8
PMEPA1 AND NOT-PCDH9	Colon	0.8889	1	0.8
PMEPA1 AND NOT-ITGA7	Colon	0.8889	1	0.8
PMEPA1 AND NOT-MPZ	Colon	0.8889	1	0.8
PMEPA1 AND NOT-ADRB2	Colon	0.8889	1	0.8
CD320 AND NOT-LPAR1	Colon	0.8889	1	0.8
CD320 AND NOT-EFNA5	Colon	0.8889	1	0.8
PMEPA1 AND NOT-ATP8B4	Colon	0.8889	1	0.8
PMEPA1 AND NOT-SLC9A9	Colon	0.8889	1	0.8
MEP1A AND NOT-SEMA4G	Colon	0.8889	1	0.8
CYBA AND NOT-LRMP	Colon	1	1	1
PMEPA1 AND NOT-ADCY6	Colon	0.8889	1	0.8
PMEPA1 AND NOT-NAALAD2	Colon	0.8889	1	0.8
CD320 AND NOT-BEST4	Colon	0.8889	1	0.8
PMEPA1 AND NOT-AGTR1	Colon	0.8889	1	0.8
PPAP2C AND NOT-ABHD6	Colon	0.8889	1	0.8
PMEPA1 AND NOT-KL	Colon	0.8889	1	0.8
PSENEN AND NOT-LRMP	Colon	0.8889	1	0.8
PMEPA1 AND NOT-CDON	Colon	0.8889	1	0.8
PMEPA1 AND NOT-SLC16A2	Colon	0.8889	1	0.8
SIGMAR1 AND NOT-SLC6A16	Colon	0.8889	1	0.8
PMEPA1 AND NOT-SFRP1	Colon	0.8889	1	0.8
PMEPA1 AND NOT-SLC41A1	Colon	0.8889	1	0.8
PMEPA1 AND NOT-CX3CL1	Colon	0.8889	1	0.8
PMEPA1 AND NOT-ATP1A2	Colon	0.8889	1	0.8
ATP10B AND NOT-SLC22A23	Colon	0.9091	0.8333	1
PMEPA1 AND NOT-VSTM4	Colon	0.8889	1	0.8
HEPH AND NOT-SLC22A23	Colon	0.9091	0.8333	1
PMEPA1 AND NOT-PEAR1	Colon	0.8889	1	0.8
PMEPA1 AND NOT-ADCY5	Colon	0.8889	1	0.8
PMEPA1 AND NOT-NTRK3	Colon	0.8889	1	0.8
PMEPA1 AND NOT-ACSL1	Colon	0.8889	1	0.8
PMEPA1 AND NOT-SLC46A1	Colon	0.8889	1	0.8
LSR AND NOT-FLVCR2	Colon	0.8889	1	0.8
PPAP2C AND NOT-SLC22A23	Colon	0.8889	1	0.8
SIGMAR1 AND NOT-KIAA0754	Colon	0.8889	1	0.8
PMEPA1 AND NOT-GPM6B	Colon	0.8889	1	0.8
EPHB4 AND NOT-SLC9A9	Colon	0.8333	0.7143	1
HEPH AND NOT-ABHD6	Colon	1	1	1
SLC26A3 AND NOT-FLVCR2	Colon	0.9091	0.8333	1
HEPH AND NOT-ANO7	Colon	0.8	0.8	0.8
LSR AND NOT-ATP6V0A1	Colon	0.8	0.8	0.8
PMEPA1 AND NOT-OR2C1	Colon	0.8889	1	0.8
LSR AND NOT-ABHD6	Colon	0.8889	1	0.8
FZD3 AND NOT-KCNJ2	Ependymoma	1	1	1
FZD3 AND NOT-KCNJ4	Ependymoma	1	1	1
SLC1A3 AND NOT-HTRA2	Ependymoma	1	1	1
CDH2 AND NOT-ADCY9	Ependymoma	1	1	1
FZD3 AND NOT-KCNK1	Ependymoma	1	1	1
FZD3 AND NOT-KCNMA1	Ependymoma	1	1	1
FZD3 AND NOT-KCNQ3	Ependymoma	1	1	1
TMEM231 AND NOT-KCNJ2	Ependymoma	1	1	1
SLC1A3 AND NOT-TNFRSF1A	Ependymoma	1	1	1
SLC1A3 AND NOT-TGFBR2	Ependymoma	1	1	1
FZD3 AND NOT-CDH12	Ependymoma	1	1	1
SLC1A3 AND NOT-LRMP	Ependymoma	1	1	1
SLC1A3 AND NOT-LTB	Ependymoma	1	1	1
FZD3 AND NOT-SCN2B	Ependymoma	1	1	1

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
SLC1A3 AND NOT-CXCR5	Ependymoma	1	1	1
SLC1A3 AND NOT-MS4A1	Ependymoma	1	1	1
SLC1A3 AND NOT-FCRL2	Ependymoma	1	1	1
SLC1A3 AND NOT-TNFRSF17	Ependymoma	1	1	1
SLC1A3 AND NOT-FCRL5	Ependymoma	1	1	1
SLC7A11 AND NOT-IL13RA1	Ependymoma	1	1	1
KCNA6 AND NOT-FOLH1	Ependymoma	1	1	1
WDR19 AND NOT-IL13RA1	Ependymoma	1	1	1
SSTR1 AND NOT-INSR	Ependymoma	1	1	1
SSTR1 AND NOT-XK	Ependymoma	1	1	1
KCNA6 AND NOT-EPCAM	Ependymoma	1	1	1
ROR2 AND NOT-STEAP1	Ependymoma	1	1	1
KCNA6 AND NOT-P2RX5	Ependymoma	1	1	1
KCNA6 AND NOT-B4GALNT1	Ependymoma	1	1	1
SSTR1 AND NOT-KCNS3	Ependymoma	1	1	1
FZD3 AND NOT-L1CAM	Ependymoma	1	1	1
SSTR1 AND NOT-ATRN	Ependymoma	1	1	1
TSPAN11 AND NOT-MET	Ependymoma	1	1	1
PCDHB10 AND NOT-MET	Ependymoma	1	1	1
KCNA6 AND NOT-CLDN1	Ependymoma	1	1	1
KCNA6 AND NOT-STEAP2	Ependymoma	1	1	1
SSTR1 AND NOT-EMB	Ependymoma	1	1	1
ZNRF3 AND NOT-CD34	Ependymoma	1	1	1
SSTR1 AND NOT-SLC31A2	Ependymoma	1	1	1
SSTR1 AND NOT-GJB2	Ependymoma	1	1	1
ROR2 AND NOT-ENG	Ependymoma	1	1	1
SSTR1 AND NOT-MRAP2	Ependymoma	1	1	1
DLL1 AND NOT-STEAP1	Ependymoma	1	1	1
KCNA6 AND NOT-EPHA3	Ependymoma	1	1	1
SSTR1 AND NOT-SLC17A5	Ependymoma	1	1	1
DLL1 AND NOT-IL2RA	Ependymoma	1	1	1
SSTR1 AND NOT-CD47	Ependymoma	1	1	1
DLL1 AND NOT-MET	Ependymoma	1	1	1
KCNA6 AND NOT-EPHB2	Ependymoma	1	1	1
DLL1 AND NOT-TRPM4	Ependymoma	1	1	1
DLL1 AND NOT-SDC1	Ependymoma	1	1	1
SSTR1 AND NOT-SLC4A7	Ependymoma	1	1	1
SSTR1 AND NOT-MTDH	Ependymoma	1	1	1
DLL1 AND NOT-CD52	Ependymoma	1	1	1
SSTR1 AND NOT-MYADM	Ependymoma	1	1	1
KCNA6 AND NOT-ERBB4	Ependymoma	1	1	1
DLL1 AND NOT-TNFRSF10A	Ependymoma	1	1	1
SSTR1 AND NOT-STX3	Ependymoma	1	1	1
KCNA6 AND NOT-CLDN12	Ependymoma	1	1	1
ZNRF3 AND NOT-AXL	Ependymoma	1	1	1
ROR2 AND NOT-CD276	Ependymoma	1	1	1
SSTR1 AND NOT-STYK1	Ependymoma	1	1	1
SSTR1 AND NOT-ANO10	Ependymoma	1	1	1
ROR2 AND NOT-CD34	Ependymoma	1	1	1
SSTR1 AND NOT-KIAA1324	Ependymoma	1	1	1
SSTR1 AND NOT-SMPD3	Ependymoma	1	1	1
SSTR1 AND NOT-NTN4	Ependymoma	1	1	1
SSTR1 AND NOT-TMEM127	Ependymoma	1	1	1
SLC39A4 AND NOT-ENPP3	Esophagus	0.7273	1	0.5714
SLC52A2 AND NOT-ENPP3	Esophagus	0.7273	1	0.5714
PTPRH AND NOT-ENPP3	Esophagus	0.7273	1	0.5714
SLC52A2 AND NOT-ABCA5	Esophagus	0.8	0.9091	0.7143
SLC52A2 AND NOT-IL3RA	Esophagus	0.6667	1	0.5
SLC52A2 AND NOT-TRPM4	Esophagus	0.7826	1	0.6429
MOK AND NOT-WDR19	Esophagus	0.6667	0.8	0.5714
SLC52A2 AND NOT-FOLR2	Esophagus	0.6667	1	0.5
MUC13 AND NOT-PIGR	Esophagus	0.7273	1	0.5714
SLC52A2 AND NOT-CLDN8	Esophagus	0.7826	1	0.6429
TM4SF5 AND NOT-ENPP3	Esophagus	0.7273	1	0.5714
SLC52A2 AND NOT-TYR	Esophagus	0.7826	1	0.6429
SLC52A2 AND NOT-TNFRSF8	Esophagus	0.7826	1	0.6429
SLC52A2 AND NOT-IL11RA	Esophagus	0.8333	1	0.7143
SLC52A2 AND NOT-FCRL1	Esophagus	0.7826	1	0.6429
SLC52A2 AND NOT-CD160	Esophagus	0.8333	1	0.7143
SLC52A2 AND NOT-HHLA2	Esophagus	0.6	1	0.4286
SLC52A2 AND NOT-CD72	Esophagus	0.6	1	0.4286
SLC52A2 AND NOT-CLEC14A	Esophagus	0.6	1	0.4286
MUC13 AND NOT-SCNN1B	Esophagus	0.6	1	0.4286
SLC52A2 AND NOT-SST	Esophagus	0.6	1	0.4286
SLC52A2 AND NOT-GPA33	Esophagus	0.7826	1	0.6429
SLC52A2 AND NOT-RAET1E	Esophagus	0.7826	1	0.6429
SLC52A2 AND NOT-PMEL	Esophagus	0.6667	1	0.5

FIG. 12 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
SLC16A2 AND NOT-LTBR	Ependymoma	1	1	1
SLC1A3 AND NOT-LY9	Ependymoma	1	1	1
SLC7A11 AND NOT-SLC12A6	Ependymoma	1	1	1
FZD3 AND NOT-RYR2	Ependymoma	1	1	1
SLC1A3 AND NOT-LY75	Ependymoma	1	1	1
SLC1A3 AND NOT-KLRB1	Ependymoma	1	1	1
SLC1A3 AND NOT-LAMP2	Ependymoma	1	1	1
SLC7A11 AND NOT-USP48	Ependymoma	1	1	1
SLC1A3 AND NOT-P2RY10	Ependymoma	1	1	1
SLC1A3 AND NOT-SELPLG	Ependymoma	1	1	1
SLC16A2 AND NOT-LEPR	Ependymoma	1	1	1
SLC1A3 AND NOT-CD2	Ependymoma	1	1	1
SLC52A2 AND NOT-SMPD1	Esophagus	0.7826	1	0.6429
SLC52A2 AND NOT-SCNN1B	Esophagus	0.7826	1	0.6429
SLC52A2 AND NOT-KLB	Esophagus	0.8333	1	0.7143
SLC52A2 AND NOT-CDHR1	Esophagus	0.8333	1	0.7143
SLC52A2 AND NOT-ST6GALNAC6	Esophagus	0.8333	1	0.7143
SLC52A2 AND NOT-VIPR1	Esophagus	0.8333	1	0.7143
GJB3 AND NOT-CDHR1	Esophagus	0.6897	0.6667	0.7143
BCAP31 AND NOT-ST6GALNAC6	Esophagus	0.7273	1	0.5714
SLC52A2 AND NOT-CLEC10A	Esophagus	0.7826	1	0.6429
SLC52A2 AND NOT-SLCO2A1	Esophagus	0.75	0.9	0.6429
SLC52A2 AND NOT-AMICA1	Esophagus	0.7826	1	0.6429
SLC52A2 AND NOT-PTGDR	Esophagus	0.8333	1	0.7143
SLC52A2 AND NOT-ABCA8	Esophagus	0.7826	1	0.6429
SLC52A2 AND NOT-GDPD2	Esophagus	0.7273	1	0.5714
SLC52A2 AND NOT-ABHD6	Esophagus	0.7826	1	0.6429
GPRC5A AND NOT-ST6GALNAC6	Esophagus	0.7826	1	0.6429
SLC52A2 AND NOT-LRMP	Esophagus	0.7273	1	0.5714
FZD5 AND NOT-PTPRD	Esophagus	0.7273	1	0.5714
SLC39A4 AND NOT-SCNN1B	Esophagus	0.6667	1	0.5
SLC52A2 AND NOT-PTPRD	Esophagus	0.8333	1	0.7143
SLC52A2 AND NOT-ABCB1	Esophagus	0.6667	1	0.5
SLC52A2 AND NOT-NEO1	Esophagus	0.6667	1	0.5
SLC52A2 AND NOT-ADTRP	Esophagus	0.6667	1	0.5
SLC52A2 AND NOT-RNF144A	Esophagus	0.6667	1	0.5
SLC52A2 AND NOT-ATP6V0A1	Esophagus	0.6667	1	0.5
SLC52A2 AND NOT-CDH19	Esophagus	0.6667	1	0.5
SLC52A2 AND NOT-ITGB7	Esophagus	0.8333	1	0.7143
SLC52A2 AND NOT-POPDC2	Esophagus	0.7273	1	0.5714
SLC52A2 AND NOT-CA12	Esophagus	0.8333	1	0.7143
SLC52A2 AND NOT-TMEM100	Esophagus	0.7826	1	0.6429
SLC52A2 AND NOT-SLC26A6	Esophagus	0.8333	1	0.7143
SLC52A2 AND NOT-NPY1R	Esophagus	0.7826	1	0.6429
SLC52A2 AND NOT-TNFRSF25	Esophagus	0.8333	1	0.7143
SLC52A2 AND NOT-SCNN1G	Esophagus	0.7826	1	0.6429
SLC52A2 AND NOT-GPR173	Esophagus	0.7826	1	0.6429
SLC52A2 AND NOT-SCN3A	Esophagus	0.7826	1	0.6429
SLC52A2 AND NOT-NCAM2	Esophagus	0.6667	1	0.5
SLC52A2 AND NOT-UNC5D	Esophagus	0.7826	1	0.6429
SLC52A2 AND NOT-NLGN1	Esophagus	0.7826	1	0.6429
SLC52A2 AND NOT-MPZ	Esophagus	0.6667	1	0.5
SLC52A2 AND NOT-MCOLN1	Esophagus	0.7826	1	0.6429
SLC52A2 AND NOT-GABRA6	Esophagus	0.6667	1	0.5
SLC52A2 AND NOT-FLT3	Esophagus	0.7273	1	0.5714
SLC52A2 AND NOT-CD101	Esophagus	0.7826	1	0.6429
SLC52A2 AND NOT-IL6R	Esophagus	0.7273	1	0.5714
SLC52A2 AND NOT-CD163L1	Esophagus	0.7273	1	0.5714
SLC52A2 AND NOT-RHOT2	Esophagus	0.7692	0.8333	0.7143
SLC52A2 AND NOT-SEMA6D	Esophagus	0.7826	1	0.6429
SLC52A2 AND NOT-MCOLN2	Esophagus	0.7273	1	0.5714
GPRC5A AND NOT-SCNN1G	Esophagus	0.6897	0.6667	0.7143
SLC52A2 AND NOT-SLC26A10	Esophagus	0.7826	1	0.6429
SLC52A2 AND NOT-KCND1	Esophagus	0.7826	1	0.6429
SLC52A2 AND NOT-AQP8	Esophagus	0.7826	1	0.6429
SLC52A2 AND NOT-SLC9A5	Esophagus	0.8333	1	0.7143
SLC52A2 AND NOT-PKD1	Esophagus	0.8333	1	0.7143
SLC52A2 AND NOT-SLC6A16	Esophagus	0.7273	1	0.5714
SLC52A2 AND NOT-EPHB3	Esophagus	0.6667	1	0.5
SLC52A2 AND NOT-PLXNA4	Esophagus	0.8333	1	0.7143
SLC52A2 AND NOT-PTGDR2	Esophagus	0.8333	1	0.7143
SLC52A2 AND NOT-COL25A1	Esophagus	0.6667	1	0.5
SLC52A2 AND NOT-EVC	Esophagus	0.6667	1	0.5
SLC52A2 AND NOT-ABCB8	Esophagus	0.7826	1	0.6429
SLC52A2 AND NOT-GPR174	Esophagus	0.6667	1	0.5
SLC52A2 AND NOT-PARM1	Esophagus	0.8333	1	0.7143
SLC52A2 AND NOT-MC1R	Esophagus	0.7273	1	0.5714

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
SLC52A2 AND NOT-NCAM1	Esophagus	0.6667	1	0.5
SLC52A2 AND NOT-SSTR5	Esophagus	0.6	1	0.4286
EPCAM AND NOT-SLC4A4	Esophagus	0.64	0.7273	0.5714
SLC6A20 AND NOT-CD160	Esophagus	0.6957	0.8889	0.5714
SLC52A2 AND NOT-EPHA3	Esophagus	0.6	1	0.4286
SLC52A2 AND NOT-CD79B	Esophagus	0.7826	1	0.6429
SLC52A2 AND NOT-ULBP3	Esophagus	0.6667	1	0.5
SLC52A2 AND NOT-CD79A	Esophagus	0.7273	1	0.5714
MARVELD2 AND NOT-ENPP3	Esophagus	0.6087	0.7778	0.5
SLC52A2 AND NOT-TNFRSF17	Esophagus	0.7826	1	0.6429
SLC52A2 AND NOT-SSTR3	Esophagus	0.6	1	0.4286
MUC13 AND NOT-SLC4A4	Esophagus	0.6957	0.8889	0.5714
SLC52A2 AND NOT-CD180	Esophagus	0.7826	1	0.6429
SLC52A2 AND NOT-SEMA5B	Esophagus	0.7826	1	0.6429
SLC6A20 AND NOT-ENPP3	Esophagus	0.6667	1	0.5
TM4SF20 AND NOT-ENPP3	Esophagus	0.6667	1	0.5
SLC52A2 AND NOT-ITGB3	Esophagus	0.7826	1	0.6429
SLC52A2 AND NOT-BCAN	Esophagus	0.8333	1	0.7143
SLC52A2 AND NOT-CXCR5	Esophagus	0.7826	1	0.6429
SLC52A2 AND NOT-DLL3	Esophagus	0.7826	1	0.6429
SLC52A2 AND NOT-ABCB5	Esophagus	0.6667	1	0.5
SLC52A2 AND NOT-CD33	Esophagus	0.8333	1	0.7143
SLC52A2 AND NOT-ERBB2	Esophagus	0.6	1	0.4286
CBX3 AND NOT-ADCY7	Esophagus	0.6667	0.8	0.5714
SLC52A2 AND NOT-CLDN23	Esophagus	0.6957	0.8889	0.5714
CBX3 AND NOT-CD69	Esophagus	0.64	0.7273	0.5714
SLC52A2 AND NOT-P2RX5	Esophagus	0.7273	1	0.5714
SLC52A2 AND NOT-DNAJB8	Esophagus	0.8333	1	0.7143
TM4SF20 AND NOT-SST	Esophagus	0.6667	1	0.5
BCAP31 AND NOT-IL11RA	Esophagus	0.7273	1	0.5714
SLC52A2 AND NOT-CLDN9	Esophagus	0.7826	1	0.6429
ST14 AND NOT-ENPP3	Esophagus	0.6667	1	0.5
PTPRZ1 AND NOT-EPCAM	Glioblastoma	0.875	0.9333	0.8235
PTPRZ1 AND NOT-RNF43	Glioblastoma	0.9091	0.9375	0.8824
PTPRZ1 AND NOT-CLDN8	Glioblastoma	0.9091	0.9375	0.8824
PTPRZ1 AND NOT-LGR5	Glioblastoma	0.875	0.9333	0.8235
PTPRZ1 AND NOT-TPBG	Glioblastoma	0.875	0.9333	0.8235
PTPRZ1 AND NOT-SSTR1	Glioblastoma	0.9091	0.9375	0.8824
PTPRZ1 AND NOT-TRPM4	Glioblastoma	0.9091	0.9375	0.8824
PTPRZ1 AND NOT-MST1R	Glioblastoma	0.9091	0.9375	0.8824
PTPRZ1 AND NOT-CD52	Glioblastoma	0.9091	0.9375	0.8824
PTPRZ1 AND NOT-IL20RA	Glioblastoma	0.9091	0.9375	0.8824
PTPRZ1 AND NOT-GPNMB	Glioblastoma	0.8	0.9231	0.7059
PTPRZ1 AND NOT-IL11RA	Glioblastoma	0.8	0.9231	0.7059
PTPRZ1 AND NOT-PROM1	Glioblastoma	0.9091	0.9375	0.8824
NRCAM AND NOT-B4GALNT1	Glioblastoma	0.7317	0.625	0.8824
PTPRZ1 AND NOT-STEAP2	Glioblastoma	0.7407	1	0.5882
AQP4 AND NOT-ERBB3	Glioblastoma	0.6897	0.8333	0.5882
NLGN4X AND NOT-L1CAM	Glioblastoma	0.6875	0.7333	0.6471
PTPRZ1 AND NOT-L1CAM	Glioblastoma	0.8276	1	0.7059
PTPRZ1 AND NOT-MET	Glioblastoma	0.8	0.9231	0.7059
JAM2 AND NOT-CD34	Glioblastoma	0.8276	1	0.7059
NRCAM AND NOT-L1CAM	Glioblastoma	0.7586	0.9167	0.6471
PON2 AND NOT-SDC1	Glioblastoma	0.7111	0.5714	0.9412
JAM2 AND NOT-EPCAM	Glioblastoma	0.8	0.9231	0.7059
JAM2 AND NOT-CLDN1	Glioblastoma	0.7586	0.9167	0.6471
PTPRZ1 AND NOT-ERBB3	Glioblastoma	0.7857	1	0.6471
ITGB8 AND NOT-ERBB3	Glioblastoma	0.64	1	0.4706
PTPRZ1 AND NOT-ERBB4	Glioblastoma	0.64	1	0.4706
PTPRZ1 AND NOT-EPCAM	Glioma	0.9855	0.9855	0.9855
PTPRZ1 AND NOT-RAET1E	Glioma	0.9853	1	0.971
PTPRZ1 AND NOT-CLDN8	Glioma	0.9781	0.9853	0.971
PTPRZ1 AND NOT-RNF43	Glioma	0.9313	0.9839	0.8841
PTPRZ1 AND NOT-MST1R	Glioma	0.9855	0.9855	0.9855
PTPRZ1 AND NOT-PMEL	Glioma	0.9706	0.9851	0.9565
PTPRZ1 AND NOT-ITGB6	Glioma	0.9855	0.9855	0.9855
PTPRZ1 AND NOT-CLDN6	Glioma	0.9781	0.9853	0.971
NRCAM AND NOT-EPCAM	Glioma	0.9242	0.9683	0.8841
PTPRZ1 AND NOT-GPA33	Glioma	0.9855	0.9855	0.9855
GOLM1 AND NOT-EPCAM	Glioma	0.9624	1	0.9275
NLGN1 AND NOT-EPCAM	Glioma	0.9385	1	0.8841
PTPRZ1 AND NOT-SLAMF7	Glioma	0.9853	1	0.971
PTPRZ1 AND NOT-CLDN1	Glioma	0.9781	0.9853	0.971
NRCAM AND NOT-CLDN11	Glioma	0.9173	0.9531	0.8841
NRCAM AND NOT-CD22	Glioma	0.9173	0.9531	0.8841
PTPRZ1 AND NOT-CLDN9	Glioma	0.9781	0.9853	0.971
PTPRZ1 AND NOT-STEAP2	Glioma	0.9855	0.9855	0.9855

FIG. 12 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
SLC52A2 AND NOT-CLEC2D	Esophagus	0.8333	1	0.7143
SLC52A2 AND NOT-C1QTNF1	Esophagus	0.7273	1	0.5714
SLC52A2 AND NOT-FLT3LG	Esophagus	0.7273	1	0.5714
SLC52A2 AND NOT-CDH24	Esophagus	0.7826	1	0.6429
SLC52A2 AND NOT-SLAMF1	Esophagus	0.8333	1	0.7143
SLC52A2 AND NOT-CHRNA3	Esophagus	0.6667	1	0.5
SLC52A2 AND NOT-ZFYVE27	Esophagus	0.7273	1	0.5714
SLC52A2 AND NOT-TMEM119	Esophagus	0.6667	1	0.5
SLC52A2 AND NOT-KIT	Esophagus	0.7826	1	0.6429
SLC39A4 AND NOT-ABHD6	Esophagus	0.72	0.8182	0.6429
GPRC5A AND NOT-SEMA6D	Esophagus	0.7826	1	0.6429
SLC52A2 AND NOT-CMKLR1	Esophagus	0.7826	1	0.6429
SLC52A2 AND NOT-TRPC5	Esophagus	0.7273	1	0.5714
SLC52A2 AND NOT-COLQ	Esophagus	0.8333	1	0.7143
SLC52A2 AND NOT-JAG1	Esophagus	0.6364	0.875	0.5
SLC52A2 AND NOT-ZP1	Esophagus	0.7826	1	0.6429
SLC52A2 AND NOT-SLC46A2	Esophagus	0.8333	1	0.7143
SLC52A2 AND NOT-SLC13A4	Esophagus	0.8333	1	0.7143
SLC52A2 AND NOT-SCNN1D	Esophagus	0.7826	1	0.6429
SLC52A2 AND NOT-CHRM1	Esophagus	0.7826	1	0.6429
SLC52A2 AND NOT-LPAR5	Esophagus	0.8	0.9091	0.7143
SLC52A2 AND NOT-SLC44A1	Esophagus	0.6667	0.8	0.5714
SLC52A2 AND NOT-GRIA3	Esophagus	0.8333	1	0.7143
SLC52A2 AND NOT-KCNK7	Esophagus	0.6667	1	0.5
SLC52A2 AND NOT-IFITM10	Esophagus	0.7826	1	0.6429
SLC52A2 AND NOT-CD1E	Esophagus	0.8333	1	0.7143
SLC52A2 AND NOT-CACNA1G	Esophagus	0.6667	1	0.5
SLC52A2 AND NOT-CLEC12B	Esophagus	0.7826	1	0.6429
SLC52A2 AND NOT-NRSN2	Esophagus	0.6667	1	0.5
SLC52A2 AND NOT-ABCG2	Esophagus	0.6667	1	0.5
SLC52A2 AND NOT-SLC6A2	Esophagus	0.7826	1	0.6429
SLC52A2 AND NOT-TACR1	Esophagus	0.7826	1	0.6429
SLC52A2 AND NOT-CD300LG	Esophagus	0.8333	1	0.7143
SLC52A2 AND NOT-JPH2	Esophagus	0.7273	1	0.5714
PTPRZ1 AND NOT-LRP11	Glioblastoma	0.875	0.9333	0.8235
PTPRZ1 AND NOT-SLC2A12	Glioblastoma	0.9091	0.9375	0.8824
PTPRZ1 AND NOT-MTUS1	Glioblastoma	0.8667	1	0.7647
PTPRZ1 AND NOT-STX8	Glioblastoma	0.875	0.9333	0.8235
PTPRZ1 AND NOT-CNST	Glioblastoma	0.875	0.9333	0.8235
PTPRZ1 AND NOT-LRFN5	Glioblastoma	0.875	0.9333	0.8235
PTPRZ1 AND NOT-VIPR1	Glioblastoma	0.9091	0.9375	0.8824
PTPRZ1 AND NOT-NRXN3	Glioblastoma	0.875	0.9333	0.8235
PTPRZ1 AND NOT-SYT1	Glioblastoma	0.875	0.9333	0.8235
PTPRZ1 AND NOT-MEP1A	Glioblastoma	0.8387	0.9286	0.7647
PTPRZ1 AND NOT-ATP1A1	Glioblastoma	0.8387	0.9286	0.7647
NRCAM AND NOT-LRFN5	Glioblastoma	0.8387	0.9286	0.7647
PTPRZ1 AND NOT-KIT	Glioblastoma	0.8387	0.9286	0.7647
PTPRZ1 AND NOT-EFNA5	Glioblastoma	0.8387	0.9286	0.7647
PTPRZ1 AND NOT-NPTN	Glioblastoma	0.8387	0.9286	0.7647
PTPRZ1 AND NOT-ENTPD3	Glioblastoma	0.8387	0.9286	0.7647
NRCAM AND NOT-NRXN3	Glioblastoma	0.8485	0.875	0.8235
PTPRZ1 AND NOT-PARM1	Glioblastoma	0.875	0.9333	0.8235
PTPRZ1 AND NOT-CDH12	Glioblastoma	0.875	0.9333	0.8235
PTPRZ1 AND NOT-ACVR2A	Glioblastoma	0.8387	0.9286	0.7647
PTPRZ1 AND NOT-KL	Glioblastoma	0.8387	0.9286	0.7647
PTPRZ1 AND NOT-ANKH	Glioblastoma	0.875	0.9333	0.8235
PTPRZ1 AND NOT-ATP2C1	Glioblastoma	0.875	0.9333	0.8235
PTPRZ1 AND NOT-GJB6	Glioblastoma	0.8276	1	0.7059
PTPRZ1 AND NOT-FLRT2	Glioblastoma	0.8276	1	0.7059
PTPRZ1 AND NOT-RNF144A	Glioblastoma	0.8387	0.9286	0.7647
PTPRZ1 AND NOT-SLC6A1	Glioblastoma	0.9032	1	0.8235
PTPRZ1 AND NOT-TRHDE	Glioblastoma	0.875	0.9333	0.8235
PTPRZ1 AND NOT-SYT13	Glioblastoma	0.875	0.9333	0.8235
PTPRZ1 AND NOT-EPHB6	Glioblastoma	0.875	0.9333	0.8235
PTPRZ1 AND NOT-SV2B	Glioblastoma	0.875	0.9333	0.8235
PTPRZ1 AND NOT-VAPA	Glioblastoma	0.875	0.9333	0.8235
NRCAM AND NOT-SYT1	Glioblastoma	0.8667	1	0.7647
PTPRZ1 AND NOT-ANO10	Glioblastoma	0.9091	0.9375	0.8824
NRCAM AND NOT-SLC2A13	Glioblastoma	0.9091	0.9375	0.8824
PTPRZ1 AND NOT-EFNA5	Glioma	0.9781	0.9853	0.971
PTPRZ1 AND NOT-CD3G	Glioma	0.9778	1	0.9565
PTPRZ1 AND NOT-KIAA1324	Glioma	0.963	0.9848	0.942
PTPRZ1 AND NOT-FAT2	Glioma	0.9855	0.9855	0.9855
PTPRZ1 AND NOT-C1orf210	Glioma	0.9927	1	0.9855
PTPRZ1 AND NOT-LRP11	Glioma	0.9855	0.9855	0.9855
PTPRZ1 AND NOT-FUT1	Glioma	0.9855	0.9855	0.9855
PTPRZ1 AND NOT-DSC3	Glioma	0.9853	1	0.971

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
PTPRZ1 AND NOT-CLDN23	Glioma	0.9855	0.9855	0.9855
PTPRZ1 AND NOT-SSTR3	Glioma	0.9855	0.9855	0.9855
PTPRZ1 AND NOT-CXCR5	Glioma	0.9624	1	0.9275
PTPRZ1 AND NOT-CLDN18	Glioma	0.9781	0.9853	0.971
PTPRZ1 AND NOT-IL3RA	Glioma	0.9855	0.9855	0.9855
PTPRZ1 AND NOT-TNFRSF10A	Glioma	0.963	0.9848	0.942
PTPRZ1 AND NOT-SSTR5	Glioma	0.9855	0.9855	0.9855
PTPRZ1 AND NOT-SDC1	Glioma	0.9855	0.9855	0.9855
NRCAM AND NOT-STEAP2	Glioma	0.9173	0.9531	0.8841
NRCAM AND NOT-ENPP3	Glioma	0.9173	0.9531	0.8841
PTPRZ1 AND NOT-ENPP3	Glioma	0.9855	0.9855	0.9855
PTPRZ1 AND NOT-MUC4	Glioma	0.9855	0.9855	0.9855
PTPRZ1 AND NOT-VTCN1	Glioma	0.9855	0.9855	0.9855
PTPRZ1 AND NOT-MSLN	Glioma	0.9706	0.9851	0.9565
PTPRZ1 AND NOT-CLDN7	Glioma	0.963	0.9848	0.942
NRCAM AND NOT-CR2	Glioma	0.9173	0.9531	0.8841
NRCAM AND NOT-CLDN8	Glioma	0.9173	0.9531	0.8841
NRCAM AND NOT-ITGB6	Glioma	0.9173	0.9531	0.8841
PTPRZ1 AND NOT-CD79B	Glioma	0.9706	0.9851	0.9565
PTPRZ1 AND NOT-GUCY2C	Glioma	0.9855	0.9855	0.9855
PTPRZ1 AND NOT-TNFRSF8	Glioma	0.9781	0.9853	0.971
NLGN1 AND NOT-CLDN6	Glioma	0.9302	1	0.8696
NLGN1 AND NOT-GPA33	Glioma	0.9385	1	0.8841
PTPRZ1 AND NOT-TNFRSF17	Glioma	0.9781	0.9853	0.971
PTPRZ1 AND NOT-FCRL5	Glioma	0.963	0.9848	0.942
PTPRZ1 AND NOT-CR2	Glioma	0.9781	0.9853	0.971
NRCAM AND NOT-CLDN1	Glioma	0.9173	0.9531	0.8841
NRCAM AND NOT-CLDN18	Glioma	0.9173	0.9531	0.8841
NRCAM AND NOT-CLDN9	Glioma	0.9173	0.9531	0.8841
NRCAM AND NOT-CXCR5	Glioma	0.9173	0.9531	0.8841
NRCAM AND NOT-CLDN6	Glioma	0.9173	0.9531	0.8841
NRCAM AND NOT-PMEL	Glioma	0.9173	0.9531	0.8841
NRCAM AND NOT-BMPR1B	Glioma	0.9173	0.9531	0.8841
NRCAM AND NOT-FCRL2	Glioma	0.9173	0.9531	0.8841
NRCAM AND NOT-MST1R	Glioma	0.9173	0.9531	0.8841
NRCAM AND NOT-CLDN2	Glioma	0.9173	0.9531	0.8841
NRCAM AND NOT-SSTR1	Glioma	0.9091	0.9524	0.8696
NRCAM AND NOT-ERBB4	Glioma	0.9091	0.9524	0.8696
NRCAM AND NOT-FOLH1	Glioma	0.9091	0.9524	0.8696
PTPRZ1 AND NOT-MUC13	Glioma	0.9781	0.9853	0.971
PTPRZ1 AND NOT-FCRL2	Glioma	0.9778	1	0.9565
NRCAM AND NOT-NCAM1	Glioma	0.9091	0.9524	0.8696
ROR2 AND NOT-IL20RA	Leiomyosarcoma	0.7619	1	0.6154
FGFR1 AND NOT-CLDN11	Leiomyosarcoma	0.7826	0.9	0.6923
DDR2 AND NOT-EPCAM	Leiomyosarcoma	0.8	0.8333	0.7692
FAP AND NOT-SGMS1	Leiomyosarcoma	0.7	1	0.5385
FGFR1 AND NOT-EPCAM	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-IL20RA	Leiomyosarcoma	0.7619	1	0.6154
ROR2 AND NOT-RNF43	Leiomyosarcoma	0.7619	1	0.6154
FGFR1 AND NOT-TRPM4	Leiomyosarcoma	0.7826	0.9	0.6923
FGFR1 AND NOT-ENPP3	Leiomyosarcoma	0.8182	1	0.6923
DDR2 AND NOT-EDNRB	Leiomyosarcoma	0.6667	0.7273	0.6154
ROR2 AND NOT-CLDN8	Leiomyosarcoma	0.6923	0.6923	0.6923
FGFR1 AND NOT-CLDN7	Leiomyosarcoma	0.7619	1	0.6154
CD276 AND NOT-LSR	Leiomyosarcoma	0.75	0.8182	0.6923
OSMR AND NOT-ITGB6	Leiomyosarcoma	0.7273	0.8889	0.6154
FGFR1 AND NOT-RNF43	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-ERBB3	Leiomyosarcoma	0.7826	0.9	0.6923
TGFB3 AND NOT-ERBB3	Leiomyosarcoma	0.6667	0.875	0.5385
OSMR AND NOT-ERBB3	Leiomyosarcoma	0.7	1	0.5385
FGFR1 AND NOT-SLC39A6	Leiomyosarcoma	0.6667	0.875	0.5385
GPR161 AND NOT-ERBB3	Leiomyosarcoma	0.8182	1	0.6923
CD276 AND NOT-CRB3	Leiomyosarcoma	0.7273	0.8889	0.6154
FGFR1 AND NOT-STEAP1	Leiomyosarcoma	0.7826	0.9	0.6923
FGFR1 AND NOT-PMEL	Leiomyosarcoma	0.7826	0.9	0.6923
FGFR1 AND NOT-SDC1	Leiomyosarcoma	0.7273	0.8889	0.6154
FGFR1 AND NOT-LGR5	Leiomyosarcoma	0.7273	0.8889	0.6154
FGFR1 AND NOT-CD34	Leiomyosarcoma	0.7826	0.9	0.6923
FGFR1 AND NOT-CLDN2	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-CLDN8	Leiomyosarcoma	0.7826	0.9	0.6923
FGFR1 AND NOT-RAET1E	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-CLDN1	Leiomyosarcoma	0.7826	0.9	0.6923
FGFR1 AND NOT-GUCY2C	Leiomyosarcoma	0.7619	1	0.6154
ROR2 AND NOT-ENPP3	Leiomyosarcoma	0.7	1	0.5385
FGFR1 AND NOT-SLC7A5	Leiomyosarcoma	0.7273	0.8889	0.6154
FGFR1 AND NOT-MUC4	Leiomyosarcoma	0.7619	1	0.6154
DDR2 AND NOT-IL20RA	Leiomyosarcoma	0.6364	0.7778	0.5385

FIG. 12 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
PTPRZ1 AND NOT-SYT2	Glioma	0.9855	0.9855	0.9855
PTPRZ1 AND NOT-ADRA2C	Glioma	0.9552	0.9846	0.9275
PTPRZ1 AND NOT-SLC26A9	Glioma	0.9781	0.9853	0.971
PTPRZ1 AND NOT-CLCA2	Glioma	0.9855	0.9855	0.9855
PTPRZ1 AND NOT-GPR68	Glioma	0.9855	0.9855	0.9855
PTPRZ1 AND NOT-SLC5A1	Glioma	0.9927	1	0.9855
PTPRZ1 AND NOT-PTGER3	Glioma	0.9853	1	0.971
PTPRZ1 AND NOT-ACVR1B	Glioma	0.9781	0.9853	0.971
PTPRZ1 AND NOT-GRM2	Glioma	0.9855	0.9855	0.9855
PTPRZ1 AND NOT-TNFSF9	Glioma	0.9853	1	0.971
PTPRZ1 AND NOT-SLC46A2	Glioma	0.9781	0.9853	0.971
PTPRZ1 AND NOT-KRT5	Glioma	0.9706	0.9851	0.9565
PTPRZ1 AND NOT-SLC28A3	Glioma	0.9855	0.9855	0.9855
PTPRZ1 AND NOT-KCNB2	Glioma	0.9855	0.9855	0.9855
PTPRZ1 AND NOT-EDAR	Glioma	0.963	0.9848	0.942
PTPRZ1 AND NOT-C15orf27	Glioma	0.9853	1	0.971
PTPRZ1 AND NOT-SLC18A2	Glioma	0.9853	1	0.971
PTPRZ1 AND NOT-SCNN1A	Glioma	0.9781	0.9853	0.971
PTPRZ1 AND NOT-HAS1	Glioma	0.9552	0.9846	0.9275
PTPRZ1 AND NOT-ATP2A3	Glioma	0.9706	0.9851	0.9565
PTPRZ1 AND NOT-ESYT3	Glioma	0.9855	0.9855	0.9855
PTPRZ1 AND NOT-PVRL4	Glioma	0.9927	1	0.9855
PTPRZ1 AND NOT-GJB5	Glioma	0.9855	0.9855	0.9855
PTPRZ1 AND NOT-SYT8	Glioma	0.9855	0.9855	0.9855
PTPRZ1 AND NOT-RHBDL2	Glioma	0.9701	1	0.942
PTPRZ1 AND NOT-MS4A2	Glioma	0.9927	1	0.9855
PTPRZ1 AND NOT-COL17A1	Glioma	0.9855	0.9855	0.9855
PTPRZ1 AND NOT-CD5	Glioma	0.9853	1	0.971
PTPRZ1 AND NOT-S1PR5	Glioma	0.9781	0.9853	0.971
PTPRZ1 AND NOT-MUC15	Glioma	0.9781	0.9853	0.971
PTPRZ1 AND NOT-CD1E	Glioma	0.9855	0.9855	0.9855
PTPRZ1 AND NOT-CFTR	Glioma	0.9855	0.9855	0.9855
PTPRZ1 AND NOT-LPAR3	Glioma	0.9474	0.9844	0.913
PTPRZ1 AND NOT-DUOXA1	Glioma	0.9701	1	0.942
PTPRZ1 AND NOT-FLT3	Glioma	0.9853	1	0.971
PTPRZ1 AND NOT-CORIN	Glioma	0.9781	0.9853	0.971
PTPRZ1 AND NOT-KIT	Glioma	0.963	0.9848	0.942
PTPRZ1 AND NOT-MARVELD2	Glioma	0.9855	0.9855	0.9855
NRCAM AND NOT-FRRS1L	Glioma	0.9385	1	0.8841
PTPRZ1 AND NOT-UPK1A	Glioma	0.9781	0.9853	0.971
PTPRZ1 AND NOT-VIPR1	Glioma	0.9781	0.9853	0.971
PTPRZ1 AND NOT-PRPH2	Glioma	0.9781	0.9853	0.971
PTPRZ1 AND NOT-GJB6	Glioma	0.9853	1	0.971
PTPRZ1 AND NOT-LY6K	Glioma	0.9706	0.9851	0.9565
PTPRZ1 AND NOT-SYT1	Glioma	0.9855	0.9855	0.9855
PTPRZ1 AND NOT-SIGLEC6	Glioma	0.9927	1	0.9855
PTPRZ1 AND NOT-GPR143	Glioma	0.9781	0.9853	0.971
NRCAM AND NOT-KCNB1	Glioma	0.9313	0.9839	0.8841
PTPRZ1 AND NOT-GPR87	Glioma	0.9855	0.9855	0.9855
PTPRZ1 AND NOT-KCNJ12	Glioma	0.9781	0.9853	0.971
PTPRZ1 AND NOT-CD40LG	Glioma	0.9781	0.9853	0.971
PTPRZ1 AND NOT-SLITRK6	Glioma	0.9706	0.9851	0.9565
PTPRZ1 AND NOT-ANO9	Glioma	0.9781	0.9853	0.971
PTPRZ1 AND NOT-LY6D	Glioma	0.9855	0.9855	0.9855
DDR2 AND NOT-PEAR1	Leiomyosarcoma	0.8182	1	0.6923
DDR2 AND NOT-F11R	Leiomyosarcoma	0.7692	0.7692	0.7692
DDR2 AND NOT-TM7SF2	Leiomyosarcoma	0.8696	1	0.7692
FGFR1 AND NOT-SGMS1	Leiomyosarcoma	0.7826	0.9	0.6923
VMP1 AND NOT-F11R	Leiomyosarcoma	0.8333	0.9091	0.7692
DDR2 AND NOT-CHL1	Leiomyosarcoma	0.7273	0.8889	0.6154
DDR2 AND NOT-EMCN	Leiomyosarcoma	0.8333	0.9091	0.7692
ROR2 AND NOT-TMEM88	Leiomyosarcoma	0.7826	0.9	0.6923
FGFR1 AND NOT-TCTN3	Leiomyosarcoma	0.7826	0.9	0.6923
DDR2 AND NOT-LIFR	Leiomyosarcoma	0.6957	0.8	0.6154
SLC4A7 AND NOT-SLC12A6	Leiomyosarcoma	0.6957	0.8	0.6154
DDR2 AND NOT-TSPAN8	Leiomyosarcoma	0.7273	0.8889	0.6154
DDR2 AND NOT-ADRA2A	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-ADRA2A	Leiomyosarcoma	0.7826	0.9	0.6923
FGFR1 AND NOT-RHBDL2	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-XG	Leiomyosarcoma	0.7826	0.9	0.6923
FGFR1 AND NOT-TSPAN31	Leiomyosarcoma	0.7619	1	0.6154
FGFR1 AND NOT-TMPRSS6	Leiomyosarcoma	0.7619	1	0.6154
FGFR1 AND NOT-TM7SF2	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-VAMP3	Leiomyosarcoma	0.7826	0.9	0.6923
FGFR1 AND NOT-EREG	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-PRR7	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-MFSD5	Leiomyosarcoma	0.7826	0.9	0.6923

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
AXL AND NOT-GPRC5C	Leiomyosarcoma	0.6364	0.7778	0.5385
FGFR1 AND NOT-MST1R	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-ERBB2	Leiomyosarcoma	0.7619	1	0.6154
OSMR AND NOT-CLDN1	Leiomyosarcoma	0.6364	0.7778	0.5385
FGFR1 AND NOT-VTCN1	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-CLDN5	Leiomyosarcoma	0.7826	0.9	0.6923
FGFR1 AND NOT-CLDN23	Leiomyosarcoma	0.7273	0.8889	0.6154
FGFR1 AND NOT-ITGB6	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-CD79B	Leiomyosarcoma	0.8182	1	0.6923
FAP AND NOT-NLGN4X	Leiomyosarcoma	0.6316	1	0.4615
AXL AND NOT-SLCO4A1	Leiomyosarcoma	0.7	1	0.5385
FGFR1 AND NOT-SSTR3	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-SSTR5	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-SLC34A2	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-MET	Leiomyosarcoma	0.7826	0.9	0.6923
OSMR AND NOT-FOLR1	Leiomyosarcoma	0.6316	1	0.4615
ROR2 AND NOT-ERBB3	Leiomyosarcoma	0.7619	1	0.6154
FGFR1 AND NOT-EDNRB	Leiomyosarcoma	0.7	1	0.5385
GPR176 AND NOT-EPCAM	Leiomyosarcoma	0.6364	0.7778	0.5385
DCHS1 AND NOT-ERBB3	Leiomyosarcoma	0.7	1	0.5385
OSMR AND NOT-RNF43	Leiomyosarcoma	0.6667	0.875	0.5385
DCHS1 AND NOT-IL20RA	Leiomyosarcoma	0.6667	0.875	0.5385
FGFR1 AND NOT-CLDN6	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-FCRL1	Leiomyosarcoma	0.7619	1	0.6154
FGFR1 AND NOT-P2RX5	Leiomyosarcoma	0.8182	1	0.6923
ROR2 AND NOT-ERBB4	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-TNFRSF13C	Leiomyosarcoma	0.7	1	0.5385
FGFR1 AND NOT-CLDN18	Leiomyosarcoma	0.7619	1	0.6154
FGFR1 AND NOT-CD22	Leiomyosarcoma	0.8182	1	0.6923
CD276 AND NOT-FXYD3	Leiomyosarcoma	0.7273	0.8889	0.6154
FAP AND NOT-BCAM	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-L1CAM	Leiomyosarcoma	0.8182	1	0.6923
AXL AND NOT-SCN7A	Leiomyosarcoma	0.6364	0.7778	0.5385
FGFR1 AND NOT-TNFRSF17	Leiomyosarcoma	0.7826	0.9	0.6923
FGFR1 AND NOT-ENG	Leiomyosarcoma	0.7826	0.9	0.6923
FAP AND NOT-FCER1A	Leiomyosarcoma	0.7	1	0.5385
FGFR1 AND NOT-IL3RA	Leiomyosarcoma	0.7826	0.9	0.6923
FGFR1 AND NOT-CD180	Leiomyosarcoma	0.7826	0.9	0.6923
FGFR1 AND NOT-FCRL5	Leiomyosarcoma	0.7826	0.9	0.6923
FGFR1 AND NOT-ULBP1	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-FOLR1	Leiomyosarcoma	0.7826	0.9	0.6923
FGFR1 AND NOT-FCRL2	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-MUC1	Leiomyosarcoma	0.7826	0.9	0.6923
FGFR1 AND NOT-CR2	Leiomyosarcoma	0.7826	0.9	0.6923
FAP AND NOT-LIFR	Leiomyosarcoma	0.7	1	0.5385
FGFR1 AND NOT-TNFRSF8	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-PROM1	Leiomyosarcoma	0.7826	0.9	0.6923
FGFR1 AND NOT-MS4A1	Leiomyosarcoma	0.7826	0.9	0.6923
FLT3 AND NOT-EDNRB	AML	0.9545	0.9914	0.9203
FLT3 AND NOT-SLAMF7	AML	0.9545	0.9914	0.9203
FLT3 AND NOT-FCRL1	AML	0.9506	0.983	0.9203
FLT3 AND NOT-CD72	AML	0.9506	0.983	0.9203
FLT3 AND NOT-AXL	AML	0.9482	0.9871	0.9124
FLT3 AND NOT-CLDN5	AML	0.9467	0.9747	0.9203
FLT3 AND NOT-CLDN12	AML	0.9465	0.9787	0.9163
FLT3 AND NOT-RNF43	AML	0.9448	0.9706	0.9203
FLT3 AND NOT-MST1R	AML	0.9429	0.9665	0.9203
FLT3 AND NOT-TNFRSF10A	AML	0.9429	0.9665	0.9203
FLT3 AND NOT-VTCN1	AML	0.9429	0.9665	0.9203
FLT3 AND NOT-IL11RA	AML	0.9429	0.9665	0.9203
FLT3 AND NOT-CD79B	AML	0.9397	0.9826	0.9004
FLT3 AND NOT-TPBG	AML	0.9385	0.9662	0.9124
FLT3 AND NOT-MUC1	AML	0.9385	0.9662	0.9124
FLT3 AND NOT-GPNMB	AML	0.9328	0.9867	0.8845
FLT3 AND NOT-ITGAV	AML	0.9325	0.991	0.8805
FLT3 AND NOT-VCAM1	AML	0.9056	0.9814	0.8406
CD37 AND NOT-HEG1	AML	0.9041	0.8885	0.9203
CD37 AND NOT-GPR171	AML	0.8919	0.8798	0.9044
CD37 AND NOT-IL7R	AML	0.8893	0.8535	0.9283
CD37 AND NOT-AMIGO2	AML	0.8872	0.8529	0.9243
P2RX1 AND NOT-IL11RA	AML	0.8971	0.875	0.9203
CD37 AND NOT-CD27	AML	0.8867	0.8697	0.9044
CD37 AND NOT-BTN3A2	AML	0.8845	0.8692	0.9004
CD37 AND NOT-CD2	AML	0.8842	0.8577	0.9124
CD37 AND NOT-P2RY10	AML	0.8975	0.8722	0.9243
CD37 AND NOT-SIT1	AML	0.8885	0.8453	0.9363
CD37 AND NOT-GPR18	AML	0.8736	0.8413	0.9084

FIG. 12 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
DDR2 AND NOT-F3	Leiomyosarcoma	0.7273	0.8889	0.6154
FGFR1 AND NOT-ERMAP	Leiomyosarcoma	0.7273	0.8889	0.6154
FGFR1 AND NOT-GCGR	Leiomyosarcoma	0.7619	1	0.6154
ROR2 AND NOT-ABCA8	Leiomyosarcoma	0.75	0.8182	0.6923
ROR2 AND NOT-DLL1	Leiomyosarcoma	0.7826	0.9	0.6923
FGFR1 AND NOT-ACVR1B	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-PAQR5	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-SCNN1B	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-F11R	Leiomyosarcoma	0.7826	0.9	0.6923
FGFR1 AND NOT-PLA2R1	Leiomyosarcoma	0.7273	0.8889	0.6154
FGFR1 AND NOT-FXYD3	Leiomyosarcoma	0.7619	1	0.6154
FGFR1 AND NOT-TACR1	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-CDH19	Leiomyosarcoma	0.8182	1	0.6923
ROR2 AND NOT-MPZ	Leiomyosarcoma	0.7826	0.9	0.6923
OSMR AND NOT-F11R	Leiomyosarcoma	0.7273	0.8889	0.6154
FGFR1 AND NOT-SMAGP	Leiomyosarcoma	0.7826	0.9	0.6923
FGFR1 AND NOT-KCNK7	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-LEPR	Leiomyosarcoma	0.7826	0.9	0.6923
FGFR1 AND NOT-CD300LB	Leiomyosarcoma	0.7619	1	0.6154
FGFR1 AND NOT-LIM2	Leiomyosarcoma	0.8182	1	0.6923
DDR2 AND NOT-ABCA8	Leiomyosarcoma	0.7273	0.8889	0.6154
CNTNAP1 AND NOT-NLGN4X	Leiomyosarcoma	0.6957	0.8	0.6154
DDR2 AND NOT-PLP1	Leiomyosarcoma	0.7	1	0.5385
FGFR1 AND NOT-AGTR1	Leiomyosarcoma	0.7273	0.8889	0.6154
FGFR1 AND NOT-SLC12A2	Leiomyosarcoma	0.6667	0.875	0.5385
FGFR1 AND NOT-SLC15A1	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-ICOSLG	Leiomyosarcoma	0.7619	1	0.6154
FGFR1 AND NOT-ATP10B	Leiomyosarcoma	0.7619	1	0.6154
FGFR1 AND NOT-P2RY2	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-MTUS1	Leiomyosarcoma	0.8182	1	0.6923
DDR2 AND NOT-TSPAN7	Leiomyosarcoma	0.6667	0.875	0.5385
FGFR1 AND NOT-LPAR3	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-ASIC5	Leiomyosarcoma	0.7619	1	0.6154
FGFR1 AND NOT-CNGA1	Leiomyosarcoma	0.7	1	0.5385
FGFR1 AND NOT-STAB2	Leiomyosarcoma	0.7619	1	0.6154
FGFR1 AND NOT-CD300LG	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-COL17A1	Leiomyosarcoma	0.8182	1	0.6923
ROR2 AND NOT-ABCB1	Leiomyosarcoma	0.7826	0.9	0.6923
FGFR1 AND NOT-CXCR1	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-ACPP	Leiomyosarcoma	0.7619	1	0.6154
FGFR1 AND NOT-NLGN3	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-KCNK15	Leiomyosarcoma	0.7619	1	0.6154
FGFR1 AND NOT-SLC44A1	Leiomyosarcoma	0.6667	0.875	0.5385
FGFR1 AND NOT-PROM2	Leiomyosarcoma	0.7619	1	0.6154
FGFR1 AND NOT-ABHD6	Leiomyosarcoma	0.7826	0.9	0.6923
PCDH18 AND NOT-ADRA2A	Leiomyosarcoma	0.6667	0.7273	0.6154
ROR2 AND NOT-CRB3	Leiomyosarcoma	0.7619	1	0.6154
FGFR1 AND NOT-IL18R1	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-IL1RL2	Leiomyosarcoma	0.7619	1	0.6154
FGFR1 AND NOT-ATP13A4	Leiomyosarcoma	0.7619	1	0.6154
FGFR1 AND NOT-LYPD6B	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-GJB3	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-FAM57A	Leiomyosarcoma	0.6667	0.875	0.5385
FGFR1 AND NOT-YIPF4	Leiomyosarcoma	0.7619	1	0.6154
FGFR1 AND NOT-DIO3	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-MMP15	Leiomyosarcoma	0.7619	1	0.6154
FGFR1 AND NOT-KCNJ1	Leiomyosarcoma	0.7619	1	0.6154
FGFR1 AND NOT-UTS2R	Leiomyosarcoma	0.7826	0.9	0.6923
FGFR1 AND NOT-GJD4	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-CRB3	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-TMPRSS5	Leiomyosarcoma	0.7619	1	0.6154
FGFR1 AND NOT-ATP12A	Leiomyosarcoma	0.7619	1	0.6154
ROR2 AND NOT-SCN7A	Leiomyosarcoma	0.7826	0.9	0.6923
FGFR1 AND NOT-NPR3	Leiomyosarcoma	0.7619	1	0.6154
FGFR1 AND NOT-BST1	Leiomyosarcoma	0.6667	0.875	0.5385
FGFR1 AND NOT-SYP	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-AQP5	Leiomyosarcoma	0.7619	1	0.6154
FGFR1 AND NOT-CLEC10A	Leiomyosarcoma	0.7826	0.9	0.6923
FGFR1 AND NOT-LY6D	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-MPZ	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-C19orf26	Leiomyosarcoma	0.7	1	0.5385
FGFR1 AND NOT-FUT1	Leiomyosarcoma	0.7619	1	0.6154
FGFR1 AND NOT-AQP3	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-OR1I1	Leiomyosarcoma	0.7619	1	0.6154
FLT3 AND NOT-HEPH	AML	0.9545	0.9914	0.9203
FLT3 AND NOT-ENPP2	AML	0.9545	0.9914	0.9203
FLT3 AND NOT-LPAR1	AML	0.9545	0.9914	0.9203

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
CD37 AND NOT-SLC44A2	AML	0.8729	0.8333	0.9163
CD37 AND NOT-ADTRP	AML	0.872	0.816	0.9363
TSPAN32 AND NOT-FCRL5	AML	0.9108	0.8537	0.9761
TSPAN32 AND NOT-CR2	AML	0.8982	0.8261	0.9841
CD37 AND NOT-IL2RB	AML	0.8716	0.8517	0.8924
TSPAN32 AND NOT-MS4A1	AML	0.9057	0.8448	0.9761
FLT3 AND NOT-SLC7A5	AML	0.8709	0.966	0.7928
TSPAN32 AND NOT-FCRL1	AML	0.8909	0.8194	0.9761
CD37 AND NOT-CCR6	AML	0.8688	0.8103	0.9363
CD37 AND NOT-FCRL3	AML	0.8688	0.8103	0.9363
CSF3R AND NOT-SLAMF7	AML	0.9006	0.8817	0.9203
ATP8B4 AND NOT-CLDN23	AML	0.8675	0.9355	0.8088
CD37 AND NOT-DPP4	AML	0.8673	0.8169	0.9243
TSPAN32 AND NOT-SLAMF7	AML	0.9017	0.8438	0.9681
CD37 AND NOT-BTLA	AML	0.8667	0.8097	0.9323
CD37 AND NOT-CLEC2D	AML	0.8818	0.8333	0.9363
CSF3R AND NOT-MS4A1	AML	0.9049	0.8826	0.9283
TSPAN32 AND NOT-CD22	AML	0.8978	0.8283	0.9801
TSPAN32 AND NOT-FCRL2	AML	0.8837	0.8019	0.9841
CD37 AND NOT-TIGIT	AML	0.864	0.802	0.9363
CD37 AND NOT-IFITM1	AML	0.8638	0.9269	0.8088
CSF3R AND NOT-FCRL1	AML	0.8927	0.8598	0.9283
CD37 AND NOT-SIRPG	AML	0.8635	0.8041	0.9323
CD37 AND NOT-CD247	AML	0.8632	0.8358	0.8924
CD37 AND NOT-ST3GAL5	AML	0.862	0.8201	0.9084
TSPAN32 AND NOT-IL11RA	AML	0.8665	0.8007	0.9442
CD37 AND NOT-LTB	AML	0.8613	0.9111	0.8167
CD37 AND NOT-PLXDC1	AML	0.8613	0.7946	0.9402
CD37 AND NOT-LY9	AML	0.8613	0.7946	0.9402
CD37 AND NOT-PTGDR	AML	0.8608	0.7966	0.9363
CD37 AND NOT-CD28	AML	0.8812	0.8487	0.9163
CD37 AND NOT-CD40LG	AML	0.8736	0.8188	0.9363
P2RX1 AND NOT-CD79B	AML	0.8663	0.868	0.8645
GYPB AND NOT-EDNRB	AML	0.859	0.9429	0.7888
CSF3R AND NOT-FCRL5	AML	0.9017	0.8731	0.9323
CSF3R AND NOT-CD22	AML	0.8851	0.8393	0.9363
CD37 AND NOT-KLRD1	AML	0.8773	0.8223	0.9402
CD37 AND NOT-CD8B	AML	0.8683	0.8125	0.9323
CD37 AND NOT-CCR7	AML	0.8577	0.8	0.9243
P2RY8 AND NOT-FCRL5	AML	0.9139	0.8622	0.9721
ATP8B4 AND NOT-CLDN12	AML	0.8571	0.9384	0.7888
TSPAN32 AND NOT-TNFRSF17	AML	0.8881	0.8231	0.9641
CD37 AND NOT-ICOS	AML	0.8555	0.8284	0.8845
CD37 AND NOT-TLR10	AML	0.8608	0.7966	0.9363
P2RY8 AND NOT-MS4A1	AML	0.9208	0.8746	0.9721
CD37 AND NOT-EPHA4	AML	0.8545	0.786	0.9363
CD37 AND NOT-STX7	AML	0.8544	0.8333	0.8765
CSF3R AND NOT-CD79B	AML	0.8543	0.8683	0.8406
P2RX1 AND NOT-CLDN5	AML	0.8541	0.7642	0.9681
P2RY8 AND NOT-SLAMF7	AML	0.8864	0.8203	0.9641
P2RY8 AND NOT-CD79B	AML	0.8533	0.8277	0.8805
CD37 AND NOT-CD8A	AML	0.8577	0.829	0.8884
CD37 AND NOT-KLRF1	AML	0.8529	0.8007	0.9124
CD37 AND NOT-IFNAR1	AML	0.8528	0.8199	0.8884
TSPAN32 AND NOT-CD79A	AML	0.8522	0.7562	0.9761
CD276 AND NOT-CXADR	Liposarcoma	0.8049	0.7174	0.9167
ADAM12 AND NOT-ITGB6	Liposarcoma	0.7294	0.6327	0.8611
ADAM12 AND NOT-CLDN7	Liposarcoma	0.7273	0.6829	0.7778
ADAM12 AND NOT-PROM1	Liposarcoma	0.7273	0.6829	0.7778
CD276 AND NOT-ATP2C2	Liposarcoma	0.6933	0.6667	0.7222
EPHB2 AND NOT-SEMA4G	Liposarcoma	0.7647	0.8125	0.7222
EPHB2 AND NOT-STYK1	Liposarcoma	0.7077	0.7931	0.6389
EPHB2 AND NOT-CLCA4	Liposarcoma	0.7467	0.7179	0.7778
ADAM12 AND NOT-SDC1	Liposarcoma	0.6739	0.5536	0.8611
EPHB2 AND NOT-IL17RB	Liposarcoma	0.7353	0.7813	0.6944
ADAM12 AND NOT-EGFR	Liposarcoma	0.7671	0.7568	0.7778
CD276 AND NOT-LY6D	Liposarcoma	0.6923	0.6429	0.75
CD276 AND NOT-FGFR2	Liposarcoma	0.7568	0.7368	0.7778
ADAM12 AND NOT-ERBB2	Liposarcoma	0.6593	0.5455	0.8333
ADAM12 AND NOT-SLC7A5	Liposarcoma	0.6588	0.5714	0.7778
ADAM12 AND NOT-CLDN23	Liposarcoma	0.6588	0.5714	0.7778
FAT4 AND NOT-EPCAM	Liposarcoma	0.8	0.8235	0.7778
FAT4 AND NOT-CLDN1	Liposarcoma	0.6765	0.7188	0.6389
EPHB2 AND NOT-ABCC3	Liposarcoma	0.6944	0.6944	0.6944
ADAM12 AND NOT-SLC34A2	Liposarcoma	0.65	0.5909	0.7222
ROR1 AND NOT-TSPAN2	Liposarcoma	0.6545	0.9474	0.5
CD276 AND NOT-TSPAN7	Liposarcoma	0.6486	0.6316	0.6667

FIG. 12 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
FLT3 AND NOT-PODXL	AML	0.9545	0.9914	0.9203
FLT3 AND NOT-KCNMA1	AML	0.9545	0.9914	0.9203
FLT3 AND NOT-ABHD6	AML	0.9545	0.9914	0.9203
FLT3 AND NOT-CD28	AML	0.9545	0.9914	0.9203
FLT3 AND NOT-PTPRM	AML	0.9526	0.9872	0.9203
FLT3 AND NOT-HEG1	AML	0.9526	0.9872	0.9203
FLT3 AND NOT-PTPRZ1	AML	0.9526	0.9872	0.9203
FLT3 AND NOT-SIT1	AML	0.9526	0.9872	0.9203
FLT3 AND NOT-LY9	AML	0.9526	0.9872	0.9203
FLT3 AND NOT-SLC9A9	AML	0.9526	0.9872	0.9203
FLT3 AND NOT-LAMP3	AML	0.9526	0.9872	0.9203
FLT3 AND NOT-TIGIT	AML	0.9524	0.9914	0.9163
FLT3 AND NOT-GOLM1	AML	0.9524	0.9914	0.9163
FLT3 AND NOT-ABCC5	AML	0.9524	0.9914	0.9163
FLT3 AND NOT-BMPR2	AML	0.9524	0.9914	0.9163
FLT3 AND NOT-PMEPA1	AML	0.9524	0.9914	0.9163
FLT3 AND NOT-P2RY12	AML	0.9522	0.9957	0.9124
FLT3 AND NOT-SLC16A4	AML	0.9506	0.983	0.9203
FLT3 AND NOT-CRIM1	AML	0.9506	0.983	0.9203
FLT3 AND NOT-PDGFRB	AML	0.9506	0.983	0.9203
FLT3 AND NOT-CD247	AML	0.9506	0.983	0.9203
FLT3 AND NOT-TSPAN9	AML	0.9506	0.983	0.9203
FLT3 AND NOT-KLRF1	AML	0.9506	0.983	0.9203
FLT3 AND NOT-SLC17A5	AML	0.9506	0.983	0.9203
FLT3 AND NOT-TSPAN6	AML	0.9506	0.983	0.9203
FLT3 AND NOT-ST6GALNAC6	AML	0.9506	0.983	0.9203
FLT3 AND NOT-ST3GAL5	AML	0.9506	0.983	0.9203
FLT3 AND NOT-P2RY6	AML	0.9506	0.983	0.9203
FLT3 AND NOT-CD274	AML	0.9506	0.983	0.9203
FLT3 AND NOT-CAV1	AML	0.9506	0.983	0.9203
FLT3 AND NOT-FPR3	AML	0.9506	0.983	0.9203
FLT3 AND NOT-AMIGO2	AML	0.9506	0.983	0.9203
FLT3 AND NOT-ITGA2	AML	0.9506	0.983	0.9203
FLT3 AND NOT-ITGB5	AML	0.9506	0.983	0.9203
FLT3 AND NOT-ITPR3	AML	0.9506	0.983	0.9203
FLT3 AND NOT-FAM73A	AML	0.9506	0.983	0.9203
FLT3 AND NOT-FAT4	AML	0.9506	0.983	0.9203
FLT3 AND NOT-KLRD1	AML	0.9506	0.983	0.9203
FLT3 AND NOT-TMEM150C	AML	0.9506	0.983	0.9203
FLT3 AND NOT-JAM3	AML	0.9506	0.983	0.9203
FLT3 AND NOT-M6PR	AML	0.9506	0.983	0.9203
FLT3 AND NOT-ZDHHC5	AML	0.9506	0.983	0.9203
FLT3 AND NOT-CD320	AML	0.9506	0.983	0.9203
FLT3 AND NOT-TM4SF1	AML	0.9506	0.983	0.9203
FLT3 AND NOT-EMCN	AML	0.9506	0.983	0.9203
FLT3 AND NOT-AGTR1	AML	0.9506	0.983	0.9203
FLT3 AND NOT-CHIC1	AML	0.9506	0.983	0.9203
FLT3 AND NOT-TEK	AML	0.9506	0.983	0.9203
FLT3 AND NOT-PROS1	AML	0.9506	0.983	0.9203
FLT3 AND NOT-MFAP3L	AML	0.9506	0.983	0.9203
FLT3 AND NOT-SLC44A2	AML	0.9506	0.983	0.9203
FLT3 AND NOT-DAG1	AML	0.9506	0.983	0.9203
FLT3 AND NOT-WLS	AML	0.9504	0.9871	0.9163
FLT3 AND NOT-IL7R	AML	0.9504	0.9871	0.9163
FLT3 AND NOT-BTLA	AML	0.9504	0.9871	0.9163
FLT3 AND NOT-APOLD1	AML	0.9502	0.9913	0.9124
FLT3 AND NOT-PPAP2B	AML	0.9502	0.9913	0.9124
FLT3 AND NOT-TRPC1	AML	0.9502	0.9913	0.9124
FLT3 AND NOT-ACVR2A	AML	0.95	0.9956	0.9084
FLT3 AND NOT-MR1	AML	0.9487	0.9788	0.9203
FLT3 AND NOT-RNF144A	AML	0.9487	0.9788	0.9203
FLT3 AND NOT-PKD2L1	AML	0.9487	0.9788	0.9203
FLT3 AND NOT-ADRA2A	AML	0.9487	0.9788	0.9203
FLT3 AND NOT-ITSN1	AML	0.9487	0.9788	0.9203
FLT3 AND NOT-APH1B	AML	0.9487	0.9788	0.9203
FLT3 AND NOT-INSR	AML	0.9487	0.9788	0.9203
FLT3 AND NOT-ABCG1	AML	0.9487	0.9788	0.9203
FLT3 AND NOT-IGF2R	AML	0.9487	0.9788	0.9203
FLT3 AND NOT-SLC25A4	AML	0.9487	0.9788	0.9203
FLT3 AND NOT-CNGA1	AML	0.9487	0.9788	0.9203
FLT3 AND NOT-SLC16A10	AML	0.9485	0.9829	0.9163
FLT3 AND NOT-IL6ST	AML	0.9485	0.9829	0.9163
FLT3 AND NOT-CDH5	AML	0.9485	0.9829	0.9163
FLT3 AND NOT-IFNAR1	AML	0.9485	0.9829	0.9163
FLT3 AND NOT-KL	AML	0.9485	0.9829	0.9163
FLT3 AND NOT-SLC2A8	AML	0.9485	0.9829	0.9163
FLT3 AND NOT-CD40	AML	0.9485	0.9829	0.9163

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
ADAM12 AND NOT-LGR5	Liposarcoma	0.6458	0.5167	0.8611
ADAM12 AND NOT-ENG	Liposarcoma	0.6444	0.537	0.8056
ROR1 AND NOT-ATP13A3	Liposarcoma	0.6429	0.9	0.5
ROR1 AND NOT-ITGA3	Liposarcoma	0.6429	0.9	0.5
FAT4 AND NOT-ERBB4	Liposarcoma	0.6984	0.8148	0.6111
CD276 AND NOT-EPHA1	Liposarcoma	0.6829	0.6087	0.7778
CD276 AND NOT-SLC12A2	Liposarcoma	0.7027	0.6842	0.7222
EPHB2 AND NOT-MARVELD2	Liposarcoma	0.6364	0.7	0.5833
EPHB2 AND NOT-ANO9	Liposarcoma	0.6364	0.7	0.5833
EPHB2 AND NOT-ACE2	Liposarcoma	0.6364	0.7	0.5833
EPHB2 AND NOT-ABCB1	Liposarcoma	0.6571	0.6765	0.6389
ROR1 AND NOT-CYP4F2	Liposarcoma	0.6316	0.8571	0.5
ROR1 AND NOT-TSPAN5	Liposarcoma	0.6316	0.8571	0.5
ROR1 AND NOT-SLC2A12	Liposarcoma	0.6316	0.8571	0.5
ROR1 AND NOT-BACE1	Liposarcoma	0.6316	0.8571	0.5
ROR1 AND NOT-LRP2	Liposarcoma	0.6316	0.8571	0.5
ROR1 AND NOT-CD274	Liposarcoma	0.6316	0.8571	0.5
ROR1 AND NOT-SLC4A7	Liposarcoma	0.6429	0.9	0.5
ROR1 AND NOT-ANO10	Liposarcoma	0.6316	0.8571	0.5
ROR1 AND NOT-SLC12A1	Liposarcoma	0.6316	0.8571	0.5
ROR1 AND NOT-SLC7A1	Liposarcoma	0.6316	0.8571	0.5
ROR1 AND NOT-KCNMA1	Liposarcoma	0.6316	0.8571	0.5
ROR1 AND NOT-AGPAT3	Liposarcoma	0.6316	0.8571	0.5
CD276 AND NOT-KCNK5	Liposarcoma	0.6301	0.6216	0.6389
CD276 AND NOT-ANO9	Liposarcoma	0.6301	0.6216	0.6389
ROR1 AND NOT-OR51B6	Liposarcoma	0.6296	0.9444	0.4722
EPHB2 AND NOT-SYT13	Liposarcoma	0.7222	0.7222	0.7222
EPHB2 AND NOT-TSPAN7	Liposarcoma	0.6557	0.8	0.5556
CD276 AND NOT-CD9	Liposarcoma	0.7952	0.7021	0.9167
MSR1 AND NOT-CLDN1	Liposarcoma	0.6429	0.5625	0.75
SLC2A10 AND NOT-MET	Liposarcoma	0.625	0.7143	0.5556
ADAM12 AND NOT-CLDN1	Liposarcoma	0.625	0.7143	0.5556
EPHB2 AND NOT-IL22RA1	Liposarcoma	0.6905	0.6042	0.8056
CD276 AND NOT-TACSTD2	Liposarcoma	0.623	0.76	0.5278
CD276 AND NOT-SEMA4G	Liposarcoma	0.6761	0.6857	0.6667
CD276 AND NOT-SLC44A3	Liposarcoma	0.7123	0.7027	0.7222
ROR1 AND NOT-GPM6A	Liposarcoma	0.6207	0.8182	0.5
ROR1 AND NOT-CDH2	Liposarcoma	0.6207	0.8182	0.5
ROR1 AND NOT-SPINT1	Liposarcoma	0.6207	0.8182	0.5
ROR1 AND NOT-DCBLD2	Liposarcoma	0.6207	0.8182	0.5
ROR1 AND NOT-LRP12	Liposarcoma	0.6207	0.8182	0.5
ROR1 AND NOT-NDRG4	Liposarcoma	0.6207	0.8182	0.5
ROR1 AND NOT-GYPE	Liposarcoma	0.6207	0.8182	0.5
ROR1 AND NOT-SLC44A5	Liposarcoma	0.6207	0.8182	0.5
ROR1 AND NOT-SORT1	Liposarcoma	0.6207	0.8182	0.5
ROR1 AND NOT-TM7SF3	Liposarcoma	0.6207	0.8182	0.5
ROR1 AND NOT-TGOLN2	Liposarcoma	0.6182	0.8947	0.4722
SLC34A2 AND NOT-FXYD1	Lung Adenocarcinoma	0.7619	0.8571	0.6857
SLC34A2 AND NOT-DUOX1	Lung Adenocarcinoma	0.7176	0.7705	0.6714
SLC34A2 AND NOT-ST6GALNAC6	Lung Adenocarcinoma	0.6861	0.7015	0.6714
SLC34A2 AND NOT-DUOXA1	Lung Adenocarcinoma	0.6765	0.697	0.6571
SLC34A2 AND NOT-PTGFR	Lung Adenocarcinoma	0.6761	0.6667	0.6857
SLC34A2 AND NOT-SLC13A4	Lung Adenocarcinoma	0.6757	0.641	0.7143
SLC34A2 AND NOT-MGAM	Lung Adenocarcinoma	0.6718	0.7213	0.6286
SLC34A2 AND NOT-SLC6A16	Lung Adenocarcinoma	0.6716	0.7031	0.6429
SLC34A2 AND NOT-ADCY9	Lung Adenocarcinoma	0.6667	0.7288	0.6143
SLC34A2 AND NOT-VSIG2	Lung Adenocarcinoma	0.6571	0.6571	0.6571
SLC34A2 AND NOT-STX2	Lung Adenocarcinoma	0.6512	0.7119	0.6
SLC34A2 AND NOT-CLCA4	Lung Adenocarcinoma	0.6536	0.6024	0.7143
SLC34A2 AND NOT-SLC52A1	Lung Adenocarcinoma	0.6452	0.5882	0.7143
SLC34A2 AND NOT-TRPV4	Lung Adenocarcinoma	0.6447	0.5976	0.7
SLC34A2 AND NOT-KCNMB2	Lung Adenocarcinoma	0.6415	0.573	0.7286
SLC34A2 AND NOT-VAPA	Lung Adenocarcinoma	0.6504	0.7547	0.5714
SLC34A2 AND NOT-ATP7B	Lung Adenocarcinoma	0.637	0.6615	0.6143
SLC34A2 AND NOT-SLC51B	Lung Adenocarcinoma	0.6364	0.6774	0.6
SLC34A2 AND NOT-SLC13A3	Lung Adenocarcinoma	0.6358	0.5926	0.6857
SLC34A2 AND NOT-ADCY2	Lung Adenocarcinoma	0.6335	0.5604	0.7286
SLC34A2 AND NOT-CNNM2	Lung Adenocarcinoma	0.6329	0.5682	0.7143
SLC34A2 AND NOT-SLC23A1	Lung Adenocarcinoma	0.6323	0.5765	0.7
SLC34A2 AND NOT-DSCAML1	Lung Adenocarcinoma	0.6323	0.5765	0.7
SLC34A2 AND NOT-SLC9C2	Lung Adenocarcinoma	0.6296	0.5543	0.7286
SLC34A2 AND NOT-CLCA2	Lung Adenocarcinoma	0.6289	0.5618	0.7143
SLC34A2 AND NOT-KCNE1	Lung Adenocarcinoma	0.6282	0.5698	0.7
SLC34A2 AND NOT-STOML3	Lung Adenocarcinoma	0.6258	0.5484	0.7286
SLC34A2 AND NOT-CELSR1	Lung Adenocarcinoma	0.625	0.5556	0.7143
SLC34A2 AND NOT-GABRP	Lung Adenocarcinoma	0.625	0.6081	0.6429
SLC34A2 AND NOT-LPAR3	Lung Adenocarcinoma	0.6242	0.5632	0.7

FIG. 12 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
FLT3 AND NOT-DST	AML	0.9485	0.9829	0.9163
FLT3 AND NOT-ENPP4	AML	0.9485	0.9829	0.9163
FLT3 AND NOT-TMEM231	AML	0.9526	0.9872	0.9203
FLT3 AND NOT-CLEC4A	AML	0.9482	0.9871	0.9124
FLT3 AND NOT-STX7	AML	0.9482	0.9871	0.9124
FLT3 AND NOT-TGFBR3	AML	0.9482	0.9871	0.9124
FLT3 AND NOT-HMOX2	AML	0.9482	0.9871	0.9124
FLT3 AND NOT-SIRPG	AML	0.9482	0.9871	0.9124
FLT3 AND NOT-GPR171	AML	0.9482	0.9871	0.9124
FLT3 AND NOT-CD27	AML	0.948	0.9913	0.9084
FLT3 AND NOT-PSEN2	AML	0.9467	0.9747	0.9203
FLT3 AND NOT-JAM2	AML	0.9467	0.9747	0.9203
FLT3 AND NOT-FZD7	AML	0.9467	0.9747	0.9203
FLT3 AND NOT-SEZ6L2	AML	0.9467	0.9747	0.9203
FLT3 AND NOT-MACF1	AML	0.9467	0.9747	0.9203
FLT3 AND NOT-SLC22A17	AML	0.9467	0.9747	0.9203
FLT3 AND NOT-CDIPT	AML	0.9467	0.9747	0.9203
FLT3 AND NOT-ZNRF3	AML	0.9467	0.9747	0.9203
FLT3 AND NOT-GPR176	AML	0.9467	0.9747	0.9203
SLC2A10 AND NOT-PERP	Liposarcoma	0.9143	0.9412	0.8889
ADAM12 AND NOT-ITGA6	Liposarcoma	0.7949	0.7381	0.8611
ADAM12 AND NOT-TACSTD2	Liposarcoma	0.8378	0.8158	0.8611
CNTNAP1 AND NOT-DPP6	Liposarcoma	0.7647	0.8125	0.7222
MSR1 AND NOT-CXADR	Liposarcoma	0.7826	0.8182	0.75
ADAM12 AND NOT-CDH3	Liposarcoma	0.7397	0.7297	0.75
ADAM12 AND NOT-DSC2	Liposarcoma	0.7368	0.7	0.7778
ADAM12 AND NOT-MUC15	Liposarcoma	0.7368	0.7	0.7778
ADAM12 AND NOT-PERP	Liposarcoma	0.8169	0.8286	0.8056
ADAM12 AND NOT-OLR1	Liposarcoma	0.75	0.6818	0.8333
ADAM12 AND NOT-LRP2	Liposarcoma	0.7229	0.6383	0.8333
ADAM12 AND NOT-ATP7B	Liposarcoma	0.7229	0.6383	0.8333
ADAM12 AND NOT-DSC3	Liposarcoma	0.72	0.6923	0.75
CNTNAP1 AND NOT-NTRK3	Liposarcoma	0.7188	0.8214	0.6389
ADAM12 AND NOT-MAL	Liposarcoma	0.725	0.6591	0.8056
ADAM12 AND NOT-PCDH10	Liposarcoma	0.7381	0.6458	0.8611
CNTNAP1 AND NOT-GRIA2	Liposarcoma	0.7879	0.8667	0.7222
ADAM12 AND NOT-MFSD2A	Liposarcoma	0.7059	0.6122	0.8333
ADAM12 AND NOT-DAG1	Liposarcoma	0.7013	0.6585	0.75
CNTNAP1 AND NOT-GRIK2	Liposarcoma	0.7188	0.8214	0.6389
ADAM12 AND NOT-STXBP2	Liposarcoma	0.6977	0.6	0.8333
ADAM12 AND NOT-DPP4	Liposarcoma	0.6977	0.6	0.8333
ADAM12 AND NOT-IL6ST	Liposarcoma	0.6933	0.6667	0.7222
ADAM12 AND NOT-P2RY14	Liposarcoma	0.6933	0.6667	0.7222
ADAM12 AND NOT-SEMA6D	Liposarcoma	0.6905	0.6042	0.8056
ADAM12 AND NOT-CSF2RB	Liposarcoma	0.6857	0.7059	0.6667
ADAM12 AND NOT-ATP12A	Liposarcoma	0.6829	0.6087	0.7778
ADAM12 AND NOT-CXADR	Liposarcoma	0.6824	0.5918	0.8056
PON2 AND NOT-ABCB1	Lung Adenocarcinoma	0.6355	0.9189	0.4857
OSMR AND NOT-LYVE1	Lung Adenocarcinoma	0.625	0.8333	0.5
OSMR AND NOT-JAM3	Lung Adenocarcinoma	0.6182	0.85	0.4857
OSMR AND NOT-SEMA6A	Lung Adenocarcinoma	0.6071	0.8095	0.4857
OSMR AND NOT-NGFR	Lung Adenocarcinoma	0.6034	0.7609	0.5
OSMR AND NOT-TGFBR3	Lung Adenocarcinoma	0.6019	0.9394	0.4429
SEMA4F AND NOT-GPM6A	Lung Carcinoma	0.6197	0.5641	0.6875
CHRNA5 AND NOT-GRIK2	Lung Carcinoma	0.608	0.623	0.5938
ADAM12 AND NOT-MPZ	Lung Carcinoma	0.6259	0.5542	0.7188
SEMA4F AND NOT-ATP1B2	Lung Carcinoma	0.6056	0.5513	0.6719
ADAM12 AND NOT-CD300LG	Lung Carcinoma	0.6138	0.464	0.9063
CXCL9 AND NOT-FCER1A	B-Cell Diffuse	0.7937	0.9615	0.6757
CXCL9 AND NOT-SLC16A10	B-Cell Diffuse	0.8	0.9286	0.7027
CXCL9 AND NOT-EREG	B-Cell Diffuse	0.7813	0.9259	0.6757
CXCL9 AND NOT-KCNJ15	B-Cell Diffuse	0.7813	0.9259	0.6757
CXCL9 AND NOT-CLEC5A	B-Cell Diffuse	0.7742	0.96	0.6486
CXCL9 AND NOT-TLR3	B-Cell Diffuse	0.7692	0.8929	0.6757
CXCL9 AND NOT-GPR35	B-Cell Diffuse	0.7692	0.8929	0.6757
CXCL9 AND NOT-SERINC5	B-Cell Diffuse	0.7692	0.8929	0.6757
CXCL9 AND NOT-S1PR5	B-Cell Diffuse	0.7647	0.8387	0.7027
CD80 AND NOT-FCER1A	B-Cell Diffuse	0.7692	0.8929	0.6757
CXCL9 AND NOT-PLB1	B-Cell Diffuse	0.7619	0.9231	0.6486
CXCL9 AND NOT-GJB2	B-Cell Diffuse	0.7619	0.9231	0.6486
MCOLN2 AND NOT-GPR35	B-Cell Diffuse	0.7619	0.9231	0.6486
CXCL9 AND NOT-SGMS2	B-Cell Diffuse	0.7576	0.8621	0.6757
CXCL9 AND NOT-ITGB5	B-Cell Diffuse	0.7576	0.8621	0.6757
CXCL9 AND NOT-ITM2B	B-Cell Diffuse	0.7536	0.8125	0.7027
CXCL9 AND NOT-GJB6	B-Cell Diffuse	0.7536	0.8125	0.7027
MCOLN2 AND NOT-MRAP2	B-Cell Diffuse	0.7536	0.8125	0.7027
CD80 AND NOT-S1PR5	B-Cell Diffuse	0.7536	0.8125	0.7027

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
SLC34A2 AND NOT-WDR19	Lung Adenocarcinoma	0.6242	0.5632	0.7
SLC34A2 AND NOT-TGFBR3	Lung Adenocarcinoma	0.6214	0.9697	0.4571
SLC34A2 AND NOT-C1orf101	Lung Adenocarcinoma	0.6296	0.5543	0.7286
SLC34A2 AND NOT-SLC51A	Lung Adenocarcinoma	0.6203	0.5568	0.7
SLC34A2 AND NOT-KRT5	Lung Adenocarcinoma	0.6187	0.6232	0.6143
SLC34A2 AND NOT-SLC27A1	Lung Adenocarcinoma	0.6164	0.5921	0.6429
SLC34A2 AND NOT-LRRC4	Lung Adenocarcinoma	0.6258	0.5484	0.7286
SLC34A2 AND NOT-BBS4	Lung Adenocarcinoma	0.6242	0.5632	0.7
SLC34A2 AND NOT-EFNB3	Lung Adenocarcinoma	0.6456	0.5795	0.7286
SLC34A2 AND NOT-DLL1	Lung Adenocarcinoma	0.6111	0.8684	0.4714
SLC34A2 AND NOT-UPK1B	Lung Adenocarcinoma	0.6111	0.5946	0.6286
SLC34A2 AND NOT-VNN3	Lung Adenocarcinoma	0.6111	0.5946	0.6286
SLC34A2 AND NOT-CDH12	Lung Adenocarcinoma	0.6104	0.5595	0.6714
SLC34A2 AND NOT-SGMS2	Lung Adenocarcinoma	0.6099	0.6056	0.6143
SLC34A2 AND NOT-TSPAN1	Lung Adenocarcinoma	0.6087	0.5385	0.7
SLC34A2 AND NOT-CDHR3	Lung Adenocarcinoma	0.6065	0.5529	0.6714
SLC34A2 AND NOT-NRG4	Lung Adenocarcinoma	0.6486	0.6154	0.6857
SLC34A2 AND NOT-GJB5	Lung Adenocarcinoma	0.6053	0.561	0.6571
SLC34A2 AND NOT-BEST4	Lung Adenocarcinoma	0.604	0.5696	0.6429
SLC34A2 AND NOT-DSC3	Lung Adenocarcinoma	0.6906	0.6957	0.6857
SLC34A2 AND NOT-APCDD1	Lung Adenocarcinoma	0.6483	0.6267	0.6714
SLC34A2 AND NOT-PRRT2	Lung Adenocarcinoma	0.6581	0.6	0.7286
FAP AND NOT-BOC	Lung Adenocarcinoma	0.6	0.825	0.4714
SLC34A2 AND NOT-SLC4A3	Lung Adenocarcinoma	0.6316	0.6667	0.6
SLC34A2 AND NOT-CDH26	Lung Adenocarcinoma	0.6	0.5333	0.6857
OSMR AND NOT-EDNRB	Lung Adenocarcinoma	0.6	0.72	0.5143
SLC34A2 AND NOT-PIGR	Lung Adenocarcinoma	0.6222	0.6462	0.6
MAGEA11 AND NOT-JAM3	Lung Carcinoma	0.6237	1	0.4531
MAGEA11 AND NOT-TRPV6	Lung Carcinoma	0.6316	0.9677	0.4688
MAGEA11 AND NOT-PRLR	Lung Carcinoma	0.625	0.9375	0.4688
MAGEA11 AND NOT-TMEM88	Lung Carcinoma	0.6383	1	0.4688
MAGEA11 AND NOT-GLDN	Lung Carcinoma	0.6237	1	0.4531
MAGEA11 AND NOT-SLC7A6	Lung Carcinoma	0.6383	1	0.4688
MAGEA11 AND NOT-LIFR	Lung Carcinoma	0.625	0.9375	0.4688
MAGEA11 AND NOT-APLNR	Lung Carcinoma	0.6316	0.9677	0.4688
MAGEA11 AND NOT-KCNK5	Lung Carcinoma	0.6383	1	0.4688
MAGEA11 AND NOT-CLEC1A	Lung Carcinoma	0.6316	0.9677	0.4688
MAGEA11 AND NOT-SLC19A3	Lung Carcinoma	0.625	0.9375	0.4688
MAGEA11 AND NOT-CADM3	Lung Carcinoma	0.6316	0.9677	0.4688
MAGEA11 AND NOT-FURIN	Lung Carcinoma	0.6316	0.9677	0.4688
MAGEA11 AND NOT-KIAA1919	Lung Carcinoma	0.6316	0.9677	0.4688
MAGEA11 AND NOT-SLC7A2	Lung Carcinoma	0.6186	0.9091	0.4688
MAGEA11 AND NOT-SYT12	Lung Carcinoma	0.617	0.9667	0.4531
MAGEA11 AND NOT-EPOR	Lung Carcinoma	0.6122	0.8824	0.4688
MAGEA11 AND NOT-SLC13A3	Lung Carcinoma	0.6122	0.8824	0.4688
MAGEA11 AND NOT-FLT4	Lung Carcinoma	0.6105	0.9355	0.4531
MAGEA11 AND NOT-PCDH12	Lung Carcinoma	0.6105	0.9355	0.4531
MAGEA11 AND NOT-C1QTNF1	Lung Carcinoma	0.6105	0.9355	0.4531
MAGEA11 AND NOT-VAMP2	Lung Carcinoma	0.6087	1	0.4375
MAGEA11 AND NOT-SLC26A6	Lung Carcinoma	0.6087	1	0.4375
MAGEA11 AND NOT-ZFYVE27	Lung Carcinoma	0.6237	1	0.4531
MAGEA11 AND NOT-SLC30A2	Lung Carcinoma	0.6122	0.8824	0.4688
MAGEA11 AND NOT-SIGLEC6	Lung Carcinoma	0.6186	0.9091	0.4688
MAGEA11 AND NOT-ADCY6	Lung Carcinoma	0.6042	0.9063	0.4531
MAGEA11 AND NOT-SPPL3	Lung Carcinoma	0.6022	0.9655	0.4375
MAGEA11 AND NOT-PTH1R	Lung Carcinoma	0.6022	0.9655	0.4375
MAGEA11 AND NOT-CRIM1	Lung Carcinoma	0.6022	0.9655	0.4375
MAGEA11 AND NOT-KCNK3	Lung Carcinoma	0.6122	0.8824	0.4688
MAGEA11 AND NOT-LRP2	Lung Carcinoma	0.6122	0.8824	0.4688
MAGEA11 AND NOT-TREML2	Lung Carcinoma	0.6122	0.8824	0.4688
MAGEA11 AND NOT-CSF3R	Lung Carcinoma	0.6122	0.8824	0.4688
MAGEA11 AND NOT-STRA6	Lung Carcinoma	0.6122	0.8824	0.4688
MAGEA11 AND NOT-TMEM150A	Lung Carcinoma	0.617	0.9667	0.4531
MAGEA11 AND NOT-KCNN4	Lung Carcinoma	0.6237	1	0.4531
MAGEA11 AND NOT-TMEM150C	Lung Carcinoma	0.617	0.9667	0.4531
MAGEA11 AND NOT-SYT7	Lung Carcinoma	0.6122	0.8824	0.4688
MAGEA11 AND NOT-BCAM	Lung Carcinoma	0.6042	0.9063	0.4531
MAGEA11 AND NOT-ADCY4	Lung Carcinoma	0.6087	1	0.4375
CD180 AND NOT-CXCR2	B-Cell Diffuse	0.7647	0.8387	0.7027
CD180 AND NOT-FCER1A	B-Cell Diffuse	0.7429	0.7879	0.7027
CXCL9 AND NOT-CLDN1	B-Cell Diffuse	0.7429	0.7879	0.7027
MCOLN2 AND NOT-ERBB3	B-Cell Diffuse	0.7429	0.7879	0.7027
CXCL9 AND NOT-CLDN11	B-Cell Diffuse	0.7353	0.8065	0.6757
CXCL9 AND NOT-RAET1E	B-Cell Diffuse	0.7324	0.7647	0.7027
CD79A AND NOT-TSPAN32	B-Cell Diffuse	0.7302	0.8846	0.6216
CXCR5 AND NOT-TSPAN32	B-Cell Diffuse	0.7397	0.75	0.7297
VCAM1 AND NOT-SLC16A2	B-Cell Diffuse	0.7273	0.8276	0.6486

FIG. 12 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
CXCL9 AND NOT-EMCN	B-Cell Diffuse	0.7536	0.8125	0.7027
CXCL9 AND NOT-STX3	B-Cell Diffuse	0.75	0.8889	0.6486
TNFRSF9 AND NOT-BEST1	B-Cell Diffuse	0.75	0.8889	0.6486
CXCL9 AND NOT-SLC22A4	B-Cell Diffuse	0.75	0.8889	0.6486
CXCL9 AND NOT-LGR6	B-Cell Diffuse	0.75	0.8889	0.6486
MCOLN2 AND NOT-SLC4A4	B-Cell Diffuse	0.75	0.7714	0.7297
MCOLN2 AND NOT-ANO5	B-Cell Diffuse	0.75	0.7714	0.7297
MCOLN2 AND NOT-SYT13	B-Cell Diffuse	0.75	0.7714	0.7297
CXCL9 AND NOT-BEST1	B-Cell Diffuse	0.7463	0.8333	0.6757
CXCL9 AND NOT-RHOT1	B-Cell Diffuse	0.7463	0.8333	0.6757
CXCL9 AND NOT-CAV2	B-Cell Diffuse	0.7463	0.8333	0.6757
CD80 AND NOT-CXCR2	B-Cell Diffuse	0.7463	0.8333	0.6757
CXCL9 AND NOT-ADCY9	B-Cell Diffuse	0.7463	0.8333	0.6757
TNFRSF9 AND NOT-CD300LB	B-Cell Diffuse	0.7463	0.8333	0.6757
CXCL9 AND NOT-PCDHB16	B-Cell Diffuse	0.7463	0.8333	0.6757
CXCL9 AND NOT-PROS1	B-Cell Diffuse	0.7463	0.8333	0.6757
MCOLN2 AND NOT-LGR6	B-Cell Diffuse	0.7463	0.8333	0.6757
FAM26F AND NOT-FCER1A	B-Cell Diffuse	0.8052	0.775	0.8378
MCOLN2 AND NOT-FCER1A	B-Cell Diffuse	0.7463	0.8333	0.6757
CXCL9 AND NOT-TNFRSF19	B-Cell Diffuse	0.7429	0.7879	0.7027
CXCL9 AND NOT-KCNJ5	B-Cell Diffuse	0.7429	0.7879	0.7027
CXCL9 AND NOT-GLDN	B-Cell Diffuse	0.7429	0.7879	0.7027
CXCL9 AND NOT-MFAP3L	B-Cell Diffuse	0.7429	0.7879	0.7027
CXCL9 AND NOT-NTRK1	B-Cell Diffuse	0.7429	0.7879	0.7027
CXCL9 AND NOT-CD300LG	B-Cell Diffuse	0.7429	0.7879	0.7027
CXCL9 AND NOT-SLC28A3	B-Cell Diffuse	0.7429	0.7879	0.7027
CXCL9 AND NOT-BDKRB1	B-Cell Diffuse	0.7429	0.7879	0.7027
CXCL9 AND NOT-NEO1	B-Cell Diffuse	0.7429	0.7879	0.7027
CXCL9 AND NOT-DUOX1	B-Cell Diffuse	0.7429	0.7879	0.7027
CXCL9 AND NOT-PTPRZ1	B-Cell Diffuse	0.7429	0.7879	0.7027
CXCL9 AND NOT-GDPD2	B-Cell Diffuse	0.7429	0.7879	0.7027
CXCL9 AND NOT-SORT1	B-Cell Diffuse	0.7419	0.92	0.6216
CXCL9 AND NOT-FCGRT	B-Cell Diffuse	0.7419	0.92	0.6216
CXCL9 AND NOT-NRP1	B-Cell Diffuse	0.7419	0.92	0.6216
CXCL9 AND NOT-SDC2	B-Cell Diffuse	0.7419	0.92	0.6216
CXCL9 AND NOT-EMP1	B-Cell Diffuse	0.7419	0.92	0.6216
MCOLN2 AND NOT-SEMA4G	B-Cell Diffuse	0.7397	0.75	0.7297
MCOLN2 AND NOT-PCDH19	B-Cell Diffuse	0.7397	0.75	0.7297
MCOLN2 AND NOT-CA4	B-Cell Diffuse	0.7397	0.75	0.7297
MCOLN2 AND NOT-IHH	B-Cell Diffuse	0.7397	0.75	0.7297
MCOLN2 AND NOT-SLC51B	B-Cell Diffuse	0.7385	0.8571	0.6486
CXCL9 AND NOT-ATP10A	B-Cell Diffuse	0.7385	0.8571	0.6486
CD80 AND NOT-SLC16A10	B-Cell Diffuse	0.7647	0.8387	0.7027
CD80 AND NOT-GPR35	B-Cell Diffuse	0.7619	0.9231	0.6486
CD80 AND NOT-HAS1	B-Cell Diffuse	0.7353	0.8065	0.6757
CXCL9 AND NOT-FZD10	B-Cell Diffuse	0.7353	0.8065	0.6757
CXCL9 AND NOT-SLC24A3	B-Cell Diffuse	0.7353	0.8065	0.6757
MCOLN2 AND NOT-SLC26A3	B-Cell Diffuse	0.7353	0.8065	0.6757
CXCL9 AND NOT-RECK	B-Cell Diffuse	0.7353	0.8065	0.6757
CXCL9 AND NOT-WLS	B-Cell Diffuse	0.7353	0.8065	0.6757
CXCL9 AND NOT-EMP2	B-Cell Diffuse	0.7353	0.8065	0.6757
CXCL9 AND NOT-LDLRAD3	B-Cell Diffuse	0.7353	0.8065	0.6757
CXCL9 AND NOT-CADM3	B-Cell Diffuse	0.7353	0.8065	0.6757
CXCL9 AND NOT-SLC18A2	B-Cell Diffuse	0.7353	0.8065	0.6757
CXCL9 AND NOT-BMP2	B-Cell Diffuse	0.7353	0.8065	0.6757
CXCL9 AND NOT-SLC1A7	B-Cell Diffuse	0.7353	0.8065	0.6757
CD80 AND NOT-ITM2B	B-Cell Diffuse	0.7536	0.8125	0.7027
CLECL1 AND NOT-SLC18A2	B-Cell Diffuse	0.7333	0.9565	0.5946
CLECL1 AND NOT-ABCA2	B-Cell Diffuse	0.7333	0.9565	0.5946
CLECL1 AND NOT-ITM2B	B-Cell Diffuse	0.7333	0.9565	0.5946
CXCL9 AND NOT-SLCO3A1	B-Cell Diffuse	0.7333	0.9565	0.5946
CXCL9 AND NOT-KCNJ13	B-Cell Diffuse	0.7324	0.7647	0.7027
CXCL9 AND NOT-ANO5	B-Cell Diffuse	0.7324	0.7647	0.7027
CXCL9 AND NOT-DSC3	B-Cell Diffuse	0.7324	0.7647	0.7027
CXCL9 AND NOT-SGCA	B-Cell Diffuse	0.7324	0.7647	0.7027
CXCL9 AND NOT-HCN3	B-Cell Diffuse	0.7324	0.7647	0.7027
CXCL9 AND NOT-CLCA2	B-Cell Diffuse	0.7324	0.7647	0.7027
CXCL9 AND NOT-AGTR1	B-Cell Diffuse	0.7324	0.7647	0.7027
CXCL9 AND NOT-ABCA8	B-Cell Diffuse	0.7324	0.7647	0.7027
CXCL9 AND NOT-SLC2A12	B-Cell Diffuse	0.7324	0.7647	0.7027
CXCL9 AND NOT-CDHR1	B-Cell Diffuse	0.7324	0.7647	0.7027
CXCL9 AND NOT-IL1RL1	B-Cell Diffuse	0.7324	0.7647	0.7027
CXCL9 AND NOT-GJD4	B-Cell Diffuse	0.7324	0.7647	0.7027
CXCL9 AND NOT-SLC15A1	B-Cell Diffuse	0.7324	0.7647	0.7027
CXCL9 AND NOT-SLC4A9	B-Cell Diffuse	0.7324	0.7647	0.7027
CXCL9 AND NOT-DPP6	B-Cell Diffuse	0.7324	0.7647	0.7027
CXCL9 AND NOT-FCER1A	Anaplastic Lymphoma	0.6667	0.8889	0.5333

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
CXCL9 AND NOT-AXL	B-Cell Diffuse	0.7246	0.7813	0.6757
CXCL9 AND NOT-IL11RA	B-Cell Diffuse	0.7246	0.7813	0.6757
CXCL9 AND NOT-BMPR1B	B-Cell Diffuse	0.7222	0.7429	0.7027
CXCL9 AND NOT-ERBB4	B-Cell Diffuse	0.7222	0.7429	0.7027
MCOLN2 AND NOT-GPA33	B-Cell Diffuse	0.7222	0.7429	0.7027
CXCL9 AND NOT-PROM1	B-Cell Diffuse	0.7222	0.7429	0.7027
MCOLN2 AND NOT-MUC1	B-Cell Diffuse	0.7164	0.8	0.6486
CD180 AND NOT-CLEC1B	B-Cell Diffuse	0.7429	0.7879	0.7027
CD79B AND NOT-RNF144A	B-Cell Diffuse	0.7143	0.7576	0.6757
CXCL9 AND NOT-ULBP2	B-Cell Diffuse	0.7143	0.7576	0.6757
CXCL9 AND NOT-ERBB3	B-Cell Diffuse	0.7143	0.7576	0.6757
CD79B AND NOT-SLC46A2	B-Cell Diffuse	0.7143	0.7576	0.6757
CD79B AND NOT-ASGR1	B-Cell Diffuse	0.7143	0.7576	0.6757
CXCL9 AND NOT-RNF43	B-Cell Diffuse	0.7143	0.7576	0.6757
CXCL9 AND NOT-CD34	B-Cell Diffuse	0.7143	0.7576	0.6757
CD79B AND NOT-CD244	B-Cell Diffuse	0.7143	0.7576	0.6757
CD79B AND NOT-SLC18A2	B-Cell Diffuse	0.7143	0.7576	0.6757
CD79B AND NOT-SLC24A1	B-Cell Diffuse	0.7143	0.7576	0.6757
CD79B AND NOT-CXCR2	B-Cell Diffuse	0.7143	0.7576	0.6757
CD79B AND NOT-AMICA1	B-Cell Diffuse	0.7143	0.7576	0.6757
CXCL9 AND NOT-IL20RA	B-Cell Diffuse	0.7123	0.7222	0.7027
CXCL9 AND NOT-CLDN18	B-Cell Diffuse	0.7123	0.7222	0.7027
MCOLN2 AND NOT-PROM1	B-Cell Diffuse	0.7123	0.7222	0.7027
CXCL9 AND NOT-MUC13	B-Cell Diffuse	0.7123	0.7222	0.7027
CD80 AND NOT-CLDN1	B-Cell Diffuse	0.7123	0.7222	0.7027
CXCL9 AND NOT-MSLN	B-Cell Diffuse	0.7123	0.7222	0.7027
CD80 AND NOT-BMPR1B	B-Cell Diffuse	0.7123	0.7222	0.7027
MCOLN2 AND NOT-MUC4	B-Cell Diffuse	0.7123	0.7222	0.7027
MCOLN2 AND NOT-ITGB6	B-Cell Diffuse	0.7123	0.7222	0.7027
MCOLN2 AND NOT-MUC13	B-Cell Diffuse	0.7105	0.6923	0.7297
CD79B AND NOT-ITSN1	B-Cell Diffuse	0.7143	0.7576	0.6757
CD79A AND NOT-FCER1A	B-Cell Diffuse	0.7188	0.8519	0.6216
CD40 AND NOT-SDC1	B-Cell Diffuse	0.7143	0.7576	0.6757
CD40 AND NOT-EDNRB	B-Cell Diffuse	0.7246	0.7813	0.6757
CD40 AND NOT-AXL	B-Cell Diffuse	0.7059	0.7742	0.6486
CD79A AND NOT-CXCR2	B-Cell Diffuse	0.7059	0.7742	0.6486
CXCL9 AND NOT-CSPG4	B-Cell Diffuse	0.7059	0.7742	0.6486
CD79B AND NOT-KLRD1	B-Cell Diffuse	0.7042	0.7353	0.6757
CD79B AND NOT-INSR	B-Cell Diffuse	0.7042	0.7353	0.6757
CXCL9 AND NOT-ITGB6	B-Cell Diffuse	0.7042	0.7353	0.6757
CXCL9 AND NOT-EDNRB	B-Cell Diffuse	0.7042	0.7353	0.6757
CD79B AND NOT-FCER1A	B-Cell Diffuse	0.7042	0.7353	0.6757
CD79B AND NOT-TLR5	B-Cell Diffuse	0.7042	0.7353	0.6757
CD79B AND NOT-CX3CR1	B-Cell Diffuse	0.7042	0.7353	0.6757
CXCL9 AND NOT-GPA33	B-Cell Diffuse	0.7042	0.7353	0.6757
CXCL9 AND NOT-FOLH1	B-Cell Diffuse	0.7042	0.7353	0.6757
CXCL9 AND NOT-EGFR	B-Cell Diffuse	0.7042	0.7353	0.6757
CD79B AND NOT-PKN2	B-Cell Diffuse	0.7042	0.7353	0.6757
CD79B AND NOT-ABCA2	B-Cell Diffuse	0.7042	0.7353	0.6757
CXCL9 AND NOT-SDC1	B-Cell Diffuse	0.7042	0.7353	0.6757
CD79B AND NOT-MFAP3L	B-Cell Diffuse	0.7042	0.7353	0.6757
CD79B AND NOT-CLEC12A	B-Cell Diffuse	0.7042	0.7353	0.6757
MCOLN2 AND NOT-ERBB2	B-Cell Diffuse	0.7042	0.7353	0.6757
CXCL9 AND NOT-CLDN23	Anaplastic Lymphoma	0.6667	0.8889	0.5333
IL2RA AND NOT-GAL3ST1	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-GPM6A	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-KCNH7	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-KCNH5	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-SCNN1D	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-GPR75	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-PKHD1	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-IL1RAPL1	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-SLC28A2	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-SGCD	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-FAT3	Anaplastic Lymphoma	0.6429	0.6923	0.6
IL2RA AND NOT-SLC29A2	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-FGFR3	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-SLC43A2	Anaplastic Lymphoma	0.6957	1	0.5333
IL2RA AND NOT-KCNJ16	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-KCNJ12	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-SHH	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-GPR20	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-TAAR2	Anaplastic Lymphoma	0.6429	0.6923	0.6
IL2RA AND NOT-FNDC5	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-VSIG2	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-TRPC4	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-OR5J2	Anaplastic Lymphoma	0.6429	0.6923	0.6

FIG. 12 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
CXCL9 AND NOT-SORT1	Anaplastic Lymphoma	0.6923	0.8182	0.6
CXCL9 AND NOT-SLC44A1	Anaplastic Lymphoma	0.75	1	0.6
CXCL9 AND NOT-STX3	Anaplastic Lymphoma	0.6667	0.75	0.6
CXCL9 AND NOT-TLR3	Anaplastic Lymphoma	0.6429	0.6923	0.6
CXCL9 AND NOT-CD9	Anaplastic Lymphoma	0.64	0.8	0.5333
CXCL9 AND NOT-SDC2	Anaplastic Lymphoma	0.64	0.8	0.5333
CXCL9 AND NOT-EFNA5	Anaplastic Lymphoma	0.6923	0.8182	0.6
CXCL9 AND NOT-F3	Anaplastic Lymphoma	0.6207	0.6429	0.6
CXCL9 AND NOT-SLC17A5	Anaplastic Lymphoma	0.6207	0.6429	0.6
CXCL9 AND NOT-KCNJ15	Anaplastic Lymphoma	0.6364	1	0.4667
CXCL9 AND NOT-TM9SF2	Anaplastic Lymphoma	0.6154	0.7273	0.5333
CXCL9 AND NOT-SERINC5	Anaplastic Lymphoma	0.6207	0.6429	0.6
CXCL9 AND NOT-ADRB2	Anaplastic Lymphoma	0.6087	0.875	0.4667
CXCL9 AND NOT-FFAR2	Anaplastic Lymphoma	0.64	0.8	0.5333
IL2RB AND NOT-TSPAN32	Anaplastic Lymphoma	0.6061	0.5556	0.6667
CXCL9 AND NOT-ADCY9	Anaplastic Lymphoma	0.6	0.6	0.6
CXCL9 AND NOT-CRIM1	Anaplastic Lymphoma	0.6	0.6	0.6
CXCL9 AND NOT-SLC18A2	Anaplastic Lymphoma	0.6	0.6	0.6
CXCL9 AND NOT-SLC34A3	Anaplastic Lymphoma	0.6	0.6	0.6
CXCL9 AND NOT-BST1	Anaplastic Lymphoma	0.6087	0.875	0.4667
CXCL9 AND NOT-JAG1	Anaplastic Lymphoma	0.6087	0.875	0.4667
CXCL9 AND NOT-B4GALT1	Anaplastic Lymphoma	0.6957	1	0.5333
CXCL9 AND NOT-TMEM127	Anaplastic Lymphoma	0.6667	0.8889	0.5333
CXCL9 AND NOT-SLC43A2	Anaplastic Lymphoma	0.6087	0.875	0.4667
LAG3 AND NOT-TMEM8B	Anaplastic Lymphoma	0.6061	0.5556	0.6667
CXCL9 AND NOT-VNN3	Anaplastic Lymphoma	0.6207	0.6429	0.6
CXCL9 AND NOT-IL22RA1	Anaplastic Lymphoma	0.64	0.8	0.5333
CXCL9 AND NOT-HCN3	Anaplastic Lymphoma	0.6	0.6	0.6
CXCL9 AND NOT-CKAP4	Anaplastic Lymphoma	0.6667	0.75	0.6
CXCL9 AND NOT-ANO6	Anaplastic Lymphoma	0.6087	0.875	0.4667
CXCL9 AND NOT-STX1B	Anaplastic Lymphoma	0.6667	0.8889	0.5333
CXCL9 AND NOT-EPHA4	Anaplastic Lymphoma	0.6429	0.6923	0.6
CLECL1 AND NOT-SLC46A2	Mantle-Cell Lymphoma	0.9737	0.9737	0.9737
CLECL1 AND NOT-PTGDR2	Mantle-Cell Lymphoma	0.9744	0.95	1
CLECL1 AND NOT-ASGR1	Mantle-Cell Lymphoma	0.9737	0.9737	0.9737
CLECL1 AND NOT-FPR2	Mantle-Cell Lymphoma	0.961	0.9487	0.9737
CLECL1 AND NOT-SLC16A10	Mantle-Cell Lymphoma	0.9737	0.9737	0.9737
CLECL1 AND NOT-S1PR5	Mantle-Cell Lymphoma	0.9737	0.9737	0.9737
CLECL1 AND NOT-ACPP	Mantle-Cell Lymphoma	0.9744	0.95	1
CLECL1 AND NOT-ADAM9	Mantle-Cell Lymphoma	0.9487	0.925	0.9737
CLECL1 AND NOT-INSR	Mantle-Cell Lymphoma	0.9474	0.9474	0.9474
CLECL1 AND NOT-GP9	Mantle-Cell Lymphoma	0.962	0.9268	1
CLECL1 AND NOT-SLC51A	Mantle-Cell Lymphoma	0.961	0.9487	0.9737
CLECL1 AND NOT-MS4A2	Mantle-Cell Lymphoma	0.95	0.9048	1
CLECL1 AND NOT-DHRS3	Mantle-Cell Lymphoma	0.961	0.9487	0.9737
CLECL1 AND NOT-ATRN	Mantle-Cell Lymphoma	0.9744	0.95	1
CLECL1 AND NOT-ITGA5	Mantle-Cell Lymphoma	0.987	0.9744	1
CLECL1 AND NOT-TMEM204	Mantle-Cell Lymphoma	0.9487	0.925	0.9737
CLECL1 AND NOT-SLC2A12	Mantle-Cell Lymphoma	1	1	1
CLECL1 AND NOT-CPD	Mantle-Cell Lymphoma	0.9737	0.9737	0.9737
CLECL1 AND NOT-MFAP3L	Mantle-Cell Lymphoma	0.9487	0.925	0.9737
CLECL1 AND NOT-FCER1A	Mantle-Cell Lymphoma	0.96	0.973	0.9474
CLECL1 AND NOT-HM13	Mantle-Cell Lymphoma	0.9744	0.95	1
CLECL1 AND NOT-VSIG2	Mantle-Cell Lymphoma	0.9383	0.8837	1
CLECL1 AND NOT-CCR3	Mantle-Cell Lymphoma	0.9744	0.95	1
CLECL1 AND NOT-SLC18A2	Mantle-Cell Lymphoma	0.987	0.9744	1
CLECL1 AND NOT-VIPR1	Mantle-Cell Lymphoma	0.9351	0.9231	0.9474
CLECL1 AND NOT-SLC24A4	Mantle-Cell Lymphoma	0.9383	0.8837	1
CLECL1 AND NOT-PTGER2	Mantle-Cell Lymphoma	0.9737	0.9737	0.9737
CLECL1 AND NOT-ANKH	Mantle-Cell Lymphoma	0.9315	0.9714	0.8947
CLECL1 AND NOT-GOLM1	Mantle-Cell Lymphoma	0.961	0.9487	0.9737
CLECL1 AND NOT-ACVR1B	Mantle-Cell Lymphoma	0.961	0.9487	0.9737
CLECL1 AND NOT-CLEC1B	Mantle-Cell Lymphoma	0.95	0.9048	1
CLECL1 AND NOT-ATP8B4	Mantle-Cell Lymphoma	0.9744	0.95	1
CLECL1 AND NOT-ADAM23	Mantle-Cell Lymphoma	0.962	0.9268	1
CLECL1 AND NOT-CLEC7A	Mantle-Cell Lymphoma	0.9296	1	0.8684
CLECL1 AND NOT-XK	Mantle-Cell Lymphoma	0.9474	0.9474	0.9474
CLECL1 AND NOT-ASPH	Mantle-Cell Lymphoma	0.961	0.9487	0.9737
CLECL1 AND NOT-CD101	Mantle-Cell Lymphoma	0.9744	0.95	1
CLECL1 AND NOT-AQP11	Mantle-Cell Lymphoma	0.961	0.9487	0.9737
CLECL1 AND NOT-TNFSF14	Mantle-Cell Lymphoma	0.9268	0.8636	1
CLECL1 AND NOT-CLEC5A	Mantle-Cell Lymphoma	0.9268	0.8636	1
CLECL1 AND NOT-SIGLEC9	Mantle-Cell Lymphoma	0.9268	0.8636	1
CLECL1 AND NOT-SLC22A4	Mantle-Cell Lymphoma	0.987	0.9744	1
CLECL1 AND NOT-KLRD1	Mantle-Cell Lymphoma	0.9737	0.9737	0.9737
CLECL1 AND NOT-TGFBR3	Mantle-Cell Lymphoma	0.9268	0.8636	1
CLECL1 AND NOT-TNFSF10	Mantle-Cell Lymphoma	0.9744	0.95	1

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
IL2RA AND NOT-CDH6	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-OR51I1	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-DPP6	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-SIGLEC6	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-SLC30A3	Anaplastic Lymphoma	0.72	0.9	0.6
IL2RA AND NOT-GRID1	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-KCNB2	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-ERVFRD-1	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-SLC4A3	Anaplastic Lymphoma	0.6923	0.8182	0.6
IL2RA AND NOT-CHRNE	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-GPR173	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-SLC9A3	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-DRD3	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-ABCB8	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-GABRB3	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-KCNJ4	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-ADRA2B	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-OR8D2	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-OR51E2	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-OR1G1	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-SLC6A9	Anaplastic Lymphoma	0.6154	0.7273	0.5333
IL2RA AND NOT-PCDHB13	Anaplastic Lymphoma	0.6154	0.7273	0.5333
IL2RA AND NOT-OR2M4	Anaplastic Lymphoma	0.6154	0.7273	0.5333
IL2RA AND NOT-TMEM8B	Anaplastic Lymphoma	0.6154	0.7273	0.5333
IL2RA AND NOT-RHD	Anaplastic Lymphoma	0.6154	0.7273	0.5333
IL2RA AND NOT-RRH	Anaplastic Lymphoma	0.6154	0.7273	0.5333
IL2RA AND NOT-SLC15A1	Anaplastic Lymphoma	0.6154	0.7273	0.5333
IL2RA AND NOT-KCNC3	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-OR10H3	Anaplastic Lymphoma	0.6154	0.7273	0.5333
IL2RA AND NOT-OR10H1	Anaplastic Lymphoma	0.6154	0.7273	0.5333
IL2RA AND NOT-KCNH2	Anaplastic Lymphoma	0.6154	0.7273	0.5333
IL2RA AND NOT-PTCHD1	Anaplastic Lymphoma	0.6154	0.7273	0.5333
IL2RA AND NOT-SLC26A10	Anaplastic Lymphoma	0.6154	0.7273	0.5333
IL2RA AND NOT-ACE2	Anaplastic Lymphoma	0.6154	0.7273	0.5333
IL2RA AND NOT-CHRNA10	Anaplastic Lymphoma	0.6154	0.7273	0.5333
IL2RA AND NOT-ENTPD2	Anaplastic Lymphoma	0.6154	0.7273	0.5333
IL2RA AND NOT-TAS2R41	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-KIRREL	Anaplastic Lymphoma	0.6154	0.7273	0.5333
IL2RA AND NOT-KCNG3	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-FFAR1	Anaplastic Lymphoma	0.6154	0.7273	0.5333
IL2RA AND NOT-IZUMO1	Anaplastic Lymphoma	0.72	0.9	0.6
IL2RA AND NOT-FLRT1	Anaplastic Lymphoma	0.6154	0.7273	0.5333
IL2RA AND NOT-CHRM5	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-OR7C2	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-OPRL1	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-KCNQ4	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-ASGR1	Anaplastic Lymphoma	0.6667	0.8889	0.5333
IL2RA AND NOT-NRXN1	Anaplastic Lymphoma	0.6154	0.7273	0.5333
IL2RA AND NOT-SGCA	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-PTGDR2	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-GP9	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-KCNJ14	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-TAS2R16	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-SGCZ	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-TAS2R13	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-SLC6A16	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-CCR3	Anaplastic Lymphoma	0.72	0.9	0.6
IL2RA AND NOT-TAS2R39	Anaplastic Lymphoma	0.6429	0.6923	0.6
IL2RA AND NOT-FGFRL1	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-CDHR5	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-WNT4	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-TAS2R50	Anaplastic Lymphoma	0.6429	0.6923	0.6
IL2RA AND NOT-KCNK7	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-BTN1A1	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-KNCN	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-FCGR2B	Anaplastic Lymphoma	0.75	1	0.6
IL2RA AND NOT-LRRTM1	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-ABCC2	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-AQP4	Anaplastic Lymphoma	0.6429	0.6923	0.6
IL2RA AND NOT-ITGA9	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-KCNA5	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-SLC13A4	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-CDH15	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-RHBDL2	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-MAS1	Anaplastic Lymphoma	0.6667	0.75	0.6
CLECL1 AND NOT-ITGB3	Mantle-Cell Lymphoma	0.987	0.9744	1
CLECL1 AND NOT-NCAM1	Mantle-Cell Lymphoma	0.9744	0.95	1

FIG. 12 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
CLECL1 AND NOT-TRAT1	Mantle-Cell Lymphoma	1	1	1
CLECL1 AND NOT-CACNA1I	Mantle-Cell Lymphoma	0.925	0.881	0.9737
CLECL1 AND NOT-CD300C	Mantle-Cell Lymphoma	0.925	0.881	0.9737
CLECL1 AND NOT-SLCO3A1	Mantle-Cell Lymphoma	0.987	0.9744	1
CLECL1 AND NOT-KCNE3	Mantle-Cell Lymphoma	0.9744	0.95	1
CLECL1 AND NOT-CLEC12B	Mantle-Cell Lymphoma	0.9383	0.8837	1
CLECL1 AND NOT-NRG1	Mantle-Cell Lymphoma	0.9367	0.9024	0.9737
CLECL1 AND NOT-HAS3	Mantle-Cell Lymphoma	0.95	0.9048	1
CLECL1 AND NOT-SGMS2	Mantle-Cell Lymphoma	0.9474	0.9474	0.9474
CLECL1 AND NOT-HS3ST3B1	Mantle-Cell Lymphoma	0.9333	0.9459	0.9211
CLECL1 AND NOT-BPI	Mantle-Cell Lymphoma	0.9189	0.9444	0.8947
CLECL1 AND NOT-TNFSF12	Mantle-Cell Lymphoma	0.9487	0.925	0.9737
CLECL1 AND NOT-IL6R	Mantle-Cell Lymphoma	0.962	0.9268	1
CLECL1 AND NOT-IL18RAP	Mantle-Cell Lymphoma	0.987	0.9744	1
CLECL1 AND NOT-CD244	Mantle-Cell Lymphoma	0.9744	0.95	1
CLECL1 AND NOT-AMICA1	Mantle-Cell Lymphoma	0.9744	0.95	1
CLECL1 AND NOT-TENM1	Mantle-Cell Lymphoma	0.987	0.9744	1
CLECL1 AND NOT-SPN	Mantle-Cell Lymphoma	0.9744	0.95	1
CLECL1 AND NOT-SULF2	Mantle-Cell Lymphoma	0.95	0.9048	1
CLECL1 AND NOT-KLRB1	Mantle-Cell Lymphoma	0.9589	1	0.9211
CLECL1 AND NOT-GPR68	Mantle-Cell Lymphoma	0.9268	0.8636	1
CLECL1 AND NOT-SLC44A1	Mantle-Cell Lymphoma	0.9744	0.95	1
CLECL1 AND NOT-SECTM1	Mantle-Cell Lymphoma	0.95	0.9048	1
CLECL1 AND NOT-TLR5	Mantle-Cell Lymphoma	0.9744	0.95	1
CLECL1 AND NOT-GPR143	Mantle-Cell Lymphoma	0.9367	0.9024	0.9737
CLECL1 AND NOT-FLT3LG	Mantle-Cell Lymphoma	0.9744	0.95	1
CLECL1 AND NOT-TSPAN2	Mantle-Cell Lymphoma	0.9737	0.9737	0.9737
CLECL1 AND NOT-SORT1	Mantle-Cell Lymphoma	0.9157	0.8444	1
CLECL1 AND NOT-SLC8A1	Mantle-Cell Lymphoma	0.9157	0.8444	1
CLECL1 AND NOT-SLC19A1	Mantle-Cell Lymphoma	0.9744	0.95	1
CLECL1 AND NOT-EGF	Mantle-Cell Lymphoma	0.9268	0.8636	1
CLECL1 AND NOT-CD226	Mantle-Cell Lymphoma	0.987	0.9744	1
CLECL1 AND NOT-C10orf54	Mantle-Cell Lymphoma	0.961	0.9487	0.9737
CLECL1 AND NOT-CD28	Mantle-Cell Lymphoma	0.987	0.9744	1
CLECL1 AND NOT-NAALAD2	Mantle-Cell Lymphoma	0.962	0.9268	1
CLECL1 AND NOT-EPHA4	Mantle-Cell Lymphoma	0.9315	0.9714	0.8947
CLECL1 AND NOT-DNAJC5	Mantle-Cell Lymphoma	0.9744	0.95	1
CLECL1 AND NOT-HTR2B	Mantle-Cell Lymphoma	0.962	0.9268	1
CLECL1 AND NOT-PTGDR	Mantle-Cell Lymphoma	0.9744	0.95	1
CLECL1 AND NOT-SCIMP	Mantle-Cell Lymphoma	0.9114	0.878	0.9474
CLECL1 AND NOT-SIGLEC7	Mantle-Cell Lymphoma	0.962	0.9268	1
CLECL1 AND NOT-NFAM1	Mantle-Cell Lymphoma	0.9744	0.95	1
CLECL1 AND NOT-NUCB2	Mantle-Cell Lymphoma	0.9737	0.9737	0.9737
CLECL1 AND NOT-ICAM4	Mantle-Cell Lymphoma	0.9091	0.8974	0.9211
CLECL1 AND NOT-DPP4	Mantle-Cell Lymphoma	0.9744	0.95	1
CLECL1 AND NOT-MPZ	Mantle-Cell Lymphoma	0.9268	0.8636	1
CLECL1 AND NOT-NT5E	Mantle-Cell Lymphoma	0.9067	0.9189	0.8947
CLECL1 AND NOT-P2RY12	Mantle-Cell Lymphoma	0.9383	0.8837	1
IL10RA AND NOT-AQP9	Mantle-Cell Lymphoma	0.9211	0.9211	0.9211
CXCL9 AND NOT-CD9	T-Cell, Peripheral	0.8462	0.9167	0.7857
CXCL9 AND NOT-EREG	T-Cell, Peripheral	0.8235	0.913	0.75
CXCL9 AND NOT-EMP2	T-Cell, Peripheral	0.7857	0.7857	0.7857
CXCL9 AND NOT-TLR3	T-Cell, Peripheral	0.7925	0.84	0.75
CXCL9 AND NOT-ITGB5	T-Cell, Peripheral	0.7843	0.8696	0.7143
CXCL9 AND NOT-ADCY9	T-Cell, Peripheral	0.7778	0.8077	0.75
CXCL9 AND NOT-CAV2	T-Cell, Peripheral	0.7778	0.8077	0.75
CXCL9 AND NOT-SGMS2	T-Cell, Peripheral	0.7857	0.7857	0.7857
CXCL9 AND NOT-EPHB4	T-Cell, Peripheral	0.7719	0.7586	0.7857
CXCL9 AND NOT-SLC6A14	T-Cell, Peripheral	0.7719	0.7586	0.7857
CXCL9 AND NOT-SLC17A5	T-Cell, Peripheral	0.8235	0.913	0.75
CXCL9 AND NOT-PTPRG	T-Cell, Peripheral	0.8	0.8148	0.7857
CXCL9 AND NOT-APLP2	T-Cell, Peripheral	0.8148	0.8462	0.7857
CXCL9 AND NOT-CRIM1	T-Cell, Peripheral	0.8148	0.8462	0.7857
CXCL9 AND NOT-SLC24A3	T-Cell, Peripheral	0.7719	0.7586	0.7857
CXCL9 AND NOT-DSC2	T-Cell, Peripheral	0.7636	0.7778	0.75
CXCL9 AND NOT-FZD5	T-Cell, Peripheral	0.7719	0.7586	0.7857
CXCL9 AND NOT-SERINC5	T-Cell, Peripheral	0.7778	0.8077	0.75
CXCL9 AND NOT-SLC44A1	T-Cell, Peripheral	0.7778	0.8077	0.75
CXCL9 AND NOT-PMP22	T-Cell, Peripheral	0.8	0.8148	0.7857
CXCL9 AND NOT-STEAP3	T-Cell, Peripheral	0.7755	0.9048	0.6786
CXCL9 AND NOT-SDC2	T-Cell, Peripheral	0.8462	0.9167	0.7857
CXCL9 AND NOT-SLC41A2	T-Cell, Peripheral	0.7692	0.8333	0.7143
CXCL9 AND NOT-SLC34A3	T-Cell, Peripheral	0.7636	0.7778	0.75
CXCL9 AND NOT-FGFR3	T-Cell, Peripheral	0.7586	0.7333	0.7857
CXCL9 AND NOT-FXYD3	T-Cell, Peripheral	0.7586	0.7333	0.7857
CXCL9 AND NOT-SORT1	T-Cell, Peripheral	0.7586	0.7333	0.7857
CXCL9 AND NOT-KCNS3	T-Cell, Peripheral	0.7586	0.7333	0.7857

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
CLECL1 AND NOT-TNFRSF8	Mantle-Cell Lymphoma	0.9367	0.9024	0.9737
MS4A1 AND NOT-ICAM4	Mantle-Cell Lymphoma	0.9167	0.9706	0.8684
MS4A1 AND NOT-KCNG2	Mantle-Cell Lymphoma	0.9383	0.8837	1
MS4A1 AND NOT-CDHR3	Mantle-Cell Lymphoma	0.9041	0.9429	0.8684
CD52 AND NOT-FPR2	Mantle-Cell Lymphoma	0.8974	0.875	0.9211
MS4A1 AND NOT-LRRTM2	Mantle-Cell Lymphoma	0.881	0.8043	0.9737
P2RX5 AND NOT-MS4A2	Mantle-Cell Lymphoma	0.878	0.8182	0.9474
MS4A1 AND NOT-GPM6B	Mantle-Cell Lymphoma	0.9136	0.8605	0.9737
CLECL1 AND NOT-CD33	Mantle-Cell Lymphoma	0.962	0.9268	1
MS4A1 AND NOT-CYBB	Mantle-Cell Lymphoma	0.8736	0.7755	1
MS4A1 AND NOT-M6PR	Mantle-Cell Lymphoma	0.8736	0.7755	1
CD52 AND NOT-PTGER2	Mantle-Cell Lymphoma	0.9211	0.9211	0.9211
P2RX5 AND NOT-SLAMF6	Mantle-Cell Lymphoma	0.8718	0.85	0.8947
P2RX5 AND NOT-SCN3A	Mantle-Cell Lymphoma	0.8718	0.85	0.8947
QSOX2 AND NOT-TNFRSF8	Mantle-Cell Lymphoma	0.9474	0.9474	0.9474
MS4A1 AND NOT-FPR2	Mantle-Cell Lymphoma	0.8706	0.7872	0.9737
CD52 AND NOT-ITGA5	Mantle-Cell Lymphoma	0.96	0.973	0.9474
P2RX5 AND NOT-FPR2	Mantle-Cell Lymphoma	0.875	0.8333	0.9211
KCNN4 AND NOT-MET	Mantle-Cell Lymphoma	0.9024	0.8409	0.9737
MS4A1 AND NOT-PTGDR2	Mantle-Cell Lymphoma	0.8736	0.7755	1
P2RX5 AND NOT-KCNG2	Mantle-Cell Lymphoma	0.9459	0.9722	0.9211
IL10RA AND NOT-CD33	Mantle-Cell Lymphoma	0.8837	0.7917	1
P2RX5 AND NOT-CYBB	Mantle-Cell Lymphoma	0.9	0.8571	0.9474
CD52 AND NOT-SLC46A2	Mantle-Cell Lymphoma	0.8642	0.814	0.9211
CD52 AND NOT-DHRS3	Mantle-Cell Lymphoma	0.8642	0.814	0.9211
MS4A1 AND NOT-MS4A2	Mantle-Cell Lymphoma	0.8636	0.76	1
MS4A1 AND NOT-GP9	Mantle-Cell Lymphoma	0.8636	0.76	1
MS4A1 AND NOT-SLC16A10	Mantle-Cell Lymphoma	0.8605	0.7708	0.9737
MS4A1 AND NOT-PTPRA	Mantle-Cell Lymphoma	0.8605	0.7708	0.9737
IL10RA AND NOT-ITGB3	Mantle-Cell Lymphoma	0.8736	0.7755	1
MS4A1 AND NOT-S1PR5	Mantle-Cell Lymphoma	0.8605	0.7708	0.9737
P2RX5 AND NOT-CLEC4A	Mantle-Cell Lymphoma	0.8571	0.8462	0.8684
QSOX2 AND NOT-SSTR5	Mantle-Cell Lymphoma	0.9024	0.8409	0.9737
MS4A1 AND NOT-CD86	Mantle-Cell Lymphoma	0.8571	0.7826	0.9474
QSOX2 AND NOT-IL20RA	Mantle-Cell Lymphoma	0.9024	0.8409	0.9737
QSOX2 AND NOT-CLDN2	Mantle-Cell Lymphoma	0.9024	0.8409	0.9737
MS4A1 AND NOT-SERINC3	Mantle-Cell Lymphoma	0.8736	0.7755	1
MS4A1 AND NOT-PSEN1	Mantle-Cell Lymphoma	0.8736	0.7755	1
QSOX2 AND NOT-ENG	Mantle-Cell Lymphoma	0.9024	0.8409	0.9737
CLECL1 AND NOT-MST1R	Mantle-Cell Lymphoma	0.8539	0.7451	1
MS4A1 AND NOT-HM13	Mantle-Cell Lymphoma	0.8539	0.7451	1
MS4A1 AND NOT-LILRB3	Mantle-Cell Lymphoma	0.8539	0.7451	1
MS4A1 AND NOT-LNPEP	Mantle-Cell Lymphoma	0.8539	0.7451	1
CD52 AND NOT-AQP9	Mantle-Cell Lymphoma	0.8533	0.8649	0.8421
CD52 AND NOT-FCER1G	Mantle-Cell Lymphoma	0.8861	0.8537	0.9211
CD52 AND NOT-CLEC7A	Mantle-Cell Lymphoma	0.9067	0.9189	0.8947
CXCL9 AND NOT-CLDN1	T-Cell, Peripheral	0.7719	0.7586	0.7857
CXCL9 AND NOT-ERBB3	T-Cell, Peripheral	0.7586	0.7333	0.7857
CXCL9 AND NOT-CLDN23	T-Cell, Peripheral	0.8462	0.9167	0.7857
CXCL9 AND NOT-ROR1	T-Cell, Peripheral	0.7586	0.7333	0.7857
CXCL9 AND NOT-SDC1	T-Cell, Peripheral	0.7458	0.7097	0.7857
CXCL9 AND NOT-CLDN7	T-Cell, Peripheral	0.7458	0.7097	0.7857
CXCL9 AND NOT-CD34	T-Cell, Peripheral	0.7458	0.7097	0.7857
CXCL9 AND NOT-RNF43	T-Cell, Peripheral	0.7368	0.7241	0.75
CXCL9 AND NOT-CLDN12	T-Cell, Peripheral	0.7333	0.6875	0.7857
CXCL9 AND NOT-CLDN8	T-Cell, Peripheral	0.7333	0.6875	0.7857
CXCL9 AND NOT-CD276	T-Cell, Peripheral	0.7241	0.7	0.75
CXCL9 AND NOT-MUC1	T-Cell, Peripheral	0.7333	0.6875	0.7857
CXCL9 AND NOT-AXL	T-Cell, Peripheral	0.7273	0.7407	0.7143
CXCL9 AND NOT-STEAP2	T-Cell, Peripheral	0.7778	0.8077	0.75
CXCL9 AND NOT-ENG	T-Cell, Peripheral	0.7368	0.7241	0.75
CXCL9 AND NOT-CLDN5	T-Cell, Peripheral	0.7333	0.6875	0.7857
CXCL9 AND NOT-STEAP1	T-Cell, Peripheral	0.7059	0.7826	0.6429
CXCL9 AND NOT-MST1R	T-Cell, Peripheral	0.7333	0.6875	0.7857
CXCL9 AND NOT-TRPM4	T-Cell, Peripheral	0.7037	0.7308	0.6786
CXCL9 AND NOT-ERBB4	T-Cell, Peripheral	0.7018	0.6897	0.7143
CXCL9 AND NOT-PROM1	T-Cell, Peripheral	0.7018	0.6897	0.7143
CXCL9 AND NOT-MUC13	T-Cell, Peripheral	0.7119	0.6774	0.75
CXCL9 AND NOT-SLC34A2	T-Cell, Peripheral	0.6786	0.6786	0.6786
CXCL9 AND NOT-EDNRB	T-Cell, Peripheral	0.6786	0.6786	0.6786
CXCL9 AND NOT-KDR	T-Cell, Peripheral	0.7018	0.6897	0.7143
CXCL9 AND NOT-BMPR1B	T-Cell, Peripheral	0.7018	0.6897	0.7143
CXCL9 AND NOT-LGR5	T-Cell, Peripheral	0.6667	0.6552	0.6786
CXCL9 AND NOT-CLDN11	T-Cell, Peripheral	0.6667	0.7391	0.6071
CXCL9 AND NOT-TPBG	T-Cell, Peripheral	0.7857	0.7857	0.7857
CXCL9 AND NOT-FOLH1	T-Cell, Peripheral	0.6545	0.6667	0.6429
CXCL9 AND NOT-EGFR	T-Cell, Peripheral	0.6545	0.6667	0.6429

FIG. 12 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
CXCL9 AND NOT-CACFD1	T-Cell, Peripheral	0.7586	0.7333	0.7857
CXCL9 AND NOT-B4GALT1	T-Cell, Peripheral	0.8235	0.913	0.75
CXCL9 AND NOT-GJB2	T-Cell, Peripheral	0.8163	0.9524	0.7143
CXCL9 AND NOT-BMP2	T-Cell, Peripheral	0.75	0.75	0.75
CXCL9 AND NOT-NTN4	T-Cell, Peripheral	0.75	0.75	0.75
CXCL9 AND NOT-MFAP3L	T-Cell, Peripheral	0.75	0.75	0.75
CXCL9 AND NOT-ATP1B1	T-Cell, Peripheral	0.75	0.75	0.75
CXCL9 AND NOT-KIT	T-Cell, Peripheral	0.75	0.75	0.75
CXCL9 AND NOT-SLC2A10	T-Cell, Peripheral	0.7778	0.8077	0.75
CXCL9 AND NOT-PERP	T-Cell, Peripheral	0.7586	0.7333	0.7857
CXCL9 AND NOT-PTPRS	T-Cell, Peripheral	0.7586	0.7333	0.7857
CXCL9 AND NOT-LY6K	T-Cell, Peripheral	0.7719	0.7586	0.7857
CXCL9 AND NOT-SLC6A8	T-Cell, Peripheral	0.7458	0.7097	0.7857
CXCL9 AND NOT-SLC3A1	T-Cell, Peripheral	0.7458	0.7097	0.7857
CXCL9 AND NOT-TSPAN9	T-Cell, Peripheral	0.7458	0.7097	0.7857
CXCL9 AND NOT-KITLG	T-Cell, Peripheral	0.7458	0.7097	0.7857
CXCL9 AND NOT-FAT4	T-Cell, Peripheral	0.7458	0.7097	0.7857
CXCL9 AND NOT-ATP10B	T-Cell, Peripheral	0.7458	0.7097	0.7857
CXCL9 AND NOT-CRB3	T-Cell, Peripheral	0.7458	0.7097	0.7857
CXCL9 AND NOT-BAMBI	T-Cell, Peripheral	0.7458	0.7097	0.7857
CXCL9 AND NOT-CYP4F12	T-Cell, Peripheral	0.7458	0.7097	0.7857
CXCL9 AND NOT-KCNK17	T-Cell, Peripheral	0.7458	0.7097	0.7857
CXCL9 AND NOT-LRP11	T-Cell, Peripheral	0.7458	0.7097	0.7857
CXCL9 AND NOT-EPHB3	T-Cell, Peripheral	0.7458	0.7097	0.7857
CXCL9 AND NOT-SPPL3	T-Cell, Peripheral	0.7458	0.7097	0.7857
CXCL9 AND NOT-TMEM231	T-Cell, Peripheral	0.7586	0.7333	0.7857
CXCL9 AND NOT-MCOLN3	T-Cell, Peripheral	0.7719	0.7586	0.7857
CXCL9 AND NOT-ANO6	T-Cell, Peripheral	0.7925	0.84	0.75
CXCL9 AND NOT-PCDHB14	T-Cell, Peripheral	0.7586	0.7333	0.7857
CXCL9 AND NOT-BACE1	T-Cell, Peripheral	0.7407	0.7692	0.7143
CXCL9 AND NOT-F2RL1	T-Cell, Peripheral	0.75	0.75	0.75
CXCL9 AND NOT-GLDN	T-Cell, Peripheral	0.75	0.75	0.75
CXCL9 AND NOT-KCNMB1	T-Cell, Peripheral	0.8148	0.8462	0.7857
CXCL9 AND NOT-WNT5A	T-Cell, Peripheral	0.7857	0.7857	0.7857
CXCL9 AND NOT-KRT5	T-Cell, Peripheral	0.7586	0.7333	0.7857
CXCL9 AND NOT-RHBG	T-Cell, Peripheral	0.7586	0.7333	0.7857
CXCL9 AND NOT-DSC3	T-Cell, Peripheral	0.7586	0.7333	0.7857
CXCL9 AND NOT-NDRG1	T-Cell, Peripheral	0.7719	0.7586	0.7857
CXCL9 AND NOT-SSPN	T-Cell, Peripheral	0.7636	0.7778	0.75
CXCL9 AND NOT-FZD7	T-Cell, Peripheral	0.7368	0.7241	0.75
CXCL9 AND NOT-ATP9B	T-Cell, Peripheral	0.7368	0.7241	0.75
CXCL9 AND NOT-MARCO	T-Cell, Peripheral	0.7368	0.7241	0.75
CXCL9 AND NOT-HAS3	T-Cell, Peripheral	0.7368	0.7241	0.75
CXCL9 AND NOT-TSPAN6	T-Cell, Peripheral	0.7368	0.7241	0.75
CXCL9 AND NOT-THSD7A	T-Cell, Peripheral	0.75	0.75	0.75
CXCL9 AND NOT-UNC5B	T-Cell, Peripheral	0.7586	0.7333	0.7857
CXCL9 AND NOT-STX3	T-Cell, Peripheral	0.7347	0.8571	0.6429
CXCL9 AND NOT-SMO	T-Cell, Peripheral	0.7333	0.6875	0.7857
CXCL9 AND NOT-APCDD1	T-Cell, Peripheral	0.7333	0.6875	0.7857
CXCL9 AND NOT-PLXNA2	T-Cell, Peripheral	0.7333	0.6875	0.7857
CXCL9 AND NOT-IL20RB	T-Cell, Peripheral	0.7333	0.6875	0.7857
CXCL9 AND NOT-ANO10	T-Cell, Peripheral	0.7333	0.6875	0.7857
CXCL9 AND NOT-KCNMB4	T-Cell, Peripheral	0.7458	0.7097	0.7857
CXCL9 AND NOT-PCDHB10	T-Cell, Peripheral	0.7333	0.6875	0.7857
CXCL9 AND NOT-MUC15	T-Cell, Peripheral	0.7333	0.6875	0.7857
CXCL9 AND NOT-ST14	T-Cell, Peripheral	0.7333	0.6875	0.7857
CXCL9 AND NOT-PTPRF	T-Cell, Peripheral	0.7333	0.6875	0.7857
CXCL9 AND NOT-ADRB2	T-Cell, Peripheral	0.7719	0.7586	0.7857
CXCL9 AND NOT-CDH1	T-Cell, Peripheral	0.7333	0.6875	0.7857
CXCL9 AND NOT-CLCN5	T-Cell, Peripheral	0.7333	0.6875	0.7857
CXCL9 AND NOT-OR2F1	T-Cell, Peripheral	0.7333	0.6875	0.7857
CXCL9 AND NOT-DSG2	T-Cell, Peripheral	0.7333	0.6875	0.7857
CXCL9 AND NOT-CELSR1	T-Cell, Peripheral	0.7333	0.6875	0.7857
CXCL9 AND NOT-JAG2	T-Cell, Peripheral	0.7333	0.6875	0.7857
CXCL9 AND NOT-FCGRT	T-Cell, Peripheral	0.7755	0.9048	0.6786
CXCL9 AND NOT-SLC22A23	T-Cell, Peripheral	0.7333	0.6875	0.7857
VANGL2 AND NOT-CADM2	Melanoma	0.9524	1	0.9091
NPC1 AND NOT-NTRK3	Melanoma	0.9091	0.9091	0.9091
VANGL2 AND NOT-TSPAN8	Melanoma	0.9	1	0.8182
VANGL2 AND NOT-ATP1A2	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-CAV3	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-KCNS2	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-TAAR2	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-KCNB1	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-SLC22A24	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-TNFSF18	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-CXCR1	Melanoma	0.9	1	0.8182

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
CXCL9 AND NOT-IL13RA1	T-Cell, Peripheral	0.6522	0.8333	0.5357
CXCL9 AND NOT-MET	T-Cell, Peripheral	0.6667	1	0.5
CXCL9 AND NOT-EPCAM	T-Cell, Peripheral	0.6415	0.68	0.6071
CXCL9 AND NOT-ERBB2	T-Cell, Peripheral	0.64	0.7273	0.5714
CXCL9 AND NOT-CD33	T-Cell, Peripheral	0.6809	0.8421	0.5714
CXCL9 AND NOT-EPHA3	T-Cell, Peripheral	0.6275	0.6957	0.5714
TNFSF11 AND NOT-FXYD1	T-Cell, Peripheral	0.6222	0.8235	0.5
CXCL9 AND NOT-MUC4	T-Cell, Peripheral	0.6182	0.6296	0.6071
CXCL10 AND NOT-CLDN12	T-Cell, Peripheral	0.6154	0.5405	0.7143
CXCL10 AND NOT-SDC1	T-Cell, Peripheral	0.6154	0.5405	0.7143
CXCL9 AND NOT-CSPG4	T-Cell, Peripheral	0.6154	0.6667	0.5714
CXCL10 AND NOT-CLDN7	T-Cell, Peripheral	0.6129	0.5588	0.6786
TNFSF11 AND NOT-NTN4	T-Cell, Peripheral	0.6087	0.7778	0.5
TNFSF11 AND NOT-ITGB5	T-Cell, Peripheral	0.6087	0.7778	0.5
TNFSF11 AND NOT-SSPN	T-Cell, Peripheral	0.619	0.9286	0.4643
CXCL10 AND NOT-CLDN23	T-Cell, Peripheral	0.6957	0.8889	0.5714
CXCL10 AND NOT-STEAP2	T-Cell, Peripheral	0.6316	0.6207	0.6429
CXCL9 AND NOT-IGF1R	T-Cell, Peripheral	0.7368	0.7241	0.75
TNFSF11 AND NOT-RAMP2	T-Cell, Peripheral	0.6087	0.7778	0.5
CXCL9 AND NOT-ABCB5	T-Cell, Peripheral	0.6038	0.64	0.5714
CXCL10 AND NOT-ROR1	T-Cell, Peripheral	0.6452	0.5882	0.7143
TNFSF11 AND NOT-OR7A10	T-Cell, Peripheral	0.6	1	0.4286
TNFSF11 AND NOT-TMEM47	T-Cell, Peripheral	0.6222	0.8235	0.5
EDNRB AND NOT-HTR2A	Melanoma	0.9	1	0.8182
EDNRB AND NOT-IL5RA	Melanoma	0.9	1	0.8182
EDNRB AND NOT-IHH	Melanoma	0.9	1	0.8182
EDNRB AND NOT-AMIGO2	Melanoma	0.9	1	0.8182
EDNRB AND NOT-IFNGR1	Melanoma	0.9	1	0.8182
EDNRB AND NOT-AQP8	Melanoma	0.9	1	0.8182
EDNRB AND NOT-SLC6A19	Melanoma	0.9	1	0.8182
EDNRB AND NOT-FAT3	Melanoma	0.9	1	0.8182
EDNRB AND NOT-ICAM4	Melanoma	0.9	1	0.8182
EDNRB AND NOT-ICAM2	Melanoma	0.9	1	0.8182
EDNRB AND NOT-HTR4	Melanoma	0.9	1	0.8182
EDNRB AND NOT-IL7R	Melanoma	0.9	1	0.8182
EDNRB AND NOT-CXCR1	Melanoma	0.9	1	0.8182
EDNRB AND NOT-AQP1	Melanoma	0.9	1	0.8182
EDNRB AND NOT-KCNB1	Melanoma	0.9	1	0.8182
EDNRB AND NOT-ABCC6	Melanoma	0.9	1	0.8182
EDNRB AND NOT-AQP9	Melanoma	0.9	1	0.8182
EDNRB AND NOT-INSRR	Melanoma	0.9	1	0.8182
EDNRB AND NOT-AQP4	Melanoma	0.9	1	0.8182
EDNRB AND NOT-ANK1	Melanoma	0.9	1	0.8182
EDNRB AND NOT-FFAR2	Melanoma	0.9	1	0.8182
EDNRB AND NOT-GPR31	Melanoma	0.9	1	0.8182
EDNRB AND NOT-GPR21	Melanoma	0.9	1	0.8182
EDNRB AND NOT-TMEM150B	Melanoma	0.9	1	0.8182
EDNRB AND NOT-CCR4	Melanoma	0.9	1	0.8182
EDNRB AND NOT-GPR20	Melanoma	0.9	1	0.8182
EDNRB AND NOT-IZUMO1	Melanoma	0.9	1	0.8182
EDNRB AND NOT-GPR15	Melanoma	0.9	1	0.8182
EDNRB AND NOT-UTS2R	Melanoma	0.9	1	0.8182
EDNRB AND NOT-PRLHR	Melanoma	0.9	1	0.8182
EDNRB AND NOT-NPBWR2	Melanoma	0.9	1	0.8182
EDNRB AND NOT-OR10A4	Melanoma	0.9	1	0.8182
EDNRB AND NOT-GRIA1	Melanoma	0.9	1	0.8182
EDNRB AND NOT-GRIK5	Melanoma	0.9	1	0.8182
EDNRB AND NOT-HCRTR1	Melanoma	0.9	1	0.8182
EDNRB AND NOT-HRH2	Melanoma	0.9	1	0.8182
EDNRB AND NOT-RXFP2	Melanoma	0.9	1	0.8182
EDNRB AND NOT-GYPE	Melanoma	0.9	1	0.8182
EDNRB AND NOT-TRHDE	Melanoma	0.9	1	0.8182
EDNRB AND NOT-GYPB	Melanoma	0.9	1	0.8182
EDNRB AND NOT-GYPA	Melanoma	0.9	1	0.8182
EDNRB AND NOT-LYVE1	Melanoma	0.9	1	0.8182
EDNRB AND NOT-GRPR	Melanoma	0.9	1	0.8182
EDNRB AND NOT-CD46	Melanoma	0.9	1	0.8182
SLC10A4 AND NOT-SSTR1	Neuroblastoma	0.9881	0.9881	0.9881
SLC10A4 AND NOT-CLDN11	Neuroblastoma	0.988	1	0.9762
SLC10A4 AND NOT-TYR	Neuroblastoma	0.9765	0.9651	0.9881
SLC10A4 AND NOT-EGFR	Neuroblastoma	0.9762	0.9762	0.9762
SLC10A4 AND NOT-BCAN	Neuroblastoma	0.9762	0.9762	0.9762
SLC10A4 AND NOT-NCAM1	Neuroblastoma	0.9762	0.9762	0.9762
SLC10A4 AND NOT-ITGB3	Neuroblastoma	0.9765	0.9651	0.9881
SLC10A4 AND NOT-ABCB5	Neuroblastoma	0.9708	0.954	0.9881
SLC10A4 AND NOT-FOLH1	Neuroblastoma	0.9708	0.954	0.9881
SLC10A4 AND NOT-CLDN18	Neuroblastoma	0.9708	0.954	0.9881

FIG. 12 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
C11orf24 AND NOT-OR51B2	Melanoma	0.9	1	0.8182
VANGL2 AND NOT-NRXN1	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-OR4D1	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-GYPB	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-CYP4F12	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-AQP2	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-AMN	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-RXFP2	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-SGMS2	Melanoma	0.9	1	0.8182
SDC3 AND NOT-VIPR2	Melanoma	0.9	1	0.8182
MC1R AND NOT-OR1J2	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-SLC7A13	Melanoma	0.9	1	0.8182
MC1R AND NOT-GPR101	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-NTRK3	Melanoma	0.9	1	0.8182
MC1R AND NOT-AQP10	Melanoma	0.9	1	0.8182
CELSR2 AND NOT-CADM2	Melanoma	0.9524	1	0.9091
CELSR2 AND NOT-ATP1A2	Melanoma	0.9524	1	0.9091
VANGL2 AND NOT-NTRK3	Melanoma	0.9	1	0.8182
TNFSF9 AND NOT-KCNQ3	Melanoma	0.9524	1	0.9091
C11orf24 AND NOT-CHRM2	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-PSENEN	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-IL17RC	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-SLC2A5	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-DCHS2	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-CD300C	Melanoma	0.9	1	0.8182
TNFSF9 AND NOT-FNDC5	Melanoma	0.9524	1	0.9091
C11orf24 AND NOT-CD300LB	Melanoma	0.9	1	0.8182
SDC3 AND NOT-SYT9	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-SCN4A	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-OR2S2	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-KCNK1	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-STX1B	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-IHH	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-GDPD2	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-OR2F2	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-GCGR	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-HCRTR1	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-TAS2R7	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-SLC1A1	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-NPC1L1	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-OXER1	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-TAAR1	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-ATP2A1	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-OR12D2	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-SLC5A5	Melanoma	0.9	1	0.8182
TNFSF9 AND NOT-FRRS1L	Melanoma	0.9524	1	0.9091
C11orf24 AND NOT-CACNG5	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-SLC36A1	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-CHRNB3	Melanoma	0.9	1	0.8182
TNFSF9 AND NOT-SYT9	Melanoma	0.9091	0.9091	0.9091
C11orf24 AND NOT-VIPR2	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-GPR101	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-AQP10	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-F2RL1	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-SLC4A1	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-ADRA1B	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-TM4SF5	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-GRIK5	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-PVRL3	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-PTPRF	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-GRPR	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-OR2F1	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-PIANP	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-SLC5A1	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-PRLHR	Melanoma	0.9	1	0.8182
TNFSF9 AND NOT-SLC7A14	Melanoma	0.9524	1	0.9091
C11orf24 AND NOT-TAAR8	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-GP2	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-SLC14A2	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-CNGB3	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-OR10A5	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-SLC6A20	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-MCHR2	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-GLP1R	Melanoma	0.9	1	0.8182
SLC10A4 AND NOT-GPR62	Neuroblastoma	0.994	1	0.9881
SLC10A4 AND NOT-PAQR6	Neuroblastoma	0.994	1	0.9881
SLC10A4 AND NOT-AQP4	Neuroblastoma	0.994	1	0.9881

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
SLC10A4 AND NOT-ITGB6	Neuroblastoma	0.9701	0.9759	0.9643
SLC10A4 AND NOT-PCYT1A	Neuroblastoma	0.9647	0.9535	0.9762
SLC10A4 AND NOT-CLDN7	Neuroblastoma	0.9639	0.9756	0.9524
SLC10A4 AND NOT-DNAJB8	Neuroblastoma	0.9595	0.9326	0.9881
SLC10A4 AND NOT-CLDN4	Neuroblastoma	0.9591	0.9425	0.9762
CHRNA3 AND NOT-TYR	Neuroblastoma	0.9565	1	0.9167
CHRNA3 AND NOT-EGFR	Neuroblastoma	0.9565	1	0.9167
SLC10A4 AND NOT-ULBP3	Neuroblastoma	0.954	0.9222	0.9881
SLC10A4 AND NOT-HHLA2	Neuroblastoma	0.9535	0.9318	0.9762
SLC10A4 AND NOT-MSLN	Neuroblastoma	0.9529	0.9419	0.9643
SLC10A4 AND NOT-ERBB3	Neuroblastoma	0.9518	0.9634	0.9405
CHRNA3 AND NOT-ULBP3	Neuroblastoma	0.9506	0.9872	0.9167
CHRNA3 AND NOT-DNAJB8	Neuroblastoma	0.9506	0.9872	0.9167
CHRNA3 AND NOT-DPEP1	Neuroblastoma	0.9506	0.9872	0.9167
CHRNA3 AND NOT-CLDN1	Neuroblastoma	0.9506	0.9872	0.9167
CHRNA3 AND NOT-HHLA2	Neuroblastoma	0.9506	0.9872	0.9167
CHRNA3 AND NOT-CLDN18	Neuroblastoma	0.9506	0.9872	0.9167
CHRNA3 AND NOT-ITGB3	Neuroblastoma	0.95	1	0.9048
CHRNA3 AND NOT-BCAN	Neuroblastoma	0.95	1	0.9048
SLC10A4 AND NOT-MST1R	Neuroblastoma	0.9486	0.9121	0.9881
SLC10A4 AND NOT-ERBB4	Neuroblastoma	0.9474	0.931	0.9643
SLC10A4 AND NOT-IL20RA	Neuroblastoma	0.9455	0.963	0.9286
CHRNA3 AND NOT-RNF43	Neuroblastoma	0.9441	0.987	0.9048
CHRNA3 AND NOT-CLDN4	Neuroblastoma	0.9441	0.987	0.9048
CHRNA3 AND NOT-IL20RA	Neuroblastoma	0.9441	0.987	0.9048
CHRNA3 AND NOT-SLC34A2	Neuroblastoma	0.9441	0.987	0.9048
CHRNA3 AND NOT-CA9	Neuroblastoma	0.9441	0.987	0.9048
CHRNA3 AND NOT-PCYT1A	Neuroblastoma	0.9441	0.987	0.9048
CHRNA3 AND NOT-MSLN	Neuroblastoma	0.9441	0.987	0.9048
CHRNA3 AND NOT-ITGB6	Neuroblastoma	0.9434	1	0.8929
CHRNA3 AND NOT-ENPP3	Neuroblastoma	0.9434	1	0.8929
SLC10A4 AND NOT-ENPP3	Neuroblastoma	0.9425	0.9111	0.9762
SLC10A4 AND NOT-RNF43	Neuroblastoma	0.9425	0.9111	0.9762
SLC10A4 AND NOT-TPBG	Neuroblastoma	0.9405	0.9405	0.9405
CHRNA3 AND NOT-CXCR5	Neuroblastoma	0.9375	0.9868	0.8929
SLC10A4 AND NOT-ULBP1	Neuroblastoma	0.9364	0.9101	0.9643
SLC10A4 AND NOT-EDNRB	Neuroblastoma	0.9349	0.9294	0.9405
SLC10A4 AND NOT-ALDH1A1	Neuroblastoma	0.9349	0.9294	0.9405
SLC10A4 AND NOT-FOLR1	Neuroblastoma	0.9325	0.962	0.9048
CHRNA3 AND NOT-ABCB5	Neuroblastoma	0.9308	0.9867	0.881
CHRNA3 AND NOT-FOLR1	Neuroblastoma	0.9308	0.9867	0.881
SLC10A4 AND NOT-TNFRSF13C	Neuroblastoma	0.9294	0.9186	0.9405
SLC10A4 AND NOT-CLDN6	Neuroblastoma	0.9286	0.9286	0.9286
SLC10A4 AND NOT-CXCR5	Neuroblastoma	0.9259	0.9615	0.8929
CHRNA3 AND NOT-EDNRB	Neuroblastoma	0.9241	0.9865	0.869
CHRNA3 AND NOT-AXL	Neuroblastoma	0.9241	0.9865	0.869
CHRNA3 AND NOT-TNFRSF13C	Neuroblastoma	0.9241	0.9865	0.869
VANGL2 AND NOT-EGFR	Neuroblastoma	0.9241	0.9865	0.869
L1CAM AND NOT-GJB6	Neuroblastoma	0.9317	0.974	0.8929
L1CAM AND NOT-CACNA1A	Neuroblastoma	0.9193	0.961	0.881
VANGL2 AND NOT-RNF43	Neuroblastoma	0.9125	0.9605	0.869
PCDHB10 AND NOT-SSTR1	Neuroblastoma	0.9091	1	0.8333
VANGL2 AND NOT-SDC1	Oligodendroglioma	0.8571	0.9231	0.8
PCDHGC3 AND NOT-IL2RA	Oligodendroglioma	0.8889	1	0.8
PCDHGC3 AND NOT-CD72	Oligodendroglioma	0.8571	0.9231	0.8
VANGL2 AND NOT-IL20RA	Oligodendroglioma	0.8571	0.9231	0.8
NLGN3 AND NOT-EPCAM	Oligodendroglioma	0.9333	0.9333	0.9333
PCDHGC3 AND NOT-MET	Oligodendroglioma	0.8889	1	0.8
VANGL2 AND NOT-CLDN1	Oligodendroglioma	0.8571	0.9231	0.8
PCDHGC3 AND NOT-ABCB5	Oligodendroglioma	0.8571	0.9231	0.8
PCDHGC3 AND NOT-ERBB2	Oligodendroglioma	0.8571	0.9231	0.8
PCDHGC3 AND NOT-TRPM4	Oligodendroglioma	0.8276	0.8571	0.8
PCDHGC3 AND NOT-ROR1	Oligodendroglioma	0.8276	0.8571	0.8
PCDHGC3 AND NOT-KDR	Oligodendroglioma	0.8276	0.8571	0.8
VANGL2 AND NOT-TRPM4	Oligodendroglioma	0.8276	0.8571	0.8
PCDHGC3 AND NOT-RAET1E	Oligodendroglioma	0.8276	0.8571	0.8
NLGN3 AND NOT-RNF43	Oligodendroglioma	0.9333	0.9333	0.9333
NLGN3 AND NOT-IL13RA1	Oligodendroglioma	0.8966	0.9286	0.8667
VANGL2 AND NOT-TPBG	Oligodendroglioma	0.8148	0.9167	0.7333
VANGL2 AND NOT-EPCAM	Oligodendroglioma	0.8148	0.9167	0.7333
NLGN3 AND NOT-CLDN8	Oligodendroglioma	0.9333	0.9333	0.9333
NLGN3 AND NOT-CLDN12	Oligodendroglioma	0.9333	0.9333	0.9333
NLGN3 AND NOT-CLDN1	Oligodendroglioma	0.9333	0.9333	0.9333
NLGN3 AND NOT-IL20RA	Oligodendroglioma	0.9333	0.9333	0.9333
NLGN3 AND NOT-CD33	Oligodendroglioma	0.8571	0.9231	0.8
NLGN3 AND NOT-CD72	Oligodendroglioma	0.9333	0.9333	0.9333
NLGN3 AND NOT-TRPM4	Oligodendroglioma	0.9333	0.9333	0.9333

FIG. 12 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
SLC10A4 AND NOT-SLC15A2	Neuroblastoma	0.994	1	0.9881
SLC10A4 AND NOT-SLC2A12	Neuroblastoma	0.994	1	0.9881
SLC10A4 AND NOT-SLC7A10	Neuroblastoma	0.994	1	0.9881
SLC10A4 AND NOT-LRP4	Neuroblastoma	0.994	1	0.9881
SLC10A4 AND NOT-SLC24A4	Neuroblastoma	0.9881	0.9881	0.9881
SLC10A4 AND NOT-SLC1A3	Neuroblastoma	0.9881	0.9881	0.9881
SLC10A4 AND NOT-KCNJ16	Neuroblastoma	0.9881	0.9881	0.9881
SLC10A4 AND NOT-GPR37	Neuroblastoma	0.9881	0.9881	0.9881
SLC10A4 AND NOT-ADRB1	Neuroblastoma	0.988	1	0.9762
SLC10A4 AND NOT-GJB6	Neuroblastoma	0.988	1	0.9762
SLC10A4 AND NOT-ABCG2	Neuroblastoma	0.988	1	0.9762
SLC10A4 AND NOT-IGSF11	Neuroblastoma	0.988	1	0.9762
SLC10A4 AND NOT-KCNN3	Neuroblastoma	0.988	1	0.9762
SLC10A4 AND NOT-CHRM5	Neuroblastoma	0.988	1	0.9762
SLC10A4 AND NOT-SLC4A4	Neuroblastoma	0.9822	0.9765	0.9881
SLC10A4 AND NOT-TRPM6	Neuroblastoma	0.9822	0.9765	0.9881
SLC10A4 AND NOT-CADM4	Neuroblastoma	0.9822	0.9765	0.9881
SLC10A4 AND NOT-AQP8	Neuroblastoma	0.9822	0.9765	0.9881
SLC10A4 AND NOT-GRIA3	Neuroblastoma	0.9822	0.9765	0.9881
SLC10A4 AND NOT-ATP1A2	Neuroblastoma	0.9822	0.9765	0.9881
SLC10A4 AND NOT-GJC2	Neuroblastoma	0.9822	0.9765	0.9881
SLC10A4 AND NOT-GPRC5B	Neuroblastoma	0.9822	0.9765	0.9881
SLC10A4 AND NOT-ATP13A4	Neuroblastoma	0.9822	0.9765	0.9881
SLC10A4 AND NOT-SLC22A23	Neuroblastoma	0.9822	0.9765	0.9881
SLC10A4 AND NOT-SLC44A1	Neuroblastoma	0.9822	0.9765	0.9881
SLC10A4 AND NOT-PRRG1	Neuroblastoma	0.9822	0.9765	0.9881
SLC10A4 AND NOT-RHBDL2	Neuroblastoma	0.9822	0.9765	0.9881
SLC10A4 AND NOT-TGFA	Neuroblastoma	0.9822	0.9765	0.9881
SLC10A4 AND NOT-SLC2A13	Neuroblastoma	0.982	0.988	0.9762
SLC10A4 AND NOT-MFAP3L	Neuroblastoma	0.982	0.988	0.9762
SLC10A4 AND NOT-CACNA1A	Neuroblastoma	0.982	0.988	0.9762
SLC10A4 AND NOT-GLDN	Neuroblastoma	0.982	0.988	0.9762
SLC10A4 AND NOT-UGT8	Neuroblastoma	0.982	0.988	0.9762
SLC10A4 AND NOT-STYK1	Neuroblastoma	0.9818	1	0.9643
SLC10A4 AND NOT-PCDHAC1	Neuroblastoma	0.9822	0.9765	0.9881
SLC10A4 AND NOT-OPN5	Neuroblastoma	0.9822	0.9765	0.9881
SLC10A4 AND NOT-CATSPERG	Neuroblastoma	0.9822	0.9765	0.9881
SLC10A4 AND NOT-ACE2	Neuroblastoma	0.9822	0.9765	0.9881
SLC10A4 AND NOT-ESYT3	Neuroblastoma	0.988	1	0.9762
SLC10A4 AND NOT-CDH26	Neuroblastoma	0.9765	0.9651	0.9881
SLC10A4 AND NOT-SLC13A3	Neuroblastoma	0.9765	0.9651	0.9881
SLC10A4 AND NOT-SLC6A1	Neuroblastoma	0.9765	0.9651	0.9881
SLC10A4 AND NOT-NINJ2	Neuroblastoma	0.9765	0.9651	0.9881
SLC10A4 AND NOT-CLCA2	Neuroblastoma	0.9765	0.9651	0.9881
SLC10A4 AND NOT-RRH	Neuroblastoma	0.9765	0.9651	0.9881
SLC10A4 AND NOT-GAL3ST1	Neuroblastoma	0.9765	0.9651	0.9881
SLC10A4 AND NOT-AGPAT3	Neuroblastoma	0.9765	0.9651	0.9881
SLC10A4 AND NOT-CRHR2	Neuroblastoma	0.9765	0.9651	0.9881
SLC10A4 AND NOT-CD99L2	Neuroblastoma	0.9765	0.9651	0.9881
SLC10A4 AND NOT-S1PR5	Neuroblastoma	0.9765	0.9651	0.9881
SLC10A4 AND NOT-FGFR2	Neuroblastoma	0.9765	0.9651	0.9881
SLC10A4 AND NOT-LRP2	Neuroblastoma	0.9765	0.9651	0.9881
SLC10A4 AND NOT-NTRK2	Neuroblastoma	0.9765	0.9651	0.9881
SLC10A4 AND NOT-SLC6A4	Neuroblastoma	0.9762	0.9762	0.9762
SLC10A4 AND NOT-UPK1B	Neuroblastoma	0.9762	0.9762	0.9762
SLC10A4 AND NOT-SLC6A20	Neuroblastoma	0.9762	0.9762	0.9762
SLC10A4 AND NOT-DIO2	Neuroblastoma	0.9762	0.9762	0.9762
SLC10A4 AND NOT-SLC26A9	Neuroblastoma	0.9762	0.9762	0.9762
SLC10A4 AND NOT-TREML1	Neuroblastoma	0.9762	0.9762	0.9762
SLC10A4 AND NOT-SLC5A1	Neuroblastoma	0.9762	0.9762	0.9762
SLC10A4 AND NOT-GABRA6	Neuroblastoma	0.9762	0.9762	0.9762
PCDHGC3 AND NOT-DSG2	Oligodendroglioma	0.8889	1	0.8
PCDHGC3 AND NOT-IL1R1	Oligodendroglioma	0.8889	1	0.8
PCDHGC3 AND NOT-IL1RL2	Oligodendroglioma	0.8889	1	0.8
PCDHGC3 AND NOT-GABRE	Oligodendroglioma	0.8889	1	0.8
PCDHGC3 AND NOT-CORIN	Oligodendroglioma	0.8889	1	0.8
PCDHGC3 AND NOT-DUOX2	Oligodendroglioma	0.8889	1	0.8
PCDHGC3 AND NOT-DSC2	Oligodendroglioma	0.8889	1	0.8
PCDHGC3 AND NOT-SGMS2	Oligodendroglioma	0.8889	1	0.8
VANGL2 AND NOT-SLITRK6	Oligodendroglioma	0.8571	0.9231	0.8
PCDHGC3 AND NOT-CLMP	Oligodendroglioma	0.8571	0.9231	0.8
PCDHGC3 AND NOT-ITGA11	Oligodendroglioma	0.8571	0.9231	0.8
PCDHGC3 AND NOT-TRPM7	Oligodendroglioma	0.8889	1	0.8
VANGL2 AND NOT-FXYD3	Oligodendroglioma	0.8571	0.9231	0.8
PCDHGC3 AND NOT-ICAM3	Oligodendroglioma	0.8571	0.9231	0.8
VANGL2 AND NOT-KRT5	Oligodendroglioma	0.8571	0.9231	0.8
VANGL2 AND NOT-PERP	Oligodendroglioma	0.8571	0.9231	0.8

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
NLGN3 AND NOT-SDC1	Oligodendroglioma	0.9333	0.9333	0.9333
VANGL2 AND NOT-CLDN8	Oligodendroglioma	0.8571	0.9231	0.8
PTPRZ1 AND NOT-EPCAM	Oligodendroglioma	0.8148	0.9167	0.7333
NLGN3 AND NOT-RAET1E	Oligodendroglioma	0.8889	1	0.8
NLGN3 AND NOT-ENPP3	Oligodendroglioma	0.9333	0.9333	0.9333
NLGN3 AND NOT-CD79A	Oligodendroglioma	0.8571	0.9231	0.8
VANGL2 AND NOT-ERBB2	Oligodendroglioma	0.8571	0.9231	0.8
NLGN3 AND NOT-SLC34A2	Oligodendroglioma	0.9333	0.9333	0.9333
NLGN3 AND NOT-ULBP2	Oligodendroglioma	0.9333	0.9333	0.9333
NLGN3 AND NOT-CLDN9	Oligodendroglioma	0.9333	0.9333	0.9333
NLGN3 AND NOT-CD34	Oligodendroglioma	0.9333	0.9333	0.9333
NLGN3 AND NOT-P2RX5	Oligodendroglioma	0.9333	0.9333	0.9333
PCDHGC3 AND NOT-CD34	Oligodendroglioma	0.8	0.8	0.8
VANGL2 AND NOT-RNF43	Oligodendroglioma	0.8	0.8	0.8
GPR173 AND NOT-ERBB2	Oligodendroglioma	0.8	1	0.6667
GPR173 AND NOT-ENG	Oligodendroglioma	0.8	1	0.6667
NLGN3 AND NOT-CD52	Oligodendroglioma	0.8	1	0.6667
GPR173 AND NOT-CLDN1	Oligodendroglioma	0.8	1	0.6667
VANGL2 AND NOT-CD52	Oligodendroglioma	0.8148	0.9167	0.7333
PTPRZ1 AND NOT-TPBG	Oligodendroglioma	0.8148	0.9167	0.7333
SLC4A4 AND NOT-EPCAM	Oligodendroglioma	0.8889	1	0.8
NLGN3 AND NOT-TNFRSF8	Oligodendroglioma	0.9333	0.9333	0.9333
NLGN3 AND NOT-GUCY2C	Oligodendroglioma	0.9333	0.9333	0.9333
NLGN3 AND NOT-CD180	Oligodendroglioma	0.9333	0.9333	0.9333
NLGN3 AND NOT-IL3RA	Oligodendroglioma	0.9333	0.9333	0.9333
NLGN3 AND NOT-STEAP1	Oligodendroglioma	0.9333	0.9333	0.9333
NLGN3 AND NOT-ENG	Oligodendroglioma	0.9333	0.9333	0.9333
NLGN3 AND NOT-STEAP2	Oligodendroglioma	0.9333	0.9333	0.9333
NLGN3 AND NOT-ITGB6	Oligodendroglioma	0.9333	0.9333	0.9333
NLGN3 AND NOT-ABCB5	Oligodendroglioma	0.9333	0.9333	0.9333
NLGN3 AND NOT-FCRL1	Oligodendroglioma	0.9333	0.9333	0.9333
NLGN3 AND NOT-SSTR4	Oligodendroglioma	0.9333	0.9333	0.9333
NLGN3 AND NOT-CD79B	Oligodendroglioma	0.9333	0.9333	0.9333
NLGN3 AND NOT-SSTR5	Oligodendroglioma	0.9333	0.9333	0.9333
NLGN3 AND NOT-SSTR3	Oligodendroglioma	0.9333	0.9333	0.9333
NLGN3 AND NOT-TNFRSF17	Oligodendroglioma	0.9333	0.9333	0.9333
NLGN3 AND NOT-VTCN1	Oligodendroglioma	0.9333	0.9333	0.9333
NLGN3 AND NOT-FCRL5	Oligodendroglioma	0.9333	0.9333	0.9333
NLGN3 AND NOT-AXL	Oligodendroglioma	0.8966	0.9286	0.8667
NLGN3 AND NOT-PMEL	Oligodendroglioma	0.9333	0.9333	0.9333
NLGN3 AND NOT-TNFRSF10A	Oligodendroglioma	0.9333	0.9333	0.9333
NLGN3 AND NOT-MSLN	Oligodendroglioma	0.8966	0.9286	0.8667
NLGN3 AND NOT-MUC1	Oligodendroglioma	0.9333	0.9333	0.9333
NLGN3 AND NOT-CLDN6	Oligodendroglioma	0.9333	0.9333	0.9333
NLGN3 AND NOT-TNFRSF13C	Oligodendroglioma	0.9333	0.9333	0.9333
NLGN3 AND NOT-SSTR1	Oligodendroglioma	0.9333	0.9333	0.9333
NLGN3 AND NOT-TNFSF11	Oligodendroglioma	0.9333	0.9333	0.9333
NLGN3 AND NOT-MUC13	Oligodendroglioma	0.9333	0.9333	0.9333
NLGN3 AND NOT-CR2	Oligodendroglioma	0.9333	0.9333	0.9333
NLGN3 AND NOT-FOLH1	Oligodendroglioma	0.9333	0.9333	0.9333
NLGN3 AND NOT-CLDN2	Oligodendroglioma	0.9333	0.9333	0.9333
NLGN3 AND NOT-ITGB3	Oligodendroglioma	0.9333	0.9333	0.9333
NLGN3 AND NOT-CLDN18	Oligodendroglioma	0.9333	0.9333	0.9333
KCNK15 AND NOT-TPBG	Ovarian	0.8571	0.75	1
KCNK15 AND NOT-EDNRB	Ovarian	0.8696	0.9091	0.8333
KCNK15 AND NOT-AXL	Ovarian	0.8889	0.8	1
KCNK15 AND NOT-IL3RA	Ovarian	0.9167	0.9167	0.9167
KCNK15 AND NOT-CD52	Ovarian	0.8276	0.7059	1
KCNK15 AND NOT-GPNMB	Ovarian	0.96	0.9231	1
KCNK15 AND NOT-VCAM1	Ovarian	0.8	0.6667	1
KCNK15 AND NOT-CLDN23	Ovarian	0.8	0.6667	1
MUC16 AND NOT-MTUS1	Ovarian	0.8	1	0.6667
KCNK15 AND NOT-ENG	Ovarian	0.8	0.6667	1
MUC16 AND NOT-CD99	Ovarian	0.8	1	0.6667
MUC16 AND NOT-PKD2	Ovarian	0.8	1	0.6667
KCNK15 AND NOT-IL20RA	Ovarian	0.7742	0.6316	1
KCNK15 AND NOT-FAP	Ovarian	0.7742	0.6316	1
KCNK15 AND NOT-CLDN8	Ovarian	0.7742	0.6316	1
KCNK15 AND NOT-PROM1	Ovarian	0.7742	0.6316	1
KCNK15 AND NOT-ROR1	Ovarian	0.7742	0.6316	1
KCNK15 AND NOT-TRPM4	Ovarian	0.7742	0.6316	1
KCNK15 AND NOT-KDR	Ovarian	0.7742	0.6316	1
KCNK15 AND NOT-CLDN1	Ovarian	0.7742	0.6316	1
KCNK15 AND NOT-CD70	Ovarian	0.7742	0.6316	1
KCNK15 AND NOT-MET	Ovarian	0.7742	0.6316	1
KCNK15 AND NOT-ULBP2	Ovarian	0.7742	0.6316	1
KCNK15 AND NOT-SLC39A6	Ovarian	0.7742	0.6316	1

FIG. 12 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
PCDHGC3 AND NOT-OR51M1	Oligodendroglioma	0.8571	0.9231	0.8
VANGL2 AND NOT-DSG2	Oligodendroglioma	0.8571	0.9231	0.8
PCDHGC3 AND NOT-SGCG	Oligodendroglioma	0.8889	1	0.8
PCDHGC3 AND NOT-SLCO2A1	Oligodendroglioma	0.8571	0.9231	0.8
PCDHGC3 AND NOT-NPR3	Oligodendroglioma	0.8889	1	0.8
PCDHGC3 AND NOT-CD40	Oligodendroglioma	0.8571	0.9231	0.8
VANGL2 AND NOT-TACSTD2	Oligodendroglioma	0.8571	0.9231	0.8
PCDHGC3 AND NOT-SPPL2A	Oligodendroglioma	0.8889	1	0.8
VANGL2 AND NOT-AQP3	Oligodendroglioma	0.8571	0.9231	0.8
VANGL2 AND NOT-DUOX1	Oligodendroglioma	0.8571	0.9231	0.8
PCDHGC3 AND NOT-LRRC32	Oligodendroglioma	0.8571	0.9231	0.8
PCDHGC3 AND NOT-AGTR1	Oligodendroglioma	0.8571	0.9231	0.8
VANGL2 AND NOT-GPR87	Oligodendroglioma	0.8571	0.9231	0.8
PTPRZ1 AND NOT-MTUS1	Oligodendroglioma	0.8889	1	0.8
PCDHGC3 AND NOT-GJA5	Oligodendroglioma	0.8889	1	0.8
PCDHGC3 AND NOT-LTA	Oligodendroglioma	0.8571	0.9231	0.8
PCDHGC3 AND NOT-JPH2	Oligodendroglioma	0.8571	0.9231	0.8
PCDHGC3 AND NOT-RTP1	Oligodendroglioma	0.8571	0.9231	0.8
PCDHGC3 AND NOT-DSC3	Oligodendroglioma	0.8571	0.9231	0.8
PCDHGC3 AND NOT-VNN3	Oligodendroglioma	0.8889	1	0.8
PCDHGC3 AND NOT-FUT1	Oligodendroglioma	0.8571	0.9231	0.8
PCDHGC3 AND NOT-ITGA9	Oligodendroglioma	0.8889	1	0.8
PCDHGC3 AND NOT-AQP7	Oligodendroglioma	0.8889	1	0.8
PCDHGC3 AND NOT-APOB	Oligodendroglioma	0.8889	1	0.8
PCDHGC3 AND NOT-LNPEP	Oligodendroglioma	0.8571	0.9231	0.8
PCDHGC3 AND NOT-KCNQ1	Oligodendroglioma	0.8571	0.9231	0.8
PCDHGC3 AND NOT-SLCO5A1	Oligodendroglioma	0.8889	1	0.8
PCDHGC3 AND NOT-P2RY2	Oligodendroglioma	0.8889	1	0.8
PCDHGC3 AND NOT-SGCA	Oligodendroglioma	0.8889	1	0.8
PCDHGC3 AND NOT-SCN5A	Oligodendroglioma	0.8571	0.9231	0.8
PCDHGC3 AND NOT-C14orf180	Oligodendroglioma	0.8889	1	0.8
PCDHGC3 AND NOT-EPHB4	Oligodendroglioma	0.8571	0.9231	0.8
PCDHGC3 AND NOT-FGG	Oligodendroglioma	0.8889	1	0.8
PCDHGC3 AND NOT-SLC5A1	Oligodendroglioma	0.8889	1	0.8
VANGL2 AND NOT-ST14	Oligodendroglioma	0.8571	0.9231	0.8
VANGL2 AND NOT-DSC3	Oligodendroglioma	0.8571	0.9231	0.8
VANGL2 AND NOT-SCNN1A	Oligodendroglioma	0.8571	0.9231	0.8
VANGL2 AND NOT-SPINT1	Oligodendroglioma	0.8571	0.9231	0.8
VANGL2 AND NOT-EMP2	Oligodendroglioma	0.8571	0.9231	0.8
PCDHGC3 AND NOT-ACVRL1	Oligodendroglioma	0.8889	1	0.8
PCDHGC3 AND NOT-LILRB1	Oligodendroglioma	0.8462	1	0.7333
PCDHGC3 AND NOT-PERP	Oligodendroglioma	0.8462	1	0.7333
PCDHGC3 AND NOT-IL7R	Oligodendroglioma	0.8462	1	0.7333
PCDHGC3 AND NOT-MYADM	Oligodendroglioma	0.8462	1	0.7333
PCDHGC3 AND NOT-ITPR3	Oligodendroglioma	0.8462	1	0.7333
PCDHGC3 AND NOT-ESYT2	Oligodendroglioma	0.8462	1	0.7333
PCDHGC3 AND NOT-PDCD1LG2	Oligodendroglioma	0.8462	1	0.7333
PCDHGC3 AND NOT-ADCY9	Oligodendroglioma	0.8462	1	0.7333
PCDHGC3 AND NOT-IL1R2	Oligodendroglioma	0.8462	1	0.7333
PCDHGC3 AND NOT-TRPV3	Oligodendroglioma	0.8462	1	0.7333
PTPRZ1 AND NOT-ATP1B1	Oligodendroglioma	0.8462	1	0.7333
PCDHGC3 AND NOT-ZDHHC5	Oligodendroglioma	0.8462	1	0.7333
VANGL2 AND NOT-IL1R1	Oligodendroglioma	0.8571	0.9231	0.8
VANGL2 AND NOT-SGMS2	Oligodendroglioma	0.8571	0.9231	0.8
PCDHGC3 AND NOT-EMP2	Oligodendroglioma	0.8462	1	0.7333
KCNK15 AND NOT-BMP2	Ovarian	0.96	0.9231	1
KCNK15 AND NOT-SLC31A2	Ovarian	0.96	0.9231	1
KCNK15 AND NOT-CAV1	Ovarian	0.96	0.9231	1
KCNK15 AND NOT-TSPAN5	Ovarian	0.9231	0.8571	1
KCNK15 AND NOT-DSC3	Ovarian	0.96	0.9231	1
KCNK15 AND NOT-SLC16A7	Ovarian	1	1	1
KCNK15 AND NOT-AQP9	Ovarian	0.9231	0.8571	1
KCNK15 AND NOT-KCNJ8	Ovarian	0.96	0.9231	1
KCNK15 AND NOT-SFRP1	Ovarian	0.96	0.9231	1
KCNK15 AND NOT-ATP10D	Ovarian	0.9167	0.9167	0.9167
KCNK15 AND NOT-MARCO	Ovarian	0.9167	0.9167	0.9167
KCNK15 AND NOT-PMP22	Ovarian	0.96	0.9231	1
KCNK15 AND NOT-PARM1	Ovarian	0.88	0.8462	0.9167
KCNK15 AND NOT-ABCA8	Ovarian	0.88	0.8462	0.9167
KCNK15 AND NOT-FCGR3B	Ovarian	0.96	0.9231	1
KCNK15 AND NOT-STX8	Ovarian	0.96	0.9231	1
KCNK15 AND NOT-TNFSF13B	Ovarian	0.96	0.9231	1
KCNK15 AND NOT-LYVE1	Ovarian	0.8889	0.8	1
KCNK15 AND NOT-LPAR5	Ovarian	0.9231	0.8571	1
KCNK15 AND NOT-PTGER3	Ovarian	0.8889	0.8	1
KCNK15 AND NOT-ABHD6	Ovarian	0.8696	0.9091	0.8333
KCNK15 AND NOT-AQP1	Ovarian	1	1	1

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
KCNK15 AND NOT-SSTR5	Ovarian	0.7692	0.7143	0.8333
MUC16 AND NOT-TMEM59	Ovarian	0.7619	0.8889	0.6667
MUC16 AND NOT-BTN3A2	Ovarian	0.7619	0.8889	0.6667
MUC16 AND NOT-F3	Ovarian	0.7619	0.8889	0.6667
KCNK15 AND NOT-TNFRSF8	Ovarian	0.7586	0.6471	0.9167
KCNK15 AND NOT-CSPG4	Ovarian	0.7407	0.6667	0.8333
KCNK15 AND NOT-CLDN11	Ovarian	0.7333	0.6111	0.9167
MUC16 AND NOT-ATP2C2	Ovarian	0.7273	0.8	0.6667
MUC16 AND NOT-DLL1	Ovarian	0.7273	0.8	0.6667
MUC16 AND NOT-PIGR	Ovarian	0.7273	0.8	0.6667
MUC16 AND NOT-KIAA1324	Ovarian	0.7273	0.8	0.6667
MUC16 AND NOT-GOLM1	Ovarian	0.7273	0.8	0.6667
MUC16 AND NOT-TSPAN8	Ovarian	0.7273	0.8	0.6667
MUC16 AND NOT-STIM2	Ovarian	0.7273	0.8	0.6667
MUC16 AND NOT-APCDD1	Ovarian	0.7273	0.8	0.6667
MUC16 AND NOT-AQP3	Ovarian	0.7273	0.8	0.6667
MUC16 AND NOT-MACF1	Ovarian	0.7273	0.8	0.6667
KCNK15 AND NOT-SSTR4	Ovarian	0.7143	0.625	0.8333
MUC16 AND NOT-LPAR6	Ovarian	0.7619	0.8889	0.6667
MUC16 AND NOT-ICAM4	Ovarian	0.7368	1	0.5833
MUC16 AND NOT-C10orf54	Ovarian	0.7619	0.8889	0.6667
MUC16 AND NOT-BAMBI	Ovarian	0.7	0.875	0.5833
MUC16 AND NOT-KIT	Ovarian	0.7	0.875	0.5833
MUC16 AND NOT-FXYD1	Ovarian	0.7	0.875	0.5833
MUC16 AND NOT-KRT5	Ovarian	0.7273	0.8	0.6667
MUC16 AND NOT-PTPRZ1	Ovarian	0.6957	0.7273	0.6667
MUC16 AND NOT-CDHR3	Ovarian	0.6957	0.7273	0.6667
MUC16 AND NOT-COMT	Ovarian	0.6957	0.7273	0.6667
MUC16 AND NOT-ITGA2	Ovarian	0.6957	0.7273	0.6667
KCNK15 AND NOT-CXCR5	Ovarian	0.6957	0.7273	0.6667
MUC16 AND NOT-DNAJC1	Ovarian	0.6957	0.7273	0.6667
MUC16 AND NOT-STIM1	Ovarian	0.6957	0.7273	0.6667
MUC16 AND NOT-ACSL1	Ovarian	0.6957	0.7273	0.6667
MUC16 AND NOT-STOM	Ovarian	0.6957	0.7273	0.6667
MUC16 AND NOT-LRP11	Ovarian	0.6957	0.7273	0.6667
MUC16 AND NOT-ATP6AP2	Ovarian	0.7273	0.8	0.6667
KCNK15 AND NOT-IL11RA	Ovarian	0.7692	0.7143	0.8333
MUC16 AND NOT-PCDHGA10	Ovarian	0.7	0.875	0.5833
MUC16 AND NOT-HLA-B	Ovarian	0.7273	0.8	0.6667
MUC16 AND NOT-CLCA2	Ovarian	0.6667	0.6667	0.6667
MUC16 AND NOT-LRRTM2	Ovarian	0.6667	1	0.5
MUC16 AND NOT-IL4R	Ovarian	0.6667	0.7778	0.5833
MUC16 AND NOT-SLC20A1	Ovarian	0.6667	0.6667	0.6667
MUC16 AND NOT-AKAP1	Ovarian	0.6667	0.6667	0.6667
MUC16 AND NOT-ENTPD3	Ovarian	0.6667	0.6667	0.6667
MUC16 AND NOT-LITAF	Ovarian	0.6667	0.6667	0.6667
MUC16 AND NOT-SERINC3	Ovarian	0.6667	0.6667	0.6667
MUC16 AND NOT-SLC41A1	Ovarian	0.6667	0.6667	0.6667
MUC16 AND NOT-ATP12A	Ovarian	0.6667	0.6667	0.6667
MUC16 AND NOT-ARL6IP5	Ovarian	0.6667	0.6667	0.6667
MUC16 AND NOT-LPAR1	Ovarian	0.6667	1	0.5
MUC16 AND NOT-CHIC2	Ovarian	0.6667	0.6667	0.6667
MUC16 AND NOT-SCN4B	Ovarian	0.6667	0.6667	0.6667
KCNK15 AND NOT-SSTR1	Ovarian	0.6667	0.6	0.75
MUC16 AND NOT-SLC16A7	Ovarian	0.8	1	0.6667
MUC16 AND NOT-AQP5	Ovarian	0.7273	0.8	0.6667
MUC16 AND NOT-CYP4F12	Ovarian	0.7273	0.8	0.6667
MUC16 AND NOT-CAV2	Ovarian	0.7368	1	0.5833
KCNK15 AND NOT-ULBP1	Ovarian	0.6429	0.5625	0.75
MUC16 AND NOT-ASPH	Ovarian	0.6667	0.6667	0.6667
MUC16 AND NOT-KCNK1	Ovarian	0.64	0.6154	0.6667
MUC16 AND NOT-CYB5R1	Ovarian	0.64	0.6154	0.6667
MUC16 AND NOT-LRMP	Ovarian	0.64	0.6154	0.6667
MUC16 AND NOT-SLC31A2	Ovarian	0.64	0.6154	0.6667
OR51E2 AND NOT-KDR	Prostate	1	1	1
OR51E2 AND NOT-AXL	Prostate	0.8889	0.8	1
OR51E2 AND NOT-TNC	Prostate	0.8889	0.8	1
OR51E2 AND NOT-CSPG4	Prostate	0.8889	0.8	1
OR51E2 AND NOT-ST8SIA1	Prostate	0.8889	0.8	1
OR51E2 AND NOT-TNFRSF17	Prostate	0.8889	0.8	1
OR51E2 AND NOT-CLDN5	Prostate	0.8889	0.8	1
OR51E2 AND NOT-ITGB6	Prostate	0.8889	0.8	1
OR51E2 AND NOT-DKK1	Prostate	0.8889	0.8	1
OR51E2 AND NOT-SLAMF7	Prostate	0.8889	0.8	1
OR51E2 AND NOT-CLDN1	Prostate	0.8889	0.8	1
OR51E2 AND NOT-MUC16	Prostate	0.8889	0.8	1
OR51E2 AND NOT-CD33	Prostate	0.8889	0.8	1

FIG. 12 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
KCNK15 AND NOT-HAS1	Ovarian	0.8696	0.9091	0.8333
KCNK15 AND NOT-CXCR2	Ovarian	0.8696	0.9091	0.8333
KCNK15 AND NOT-EMB	Ovarian	0.8889	0.8	1
KCNK15 AND NOT-JAM3	Ovarian	0.9231	0.8571	1
KCNK15 AND NOT-IL22RA1	Ovarian	0.9167	0.9167	0.9167
KCNK15 AND NOT-PTPRZ1	Ovarian	0.96	0.9231	1
KCNK15 AND NOT-LRFN5	Ovarian	0.8571	0.75	1
KCNK15 AND NOT-TNFSF12	Ovarian	0.8571	1	0.75
KCNK15 AND NOT-PTGDR	Ovarian	0.8571	0.75	1
KCNK15 AND NOT-STIM2	Ovarian	0.8571	0.75	1
KCNK15 AND NOT-ADRA2A	Ovarian	0.8571	0.75	1
KCNK15 AND NOT-TMEM150C	Ovarian	0.8571	0.75	1
KCNK15 AND NOT-CNIH3	Ovarian	0.8571	0.75	1
KCNK15 AND NOT-PPAP2A	Ovarian	0.8571	0.75	1
KCNK15 AND NOT-SLC9A9	Ovarian	0.8889	0.8	1
KCNK15 AND NOT-DDR2	Ovarian	0.9231	0.8571	1
KCNK15 AND NOT-CSF2RB	Ovarian	0.8696	0.9091	0.8333
KCNK15 AND NOT-IL6ST	Ovarian	0.8889	0.8	1
KCNK15 AND NOT-C10orf54	Ovarian	1	1	1
KCNK15 AND NOT-SDC2	Ovarian	0.8571	0.75	1
KCNK15 AND NOT-ITM2A	Ovarian	0.96	0.9231	1
KCNK15 AND NOT-CMKLR1	Ovarian	0.9091	1	0.8333
KCNK15 AND NOT-SELL	Ovarian	0.9167	0.9167	0.9167
KCNK15 AND NOT-CLEC10A	Ovarian	0.8462	0.7857	0.9167
KCNK15 AND NOT-SLC51A	Ovarian	0.8462	0.7857	0.9167
KCNK15 AND NOT-PDPN	Ovarian	0.8571	0.75	1
KCNK15 AND NOT-AMICA1	Ovarian	0.9167	0.9167	0.9167
KCNK15 AND NOT-NAALADL1	Ovarian	0.8462	0.7857	0.9167
KCNK15 AND NOT-TLR2	Ovarian	0.96	0.9231	1
KCNK15 AND NOT-ACSL1	Ovarian	0.96	0.9231	1
KCNK15 AND NOT-DST	Ovarian	0.8889	0.8	1
KCNK15 AND NOT-P2RY14	Ovarian	0.9167	0.9167	0.9167
KCNK15 AND NOT-EMCN	Ovarian	0.9231	0.8571	1
KCNK15 AND NOT-LPAR1	Ovarian	1	1	1
KCNK15 AND NOT-ITGAM	Ovarian	0.96	0.9231	1
KCNK15 AND NOT-C3AR1	Ovarian	0.96	0.9231	1
KCNK15 AND NOT-FPR1	Ovarian	1	1	1
KCNK15 AND NOT-KIT	Ovarian	0.96	0.9231	1
KCNK15 AND NOT-APH1B	Ovarian	0.9231	0.8571	1
KCNK15 AND NOT-PCDH17	Ovarian	0.8333	0.8333	0.8333
KCNK15 AND NOT-IL7R	Ovarian	0.96	0.9231	1
KCNK15 AND NOT-CCR1	Ovarian	0.9167	0.9167	0.9167
KCNK15 AND NOT-JAM2	Ovarian	0.9091	1	0.8333
KCNK15 AND NOT-SULF2	Ovarian	0.8276	0.7059	1
KCNK15 AND NOT-RECK	Ovarian	0.8276	0.7059	1
KCNK15 AND NOT-CNST	Ovarian	0.8276	0.7059	1
KCNK15 AND NOT-COLEC12	Ovarian	0.8276	0.7059	1
KCNK15 AND NOT-STX7	Ovarian	0.8276	0.7059	1
KCNK15 AND NOT-SEMA5A	Ovarian	0.8276	0.7059	1
KCNK15 AND NOT-SLC41A2	Ovarian	0.8276	0.7059	1
KCNK15 AND NOT-BDKRB1	Ovarian	0.8276	0.7059	1
KCNK15 AND NOT-STX2	Ovarian	0.8276	0.7059	1
KCNK15 AND NOT-NT5E	Ovarian	0.8276	0.7059	1
KCNK15 AND NOT-PROS1	Ovarian	0.8276	0.7059	1
KCNK15 AND NOT-CAV2	Ovarian	0.8276	0.7059	1
KCNK15 AND NOT-HMOX1	Ovarian	0.8889	0.8	1
KCNK15 AND NOT-BDKRB2	Ovarian	0.8696	0.9091	0.8333
KCNK15 AND NOT-PDGFRA	Ovarian	0.96	0.9231	1
KCNK15 AND NOT-HCST	Ovarian	0.88	0.8462	0.9167
KCNK15 AND NOT-FPR2	Ovarian	0.88	0.8462	0.9167
KCNK15 AND NOT-ATP8B4	Ovarian	0.8333	0.8333	0.8333
KCNK15 AND NOT-HVCN1	Ovarian	0.8276	0.7059	1
KCNK15 AND NOT-VAMP5	Ovarian	0.8889	0.8	1
KCNK15 AND NOT-GPR34	Ovarian	0.8276	0.7059	1
KCNK15 AND NOT-VNN2	Ovarian	0.8889	0.8	1
KCNK15 AND NOT-PTPRO	Ovarian	0.8571	0.75	1
KCNK15 AND NOT-LRRTM2	Ovarian	0.9565	1	0.9167
KCNK15 AND NOT-VSTM4	Ovarian	0.88	0.8462	0.9167
KCNK15 AND NOT-C5AR1	Ovarian	0.9565	1	0.9167
KCNK15 AND NOT-GLIPR1	Ovarian	0.96	0.9231	1
KCNK15 AND NOT-HEPH	Ovarian	0.8571	1	0.75
KCNK15 AND NOT-VAMP7	Ovarian	0.8571	0.75	1
KCNK15 AND NOT-ENPP2	Ovarian	0.96	0.9231	1
KCNK15 AND NOT-CD53	Ovarian	0.96	0.9231	1
KCNK15 AND NOT-GPR171	Ovarian	0.96	0.9231	1
OR51E2 AND NOT-LSAMP	Prostate	1	1	1
OR51E2 AND NOT-SGCD	Prostate	1	1	1

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
OR51E2 AND NOT-CD72	Prostate	0.8889	0.8	1
OR51E2 AND NOT-CD79B	Prostate	0.8889	0.8	1
STEAP2 AND NOT-SLC52A3	Prostate	0.8889	0.8	1
STEAP2 AND NOT-FZD1	Prostate	0.8889	0.8	1
OR51E2 AND NOT-CD180	Prostate	0.8889	0.8	1
OR51E2 AND NOT-MOK	Prostate	0.8889	0.8	1
STEAP2 AND NOT-FADS2	Prostate	0.8889	0.8	1
OR51E2 AND NOT-LGR5	Prostate	0.8889	0.8	1
OR51E2 AND NOT-L1CAM	Prostate	0.8889	0.8	1
OR51E2 AND NOT-SLC34A2	Prostate	0.8889	0.8	1
OR51E2 AND NOT-MST1R	Prostate	0.8889	0.8	1
OR51E2 AND NOT-HLA-DOB	Prostate	0.8889	0.8	1
OR51E2 AND NOT-IGF1R	Prostate	0.8889	0.8	1
OR51E2 AND NOT-IL11RA	Prostate	0.8889	0.8	1
OR51E2 AND NOT-CLDN11	Prostate	0.8889	0.8	1
OR51E2 AND NOT-FOLR1	Prostate	0.8889	0.8	1
OR51E2 AND NOT-MUC1	Prostate	0.8889	0.8	1
STEAP2 AND NOT-SLC2A5	Prostate	1	1	1
OR51E2 AND NOT-MUC4	Prostate	0.8889	0.8	1
OR51E2 AND NOT-VTCN1	Prostate	0.8889	0.8	1
OR51E2 AND NOT-CEACAM6	Prostate	0.8889	0.8	1
STEAP2 AND NOT-PRRT2	Prostate	0.8571	1	0.75
OR51E1 AND NOT-CSPG4	Prostate	0.8571	1	0.75
STEAP2 AND NOT-SGCD	Prostate	1	1	1
OR51E2 AND NOT-PMEL	Prostate	0.8571	1	0.75
STEAP2 AND NOT-PCDH9	Prostate	0.8571	1	0.75
STEAP2 AND NOT-PTPRF	Prostate	0.8571	1	0.75
STEAP2 AND NOT-IL6ST	Prostate	0.8571	1	0.75
STEAP2 AND NOT-KCNK3	Prostate	0.8571	1	0.75
OR51E2 AND NOT-DPEP1	Prostate	0.8571	1	0.75
OR51E2 AND NOT-ITGB3	Prostate	0.8571	1	0.75
OR51E2 AND NOT-B4GALNT1	Prostate	0.8571	1	0.75
OR51E1 AND NOT-CLDN5	Prostate	0.8571	1	0.75
OR51E2 AND NOT-CXCR5	Prostate	0.8889	0.8	1
OR51E2 AND NOT-CEACAM5	Prostate	0.8889	0.8	1
OR51E2 AND NOT-SSTR3	Prostate	0.8889	0.8	1
OR51E2 AND NOT-AFP	Prostate	0.8889	0.8	1
OR51E2 AND NOT-CA9	Prostate	0.8889	0.8	1
OR51E2 AND NOT-SSTR4	Prostate	0.8889	0.8	1
OR51E2 AND NOT-CR2	Prostate	0.8889	0.8	1
OR51E2 AND NOT-SSX1	Prostate	0.8889	0.8	1
OR51E2 AND NOT-ROR1	Prostate	0.8889	0.8	1
OR51E2 AND NOT-BCAN	Prostate	0.8889	0.8	1
OR51E2 AND NOT-DNAJB8	Prostate	0.8889	0.8	1
OR51E2 AND NOT-DLL3	Prostate	0.8889	0.8	1
OR51E2 AND NOT-SSTR2	Prostate	0.8889	0.8	1
OR51E2 AND NOT-VCAM1	Prostate	0.8889	0.8	1
OR51E2 AND NOT-SSTR1	Prostate	0.8889	0.8	1
OR51E2 AND NOT-SST	Prostate	0.8889	0.8	1
OR51E2 AND NOT-FCRL2	Prostate	0.8889	0.8	1
OR51E2 AND NOT-GAGE1	Prostate	0.8889	0.8	1
OR51E2 AND NOT-CLDN4	Prostate	0.8889	0.8	1
OR51E2 AND NOT-IL2RA	Prostate	0.8889	0.8	1
OR51E2 AND NOT-TNFRSF13C	Prostate	0.8889	0.8	1
OR51E2 AND NOT-ABCB5	Prostate	0.8889	0.8	1
OR51E2 AND NOT-WT1	Prostate	0.8889	0.8	1
OR51E2 AND NOT-BIRC5	Prostate	0.8889	0.8	1
OR51E2 AND NOT-ULBP3	Prostate	0.8889	0.8	1
OR51E2 AND NOT-CTAG2	Prostate	0.8889	0.8	1
OR51E2 AND NOT-SEMA5B	Prostate	0.8889	0.8	1
OR51E2 AND NOT-GUCY2C	Prostate	0.8889	0.8	1
OR51E2 AND NOT-GPA33	Prostate	0.8889	0.8	1
OR51E2 AND NOT-RAET1E	Prostate	0.8889	0.8	1
OR51E2 AND NOT-MSLN	Prostate	0.8889	0.8	1
OR51E2 AND NOT-THY1	Prostate	0.8889	0.8	1
OR51E2 AND NOT-CD52	Prostate	0.8889	0.8	1
STEAP2 AND NOT-KL	Prostate	0.8571	1	0.75
OR51E2 AND NOT-TYR	Prostate	0.8889	0.8	1
OR51E2 AND NOT-FCRL5	Prostate	0.8889	0.8	1
OR51E2 AND NOT-CD22	Prostate	0.8889	0.8	1
STEAP2 AND NOT-ANTXR1	Prostate	0.8571	1	0.75
OR51E2 AND NOT-ULBP2	Prostate	0.8889	0.8	1
STEAP2 AND NOT-IL6R	Prostate	0.8571	1	0.75
OR51E2 AND NOT-CLDN7	Prostate	0.8889	0.8	1
OR51E2 AND NOT-CD79A	Prostate	0.8889	0.8	1
OR51E2 AND NOT-CD70	Prostate	0.8889	0.8	1
OR51E2 AND NOT-CD37	Prostate	0.8889	0.8	1

FIG. 12 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
OR51E2 AND NOT-SLC2A5	Prostate	1	1	1
OR51E2 AND NOT-ACE	Prostate	1	1	1
OR51E2 AND NOT-SI	Prostate	1	1	1
OR51E2 AND NOT-DAGLA	Prostate	1	1	1
OR51E2 AND NOT-CDH6	Prostate	0.8889	0.8	1
OR51E2 AND NOT-TSPAN11	Prostate	0.8889	0.8	1
OR51E2 AND NOT-STX2	Prostate	0.8889	0.8	1
OR51E2 AND NOT-SDC3	Prostate	0.8889	0.8	1
OR51E2 AND NOT-CD69	Prostate	0.8889	0.8	1
OR51E2 AND NOT-GPR65	Prostate	0.8889	0.8	1
OR51E2 AND NOT-SLC5A4	Prostate	0.8889	0.8	1
OR51E2 AND NOT-GPR87	Prostate	0.8889	0.8	1
OR51E2 AND NOT-FAT4	Prostate	0.8889	0.8	1
OR51E2 AND NOT-TMEM67	Prostate	0.8889	0.8	1
OR51E2 AND NOT-CD1C	Prostate	0.8889	0.8	1
OR51E2 AND NOT-SLC5A3	Prostate	0.8889	0.8	1
OR51E2 AND NOT-MYOF	Prostate	0.8889	0.8	1
OR51E2 AND NOT-SV2B	Prostate	0.8889	0.8	1
OR51E2 AND NOT-LPAR3	Prostate	0.8889	0.8	1
OR51E2 AND NOT-KCNE4	Prostate	0.8889	0.8	1
OR51E2 AND NOT-CEACAM1	Prostate	0.8889	0.8	1
OR51E2 AND NOT-BMP2	Prostate	0.8889	0.8	1
OR51E2 AND NOT-CHL1	Prostate	0.8889	0.8	1
OR51E2 AND NOT-SLC23A1	Prostate	0.8889	0.8	1
OR51E2 AND NOT-TM7SF3	Prostate	0.8889	0.8	1
OR51E2 AND NOT-DUOXA1	Prostate	1	1	1
OR51E2 AND NOT-CLCA4	Prostate	0.8889	0.8	1
OR51E2 AND NOT-LEPR	Prostate	0.8889	0.8	1
ANO7 AND NOT-CAV1	Prostate	0.8889	0.8	1
OR51E2 AND NOT-ADCY3	Prostate	0.8889	0.8	1
ANO7 AND NOT-ATP1A2	Prostate	0.8889	0.8	1
OR51E2 AND NOT-ITGB5	Prostate	0.8889	0.8	1
OR51E2 AND NOT-SGCA	Prostate	0.8889	0.8	1
OR51E2 AND NOT-SLC47A1	Prostate	0.8889	0.8	1
OR51E2 AND NOT-PRPH2	Prostate	0.8889	0.8	1
OR51E2 AND NOT-LRFN4	Prostate	0.8889	0.8	1
OR51E2 AND NOT-TRPM6	Prostate	0.8889	0.8	1
OR51E2 AND NOT-NKG7	Prostate	0.8889	0.8	1
OR51E2 AND NOT-WDR19	Prostate	0.8889	0.8	1
OR51E2 AND NOT-GGT7	Prostate	0.8889	0.8	1
OR51E2 AND NOT-SEMA4F	Prostate	0.8889	0.8	1
OR51E2 AND NOT-SLC38A4	Prostate	0.8889	0.8	1
OR51E2 AND NOT-LRRC8A	Prostate	0.8889	0.8	1
OR51E2 AND NOT-PMEPA1	Prostate	0.8889	0.8	1
OR51E2 AND NOT-ITGA2	Prostate	0.8889	0.8	1
OR51E2 AND NOT-ATP1A2	Prostate	0.8889	0.8	1
OR51E2 AND NOT-ADRB1	Prostate	0.8889	0.8	1
OR51E2 AND NOT-CELSR2	Prostate	0.8889	0.8	1
OR51E2 AND NOT-ENPP2	Prostate	0.8889	0.8	1
OR51E2 AND NOT-FCGRT	Prostate	0.8889	0.8	1
ANO7 AND NOT-FADS2	Prostate	0.8889	0.8	1
OR51E2 AND NOT-SLC18A2	Prostate	0.8889	0.8	1
OR51E2 AND NOT-DNAJC5	Prostate	0.8889	0.8	1
OR51E2 AND NOT-PANX1	Prostate	0.8889	0.8	1
OR51E2 AND NOT-TENM2	Prostate	0.8889	0.8	1
OR51E2 AND NOT-SLC52A3	Prostate	0.8889	0.8	1
OR51E2 AND NOT-SLC8A1	Prostate	0.8889	0.8	1
ANO7 AND NOT-SLC52A3	Prostate	0.8889	0.8	1
OR51E2 AND NOT-SHISA6	Prostate	0.8889	0.8	1
ANO7 AND NOT-LSAMP	Prostate	1	1	1
OR51E2 AND NOT-SLC16A14	Prostate	0.8889	0.8	1
OR51E2 AND NOT-STX4	Prostate	0.8889	0.8	1
OR51E2 AND NOT-AQP1	Prostate	0.8889	0.8	1
OR51E2 AND NOT-SLC24A3	Prostate	0.8889	0.8	1
OR51E2 AND NOT-SEMA6D	Prostate	0.8889	0.8	1
OR51E2 AND NOT-DAGLB	Prostate	0.8889	0.8	1
OR51E2 AND NOT-SDC4	Prostate	0.8889	0.8	1
OR51E2 AND NOT-ADRA2C	Prostate	0.8889	0.8	1
OR51E2 AND NOT-CAV2	Prostate	0.8889	0.8	1
OR51E2 AND NOT-SLC6A16	Prostate	0.8889	0.8	1
OR51E2 AND NOT-CTSZ	Prostate	0.8889	0.8	1
OR51E2 AND NOT-EMP2	Prostate	0.8889	0.8	1
OR51E2 AND NOT-CYP4F2	Prostate	1	1	1
OR51E2 AND NOT-BCAM	Prostate	0.8889	0.8	1
OR51E2 AND NOT-JPH1	Prostate	0.8889	0.8	1
OR51E2 AND NOT-ABHD6	Prostate	0.8889	0.8	1
OR51E2 AND NOT-VAMP1	Prostate	0.8889	0.8	1

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
OR51E2 AND NOT-TNFRSF8	Prostate	0.8889	0.8	1
OR51E2 AND NOT-MS4A1	Prostate	0.8889	0.8	1
ADAM12 AND NOT-ERBB3	Sarcoma	0.8333	0.7895	0.8824
FAP AND NOT-SLC8A1	Sarcoma	0.8276	1	0.7059
FAP AND NOT-SLC25A4	Sarcoma	0.8276	1	0.7059
FAP AND NOT-TSPAN12	Sarcoma	0.8276	1	0.7059
FAP AND NOT-ABCB1	Sarcoma	0.8667	1	0.7647
FAP AND NOT-SCN7A	Sarcoma	0.8667	1	0.7647
FAP AND NOT-SCN3A	Sarcoma	0.8276	1	0.7059
FAP AND NOT-SLMAP	Sarcoma	0.8667	1	0.7647
FAP AND NOT-SLC4A3	Sarcoma	0.8667	1	0.7647
FAP AND NOT-NAALAD2	Sarcoma	0.8276	1	0.7059
FAP AND NOT-PRIMA1	Sarcoma	0.8276	1	0.7059
FAP AND NOT-NPY1R	Sarcoma	0.8276	1	0.7059
FAP AND NOT-SIGLEC6	Sarcoma	0.8667	1	0.7647
FAP AND NOT-PRLR	Sarcoma	0.8276	1	0.7059
FAP AND NOT-MRAP2	Sarcoma	0.8276	1	0.7059
ADAM12 AND NOT-RNF43	Sarcoma	0.8421	0.7619	0.9412
FAP AND NOT-SLC2A12	Sarcoma	0.7857	1	0.6471
FAP AND NOT-EMCN	Sarcoma	0.7857	1	0.6471
FAP AND NOT-BCAM	Sarcoma	0.7857	1	0.6471
FAP AND NOT-PCDH20	Sarcoma	0.7857	1	0.6471
FAP AND NOT-TSPAN2	Sarcoma	0.7857	1	0.6471
FAP AND NOT-KCNMB1	Sarcoma	0.7857	1	0.6471
FAP AND NOT-PARM1	Sarcoma	0.7857	1	0.6471
FAP AND NOT-PLA2R1	Sarcoma	0.7857	1	0.6471
FAP AND NOT-SLCO2A1	Sarcoma	0.7857	1	0.6471
FAP AND NOT-HTR2B	Sarcoma	0.7857	1	0.6471
FAP AND NOT-SLC3A1	Sarcoma	0.7857	1	0.6471
FAP AND NOT-GDPD2	Sarcoma	0.8667	1	0.7647
FAP AND NOT-CDH3	Sarcoma	0.7857	1	0.6471
FAP AND NOT-TSPAN7	Sarcoma	0.8667	1	0.7647
FAP AND NOT-GPM6A	Sarcoma	0.8276	1	0.7059
ADAM12 AND NOT-PROM1	Sarcoma	0.8095	0.68	1
FAP AND NOT-CLSTN2	Sarcoma	0.8276	1	0.7059
OSMR AND NOT-FOLR1	Sarcoma	0.8387	0.9286	0.7647
FAP AND NOT-CACNA1D	Sarcoma	0.8667	1	0.7647
FAP AND NOT-KCNQ4	Sarcoma	0.8667	1	0.7647
FAP AND NOT-KCNC4	Sarcoma	0.8276	1	0.7059
FAP AND NOT-SLC6A2	Sarcoma	0.7857	1	0.6471
FAP AND NOT-SLC8A2	Sarcoma	0.8276	1	0.7059
FAP AND NOT-CD9	Sarcoma	0.7857	1	0.6471
FAP AND NOT-ATP7B	Sarcoma	0.8667	1	0.7647
FAP AND NOT-P2RX1	Sarcoma	0.7857	1	0.6471
FAP AND NOT-TACR1	Sarcoma	0.7857	1	0.6471
FAP AND NOT-PCDHA10	Sarcoma	0.8276	1	0.7059
FAP AND NOT-PERP	Sarcoma	0.7857	1	0.6471
FAP AND NOT-SLC6A16	Sarcoma	0.7857	1	0.6471
FAP AND NOT-SGCA	Sarcoma	0.8276	1	0.7059
FAP AND NOT-JPH2	Sarcoma	0.8667	1	0.7647
FAP AND NOT-TACSTD2	Sarcoma	0.7857	1	0.6471
FAP AND NOT-VIPR2	Sarcoma	0.7857	1	0.6471
OSMR AND NOT-ITGB6	Sarcoma	0.7586	0.9167	0.6471
FAP AND NOT-PTPRS	Sarcoma	0.7857	1	0.6471
FAP AND NOT-PIGR	Sarcoma	0.8276	1	0.7059
FAP AND NOT-KCNH2	Sarcoma	0.7857	1	0.6471
FAP AND NOT-ATP1B2	Sarcoma	0.7857	1	0.6471
FAP AND NOT-TRPC4	Sarcoma	0.8667	1	0.7647
FAP AND NOT-SLC47A1	Sarcoma	0.7857	1	0.6471
FAP AND NOT-CXADR	Sarcoma	0.7857	1	0.6471
FAP AND NOT-GPR162	Sarcoma	0.7857	1	0.6471
ADAM12 AND NOT-ITGB6	Sarcoma	0.8235	0.8235	0.8235
FAP AND NOT-NTRK3	Sarcoma	0.7857	1	0.6471
FAP AND NOT-GPR37	Sarcoma	0.8667	1	0.7647
FAP AND NOT-KCNB1	Sarcoma	0.8276	1	0.7059
FAP AND NOT-ITM2B	Sarcoma	0.7857	1	0.6471
FAP AND NOT-ZDHHC21	Sarcoma	0.8276	1	0.7059
FAP AND NOT-MCHR1	Sarcoma	0.8276	1	0.7059
FAP AND NOT-RAMP1	Sarcoma	0.7857	1	0.6471
FAP AND NOT-NAALADL1	Sarcoma	0.7857	1	0.6471
FAP AND NOT-ART4	Sarcoma	0.7857	1	0.6471
FAP AND NOT-ADCY4	Sarcoma	0.8276	1	0.7059
FAP AND NOT-ISLR2	Sarcoma	0.7857	1	0.6471
FAP AND NOT-KCNN2	Sarcoma	0.7857	1	0.6471
FAP AND NOT-DPP6	Sarcoma	0.8276	1	0.7059
FAP AND NOT-SLC27A5	Sarcoma	0.8276	1	0.7059
OSMR AND NOT-RNF43	Sarcoma	0.7586	0.9167	0.6471

FIG. 12 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
OR51E2 AND NOT-ATP10A	Prostate	0.8889	0.8	1
OR51E2 AND NOT-NRSN2	Prostate	0.8889	0.8	1
OR51E2 AND NOT-FAM57A	Prostate	0.8889	0.8	1
OR51E2 AND NOT-TMEM97	Prostate	0.8889	0.8	1
OR51E2 AND NOT-DLL1	Prostate	0.8889	0.8	1
ADAM12 AND NOT-CDH1	Sarcoma	0.8333	0.7895	0.8824
OSMR AND NOT-SCN7A	Sarcoma	0.9032	1	0.8235
OSMR AND NOT-F11R	Sarcoma	0.875	0.9333	0.8235
ADAM12 AND NOT-F11R	Sarcoma	0.9444	0.8947	1
DDR2 AND NOT-SLC25A4	Sarcoma	0.8276	1	0.7059
ADAM12 AND NOT-LSR	Sarcoma	0.8387	0.9286	0.7647
ADAM12 AND NOT-TACSTD2	Sarcoma	0.8333	0.7895	0.8824
SLC2A10 AND NOT-F11R	Sarcoma	0.8276	1	0.7059
OSMR AND NOT-CDH1	Sarcoma	0.8	0.9231	0.7059
ADAM12 AND NOT-PARM1	Sarcoma	0.8108	0.75	0.8824
ADAM12 AND NOT-NAALAD2	Sarcoma	0.7805	0.6667	0.9412
ADAM12 AND NOT-PERP	Sarcoma	0.8	0.7778	0.8235
ADAM12 AND NOT-FXYD3	Sarcoma	0.8333	0.7895	0.8824
OSMR AND NOT-TACSTD2	Sarcoma	0.7742	0.8571	0.7059
ADAM12 AND NOT-SCN7A	Sarcoma	0.7727	0.6296	1
ADAM12 AND NOT-ATP7B	Sarcoma	0.8205	0.7273	0.9412
DDR2 AND NOT-SCN7A	Sarcoma	0.8	0.9231	0.7059
ADAM12 AND NOT-CLEC1A	Sarcoma	0.8205	0.7273	0.9412
OSMR AND NOT-FXYD3	Sarcoma	0.8	0.9231	0.7059
DDR2 AND NOT-TSPAN7	Sarcoma	0.7586	0.9167	0.6471
OSMR AND NOT-BCAM	Sarcoma	0.7586	0.9167	0.6471
OSMR AND NOT-LSR	Sarcoma	0.7586	0.9167	0.6471
DDR2 AND NOT-NAALAD2	Sarcoma	0.7586	0.9167	0.6471
VMP1 AND NOT-F11R	Sarcoma	0.875	0.9333	0.8235
OSMR AND NOT-LRP5	Sarcoma	0.8276	1	0.7059
ADAM12 AND NOT-TM7SF2	Sarcoma	0.7778	0.7368	0.8235
OSMR AND NOT-ABCB1	Sarcoma	0.875	0.9333	0.8235
OSMR AND NOT-SLC19A3	Sarcoma	0.8485	0.875	0.8235
OSMR AND NOT-MUC15	Sarcoma	0.8387	0.9286	0.7647
OSMR AND NOT-CRB3	Sarcoma	0.875	0.9333	0.8235
OSMR AND NOT-TMEM88	Sarcoma	0.7586	0.9167	0.6471
OSMR AND NOT-SCNN1B	Sarcoma	0.875	0.9333	0.8235
OSMR AND NOT-SPINT1	Sarcoma	0.875	0.9333	0.8235
OSMR AND NOT-PCDH1	Sarcoma	0.8	0.9231	0.7059
OSMR AND NOT-PARM1	Sarcoma	0.8387	0.9286	0.7647
OSMR AND NOT-CLEC1A	Sarcoma	0.875	0.9333	0.8235
ADAM12 AND NOT-NRCAM	Sarcoma	0.8	0.7778	0.8235
OSMR AND NOT-SEMA6A	Sarcoma	0.8667	1	0.7647
ADAM12 AND NOT-SPINT1	Sarcoma	0.8718	0.7727	1
DDR2 AND NOT-PARM1	Sarcoma	0.7407	1	0.5882
DDR2 AND NOT-TM7SF2	Sarcoma	0.7407	1	0.5882
ADAM12 AND NOT-TSPAN13	Sarcoma	0.7907	0.6538	1
ADAM12 AND NOT-SLC24A3	Sarcoma	0.7857	1	0.6471
OSMR AND NOT-KCNK3	Sarcoma	0.8667	1	0.7647
OSMR AND NOT-PERP	Sarcoma	0.7586	0.9167	0.6471
OSMR AND NOT-TM7SF2	Sarcoma	0.7407	1	0.5882
OSMR AND NOT-ACPP	Sarcoma	0.875	0.9333	0.8235
OSMR AND NOT-CSF3R	Sarcoma	0.8125	0.8667	0.7647
ADAM12 AND NOT-TMEM150C	Sarcoma	0.7368	0.6667	0.8235
DDR2 AND NOT-ENPP4	Sarcoma	0.7586	0.9167	0.6471
OSMR AND NOT-CEACAM1	Sarcoma	0.7586	0.9167	0.6471
OSMR AND NOT-SLC6A2	Sarcoma	0.7857	1	0.6471
DDR2 AND NOT-TSPAN8	Sarcoma	0.7333	0.8462	0.6471
OSMR AND NOT-MGST2	Sarcoma	0.7333	0.8462	0.6471
OSMR AND NOT-SLC25A4	Sarcoma	0.7586	0.9167	0.6471
ADAM12 AND NOT-AQP3	Sarcoma	0.7333	0.8462	0.6471
OSMR AND NOT-MTUS1	Sarcoma	0.7857	1	0.6471
OSMR AND NOT-TSPAN31	Sarcoma	0.7742	0.8571	0.7059
ADAM12 AND NOT-CEACAM1	Sarcoma	0.7429	0.7222	0.7647
ADAM12 AND NOT-SLC19A3	Sarcoma	0.7317	0.625	0.8824
ADAM12 AND NOT-SLC6A2	Sarcoma	0.8485	0.875	0.8235
ADAM12 AND NOT-SIGLEC6	Sarcoma	0.85	0.7391	1
OSMR AND NOT-TSPAN12	Sarcoma	0.875	0.9333	0.8235
OSMR AND NOT-TMPRSS2	Sarcoma	0.8	0.9231	0.7059
ADAM12 AND NOT-CDH3	Sarcoma	0.8108	0.75	0.8824
OSMR AND NOT-FZD5	Sarcoma	0.8125	0.8667	0.7647
OSMR AND NOT-HAS3	Sarcoma	0.8387	0.9286	0.7647
ADAM12 AND NOT-CNGA1	Sarcoma	0.7273	0.75	0.7059
OSMR AND NOT-LRP2	Sarcoma	0.8387	0.9286	0.7647
OSMR AND NOT-TMEM150C	Sarcoma	0.7586	0.9167	0.6471
ADAM12 AND NOT-ADRA2C	Sarcoma	0.7273	0.75	0.7059
OSMR AND NOT-PIGR	Sarcoma	0.8667	1	0.7647

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
OSMR AND NOT-EDNRB	Sarcoma	0.875	0.9333	0.8235
FAP AND NOT-NLGN3	Sarcoma	0.8276	1	0.7059
FAP AND NOT-ABCG2	Sarcoma	0.7857	1	0.6471
FAP AND NOT-CD101	Sarcoma	0.8667	1	0.7647
FAP AND NOT-DIO3	Sarcoma	0.7857	1	0.6471
FAP AND NOT-ADRA2A	Sarcoma	0.7857	1	0.6471
FAP AND NOT-MPZ	Sarcoma	0.7857	1	0.6471
FAP AND NOT-TMEM123	Sarcoma	0.8667	1	0.7647
FAP AND NOT-TMEM198	Sarcoma	0.7857	1	0.6471
FAP AND NOT-STYK1	Sarcoma	0.8276	1	0.7059
FAP AND NOT-AVPR1A	Sarcoma	0.8276	1	0.7059
FAP AND NOT-CORIN	Sarcoma	0.8667	1	0.7647
FAP AND NOT-SLC26A10	Sarcoma	0.7857	1	0.6471
ADAM12 AND NOT-EPCAM	Sarcoma	0.85	0.7391	1
FAP AND NOT-TMPRSS6	Sarcoma	0.7857	1	0.6471
FAP AND NOT-SYT2	Sarcoma	0.8276	1	0.7059
FAP AND NOT-SLC6A5	Sarcoma	0.7407	1	0.5882
FAP AND NOT-MYADM	Sarcoma	0.7407	1	0.5882
FAP AND NOT-TSPAN8	Sarcoma	0.7407	1	0.5882
FAP AND NOT-SUSD3	Sarcoma	0.7407	1	0.5882
FAP AND NOT-CNR1	Sarcoma	0.7407	1	0.5882
FAP AND NOT-CELSR2	Sarcoma	0.7407	1	0.5882
CLDN18 AND NOT-SCNN1G	Stomach	0.809	0.7826	0.8372
CLDN18 AND NOT-SCNN1B	Stomach	0.7912	0.75	0.8372
MUC13 AND NOT-BEST4	Stomach	0.8718	0.9714	0.7907
CLDN18 AND NOT-ABCA3	Stomach	0.7912	0.75	0.8372
CLDN18 AND NOT-GPM6A	Stomach	0.7865	0.7609	0.814
CLDN18 AND NOT-MARCO	Stomach	0.7912	0.75	0.8372
CLDN18 AND NOT-NPR3	Stomach	0.7692	0.8571	0.6977
MUC13 AND NOT-SLC51B	Stomach	0.8831	1	0.7907
CLDN18 AND NOT-CD300LG	Stomach	0.7692	0.7292	0.814
CLDN18 AND NOT-SGCG	Stomach	0.8235	0.8333	0.814
CLDN18 AND NOT-ADRB2	Stomach	0.7674	0.7674	0.7674
CLDN18 AND NOT-SLC6A4	Stomach	0.7674	0.7674	0.7674
CLDN18 AND NOT-CADM1	Stomach	0.7865	0.7609	0.814
CLDN18 AND NOT-EVA1A	Stomach	0.766	0.7059	0.8372
CLDN18 AND NOT-SLCO4C1	Stomach	0.766	0.7059	0.8372
CLDN18 AND NOT-ADRB1	Stomach	0.7865	0.7609	0.814
CLDN18 AND NOT-DUOX1	Stomach	0.7609	0.7143	0.814
MUC13 AND NOT-SLC16A9	Stomach	0.9136	0.9737	0.8605
CLDN18 AND NOT-SLC11A1	Stomach	0.7778	0.7447	0.814
CLDN18 AND NOT-ESYT3	Stomach	0.7586	0.75	0.7674
MUC13 AND NOT-SCNN1B	Stomach	0.9882	1	0.9767
CLDN18 AND NOT-GLDN	Stomach	0.7529	0.7619	0.7442
CLDN18 AND NOT-SLC15A2	Stomach	0.7527	0.7	0.814
CLDN18 AND NOT-GPM6B	Stomach	0.7857	0.8049	0.7674
CLDN18 AND NOT-GJA5	Stomach	0.75	0.7333	0.7674
CLDN18 AND NOT-SUSD2	Stomach	0.75	0.7333	0.7674
CLDN18 AND NOT-KLRF1	Stomach	0.7826	0.7347	0.8372
CLDN18 AND NOT-CAV2	Stomach	0.764	0.7391	0.7907
CLDN18 AND NOT-AMICA1	Stomach	0.7556	0.7234	0.7907
CLDN18 AND NOT-TREM1	Stomach	0.7473	0.7083	0.7907
CLDN18 AND NOT-DUOXA1	Stomach	0.747	0.775	0.7209
CLDN18 AND NOT-ICAM4	Stomach	0.7714	1	0.6279
CLDN18 AND NOT-CLEC12A	Stomach	0.7692	0.7292	0.814
CLDN18 AND NOT-CAV1	Stomach	0.7865	0.7609	0.814
CLDN18 AND NOT-OLR1	Stomach	0.75	0.7333	0.7674
MST1R AND NOT-SCNN1B	Stomach	0.9318	0.9111	0.9535
CLDN7 AND NOT-SCNN1B	Stomach	0.9195	0.9091	0.9302
CLDN18 AND NOT-SCN7A	Stomach	0.7619	0.7805	0.7442
CLDN18 AND NOT-ST3GAL5	Stomach	0.7556	0.7234	0.7907
CLDN18 AND NOT-COLEC12	Stomach	0.7391	0.6939	0.7907
CLDN18 AND NOT-TMEM100	Stomach	0.7955	0.7778	0.814
MUC13 AND NOT-TSPAN1	Stomach	0.866	0.7778	0.9767
CLDN18 AND NOT-LIFR	Stomach	0.8182	0.8	0.8372
CLDN18 AND NOT-ECSCR	Stomach	0.7356	0.7273	0.7442
CLDN18 AND NOT-KL	Stomach	0.7356	0.7273	0.7442
CLDN18 AND NOT-CDH5	Stomach	0.7586	0.75	0.7674
CLDN18 AND NOT-EMCN	Stomach	0.7333	0.7021	0.7674
CLDN18 AND NOT-SIRPB1	Stomach	0.7333	0.7021	0.7674
CLDN18 AND NOT-RAMP3	Stomach	0.7356	0.7273	0.7442
CEACAM5 AND NOT-SCNN1B	Stomach	0.878	0.9231	0.8372
CLDN18 AND NOT-LYVE1	Stomach	0.7674	0.7674	0.7674
CLDN18 AND NOT-CSF3R	Stomach	0.7391	0.6939	0.7907
CLDN18 AND NOT-CA4	Stomach	0.7273	0.8235	0.6512
CLDN18 AND NOT-MSR1	Stomach	0.7273	0.7111	0.7442
CLDN18 AND NOT-CD300LF	Stomach	0.7416	0.7174	0.7674

FIG. 12 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
ADAM12 AND NOT-PROM2	Sarcoma	0.8649	0.8	0.9412
ADAM12 AND NOT-NPY1R	Sarcoma	0.7222	0.6842	0.7647
OSMR AND NOT-KCNK5	Sarcoma	0.7586	0.9167	0.6471
OSMR AND NOT-F3	Sarcoma	0.8	0.9231	0.7059
MMP14 AND NOT-BCAM	Sarcoma	0.7407	1	0.5882
MMP14 AND NOT-LSR	Sarcoma	0.7407	1	0.5882
MMP14 AND NOT-SIGLEC6	Sarcoma	0.7407	1	0.5882
SLC2A10 AND NOT-PIGR	Sarcoma	0.8276	1	0.7059
OSMR AND NOT-SLCO4A1	Sarcoma	0.8387	0.9286	0.7647
ADAM12 AND NOT-SLC44A5	Sarcoma	0.7273	0.5926	0.9412
OSMR AND NOT-TSPAN8	Sarcoma	0.7857	1	0.6471
OSMR AND NOT-SLC6A4	Sarcoma	0.8387	0.9286	0.7647
GPRC5A AND NOT-SCNN1B	Stomach	0.9535	0.9535	0.9535
ST14 AND NOT-SCNN1B	Stomach	0.9111	0.8723	0.9535
GPRC5A AND NOT-SCNN1G	Stomach	0.7342	0.8056	0.6744
TSPAN1 AND NOT-SCNN1B	Stomach	0.75	0.6087	0.9767
LSR AND NOT-SCNN1B	Stomach	0.9383	1	0.8837
LSR AND NOT-ABHD6	Stomach	0.7536	1	0.6047
SLC39A4 AND NOT-CLCA4	Stomach	0.8235	0.8333	0.814
LSR AND NOT-SLC16A9	Stomach	0.9136	0.9737	0.8605
B3GNT3 AND NOT-SCNN1B	Stomach	0.8958	0.8113	1
LSR AND NOT-CLCA4	Stomach	0.8378	1	0.7209
LSR AND NOT-BEST4	Stomach	0.9268	0.9744	0.8837
VSIG1 AND NOT-GP1BA	Stomach	0.6866	0.9583	0.5349
SMAGP AND NOT-SCNN1B	Stomach	0.9767	0.9767	0.9767
LSR AND NOT-ST6GALNAC6	Stomach	0.8571	0.9706	0.7674
B3GNT3 AND NOT-CLCA4	Stomach	0.7353	1	0.5814
NOT-LIFR AND PODXL	Colon	0.8889	1	0.8
NOT-LIFR AND TMEM47	Colon	0.8889	1	0.8
NOT-SLC16A7 AND EDAR	Colon	0.8889	1	0.8
NOT-LIFR AND EVA1A	Colon	0.8889	1	0.8
NOT-MUC15 AND CDH3	Colon	0.9091	0.8333	1
NOT-LIFR AND PCDH18	Colon	0.8889	1	0.8
NOT-LIFR AND EPHB4	Colon	0.8333	0.7143	1
NOT-SLC16A7 AND TMEM97	Colon	1	1	1
NOT-LIFR AND PDGFRB	Colon	0.8333	0.7143	1
NOT-SLC7A2 AND CDH3	Colon	0.8333	0.7143	1
NOT-LIFR AND COLEC12	Colon	0.8	0.8	0.8
NOT-AQP3 AND SLC10A3	Leiomyosarcoma	0.7273	0.8889	0.6154
NOT-CXADR AND DCBLD2	Liposarcoma	0.7143	0.7353	0.6944
NOT-PERP AND FAM57A	Liposarcoma	0.7813	0.8929	0.6944
NOT-CXADR AND SMO	Liposarcoma	0.6885	0.84	0.5833
NOT-NTRK2 AND NTM	Lung Adenocarcinoma	0.7287	0.7966	0.6714
NOT-CLCA4 AND PVRL4	Lung Adenocarcinoma	0.6512	0.7119	0.6
NOT-PLP1 AND NTM	Lung Adenocarcinoma	0.7397	0.7105	0.7714
NOT-NTRK2 AND SDC3	Lung Adenocarcinoma	0.6202	0.678	0.5714
NOT-CLCA4 AND TLCD1	Lung Adenocarcinoma	0.6195	0.814	0.5
NOT-ABHD6 AND MARVELD2	Lung Adenocarcinoma	0.6069	0.5867	0.6286
NOT-TENM2 AND PVRL4	Lung Adenocarcinoma	0.6015	0.6349	0.5714
NOT-DLL1 AND MARVELD2	Lung Adenocarcinoma	0.6104	0.5595	0.6714
NOT-SLC5A1 AND SLC6A14	Lung Adenocarcinoma	0.631	0.5408	0.7571
NOT-ADCY4 AND NOX4	Lung Carcinoma	0.6102	0.4779	0.8438
NOT-ADCY4 AND ADAM12	Lung Carcinoma	0.6974	0.6023	0.8281
NOT-ABCA8 AND CDH5	Anaplastic Lymphoma	0.6667	0.6667	0.6667
NOT-FCER1A AND CXCL10	Anaplastic Lymphoma	0.72	0.9	0.6
NOT-FCER1A AND LILRB2	Anaplastic Lymphoma	0.6	0.6	0.6
NOT-ASPH AND ST14	Mantle-Cell Lymphoma	0.9444	1	0.8947
NOT-ASPH AND CELSR1	Mantle-Cell Lymphoma	0.9867	1	0.9737
NOT-SLC2A12 AND CELSR1	Mantle-Cell Lymphoma	1	1	1
NOT-DHRS3 AND ST14	Mantle-Cell Lymphoma	0.9167	0.9706	0.8684
NOT-MTUS1 AND ST14	Mantle-Cell Lymphoma	0.9444	1	0.8947
NOT-GPM6B AND CELSR1	Mantle-Cell Lymphoma	0.9487	0.925	0.9737
NOT-TMEM150B AND SLC16A4	Melanoma	0.9524	1	0.9091
NOT-TM4SF5 AND ADAM12	Melanoma	0.9524	1	0.9091
NOT-TM4SF5 AND SGCD	Melanoma	0.9524	1	0.9091
NOT-CDH1 AND FAM57A	Sarcoma	0.8667	1	0.7647
NOT-CDH1 AND SLC2A10	Sarcoma	0.7857	1	0.6471
NOT-LSR AND SLC2A10	Sarcoma	0.7857	1	0.6471
NOT-F11R AND MPZL1	Sarcoma	0.8125	0.8667	0.7647
NOT-F11R AND FAM57A	Sarcoma	0.8	0.9231	0.7059
NKAIN4 AND NOT-CNTNAP2	Astrocytoma	0.825	0.9706	0.7174
NKAIN4 AND NOT-CDH8	Astrocytoma	0.8148	0.9429	0.7174
NKAIN4 AND NOT-HTR2C	Astrocytoma	0.8101	0.9697	0.6957
NKAIN4 AND NOT-KIAA0319	Astrocytoma	0.8049	0.9167	0.7174
NKAIN4 AND NOT-SCN2A	Astrocytoma	0.7952	0.8919	0.7174
NKAIN4 AND NOT-KCNJ6	Astrocytoma	0.7895	1	0.6522
NKAIN4 AND NOT-SCN8A	Astrocytoma	0.7895	1	0.6522

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
CLDN18 AND NOT-TYROBP	Stomach	0.7692	0.7292	0.814
CLDN18 AND NOT-CLEC1A	Stomach	0.7436	0.8286	0.6744
CLDN18 AND NOT-NINJ2	Stomach	0.7253	0.6875	0.7674
CLDN18 AND NOT-GRIA1	Stomach	0.7778	0.9655	0.6512
NOT-SDC1 AND ATP13A4	Astrocytoma	0.7654	0.8857	0.6739
NOT-TPBG AND PTPRF	Liver	0.8333	0.8333	0.8333
NOT-SLC7A2 AND CLDN1	Colon	0.9091	0.8333	1
NOT-SLC33A1 AND SLC39A6	Ependymoma	1	1	1
NOT-LTBR AND ERBB2	Ependymoma	1	1	1
NOT-IL11RA AND EPHB4	Esophagus	0.6	0.5625	0.6429
NOT-GYPE AND CLDN18	Esophagus	0.6	1	0.4286
NOT-IL11RA AND F2RL2	Esophagus	0.6667	0.5789	0.7857
NOT-IL11RA AND OSMR	Esophagus	0.6087	0.7778	0.5
NOT-COL25A1 AND CLDN18	Esophagus	0.6	1	0.4286
NOT-ENPP3 AND TSPAN1	Esophagus	0.6364	0.875	0.5
NOT-SDC1 AND FAM57A	Glioblastoma	0.7143	0.9091	0.5882
NOT-ERBB3 AND ATP2A2	Leiomyosarcoma	0.8696	1	0.7692
NOT-AQP3 AND CD276	Leiomyosarcoma	0.6923	0.6923	0.6923
NOT-ERBB3 AND EVC2	Leiomyosarcoma	0.7273	0.8889	0.6154
NOT-ERBB3 AND PCDH7	Leiomyosarcoma	0.6429	0.6	0.6923
NOT-AQP3 AND SDC1	Leiomyosarcoma	0.64	0.6667	0.6154
NOT-LSR AND SDC1	Leiomyosarcoma	0.8462	0.8462	0.8462
NOT-ERBB3 AND PKD1	Leiomyosarcoma	0.8696	1	0.7692
NOT-CRB3 AND AXL	Leiomyosarcoma	0.6957	0.8	0.6154
NOT-ERBB3 AND PCDHB7	Leiomyosarcoma	0.7619	1	0.6154
NOT-EPCAM AND FAM57A	Leiomyosarcoma	0.6667	0.875	0.5385
NOT-ERBB3 AND HEPH	Leiomyosarcoma	0.6957	0.8	0.6154
NOT-ERBB3 AND PCDHGC3	Leiomyosarcoma	0.6667	0.7273	0.6154
NOT-PTPRK AND SLC7A5	AML	0.8925	0.8111	0.992
NOT-WLS AND CD37	AML	0.8778	0.8201	0.9442
NOT-PTPRK AND CD70	AML	0.8737	0.7806	0.992
NOT-TGFBR3 AND CD37	AML	0.8722	0.8884	0.8566
NOT-HEG1 AND SLC7A5	AML	0.8745	0.861	0.8884
NOT-WLS AND CD70	AML	0.8689	0.7908	0.9641
NOT-CADM1 AND SLC7A5	AML	0.8801	0.8304	0.9363
NOT-HEG1 AND CD70	AML	0.8862	0.888	0.8845
NOT-PODXL AND SLC7A5	AML	0.8592	0.7697	0.9721
NOT-ATP9A AND SLC7A5	AML	0.8664	0.7921	0.9562
NOT-PTPRK AND CD37	AML	0.8566	0.7632	0.9761
NOT-SGCE AND SLC7A5	AML	0.8557	0.7593	0.9801
NOT-WLS AND SLC7A5	AML	0.8531	0.7675	0.9602
NOT-CXADR AND EPHB2	Liposarcoma	0.7647	0.8125	0.7222
NOT-FGFR2 AND PTK7	Liposarcoma	0.6575	0.6486	0.6667
NOT-XK AND EPHB2	Liposarcoma	0.6494	0.6098	0.6944
NOT-SORT1 AND GPNMB	B-Cell Diffuse	0.7742	0.96	0.6486
NOT-MTUS1 AND GPNMB	B-Cell Diffuse	0.7105	0.6923	0.7297
NOT-JAG1 AND GPNMB	B-Cell Diffuse	0.7097	0.88	0.5946
NOT-ITGB6 AND ST14	Mantle-Cell Lymphoma	0.9143	1	0.8421
NOT-TPBG AND ST14	Mantle-Cell Lymphoma	0.9315	0.9714	0.8947
NOT-ERBB3 AND ST14	Mantle-Cell Lymphoma	0.9014	0.9697	0.8421
NOT-STEAP2 AND ST14	Mantle-Cell Lymphoma	0.8657	1	0.7632
NOT-IL20RA AND ST14	Mantle-Cell Lymphoma	0.9315	0.9714	0.8947
NOT-ITGB6 AND CELSR1	Mantle-Cell Lymphoma	0.8706	0.7872	0.9737
NOT-STEAP2 AND CELSR1	Mantle-Cell Lymphoma	0.8608	0.8293	0.8947
NOT-TPBG AND CELSR1	Mantle-Cell Lymphoma	0.8539	0.7451	1
NOT-MST1R AND ST14	Mantle-Cell Lymphoma	0.8529	0.9667	0.7632
NOT-CD9 AND VCAM1	T-Cell, Peripheral	0.6939	0.8095	0.6071
NOT-ITGB5 AND GPNMB	T-Cell, Peripheral	0.6792	0.72	0.6429
NOT-APCDD1 AND GPNMB	T-Cell, Peripheral	0.6579	0.5208	0.8929
NOT-NTN4 AND GPNMB	T-Cell, Peripheral	0.6575	0.5333	0.8571
NOT-TMEM59 AND RNF43	Ovarian	0.7273	0.8	0.6667
NOT-SLC16A7 AND IL2RA	Ovarian	0.7368	1	0.5833
NOT-MTUS1 AND RNF43	Ovarian	0.6957	0.7273	0.6667
NOT-EPCAM AND FAM57A	Sarcoma	0.8	0.7778	0.8235
NOT-RNF43 AND FAM57A	Sarcoma	0.7586	0.9167	0.6471
FAP AND NOT-FOLR2	Breast	0.7879	0.8667	0.7222
FAP AND NOT-IL11RA	Breast	0.8317	0.8936	0.7778
FAP AND NOT-ALDH1A1	Breast	0.8163	0.9091	0.7407
FAP AND NOT-PMEL	Breast	0.8381	0.8627	0.8148
FAP AND NOT-CLDN1	Breast	0.8515	0.9149	0.7963
FAP AND NOT-EGFR	Breast	0.8269	0.86	0.7963
FAP AND NOT-LGR5	Breast	0.8269	0.86	0.7963
FAP AND NOT-EDNRB	Breast	0.7556	0.9444	0.6296
FAP AND NOT-ABCA5	Breast	0.8155	0.8571	0.7778
FAP AND NOT-CD34	Breast	0.8381	0.8627	0.8148
FAP AND NOT-TNC	Breast	0.7579	0.878	0.6667
FAP AND NOT-NCAM1	Breast	0.8776	0.9773	0.7963

FIG. 12 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
NKAIN4 AND NOT-GRM1	Astrocytoma	0.7895	1	0.6522
NKAIN4 AND NOT-LRFN2	Astrocytoma	0.7895	1	0.6522
CRB1 AND NOT-GRM3	Astrocytoma	0.7857	0.8684	0.7174
NKAIN4 AND NOT-SLC13A5	Astrocytoma	0.7805	0.8889	0.6957
NKAIN4 AND NOT-GRM3	Astrocytoma	0.7792	0.9677	0.6522
NKAIN4 AND NOT-KIRREL3	Astrocytoma	0.7792	0.9677	0.6522
NKAIN4 AND NOT-GPR26	Astrocytoma	0.7792	0.9677	0.6522
NKAIN4 AND NOT-CACNA1E	Astrocytoma	0.7792	0.9677	0.6522
NKAIN4 AND NOT-CDH22	Astrocytoma	0.7733	1	0.6304
NKAIN4 AND NOT-CACNA1B	Astrocytoma	0.7733	1	0.6304
NKAIN4 AND NOT-GPR6	Astrocytoma	0.7654	0.8857	0.6739
NKAIN4 AND NOT-CNIH2	Astrocytoma	0.7654	0.8857	0.6739
NKAIN4 AND NOT-KCNV1	Astrocytoma	0.7654	0.8857	0.6739
NKAIN4 AND NOT-CCKBR	Astrocytoma	0.7654	0.8857	0.6739
NKAIN4 AND NOT-FXYD7	Astrocytoma	0.7654	0.8857	0.6739
NKAIN4 AND NOT-CDH9	Astrocytoma	0.7654	0.8857	0.6739
NKAIN4 AND NOT-PTPRR	Astrocytoma	0.7632	0.9667	0.6304
NKAIN4 AND NOT-GPR22	Astrocytoma	0.7595	0.9091	0.6522
NKAIN4 AND NOT-DGKE	Astrocytoma	0.7595	0.9091	0.6522
NKAIN4 AND NOT-CDH18	Astrocytoma	0.75	0.8824	0.6522
NKAIN4 AND NOT-GRIN2A	Astrocytoma	0.75	0.8824	0.6522
NKAIN4 AND NOT-CHRND	Astrocytoma	0.75	0.8824	0.6522
CRB1 AND NOT-KCNC1	Astrocytoma	0.7447	0.7292	0.7609
NKAIN4 AND NOT-GABRA1	Astrocytoma	0.7436	0.9063	0.6304
SLCO1C1 AND NOT-GRM3	Astrocytoma	0.7407	0.8571	0.6522
NKAIN4 AND NOT-HCN1	Astrocytoma	0.7397	1	0.587
NKAIN4 AND NOT-SLC6A15	Astrocytoma	0.7397	1	0.587
NKAIN4 AND NOT-KCNA4	Astrocytoma	0.7342	0.8788	0.6304
NKAIN4 AND NOT-CACNG8	Astrocytoma	0.7342	0.8788	0.6304
NKAIN4 AND NOT-JPH3	Astrocytoma	0.7342	0.8788	0.6304
NKAIN4 AND NOT-KCNJ9	Astrocytoma	0.7342	0.8788	0.6304
NKAIN4 AND NOT-DRD1	Astrocytoma	0.7342	0.8788	0.6304
CRB1 AND NOT-SCN8A	Astrocytoma	0.7317	0.8333	0.6522
CRB1 AND NOT-GABRA2	Astrocytoma	0.7317	0.8333	0.6522
NKAIN4 AND NOT-CLSTN3	Astrocytoma	0.7273	0.9032	0.6087
SLCO1C1 AND NOT-GABRA2	Astrocytoma	0.7273	0.7619	0.6957
CRB1 AND NOT-KIAA0319	Astrocytoma	0.7229	0.8108	0.6522
NKAIN4 AND NOT-EPHA8	Astrocytoma	0.7222	1	0.5652
NKAIN4 AND NOT-GPR83	Astrocytoma	0.7222	1	0.5652
NKAIN4 AND NOT-ABCG4	Astrocytoma	0.72	0.931	0.587
NKAIN4 AND NOT-NKAIN2	Astrocytoma	0.72	0.931	0.587
NKAIN4 AND NOT-CACNG3	Astrocytoma	0.7179	0.875	0.6087
NKAIN4 AND NOT-GABRA5	Astrocytoma	0.7179	0.875	0.6087
NKAIN4 AND NOT-GABRA4	Astrocytoma	0.7179	0.875	0.6087
NKAIN4 AND NOT-GABRB2	Astrocytoma	0.7179	0.875	0.6087
CRB1 AND NOT-CDH8	Astrocytoma	0.7111	0.7273	0.6957
NKAIN4 AND NOT-OPRK1	Astrocytoma	0.7105	0.9	0.587
BEST3 AND NOT-AOC3	Astrocytoma	0.7089	0.8485	0.6087
CRB1 AND NOT-SYT4	Astrocytoma	0.7073	0.8056	0.6304
CRB1 AND NOT-GPR83	Astrocytoma	0.7073	0.8056	0.6304
CRB1 AND NOT-DGKE	Astrocytoma	0.7059	0.7692	0.6522
CRB1 AND NOT-ATP8A2	Astrocytoma	0.7059	0.7692	0.6522
PPAPDC1A AND NOT-TMEFF2	Breast	0.7755	0.8636	0.7037
PPAPDC1A AND NOT-GRM3	Breast	0.7677	0.8444	0.7037
PPAPDC1A AND NOT-TRPM3	Breast	0.76	0.8261	0.7037
PPAPDC1A AND NOT-GABRG1	Breast	0.76	0.8261	0.7037
PPAPDC1A AND NOT-ASTN1	Breast	0.76	0.8261	0.7037
PPAPDC1A AND NOT-OMG	Breast	0.7917	0.9048	0.7037
PPAPDC1A AND NOT-KCNJ10	Breast	0.8132	1	0.6852
PPAPDC1A AND NOT-CALN1	Breast	0.7451	0.7917	0.7037
PPAPDC1A AND NOT-GABRB1	Breast	0.7551	0.8409	0.6852
PPAPDC1A AND NOT-TMEM235	Breast	0.7677	0.8444	0.7037
PPAPDC1A AND NOT-MLC1	Breast	0.7677	0.8444	0.7037
PPAPDC1A AND NOT-CNTNAP4	Breast	0.7755	0.8636	0.7037
PPAPDC1A AND NOT-SLC24A2	Breast	0.7451	0.7917	0.7037
PPAPDC1A AND NOT-OPALIN	Breast	0.7677	0.8444	0.7037
PPAPDC1A AND NOT-NTSR2	Breast	0.7525	0.8085	0.7037
PPAPDC1A AND NOT-DTNA	Breast	0.7308	0.76	0.7037
PPAPDC1A AND NOT-ANO4	Breast	0.7238	0.7451	0.7037
PPAPDC1A AND NOT-MOG	Breast	0.76	0.8261	0.7037
LRRC8E AND NOT-CLDN10	Breast	0.729	0.7358	0.7222
PPAPDC1A AND NOT-ACSL6	Breast	0.7525	0.8085	0.7037
PPAPDC1A AND NOT-OPCML	Breast	0.717	0.7308	0.7037
PPAPDC1A AND NOT-FXYD7	Breast	0.7238	0.7451	0.7037
PPAPDC1A AND NOT-SLC1A2	Breast	0.7347	0.8182	0.6667
PPAPDC1A AND NOT-MAG	Breast	0.76	0.8261	0.7037
PPAPDC1A AND NOT-ATP2B2	Breast	0.7115	0.74	0.6852

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
MUC1 AND NOT-ALDH1A1	Breast	0.7347	0.8182	0.6667
FAP AND NOT-CD160	Breast	0.7579	0.878	0.6667
FAP AND NOT-RAET1E	Breast	0.7755	0.8636	0.7037
FAP AND NOT-CLDN11	Breast	0.8269	0.86	0.7963
FAP AND NOT-CLDN8	Breast	0.7158	0.8293	0.6296
CLDN7 AND NOT-ALDH1A1	Breast	0.7143	0.6897	0.7407
FAP AND NOT-SSTR1	Breast	0.8381	0.8627	0.8148
FAP AND NOT-PCYT1A	Breast	0.8269	0.86	0.7963
FAP AND NOT-GPC3	Breast	0.78	0.8478	0.7222
FAP AND NOT-GUCY2C	Breast	0.8544	0.898	0.8148
FAP AND NOT-GAGE1	Breast	0.8317	0.8936	0.7778
FAP AND NOT-CLDN2	Breast	0.8269	0.86	0.7963
FAP AND NOT-CLDN12	Breast	0.7158	0.8293	0.6296
FAP AND NOT-DDX3X	Breast	0.8381	0.8627	0.8148
FAP AND NOT-CSPG4	Breast	0.75	0.8571	0.6667
FAP AND NOT-CLDN18	Breast	0.8381	0.8627	0.8148
FAP AND NOT-IL20RA	Breast	0.7158	0.8293	0.6296
FAP AND NOT-EPHA3	Breast	0.6977	0.9375	0.5556
FAP AND NOT-CD33	Breast	0.7579	0.878	0.6667
FAP AND NOT-VCAM1	Breast	0.7423	0.8372	0.6667
FAP AND NOT-KDR	Breast	0.7755	0.8636	0.7037
FAP AND NOT-TNFRSF8	Breast	0.8081	0.8889	0.7407
FAP AND NOT-MAGEA11	Breast	0.8381	0.8627	0.8148
FAP AND NOT-CLDN23	Breast	0.8381	0.8627	0.8148
FAP AND NOT-SSTR5	Breast	0.8462	0.88	0.8148
FAP AND NOT-SST	Breast	0.8544	0.898	0.8148
FAP AND NOT-STEAP2	Breast	0.8269	0.86	0.7963
VTCN1 AND NOT-MUC13	Breast	0.6905	0.9667	0.537
VTCN1 AND NOT-MUC4	Breast	0.6905	0.9667	0.537
VTCN1 AND NOT-MOK	Breast	0.6905	0.9667	0.537
VTCN1 AND NOT-CLDN23	Breast	0.6905	0.9667	0.537
VTCN1 AND NOT-PMEL	Breast	0.6905	0.9667	0.537
VTCN1 AND NOT-HHLA2	Breast	0.6905	0.9667	0.537
VTCN1 AND NOT-SSTR1	Breast	0.6905	0.9667	0.537
VTCN1 AND NOT-IL11RA	Breast	0.6905	0.9667	0.537
VTCN1 AND NOT-MAGEA11	Breast	0.6905	0.9667	0.537
FAP AND NOT-IL3RA	Breast	0.8544	0.898	0.8148
FAP AND NOT-ITGB3	Breast	0.8571	0.9545	0.7778
VTCN1 AND NOT-MUC16	Breast	0.6905	0.9667	0.537
VTCN1 AND NOT-TNC	Breast	0.6905	0.9667	0.537
VTCN1 AND NOT-GPA33	Breast	0.6905	0.9667	0.537
VTCN1 AND NOT-GUCY2C	Breast	0.6905	0.9667	0.537
VTCN1 AND NOT-CLDN9	Breast	0.6905	0.9667	0.537
FAP AND NOT-MOK	Breast	0.8039	0.8542	0.7593
FAP AND NOT-SSX1	Breast	0.8	0.8696	0.7407
TPBG AND NOT-NCAM1	Breast	0.6813	0.8378	0.5741
FAP AND NOT-CD52	Breast	0.7677	0.8444	0.7037
FAP AND NOT-CLEC14A	Breast	0.8283	0.9111	0.7593
VTCN1 AND NOT-EDNRB	Breast	0.6747	0.9655	0.5185
VTCN1 AND NOT-GAGE1	Breast	0.6747	0.9655	0.5185
VTCN1 AND NOT-HSPA5	Breast	0.6747	0.9655	0.5185
VTCN1 AND NOT-SLC34A2	Breast	0.6905	0.9667	0.537
VTCN1 AND NOT-NCAM1	Breast	0.6905	0.9667	0.537
TPBG AND NOT-ALK	Breast	0.6739	0.8158	0.5741
VTCN1 AND NOT-CLDN1	Breast	0.6747	0.9655	0.5185
CLDN12 AND NOT-EPCAM	Liver	0.8	1	0.6667
CLDN12 AND NOT-TPBG	Liver	0.9091	1	0.8333
SDC1 AND NOT-TPBG	Liver	0.8	0.6667	1
CLDN1 AND NOT-TPBG	Liver	0.8	1	0.6667
CLDN1 AND NOT-IL20RA	Liver	0.8	1	0.6667
CLDN12 AND NOT-IL20RA	Liver	0.9231	0.8571	1
CLDN12 AND NOT-ITGB6	Liver	1	1	1
CLDN2 AND NOT-FOLR1	Liver	0.625	0.5	0.8333
CLDN12 AND NOT-PROM1	Liver	0.9091	1	0.8333
CLDN12 AND NOT-MST1R	Liver	1	1	1
CLDN12 AND NOT-CLDN8	Liver	0.9231	0.8571	1
CLDN12 AND NOT-ENPP3	Liver	0.6667	1	0.5
ERBB3 AND NOT-FOLR1	Liver	0.7692	0.7143	0.8333
FAP AND NOT-CSPG4	Pancreas	0.8421	1	0.7273
FAP AND NOT-RAET1E	Pancreas	0.7059	1	0.5455
FAP AND NOT-IL11RA	Pancreas	0.7778	1	0.6364
FAP AND NOT-NCAM1	Pancreas	0.8421	1	0.7273
FAP AND NOT-CLDN8	Pancreas	0.8421	1	0.7273
MUC13 AND NOT-ENPP3	Pancreas	0.6154	0.5333	0.7273
FAP AND NOT-KDR	Pancreas	0.625	1	0.4545
EPHA3 AND NOT-CLDN8	Pancreas	0.6667	0.8571	0.5455
CLDN2 AND NOT-ENPP3	Pancreas	0.6316	0.75	0.5455

FIG. 12 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
PPAPDC1A AND NOT-NKAIN2	Breast	0.7097	0.8462	0.6111
PPAPDC1A AND NOT-SLC5A11	Breast	0.7527	0.8974	0.6481
PPAPDC1A AND NOT-SLC6A12	Breast	0.717	0.7308	0.7037
PPAPDC1A AND NOT-SLCO1A2	Breast	0.76	0.8261	0.7037
PPAPDC1A AND NOT-NKAIN4	Breast	0.7379	0.7755	0.7037
PPAPDC1A AND NOT-SLC12A5	Breast	0.7037	0.7037	0.7037
PPAPDC1A AND NOT-NFASC	Breast	0.7379	0.7755	0.7037
PPAPDC1A AND NOT-CSPG5	Breast	0.701	0.7907	0.6296
PPAPDC1A AND NOT-SCN2A	Breast	0.7048	0.7255	0.6852
PPAPDC1A AND NOT-KCNK10	Breast	0.6981	0.7115	0.6852
PPAPDC1A AND NOT-GABRA2	Breast	0.6981	0.7115	0.6852
PPAPDC1A AND NOT-CSMD3	Breast	0.6972	0.6909	0.7037
PPAPDC1A AND NOT-SCN1A	Breast	0.6972	0.6909	0.7037
PPAPDC1A AND NOT-GABRG2	Breast	0.6972	0.6909	0.7037
PPAPDC1A AND NOT-CALY	Breast	0.6972	0.6909	0.7037
PPAPDC1A AND NOT-SYT4	Breast	0.6916	0.6981	0.6852
PPAPDC1A AND NOT-GRIN2C	Breast	0.6909	0.6786	0.7037
PPAPDC1A AND NOT-CLDN10	Breast	0.6857	0.7059	0.6667
PPAPDC1A AND NOT-SLC39A12	Breast	0.6847	0.6667	0.7037
PPAPDC1A AND NOT-OR1Q1	Breast	0.7835	0.8837	0.7037
PPAPDC1A AND NOT-HCN2	Breast	0.7451	0.7917	0.7037
PPAPDC1A AND NOT-ATP8A1	Breast	0.68	0.7391	0.6296
PPAPDC1A AND NOT-DPP10	Breast	0.6792	0.6923	0.6667
ENPP1 AND NOT-CLDN10	Breast	0.6789	0.6727	0.6852
PPAPDC1A AND NOT-TTYH2	Breast	0.7021	0.825	0.6111
PPAPDC1A AND NOT-KCNA2	Breast	0.6739	0.8158	0.5741
LRRC8E AND NOT-SLC39A2	Breast	0.7407	0.7407	0.7407
PPAPDC1A AND NOT-CDH18	Breast	0.6727	0.6607	0.6852
PPAPDC1A AND NOT-GRIN2A	Breast	0.6726	0.6441	0.7037
PPAPDC1A AND NOT-PIRT	Breast	0.6726	0.6441	0.7037
PPAPDC1A AND NOT-ADCY8	Breast	0.6726	0.6441	0.7037
PPAPDC1A AND NOT-GRIA4	Breast	0.6667	0.6333	0.7037
PPAPDC1A AND NOT-SLCO1C1	Breast	0.6667	0.6491	0.6852
LRRC8E AND NOT-TMPRSS11E	Breast	0.7879	0.8667	0.7222
PPAPDC1A AND NOT-SEMA4D	Breast	0.6609	0.623	0.7037
PPAPDC1A AND NOT-DISP2	Breast	0.6604	0.6731	0.6481
PPAPDC1A AND NOT-CACNA1B	Breast	0.6496	0.6032	0.7037
ENPP1 AND NOT-MUC17	Breast	0.6613	0.5857	0.7593
PPAPDC1A AND NOT-GPR85	Breast	0.6441	0.5938	0.7037
CLDN15 AND NOT-SLC16A5	Liver	0.8333	0.8333	0.8333
SLCO1B1 AND NOT-C8B	Liver	0.75	0.6	1
SLC30A8 AND NOT-NPHS1	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-SLC6A11	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-DPP10	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-ENPP1	Pancreas	0.7059	1	0.5455
SLC30A8 AND NOT-HCN1	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-KCNK16	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-CNIH2	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-SLC1A2	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-GPR50	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-CHRNB2	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-SLC5A2	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-SLC28A1	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-DTNA	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-FSHR	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-SLC39A8	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-SHISA9	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-CDH9	Pancreas	0.7059	1	0.5455
SLC30A8 AND NOT-FAM26D	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-CCKAR	Pancreas	0.7059	1	0.5455
SLC30A8 AND NOT-PPAPDC1B	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-LRRC55	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-CCKBR	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-LCT	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-KIRREL2	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-KCNF1	Pancreas	0.7059	1	0.5455
SLC30A8 AND NOT-GPR22	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-CYP4A11	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-KCNK10	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-TMPRSS11E	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-ATP8A2	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-LRRC8E	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-SLC2A2	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-PAQR7	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-SLC17A2	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-P2RX3	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-MRAP	Pancreas	0.7778	1	0.6364

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
FAP AND NOT-AXL	Pancreas	0.7778	1	0.6364
FAP AND NOT-CD34	Pancreas	0.7059	1	0.5455
FAP AND NOT-PMEL	Pancreas	0.8421	1	0.7273
EPHA3 AND NOT-CSPG4	Pancreas	0.6	0.6667	0.5455
FAP AND NOT-ROR1	Pancreas	0.625	1	0.4545
FAP AND NOT-EDNRB	Pancreas	0.7059	1	0.5455
FAP AND NOT-CLDN6	Pancreas	0.8421	1	0.7273
FAP AND NOT-ENPP3	Pancreas	0.7778	1	0.6364
FAP AND NOT-ITGB3	Pancreas	0.7778	1	0.6364
FAP AND NOT-TNFRSF8	Pancreas	0.8421	1	0.7273
FAP AND NOT-CLDN11	Pancreas	0.8421	1	0.7273
VCAM1 AND NOT-CLDN11	Renal	0.8	1	0.6667
VCAM1 AND NOT-IL11RA	Renal	0.7368	1	0.5833
CLDN2 AND NOT-LGR5	Renal	0.7619	0.8889	0.6667
CLDN2 AND NOT-IL11RA	Renal	0.6667	0.7778	0.5833
CLDN2 AND NOT-SSTR4	Renal	0.6667	1	0.5
VCAM1 AND NOT-BMPR1B	Renal	0.6667	1	0.5
VCAM1 AND NOT-LGR5	Renal	0.8	1	0.6667
CLDN2 AND NOT-B4GALNT1	Renal	0.6667	1	0.5
VCAM1 AND NOT-FAP	Renal	0.6667	0.6	0.75
CLDN2 AND NOT-CD19	Renal	0.6087	0.6364	0.5833
CLDN2 AND NOT-FCRL1	Renal	0.8	1	0.6667
DPEP1 AND NOT-AFP	Colon	1	1	1
EPHB2 AND NOT-MUC4	Colon	0.8889	1	0.8
GUCY2C AND NOT-GPA33	Colon	0.8889	1	0.8
DPEP1 AND NOT-SSTR4	Colon	0.8889	1	0.8
DPEP1 AND NOT-MAGEA1	Colon	1	1	1
DPEP1 AND NOT-MAGEA11	Colon	1	1	1
DPEP1 AND NOT-CLDN18	Colon	1	1	1
MUC13 AND NOT-GPA33	Colon	0.8889	1	0.8
DPEP1 AND NOT-FCRL1	Colon	0.8889	1	0.8
DPEP1 AND NOT-CLDN9	Colon	0.9091	0.8333	1
EPHB2 AND NOT-SST	Colon	0.8889	1	0.8
EPHB2 AND NOT-CLDN3	Colon	0.8	0.8	0.8
DPEP1 AND NOT-MAGEA4	Colon	1	1	1
DPEP1 AND NOT-TNFRSF13C	Colon	1	1	1
DPEP1 AND NOT-ITGB3	Colon	0.8889	1	0.8
EPHB2 AND NOT-CLDN23	Colon	0.8889	1	0.8
EPHB2 AND NOT-IGF1R	Colon	0.8889	1	0.8
BIRC5 AND NOT-IL11RA	Colon	1	1	1
CLDN2 AND NOT-AFP	Colon	0.8889	1	0.8
EPHB2 AND NOT-TNFRSF13C	Colon	0.8889	1	0.8
GUCY2C AND NOT-SSTR1	Colon	0.8889	1	0.8
RNF43 AND NOT-FLOT2	Colon	0.75	1	0.6
GUCY2C AND NOT-HHLA2	Colon	0.75	1	0.6
RNF43 AND NOT-ERBB4	Colon	0.75	1	0.6
RNF43 AND NOT-CSPG4	Colon	0.75	1	0.6
RNF43 AND NOT-EGFR	Colon	0.75	1	0.6
DPEP1 AND NOT-CXCR5	Colon	0.75	1	0.6
EPHB2 AND NOT-TRPM4	Colon	0.75	1	0.6
RNF43 AND NOT-CD160	Colon	0.75	1	0.6
RNF43 AND NOT-CLDN11	Colon	0.75	1	0.6
CLDN2 AND NOT-TYR	Colon	0.75	1	0.6
RNF43 AND NOT-PCYT1A	Colon	0.75	1	0.6
RNF43 AND NOT-MOK	Colon	0.75	1	0.6
RNF43 AND NOT-ST8SIA1	Colon	0.75	1	0.6
RNF43 AND NOT-IL11RA	Colon	0.75	1	0.6
CLDN2 AND NOT-SSTR4	Colon	0.75	1	0.6
RNF43 AND NOT-TNC	Colon	0.75	1	0.6
RNF43 AND NOT-CD34	Colon	0.75	1	0.6
EPHB2 AND NOT-GPA33	Colon	0.75	1	0.6
DPEP1 AND NOT-ULBP1	Colon	0.8889	1	0.8
DPEP1 AND NOT-TYR	Colon	0.8889	1	0.8
DPEP1 AND NOT-PMEL	Colon	0.75	1	0.6
BIRC5 AND NOT-CD79B	Colon	0.8333	0.7143	1
DPEP1 AND NOT-B4GALNT1	Colon	0.8889	1	0.8
EPHB2 AND NOT-MST1R	Colon	0.8	0.8	0.8
RNF43 AND NOT-NCAM1	Colon	0.75	1	0.6
RNF43 AND NOT-TRPM4	Colon	0.75	1	0.6
SSTR1 AND NOT-EPCAM	Ependymoma	1	1	1
SSTR1 AND NOT-CLDN12	Ependymoma	1	1	1
MUC13 AND NOT-ENPP3	Esophagus	0.6667	1	0.5
EPCAM AND NOT-ENPP3	Esophagus	0.6667	1	0.5
MST1R AND NOT-ENPP3	Esophagus	0.6	1	0.4286
EPHB2 AND NOT-ENPP3	Esophagus	0.7647	0.65	0.9286
CBX3 AND NOT-CD19	Esophagus	0.64	0.7273	0.5714
CBX3 AND NOT-IL11RA	Esophagus	0.6667	1	0.5

FIG. 12 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
SLC30A8 AND NOT-IL13	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-SLC17A3	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-HTR3C	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-SLC6A12	Pancreas	0.7059	1	0.5455
SLC30A8 AND NOT-SLC43A1	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-TMPRSS11D	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-CSPG5	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-GRIK1	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-ADAM2	Pancreas	0.7059	1	0.5455
SLC30A8 AND NOT-GABRG1	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-FCAMR	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-NTSR2	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-SLC22A12	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-GPR83	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-SLC12A3	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-IGDCC3	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-KREMEN2	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-SLC34A1	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-DRD5	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-MIP	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-SLC22A2	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-SMPD2	Pancreas	0.7059	1	0.5455
SLC30A8 AND NOT-PCDH11X	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-CHRNB4	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-KCNQ2	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-OR4N4	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-PTGER4	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-ATP13A5	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-ADAM20	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-ATP6V0A4	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-MMP24	Pancreas	0.7059	1	0.5455
SLC30A8 AND NOT-TAS2R1	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-CNTNAP2	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-AQP6	Pancreas	0.7059	1	0.5455
SLC30A8 AND NOT-CNGA4	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-KIAA0319	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-GRM1	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-KCNA4	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-SLC22A13	Pancreas	0.7778	1	0.6364
PPAPDC1A AND NOT-CSPG5	Pancreas	0.6	0.6667	0.5455
SLC30A8 AND NOT-SLC22A6	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-SLC5A10	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-PORCN	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-FGF6	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-GRM3	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-ATP4B	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-SLC22A11	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-TRPM3	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-SLC22A14	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-GUCY2D	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-HRH3	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-NMBR	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-GJA3	Pancreas	0.7059	1	0.5455
SLC30A8 AND NOT-CHRNA1	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-HTR1D	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-CSMD3	Pancreas	0.625	1	0.4545
SLC30A8 AND NOT-KCNU1	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-SLC5A11	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-TRPC7	Pancreas	0.7778	1	0.6364
SLC22A2 AND NOT-SLC16A5	Renal	0.7368	1	0.5833
SLC22A2 AND NOT-ATP6V0A4	Renal	0.7368	1	0.5833
SLC22A2 AND NOT-UMOD	Renal	0.7368	1	0.5833
GGTLC1 AND NOT-SLC16A5	Renal	0.75	0.75	0.75
TNFRSF12A AND NOT-SCARA5	Renal	0.75	0.75	0.75
SLC13A1 AND NOT-SLC5A11	Renal	0.6667	1	0.5
SLC13A1 AND NOT-SLC16A5	Renal	0.6667	1	0.5
SLC22A2 AND NOT-CASR	Renal	0.6667	0.7778	0.5833
SLC13A1 AND NOT-SLC13A2	Renal	0.6667	1	0.5
SLC22A2 AND NOT-DIO1	Renal	0.6667	0.7778	0.5833
SLC17A1 AND NOT-DIO1	Renal	0.6667	1	0.5
SLC6A13 AND NOT-CLDN19	Renal	0.6667	1	0.5
SLC22A2 AND NOT-SLC12A3	Renal	0.6667	0.7778	0.5833
SLC22A2 AND NOT-SLC13A2	Renal	0.6667	0.7778	0.5833
SLC6A13 AND NOT-SYT6	Renal	0.6667	1	0.5
SLC22A2 AND NOT-SLC22A8	Renal	0.6667	0.7778	0.5833
SLC22A2 AND NOT-SYT6	Renal	0.6667	1	0.5
SLC22A2 AND NOT-SLC5A11	Renal	0.6667	0.7778	0.5833

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
EPHB2 AND NOT-ABCA5	Esophagus	0.6875	0.6111	0.7857
CLDN18 AND NOT-FOLR1	Esophagus	0.6	0.5625	0.6429
EPHB2 AND NOT-GPA33	Esophagus	0.7059	0.6	0.8571
MUC13 AND NOT-ABCA5	Esophagus	0.6897	0.6667	0.7143
BIRC5 AND NOT-CD33	Esophagus	0.6923	0.75	0.6429
BIRC5 AND NOT-CD160	Esophagus	0.6923	0.75	0.6429
BIRC5 AND NOT-SLC39A6	Esophagus	0.6	1	0.4286
ITGAV AND NOT-RNF43	Glioma	0.9037	0.9242	0.8841
ITGAV AND NOT-CLDN1	Glioma	0.8936	0.875	0.913
ITGAV AND NOT-EPCAM	Glioma	0.8966	0.8553	0.942
ITGAV AND NOT-TRPM4	Glioma	0.8837	0.95	0.8261
ITGAV AND NOT-SLAMF7	Glioma	0.9489	0.9559	0.942
ITGAV AND NOT-TPBG	Glioma	0.8321	0.8382	0.8261
SLC39A6 AND NOT-CLDN1	Glioma	0.7941	0.806	0.7826
ITGAV AND NOT-CLDN8	Glioma	0.9429	0.9296	0.9565
ITGAV AND NOT-ITGB6	Glioma	0.942	0.942	0.942
SLC39A6 AND NOT-RNF43	Glioma	0.7786	0.8226	0.7391
ITGAV AND NOT-RAET1E	Glioma	0.942	0.942	0.942
EDNRB AND NOT-EPCAM	Glioma	0.7704	0.7879	0.7536
ITGAV AND NOT-CLDN23	Glioma	0.8889	0.9091	0.8696
ITGAV AND NOT-GPNMB	Glioma	0.8148	0.8333	0.7971
ITGAV AND NOT-CLDN7	Glioma	0.942	0.942	0.942
ITGAV AND NOT-IL20RA	Glioma	0.8702	0.9194	0.8261
ITGAV AND NOT-CD34	Glioma	0.8769	0.9344	0.8261
ITGAV AND NOT-SDC1	Glioma	0.8966	0.8553	0.942
SLC39A6 AND NOT-TRPM4	Glioma	0.75	0.8136	0.6957
SLC39A6 AND NOT-ITGB6	Glioma	0.8209	0.8462	0.7971
NCAM1 AND NOT-RAET1E	Glioma	0.7453	0.6522	0.8696
ITGAV AND NOT-TNFRSF10A	Glioma	0.9209	0.9143	0.9275
SLC39A6 AND NOT-SDC1	Glioma	0.7424	0.7778	0.7101
ITGAV AND NOT-ROR1	Glioma	0.9091	0.9524	0.8696
ITGAV AND NOT-PMEL	Glioma	0.9362	0.9167	0.9565
SLC39A6 AND NOT-IL20RA	Glioma	0.7385	0.7869	0.6957
ITGAV AND NOT-CD52	Glioma	0.791	0.8154	0.7681
ITGAV AND NOT-VTCN1	Glioma	0.9037	0.9242	0.8841
SLC39A6 AND NOT-CLDN8	Glioma	0.8116	0.8116	0.8116
ITGAV AND NOT-IL3RA	Glioma	0.8971	0.9104	0.8841
ITGAV AND NOT-CD79A	Glioma	0.8438	0.9153	0.7826
ITGAV AND NOT-ENPP3	Glioma	0.9265	0.9403	0.913
SLC39A6 AND NOT-RAET1E	Glioma	0.8346	0.9138	0.7681
SLC39A6 AND NOT-CLDN7	Glioma	0.8209	0.8462	0.7971
EDNRB AND NOT-ITGB6	Glioma	0.7761	0.8	0.7536
EDNRB AND NOT-RNF43	Glioma	0.7231	0.7705	0.6812
NCAM1 AND NOT-GPA33	Glioma	0.7229	0.6186	0.8696
ITGAV AND NOT-MST1R	Glioma	0.9343	0.9412	0.9275
EDNRB AND NOT-CLDN1	Glioma	0.7805	0.8889	0.6957
EDNRB AND NOT-SLAMF7	Glioma	0.8254	0.9123	0.7536
EDNRB AND NOT-CLDN7	Glioma	0.7934	0.9231	0.6957
ITGAV AND NOT-MUC13	Glioma	0.8855	0.9355	0.8406
EDNRB AND NOT-CLDN23	Glioma	0.7133	0.6892	0.7391
ITGAV AND NOT-CD79B	Glioma	0.8788	0.9206	0.8406
ITGAV AND NOT-FAP	Glioma	0.768	0.8571	0.6957
ITGAV AND NOT-MET	Glioma	0.708	0.9091	0.5797
SLC39A6 AND NOT-CLDN23	Glioma	0.7778	0.8596	0.7101
ITGAV AND NOT-ITGB3	Glioma	0.8254	0.9123	0.7536
SLC39A6 AND NOT-TPBG	Glioma	0.704	0.7857	0.6377
EDNRB AND NOT-TRPM4	Glioma	0.7023	0.7419	0.6667
SLC39A6 AND NOT-SLAMF7	Glioma	0.8661	0.9483	0.7971
SLC7A5 AND NOT-CLDN1	Glioma	0.7	0.6154	0.8116
SLC39A6 AND NOT-CD52	Glioma	0.7	0.8235	0.6087
ITGAV AND NOT-PTK7	Glioma	0.8429	0.831	0.8551
SLC39A6 AND NOT-TNFRSF10A	Glioma	0.813	0.9259	0.7246
ITGAV AND NOT-SLC34A2	Glioma	0.913	0.913	0.913
ITGAV AND NOT-AXL	Glioma	0.8	0.8182	0.7826
ITGAV AND NOT-MS4A1	Glioma	0.9275	0.9275	0.9275
EDNRB AND NOT-TNFRSF10A	Glioma	0.7634	0.8065	0.7246
ITGAV AND NOT-LGR5	Glioma	0.8671	0.8378	0.8986
ITGAV AND NOT-CLDN6	Glioma	0.942	0.942	0.942
ITGAV AND NOT-ABCB5	Glioma	0.9104	0.9385	0.8841
ITGAV AND NOT-TNFRSF8	Glioma	0.913	0.913	0.913
ITGAV AND NOT-IL11RA	Glioma	0.8777	0.8714	0.8841
EDNRB AND NOT-CLDN8	Glioma	0.7259	0.7424	0.7101
ITGAV AND NOT-TNFRSF17	Glioma	0.9286	0.9155	0.942
EDNRB AND NOT-IL20RA	Glioma	0.7097	0.8	0.6377
EDNRB AND NOT-IL3RA	Glioma	0.7619	0.8421	0.6957
ITGAV AND NOT-GPA33	Glioma	0.9275	0.9275	0.9275
CD276 AND NOT-RNF43	Leiomyosarcoma	0.7826	0.9	0.6923

FIG. 12 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
SLC22A2 AND NOT-GABRB1	Renal	0.6667	1	0.5
SLC6A13 AND NOT-GPRC6A	Renal	0.6667	1	0.5
SLC22A2 AND NOT-GHRHR	Renal	0.6667	0.7778	0.5833
SLC22A2 AND NOT-NPHS2	Renal	0.6667	0.7778	0.5833
SLC22A2 AND NOT-OPALIN	Renal	0.6364	0.7	0.5833
SLC22A2 AND NOT-SCN1A	Renal	0.6364	0.7	0.5833
SLC22A2 AND NOT-SLC22A25	Renal	0.6364	0.7	0.5833
SLC22A2 AND NOT-CD207	Renal	0.6364	0.7	0.5833
SLC22A2 AND NOT-PTPRT	Renal	0.6364	0.7	0.5833
SLC6A13 AND NOT-AQP6	Renal	0.6316	0.8571	0.5
SLC6A13 AND NOT-SLC5A11	Renal	0.6316	0.8571	0.5
SLC6A13 AND NOT-ATP6V0A4	Renal	0.6316	0.8571	0.5
SLC6A13 AND NOT-SLC22A8	Renal	0.6316	0.8571	0.5
SLC6A13 AND NOT-KCNJ10	Renal	0.6316	0.8571	0.5
SLC6A13 AND NOT-SLC16A5	Renal	0.6316	0.8571	0.5
SLC6A13 AND NOT-NPHS2	Renal	0.6316	0.8571	0.5
SLC6A13 AND NOT-SLC13A2	Renal	0.6316	0.8571	0.5
SLC6A13 AND NOT-SLC12A3	Renal	0.6316	0.8571	0.5
MMP24 AND NOT-LAMP5	Renal	0.6316	0.8571	0.5
SLC22A2 AND NOT-KCNK10	Renal	0.6087	0.6364	0.5833
SLC22A2 AND NOT-OR2L2	Renal	0.6087	0.6364	0.5833
SLC22A2 AND NOT-TRHR	Renal	0.6087	0.6364	0.5833
SLC22A2 AND NOT-CHRND	Renal	0.6087	0.6364	0.5833
TNFRSF12A AND NOT-STEAP4	Renal	0.8182	0.9	0.75
SLC22A2 AND NOT-MRGPRX2	Renal	0.6	0.75	0.5
SLC22A2 AND NOT-SLC22A12	Renal	0.6	0.75	0.5
SLC22A2 AND NOT-SLC22A6	Renal	0.6	0.75	0.5
SLC22A2 AND NOT-GPR6	Renal	0.6	0.75	0.5
FAT1 AND NOT-SLC16A5	Renal	0.6	0.75	0.5
SLC22A2 AND NOT-USH2A	Renal	0.6	0.75	0.5
SLC6A13 AND NOT-DIO1	Renal	0.6	0.75	0.5
SLC6A13 AND NOT-GHRHR	Renal	0.6	0.75	0.5
SLC17A1 AND NOT-CALHM1	Renal	0.6	0.75	0.5
SLC6A13 AND NOT-NPHS1	Renal	0.6	0.75	0.5
SLC17A1 AND NOT-CATSPERD	Renal	0.6	0.75	0.5
SLC17A1 AND NOT-SLC5A11	Renal	0.6667	1	0.5
SLC17A1 AND NOT-NPHS2	Renal	0.6667	1	0.5
SLC17A1 AND NOT-SLC12A3	Renal	0.6667	1	0.5
SLC17A1 AND NOT-OR4N4	Renal	0.6667	1	0.5
SLC17A1 AND NOT-KCNK10	Renal	0.6	0.75	0.5
NOX1 AND NOT-SLC9A1	Colon	1	1	1
CDH17 AND NOT-SLC9A1	Colon	1	1	1
NOX1 AND NOT-SLC30A10	Colon	0.8889	1	0.8
CDH17 AND NOT-SLC30A10	Colon	0.8889	1	0.8
IFI6 AND NOT-SLC1A2	Colon	0.9091	0.8333	1
IFI6 AND NOT-GPR26	Colon	0.8333	0.7143	1
FAT1 AND NOT-ENPP1	Colon	0.8	0.8	0.8
IFI6 AND NOT-STEAP4	Colon	0.8333	0.7143	1
CDH17 AND NOT-EPHA10	Colon	0.8333	0.7143	1
IFI6 AND NOT-MLC1	Colon	0.9091	0.8333	1
NOX1 AND NOT-CHRND	Colon	0.7692	0.625	1
FAT1 AND NOT-SLC9A1	Colon	0.75	1	0.6
NOX1 AND NOT-BEST2	Colon	0.75	1	0.6
IFI6 AND NOT-TTYH2	Colon	0.8	0.8	0.8
CDH17 AND NOT-CHRND	Colon	0.7692	0.625	1
IFI6 AND NOT-KCNV2	Colon	0.9091	0.8333	1
IFI6 AND NOT-S1PR1	Colon	0.7273	0.6667	0.8
IFI6 AND NOT-CLDN10	Colon	0.75	1	0.6
IFI6 AND NOT-SLC23A2	Colon	0.75	1	0.6
FAT1 AND NOT-SLC16A5	Colon	0.8	0.8	0.8
IFI6 AND NOT-CD36	Colon	0.8333	0.7143	1
NOX1 AND NOT-EPHA10	Colon	0.7143	0.5556	1
IFI6 AND NOT-GYPC	Colon	0.7273	0.6667	0.8
NOX1 AND NOT-GABRB2	Colon	1	1	1
CDH17 AND NOT-GABRB2	Colon	0.9091	0.8333	1
CDH17 AND NOT-DISP2	Colon	1	1	1
NOX1 AND NOT-SLC16A5	Colon	0.8889	1	0.8
CDH17 AND NOT-SLC16A5	Colon	0.8889	1	0.8
FAT1 AND NOT-SLC22A5	Colon	0.8	0.8	0.8
IFI6 AND NOT-OXTR	Colon	0.8889	1	0.8
IFI6 AND NOT-SLC6A15	Colon	0.7273	0.6667	0.8
IFI6 AND NOT-PAQR7	Colon	0.9091	0.8333	1
NOX1 AND NOT-DISP2	Colon	1	1	1
IFI6 AND NOT-CHRND	Colon	0.8333	0.7143	1
NOX1 AND NOT-ATP8A1	Colon	0.8889	1	0.8
IFI6 AND NOT-SLCO6A1	Colon	0.9091	0.8333	1
FAT1 AND NOT-MMP24	Colon	0.6667	0.5714	0.8

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
CD276 AND NOT-IL20RA	Leiomyosarcoma	0.6364	0.7778	0.5385
CD276 AND NOT-EPCAM	Leiomyosarcoma	0.75	0.8182	0.6923
FAP AND NOT-CLDN1	Leiomyosarcoma	0.6316	1	0.4615
FAP AND NOT-ERBB3	Leiomyosarcoma	0.7619	1	0.6154
FAP AND NOT-RNF43	Leiomyosarcoma	0.7619	1	0.6154
FAP AND NOT-CLDN6	Leiomyosarcoma	0.8182	1	0.6923
FAP AND NOT-IL20RA	Leiomyosarcoma	0.6316	1	0.4615
FAP AND NOT-CD79A	Leiomyosarcoma	0.8182	1	0.6923
FAP AND NOT-EDNRB	Leiomyosarcoma	0.7619	1	0.6154
FAP AND NOT-IL3RA	Leiomyosarcoma	0.8182	1	0.6923
CD276 AND NOT-CLDN8	Leiomyosarcoma	0.75	0.8182	0.6923
FAP AND NOT-CLDN8	Leiomyosarcoma	0.7619	1	0.6154
FAP AND NOT-CLDN11	Leiomyosarcoma	0.7619	1	0.6154
FAP AND NOT-FCRL2	Leiomyosarcoma	0.8182	1	0.6923
FAP AND NOT-KDR	Leiomyosarcoma	0.7619	1	0.6154
CD37 AND NOT-SLAMF7	AML	0.8783	0.84	0.9203
CD37 AND NOT-MS4A1	AML	0.8743	0.8262	0.9283
CD37 AND NOT-FCRL5	AML	0.8731	0.8211	0.9323
CD37 AND NOT-CR2	AML	0.864	0.802	0.9363
CD37 AND NOT-TNFRSF17	AML	0.8614	0.8127	0.9163
CD37 AND NOT-FCRL1	AML	0.8519	0.7872	0.9283
CD37 AND NOT-CD22	AML	0.8499	0.7781	0.9363
CD37 AND NOT-FCRL2	AML	0.8484	0.7756	0.9363
CD37 AND NOT-CD19	AML	0.8593	0.8121	0.9124
CD37 AND NOT-CD72	AML	0.8412	0.769	0.9283
CD37 AND NOT-CD79A	AML	0.8271	0.7484	0.9243
CD37 AND NOT-CD79B	AML	0.8124	0.7895	0.8367
CD37 AND NOT-IL11RA	AML	0.8078	0.7879	0.8287
SLC7A5 AND NOT-CLDN11	AML	0.8053	0.6873	0.9721
CD37 AND NOT-CXCR5	AML	0.7516	0.626	0.9402
CD37 AND NOT-CLDN5	AML	0.7433	0.6146	0.9402
CD37 AND NOT-ITGAV	AML	0.7422	0.6313	0.9004
CD37 AND NOT-KDR	AML	0.741	0.6114	0.9402
CD37 AND NOT-EDNRB	AML	0.7353	0.6105	0.9243
CD37 AND NOT-STEAP2	AML	0.7352	0.6036	0.9402
CD37 AND NOT-SDC1	AML	0.7352	0.6036	0.9402
CD37 AND NOT-GPNMB	AML	0.734	0.6139	0.9124
CD37 AND NOT-AXL	AML	0.7335	0.6047	0.9323
CD37 AND NOT-STEAP1	AML	0.7316	0.6036	0.9283
CD37 AND NOT-ROR1	AML	0.7284	0.5945	0.9402
CD37 AND NOT-CLDN11	AML	0.725	0.59	0.9402
CD38 AND NOT-SLAMF7	AML	0.771	0.7399	0.8048
CD37 AND NOT-CLDN7	AML	0.7217	0.5856	0.9402
CD37 AND NOT-ERBB3	AML	0.7206	0.5842	0.9402
CD37 AND NOT-MET	AML	0.7205	0.5903	0.9243
CD37 AND NOT-VCAM1	AML	0.7204	0.6134	0.8725
CD37 AND NOT-ERBB2	AML	0.7195	0.5827	0.9402
CD37 AND NOT-SLC34A2	AML	0.7173	0.5799	0.9402
CD37 AND NOT-IL13RA1	AML	0.6968	0.5854	0.8606
SLC7A5 AND NOT-FOLH1	AML	0.7579	0.626	0.9602
CD38 AND NOT-CLDN23	AML	0.6787	0.5787	0.8207
CD38 AND NOT-CLDN5	AML	0.6732	0.5687	0.8247
CD38 AND NOT-ERBB2	AML	0.6635	0.555	0.8247
P2RX5 AND NOT-SLAMF7	AML	0.6616	0.5358	0.8645
SLC7A5 AND NOT-BMPR1B	AML	0.7397	0.5985	0.9681
CD38 AND NOT-CLDN7	AML	0.6551	0.5433	0.8247
CD38 AND NOT-CLDN8	AML	0.6551	0.5433	0.8247
CD38 AND NOT-RNF43	AML	0.653	0.5405	0.8247
CD38 AND NOT-CD19	AML	0.7034	0.6201	0.8127
CD38 AND NOT-EGFR	AML	0.6509	0.5377	0.8247
CD38 AND NOT-ITGB6	AML	0.6479	0.5335	0.8247
CD38 AND NOT-EPCAM	AML	0.6455	0.537	0.8088
CD38 AND NOT-CLDN1	AML	0.6419	0.5254	0.8247
CD38 AND NOT-IL20RA	AML	0.6409	0.5241	0.8247
SLC7A5 AND NOT-EGFR	AML	0.6404	0.4775	0.9721
CD38 AND NOT-TRPM4	AML	0.6403	0.5316	0.8048
SLC7A5 AND NOT-CLDN23	AML	0.6402	0.4792	0.9641
CD276 AND NOT-CLDN1	Liposarcoma	0.7429	0.7647	0.7222
ROR1 AND NOT-MET	Liposarcoma	0.6545	0.9474	0.5
ROR1 AND NOT-SLC7A5	Liposarcoma	0.6316	0.8571	0.5
EPHB2 AND NOT-MUC13	Liposarcoma	0.6757	0.6579	0.6944
ROR1 AND NOT-EPCAM	Liposarcoma	0.6207	0.8182	0.5
ROR1 AND NOT-SLC34A2	Liposarcoma	0.6182	0.8947	0.4722
ROR1 AND NOT-CD52	Liposarcoma	0.6	0.75	0.5
EPHB2 AND NOT-GPA33	Liposarcoma	0.6301	0.6216	0.6389
EPHB2 AND NOT-CLDN7	Liposarcoma	0.6667	0.5686	0.8056
CD276 AND NOT-PROM1	Liposarcoma	0.7778	0.7778	0.7778

FIG. 12 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
NOX1 AND NOT-CDH10	Colon	0.6667	0.5	1
IFI6 AND NOT-GPR85	Colon	0.75	1	0.6
NOX1 AND NOT-CLDN10	Colon	0.6667	0.75	0.6
MUC12 AND NOT-CLDN10	Colon	0.6667	0.75	0.6
MUC12 AND NOT-EPHA10	Colon	0.6667	0.75	0.6
CDH17 AND NOT-CLDN10	Colon	0.6667	0.75	0.6
FUT3 AND NOT-CLDN10	Colon	0.6667	0.75	0.6
IFI6 AND NOT-MOG	Colon	0.8333	0.7143	1
IFI6 AND NOT-MDGA2	Colon	0.9091	0.8333	1
IFI6 AND NOT-TSHR	Colon	0.8333	0.7143	1
IFI6 AND NOT-OPRK1	Colon	0.8333	0.7143	1
IFI6 AND NOT-GRIN2C	Colon	0.7692	0.625	1
CDH17 AND NOT-ATP8A1	Colon	0.8889	1	0.8
FAT1 AND NOT-GABRA2	Colon	0.75	1	0.6
IFI6 AND NOT-LRRTM4	Colon	0.75	1	0.6
IFI6 AND NOT-GABRA5	Colon	0.8333	0.7143	1
NOX1 AND NOT-PCDHA6	Colon	1	1	1
IFI6 AND NOT-SLC9B1	Colon	0.9091	0.8333	1
CDH17 AND NOT-KIRREL3	Colon	0.75	1	0.6
IFI6 AND NOT-DYSF	Colon	0.7273	0.6667	0.8
NOX1 AND NOT-KIRREL3	Colon	0.75	1	0.6
FAT1 AND NOT-SLC13A2	Colon	0.8	0.8	0.8
FAT1 AND NOT-ANPEP	Colon	0.75	1	0.6
IFI6 AND NOT-DSC1	Colon	0.9091	0.8333	1
CDH17 AND NOT-ABHD3	Colon	0.8889	1	0.8
CDH17 AND NOT-CD1B	Colon	0.625	0.4545	1
IFI6 AND NOT-PIRT	Colon	0.8333	0.7143	1
NOX1 AND NOT-CD1B	Colon	0.625	0.4545	1
FUT3 AND NOT-CHRND	Colon	0.625	0.4545	1
GRIN2A AND NOT-ATP2B2	Ependymoma	1	1	1
GRIN2A AND NOT-ANO4	Ependymoma	1	1	1
GRIN2A AND NOT-SYT4	Ependymoma	1	1	1
GRIN2A AND NOT-NKAIN2	Ependymoma	1	1	1
GRIN2A AND NOT-GRM5	Ependymoma	1	1	1
GRIN2A AND NOT-SCN2A	Ependymoma	1	1	1
GRIN2A AND NOT-SCN1A	Ependymoma	1	1	1
GRIN2A AND NOT-PTPRR	Ependymoma	1	1	1
GRIN2A AND NOT-GABRA1	Ependymoma	1	1	1
GRIN2A AND NOT-GABRA2	Ependymoma	1	1	1
GRIN2A AND NOT-GABRA4	Ependymoma	1	1	1
GRIN2A AND NOT-GABRB2	Ependymoma	1	1	1
GRIN2A AND NOT-SLC12A5	Ependymoma	1	1	1
GRIN2A AND NOT-SLC24A2	Ependymoma	1	1	1
GRIN2A AND NOT-CNTNAP2	Ependymoma	1	1	1
GRIN2A AND NOT-JPH3	Ependymoma	1	1	1
GRIN2A AND NOT-ATP8A2	Ependymoma	1	1	1
GRIN2A AND NOT-PCDH8	Ependymoma	1	1	1
GRIN2A AND NOT-TMEM235	Ependymoma	1	1	1
GRIN2A AND NOT-KCNC1	Ependymoma	1	1	1
GRIN2A AND NOT-GRM7	Ependymoma	1	1	1
GRIN2A AND NOT-SLC32A1	Ependymoma	1	1	1
GRIN2A AND NOT-SCN8A	Ependymoma	1	1	1
GRIN2A AND NOT-GRM3	Ependymoma	1	1	1
GRIN2A AND NOT-CSMD2	Ependymoma	1	1	1
GRIN2A AND NOT-CACNG3	Ependymoma	1	1	1
GRIN2A AND NOT-UNC5A	Ependymoma	1	1	1
GRIN2A AND NOT-CDH18	Ependymoma	1	1	1
GRIN2A AND NOT-KCNV1	Ependymoma	1	1	1
GRIN2A AND NOT-OPCML	Ependymoma	1	1	1
GRIN2A AND NOT-KCNJ6	Ependymoma	0.6667	0.5	1
GRIN2A AND NOT-CACNG8	Ependymoma	0.6667	0.5	1
GRIN2A AND NOT-HCN1	Ependymoma	0.6667	0.5	1
GRIN2A AND NOT-OPALIN	Ependymoma	1	1	1
MEGF10 AND NOT-NKAIN2	Ependymoma	1	1	1
GRIN2A AND NOT-CNTNAP4	Ependymoma	1	1	1
GRIN2A AND NOT-GABRG2	Ependymoma	1	1	1
MEGF10 AND NOT-SEMA4D	Ependymoma	1	1	1
TNFRSF10B AND NOT-KCNK6	Ependymoma	1	1	1
GRIN2A AND NOT-ASTN1	Ependymoma	1	1	1
TNFRSF10B AND NOT-IFNAR2	Ependymoma	1	1	1
GRIN2A AND NOT-GPR158	Ependymoma	1	1	1
TNFRSF10B AND NOT-SLC50A1	Ependymoma	0.6667	0.5	1
TNFRSF10B AND NOT-CD58	Ependymoma	1	1	1
GRIN2A AND NOT-CDH10	Ependymoma	1	1	1
SLCO1B3 AND NOT-SLC1A2	Esophagus	0.6667	1	0.5
SLCO1B3 AND NOT-ABCG8	Esophagus	0.6667	1	0.5
SLCO1B3 AND NOT-SLC23A2	Esophagus	0.6	1	0.4286

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
CD276 AND NOT-ERBB2	Liposarcoma	0.6471	0.6875	0.6111
EPHB2 AND NOT-TRPM4	Liposarcoma	0.6024	0.5319	0.6944
CD276 AND NOT-EGFR	Liposarcoma	0.8533	0.8205	0.8889
CD276 AND NOT-SDC1	Liposarcoma	0.6757	0.6579	0.6944
CD276 AND NOT-MET	Liposarcoma	0.6667	0.7778	0.5833
CD276 AND NOT-IGF1R	Liposarcoma	0.8312	0.7805	0.8889
CD276 AND NOT-SLC34A2	Liposarcoma	0.7674	0.66	0.9167
CD276 AND NOT-SLC7A5	Liposarcoma	0.6133	0.5897	0.6389
SLC34A2 AND NOT-ERBB4	Lung Adenocarcinoma	0.6179	0.717	0.5429
SLC34A2 AND NOT-BMPR1B	Lung Adenocarcinoma	0.6125	0.5444	0.7
SLC34A2 AND NOT-CLDN8	Lung Adenocarcinoma	0.6316	0.5854	0.6857
SLC34A2 AND NOT-ABCA5	Lung Adenocarcinoma	0.6093	0.5679	0.6571
SLC34A2 AND NOT-MUC16	Lung Adenocarcinoma	0.6	0.5333	0.6857
SLC34A2 AND NOT-PSCA	Lung Adenocarcinoma	0.6093	0.5679	0.6571
SLC34A2 AND NOT-SSTR4	Lung Adenocarcinoma	0.6667	0.6145	0.7286
SLC34A2 AND NOT-GPA33	Lung Adenocarcinoma	0.6508	0.7321	0.5857
CD79A AND NOT-ITGB3	B-Cell Diffuse	0.6957	0.75	0.6486
CD79B AND NOT-ITGB3	B-Cell Diffuse	0.6944	0.7143	0.6757
CD79B AND NOT-CD33	B-Cell Diffuse	0.6765	0.7419	0.6216
CXCR5 AND NOT-GPA33	B-Cell Diffuse	0.6667	0.6579	0.6757
IL2RA AND NOT-GPA33	B-Cell Diffuse	0.6984	0.8462	0.5946
CD79A AND NOT-CD33	B-Cell Diffuse	0.6571	0.697	0.6216
CD180 AND NOT-CD33	B-Cell Diffuse	0.6479	0.6765	0.6216
P2RX5 AND NOT-ITGB3	B-Cell Diffuse	0.6829	0.6222	0.7568
IL2RA AND NOT-NCAM1	B-Cell Diffuse	0.6667	0.6857	0.6486
FCRL5 AND NOT-NCAM1	B-Cell Diffuse	0.6575	0.6667	0.6486
IL2RA AND NOT-ITGB3	B-Cell Diffuse	0.6774	0.84	0.5676
FCRL5 AND NOT-ITGB3	B-Cell Diffuse	0.6933	0.6842	0.7027
VCAM1 AND NOT-CLDN1	B-Cell Diffuse	0.6329	0.5952	0.6757
CD79B AND NOT-NCAM1	B-Cell Diffuse	0.6301	0.6389	0.6216
P2RX5 AND NOT-CD33	B-Cell Diffuse	0.6585	0.6	0.7297
CD180 AND NOT-ITGB3	B-Cell Diffuse	0.6269	0.7	0.5676
CD79A AND NOT-GPA33	B-Cell Diffuse	0.623	0.7917	0.5135
CD72 AND NOT-CD33	B-Cell Diffuse	0.6176	0.6774	0.5676
CD79B AND NOT-GPA33	B-Cell Diffuse	0.6176	0.6774	0.5676
CXCR5 AND NOT-ITGB3	B-Cell Diffuse	0.6897	0.6	0.8108
VCAM1 AND NOT-SDC1	B-Cell Diffuse	0.6098	0.5556	0.6757
VCAM1 AND NOT-CLDN11	B-Cell Diffuse	0.6667	0.6579	0.6757
CD72 AND NOT-NCAM1	B-Cell Diffuse	0.6027	0.6111	0.5946
IL2RA AND NOT-IL20RA	B-Cell Diffuse	0.6582	0.619	0.7027
IL2RA AND NOT-MSLN	B-Cell Diffuse	0.6667	0.6136	0.7297
IL2RA AND NOT-CLDN9	Anaplastic Lymphoma	0.6923	0.8182	0.6
IL2RA AND NOT-ERBB3	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-ITGB3	Anaplastic Lymphoma	0.6667	0.8889	0.5333
IL2RA AND NOT-CLDN2	Anaplastic Lymphoma	0.6923	0.8182	0.6
IL2RA AND NOT-CLDN6	Anaplastic Lymphoma	0.6154	0.7273	0.5333
IL2RA AND NOT-GUCY2C	Anaplastic Lymphoma	0.6154	0.7273	0.5333
IL2RA AND NOT-ULBP1	Anaplastic Lymphoma	0.6154	0.7273	0.5333
IL2RA AND NOT-SSTR1	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-CLDN18	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-IL20RA	Anaplastic Lymphoma	0.6154	0.7273	0.5333
IL2RA AND NOT-CSPG4	Anaplastic Lymphoma	0.6154	0.7273	0.5333
IL2RA AND NOT-LGR5	Anaplastic Lymphoma	0.6429	0.6923	0.6
IL2RA AND NOT-MUC4	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-SSTR3	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-TNFRSF13C	Anaplastic Lymphoma	0.6154	0.7273	0.5333
IL2RA AND NOT-MSLN	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-CLDN23	Anaplastic Lymphoma	0.6957	1	0.5333
IL2RA AND NOT-GPA33	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-BMPR1B	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-SLC34A2	Anaplastic Lymphoma	0.6429	0.6923	0.6
IL2RA AND NOT-SSTR5	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-MUC16	Anaplastic Lymphoma	0.6154	0.7273	0.5333
IL2RA AND NOT-ITGB6	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-PMEL	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-IL13RA1	Anaplastic Lymphoma	0.6154	0.7273	0.5333
IL2RA AND NOT-ERBB2	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-CXCR5	Anaplastic Lymphoma	0.6429	0.6923	0.6
IL2RA AND NOT-TNFRSF10A	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-FCRL1	Anaplastic Lymphoma	0.6429	0.6923	0.6
IL2RA AND NOT-CD22	Anaplastic Lymphoma	0.6429	0.6923	0.6
IL2RA AND NOT-CA9	Anaplastic Lymphoma	0.6429	0.6923	0.6
IL2RA AND NOT-SSTR4	Anaplastic Lymphoma	0.6429	0.6923	0.6
IL2RA AND NOT-NCAM1	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-ABCB5	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-VTCN1	Anaplastic Lymphoma	0.6429	0.6923	0.6
IL2RA AND NOT-EGFR	Anaplastic Lymphoma	0.6667	0.75	0.6

FIG. 12 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
SLCO1B3 AND NOT-ATP2B2	Esophagus	0.6	1	0.4286
SLCO1B3 AND NOT-DIO1	Esophagus	0.6667	1	0.5
SLCO1B3 AND NOT-SLC30A10	Esophagus	0.6667	1	0.5
SLCO1B3 AND NOT-SLC22A9	Esophagus	0.6	1	0.4286
SLCO1B3 AND NOT-SLC17A2	Esophagus	0.6667	1	0.5
SLCO1B3 AND NOT-SLC10A1	Esophagus	0.6667	1	0.5
SLCO1B3 AND NOT-TRPM8	Esophagus	0.6364	0.875	0.5
MUC17 AND NOT-FXYD7	Esophagus	0.6364	0.875	0.5
SLCO1B3 AND NOT-SLC28A1	Esophagus	0.6	1	0.4286
SLCO1B3 AND NOT-CYP4A11	Esophagus	0.6667	1	0.5
SLCO1B3 AND NOT-GHR	Esophagus	0.6364	0.875	0.5
MUC17 AND NOT-ABHD3	Esophagus	0.6364	0.875	0.5
MUC17 AND NOT-SLC30A10	Esophagus	0.88	1	0.7857
MUC17 AND NOT-SLC22A5	Esophagus	0.7273	1	0.5714
MUC17 AND NOT-ABCG8	Esophagus	0.75	0.9	0.6429
SLCO1B3 AND NOT-SLC13A5	Esophagus	0.6667	1	0.5
MUC17 AND NOT-SLC5A11	Esophagus	0.6957	0.8889	0.5714
MUC17 AND NOT-CD36	Esophagus	0.7273	1	0.5714
LYPD1 AND NOT-CNTNAP2	Glioblastoma	0.8	0.7778	0.8235
LYPD1 AND NOT-GRM5	Glioblastoma	0.7368	0.6667	0.8235
LYPD1 AND NOT-KIAA0319	Glioblastoma	0.7742	0.8571	0.7059
LYPD1 AND NOT-CDH8	Glioblastoma	0.75	0.8	0.7059
LYPD1 AND NOT-ATP8A2	Glioblastoma	0.7273	0.75	0.7059
LYPD1 AND NOT-GABRG1	Glioblastoma	0.8125	0.8667	0.7647
LYPD1 AND NOT-GRM3	Glioblastoma	0.7273	0.75	0.7059
LYPD1 AND NOT-NKAIN2	Glioblastoma	0.7059	0.7059	0.7059
LYPD1 AND NOT-SYT4	Glioblastoma	0.7568	0.7	0.8235
LYPD1 AND NOT-SLC6A15	Glioblastoma	0.7027	0.65	0.7647
LYPD1 AND NOT-CACNA1B	Glioblastoma	0.7027	0.65	0.7647
LYPD1 AND NOT-DGKE	Glioblastoma	0.7027	0.65	0.7647
LYPD1 AND NOT-CALN1	Glioblastoma	0.6875	0.7333	0.6471
LYPD1 AND NOT-ATP2B2	Glioblastoma	0.6857	0.6667	0.7059
LYPD1 AND NOT-SCN2A	Glioblastoma	0.6857	0.6667	0.7059
LYPD1 AND NOT-SCN8A	Glioblastoma	0.6857	0.6667	0.7059
LYPD1 AND NOT-TRPM3	Glioblastoma	0.7179	0.6364	0.8235
LYPD1 AND NOT-AJAP1	Glioblastoma	0.6842	0.619	0.7647
LYPD1 AND NOT-PPAPDC1B	Glioblastoma	0.6829	0.5833	0.8235
LYPD1 AND NOT-GABRA2	Glioblastoma	0.6667	0.5909	0.7647
LYPD1 AND NOT-ANO4	Glioblastoma	0.6667	0.6875	0.6471
LYPD1 AND NOT-SLC6A6	Glioblastoma	0.6667	0.56	0.8235
LYPD1 AND NOT-PPAPDC1A	Glioblastoma	0.6667	0.56	0.8235
LYPD1 AND NOT-NFASC	Glioblastoma	0.6667	0.56	0.8235
LYPD1 AND NOT-ATP8B1	Glioblastoma	0.6667	0.56	0.8235
SYT11 AND NOT-SLC6A15	Glioma	0.9481	0.9697	0.9275
SYT11 AND NOT-CDH8	Glioma	0.9645	0.9444	0.9855
SYT11 AND NOT-DGKE	Glioma	0.9781	0.9853	0.971
SYT11 AND NOT-SCN8A	Glioma	0.964	0.9571	0.971
SYT11 AND NOT-ATP8A2	Glioma	0.9853	1	0.971
SYT11 AND NOT-SLC32A1	Glioma	0.9496	0.9429	0.9565
SYT11 AND NOT-GRM1	Glioma	0.9496	0.9429	0.9565
SYT11 AND NOT-KIAA0319	Glioma	0.9559	0.9701	0.942
SYT11 AND NOT-ABCG4	Glioma	0.9781	0.9853	0.971
SYT11 AND NOT-KCNA2	Glioma	0.9714	0.9577	0.9855
SYT11 AND NOT-CACNA1B	Glioma	0.971	0.971	0.971
SYT11 AND NOT-NKAIN2	Glioma	0.9853	1	0.971
SYT11 AND NOT-GPR26	Glioma	0.9571	0.9437	0.971
SYT11 AND NOT-HCN1	Glioma	0.9781	0.9853	0.971
SYT11 AND NOT-KCNA1	Glioma	0.9853	1	0.971
SYT11 AND NOT-CACNA1E	Glioma	0.9496	0.9429	0.9565
SYT11 AND NOT-ATP2B2	Glioma	0.9571	0.9437	0.971
SYT11 AND NOT-KCNJ6	Glioma	0.9571	0.9437	0.971
SYT11 AND NOT-CNTNAP2	Glioma	0.971	0.971	0.971
SYT11 AND NOT-GRM3	Glioma	0.9645	0.9444	0.9855
SYT11 AND NOT-SLC12A5	Glioma	0.942	0.942	0.942
SYT11 AND NOT-KIRREL3	Glioma	0.9571	0.9437	0.971
SYT11 AND NOT-CACNG3	Glioma	0.9481	0.9697	0.9275
SYT11 AND NOT-GPR83	Glioma	0.9853	1	0.971
SYT11 AND NOT-GPR22	Glioma	0.9645	0.9444	0.9855
SYT11 AND NOT-KCNK9	Glioma	0.964	0.9571	0.971
SYT11 AND NOT-HTR1E	Glioma	0.9714	0.9577	0.9855
SYT11 AND NOT-GRIN1	Glioma	0.9565	0.9565	0.9565
SYT11 AND NOT-GABRA1	Glioma	0.9496	0.9429	0.9565
SYT11 AND NOT-GABRG1	Glioma	0.9265	0.9403	0.913
SYT11 AND NOT-CCKBR	Glioma	0.9645	0.9444	0.9855
SYT11 AND NOT-CALY	Glioma	0.9701	1	0.942
SYT11 AND NOT-SYT4	Glioma	0.9481	0.9697	0.9275
SYT11 AND NOT-KCNJ9	Glioma	0.964	0.9571	0.971

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
IL2RA AND NOT-CD180	Anaplastic Lymphoma	0.6429	0.6923	0.6
IL2RA AND NOT-EDNRB	Anaplastic Lymphoma	0.6429	0.6923	0.6
IL2RA AND NOT-EPHB2	Anaplastic Lymphoma	0.6207	0.6429	0.6
IL2RA AND NOT-ENPP3	Anaplastic Lymphoma	0.6923	0.8182	0.6
IL2RA AND NOT-CLDN1	Anaplastic Lymphoma	0.6207	0.6429	0.6
IL2RA AND NOT-CD19	Anaplastic Lymphoma	0.6207	0.6429	0.6
IL2RA AND NOT-MET	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-EPHA3	Anaplastic Lymphoma	0.6207	0.6429	0.6
IL2RA AND NOT-MUC1	Anaplastic Lymphoma	0.6207	0.6429	0.6
IL2RA AND NOT-IL11RA	Anaplastic Lymphoma	0.6429	0.6923	0.6
IL2RA AND NOT-ROR1	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-CLDN8	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-L1CAM	Anaplastic Lymphoma	0.6207	0.6429	0.6
TNFRSF8 AND NOT-ULBP1	Anaplastic Lymphoma	0.7586	0.7857	0.7333
IL2RA AND NOT-RAET1E	Anaplastic Lymphoma	0.6087	0.875	0.4667
TNFRSF8 AND NOT-SSTR4	Anaplastic Lymphoma	0.875	0.8235	0.9333
TNFRSF8 AND NOT-CLDN9	Anaplastic Lymphoma	0.8	0.8	0.8
TNFRSF8 AND NOT-TNFRSF13C	Anaplastic Lymphoma	0.8387	0.8125	0.8667
TNFRSF8 AND NOT-CLDN6	Anaplastic Lymphoma	0.875	0.8235	0.9333
TNFRSF8 AND NOT-RAET1E	Anaplastic Lymphoma	0.8	0.8	0.8
TNFRSF8 AND NOT-ERBB3	Anaplastic Lymphoma	0.875	0.8235	0.9333
TNFRSF8 AND NOT-SLC34A2	Anaplastic Lymphoma	0.8387	0.8125	0.8667
TNFRSF8 AND NOT-GUCY2C	Anaplastic Lymphoma	0.8387	0.8125	0.8667
TNFRSF8 AND NOT-IGF1R	Anaplastic Lymphoma	0.8387	0.8125	0.8667
TNFRSF8 AND NOT-CD33	Anaplastic Lymphoma	0.8889	1	0.8
TNFRSF8 AND NOT-SSTR1	Anaplastic Lymphoma	0.875	0.8235	0.9333
TNFRSF8 AND NOT-CD180	Anaplastic Lymphoma	0.9655	1	0.9333
TNFRSF8 AND NOT-ITGB3	Anaplastic Lymphoma	0.9286	1	0.8667
TNFRSF8 AND NOT-ERBB2	Anaplastic Lymphoma	0.875	0.8235	0.9333
TNFRSF8 AND NOT-ENPP3	Anaplastic Lymphoma	0.9655	1	0.9333
TNFRSF8 AND NOT-IL13RA1	Anaplastic Lymphoma	0.8387	0.8125	0.8667
TNFRSF8 AND NOT-TNFRSF10A	Anaplastic Lymphoma	0.875	0.8235	0.9333
TNFRSF8 AND NOT-CD79B	Anaplastic Lymphoma	0.9333	0.9333	0.9333
TNFRSF8 AND NOT-SLAMF7	Anaplastic Lymphoma	0.7857	0.8462	0.7333
MS4A1 AND NOT-SSTR2	Mantle-Cell Lymphoma	0.8941	0.8085	1
MS4A1 AND NOT-TNFRSF8	Mantle-Cell Lymphoma	0.8506	0.7551	0.9737
MS4A1 AND NOT-CLDN8	Mantle-Cell Lymphoma	0.8315	0.7255	0.9737
CD52 AND NOT-CD33	Mantle-Cell Lymphoma	0.8675	0.8	0.9474
MS4A1 AND NOT-MST1R	Mantle-Cell Lymphoma	0.814	0.7292	0.9211
MS4A1 AND NOT-SLAMF7	Mantle-Cell Lymphoma	0.809	0.7059	0.9474
CD52 AND NOT-NCAM1	Mantle-Cell Lymphoma	0.8537	0.7955	0.9211
CXCR5 AND NOT-TNFRSF8	Mantle-Cell Lymphoma	0.7937	1	0.6579
CXCR5 AND NOT-ITGB3	Mantle-Cell Lymphoma	0.7937	1	0.6579
CXCR5 AND NOT-NCAM1	Mantle-Cell Lymphoma	0.7937	1	0.6579
CXCR5 AND NOT-ITGB6	Mantle-Cell Lymphoma	0.7937	1	0.6579
CXCR5 AND NOT-MUC4	Mantle-Cell Lymphoma	0.7937	1	0.6579
MS4A1 AND NOT-TRPM4	Mantle-Cell Lymphoma	0.7912	0.6792	0.9474
MS4A1 AND NOT-CLDN6	Mantle-Cell Lymphoma	0.7901	0.7442	0.8421
P2RX5 AND NOT-TNFRSF17	Mantle-Cell Lymphoma	0.7865	0.6863	0.9211
TNFRSF13C AND NOT-TNFRSF8	Mantle-Cell Lymphoma	0.8132	0.6981	0.9737
CXCR5 AND NOT-CLDN2	Mantle-Cell Lymphoma	0.7813	0.9615	0.6579
CXCR5 AND NOT-ULBP1	Mantle-Cell Lymphoma	0.7813	0.9615	0.6579
CXCR5 AND NOT-SSTR5	Mantle-Cell Lymphoma	0.7813	0.9615	0.6579
CD70 AND NOT-ITGB3	Mantle-Cell Lymphoma	0.8986	1	0.8158
CD70 AND NOT-ITGB6	Mantle-Cell Lymphoma	0.8986	1	0.8158
CD70 AND NOT-ENG	Mantle-Cell Lymphoma	0.8986	1	0.8158
CD70 AND NOT-SLC39A6	Mantle-Cell Lymphoma	0.8986	1	0.8158
CD70 AND NOT-IL13RA1	Mantle-Cell Lymphoma	0.8986	1	0.8158
CD79B AND NOT-TNFRSF8	Mantle-Cell Lymphoma	0.7778	0.8235	0.7368
CD70 AND NOT-BMPR1B	Mantle-Cell Lymphoma	0.8986	1	0.8158
CD70 AND NOT-IL20RA	Mantle-Cell Lymphoma	0.8986	1	0.8158
CD72 AND NOT-ITGB3	Mantle-Cell Lymphoma	0.7733	0.7838	0.7632
TNFRSF13C AND NOT-CLDN2	Mantle-Cell Lymphoma	0.8352	0.717	1
CD70 AND NOT-NCAM1	Mantle-Cell Lymphoma	0.8986	1	0.8158
CD70 AND NOT-CSPG4	Mantle-Cell Lymphoma	0.8824	1	0.7895
MS4A1 AND NOT-MET	Mantle-Cell Lymphoma	0.7692	0.75	0.7895
TNFRSF13C AND NOT-ITGB3	Mantle-Cell Lymphoma	0.8444	0.7308	1
CD70 AND NOT-IGF1R	Mantle-Cell Lymphoma	0.8986	1	0.8158
TNFRSF13C AND NOT-SSTR5	Mantle-Cell Lymphoma	0.8172	0.6909	1
CD70 AND NOT-EPHA3	Mantle-Cell Lymphoma	0.8986	1	0.8158
CD70 AND NOT-SSTR1	Mantle-Cell Lymphoma	0.8986	1	0.8158
FCRL1 AND NOT-TNFRSF8	Mantle-Cell Lymphoma	0.7647	0.8667	0.6842
TNFRSF13C AND NOT-SSTR1	Mantle-Cell Lymphoma	0.8261	0.7037	1
TNFRSF13C AND NOT-ITGB6	Mantle-Cell Lymphoma	0.8222	0.7115	0.9737
CD52 AND NOT-TNFRSF8	Mantle-Cell Lymphoma	0.7609	0.6481	0.9211
CD79B AND NOT-TNFRSF17	Mantle-Cell Lymphoma	0.7606	0.8182	0.7105
CD70 AND NOT-KDR	Mantle-Cell Lymphoma	0.8986	1	0.8158

FIG. 12 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
SYT11 AND NOT-SLC26A8	Glioma	0.9714	0.9577	0.9855
SYT11 AND NOT-GPR6	Glioma	0.9489	0.9559	0.942
SYT11 AND NOT-OPRK1	Glioma	0.9624	1	0.9275
SYT11 AND NOT-KCNC1	Glioma	0.9645	0.9444	0.9855
SYT11 AND NOT-UNC5A	Glioma	0.9496	0.9429	0.9565
SYT11 AND NOT-GABRA4	Glioma	0.9496	0.9429	0.9565
SYT11 AND NOT-GABRA2	Glioma	0.942	0.942	0.942
SYT11 AND NOT-AJAP1	Glioma	0.9008	0.9516	0.8551
SYT11 AND NOT-KCNV1	Glioma	0.9645	0.9444	0.9855
SYT11 AND NOT-GABRB2	Glioma	0.9185	0.9394	0.8986
ATP8B2 AND NOT-SELP	Leiomyosarcoma	0.75	0.8182	0.6923
ATP8B2 AND NOT-SLC6A6	Leiomyosarcoma	0.7	1	0.5385
ATP8B2 AND NOT-CD1A	Leiomyosarcoma	0.6923	0.6923	0.6923
ATP8B2 AND NOT-SEMA4D	Leiomyosarcoma	0.8333	0.9091	0.7692
ATP8B2 AND NOT-DGKE	Leiomyosarcoma	0.7619	1	0.6154
ATP8B2 AND NOT-SLC12A9	Leiomyosarcoma	0.8333	0.9091	0.7692
ATP8B2 AND NOT-MPL	Leiomyosarcoma	0.6667	0.7273	0.6154
EMP3 AND NOT-SORL1	Leiomyosarcoma	0.6667	0.7273	0.6154
ATP8B2 AND NOT-CD44	Leiomyosarcoma	0.6667	0.875	0.5385
ATP8B2 AND NOT-KCNK6	Leiomyosarcoma	0.6	0.8571	0.4615
ATP8B2 AND NOT-MLC1	Leiomyosarcoma	0.7407	0.7143	0.7692
ATP8B2 AND NOT-SORL1	Leiomyosarcoma	0.7619	1	0.6154
ATP8B2 AND NOT-LAT	Leiomyosarcoma	0.6	0.8571	0.4615
ATP8B2 AND NOT-BTN3A1	Leiomyosarcoma	0.6667	0.875	0.5385
EMP3 AND NOT-SELP	Leiomyosarcoma	0.8333	0.9091	0.7692
SLC22A16 AND NOT-EFNB2	AML	0.9299	0.9355	0.9243
SLC22A16 AND NOT-STEAP4	AML	0.9256	0.935	0.9163
SLC22A16 AND NOT-EPHA2	AML	0.9231	0.9141	0.9323
SLC22A16 AND NOT-FAT1	AML	0.9215	0.9309	0.9124
SLC22A16 AND NOT-ATP8B1	AML	0.9209	0.9137	0.9283
SLC22A16 AND NOT-LRIG3	AML	0.9176	0.9035	0.9323
SLC22A16 AND NOT-DIO1	AML	0.9052	0.8797	0.9323
SLC22A16 AND NOT-SLC4A8	AML	0.9035	0.8764	0.9323
SLC22A16 AND NOT-PHLDB2	AML	0.9409	0.9625	0.9203
SLC22A16 AND NOT-CATSPERD	AML	0.9017	0.8731	0.9323
SLC22A16 AND NOT-SLC22A5	AML	0.9299	0.9355	0.9243
SLC22A16 AND NOT-KIAA0319	AML	0.9	0.8699	0.9323
SLC22A16 AND NOT-CACNG6	AML	0.9	0.8699	0.9323
SLC22A16 AND NOT-LRRC8E	AML	0.8983	0.8667	0.9323
SLC22A16 AND NOT-OXTR	AML	0.8983	0.8667	0.9323
SLC22A16 AND NOT-GRM7	AML	0.8983	0.8667	0.9323
SLC22A16 AND NOT-SLC31A1	AML	0.8975	0.8722	0.9243
SLC22A16 AND NOT-CNGA4	AML	0.8966	0.8635	0.9323
SLC22A16 AND NOT-ATP6V0A4	AML	0.8966	0.8635	0.9323
SLC22A16 AND NOT-PTPRT	AML	0.8966	0.8635	0.9323
SLC22A16 AND NOT-SMPD2	AML	0.8953	0.8717	0.9203
SLC22A16 AND NOT-GPR137B	AML	0.8941	0.9548	0.8406
SLC22A16 AND NOT-PPAPDC1B	AML	0.8854	0.8784	0.8924
SLC22A16 AND NOT-SLC16A5	AML	0.8789	0.8621	0.8964
SLC22A16 AND NOT-FLVCR1	AML	0.8624	0.8898	0.8367
SLC22A16 AND NOT-CLDN10	AML	0.8571	0.8538	0.8606
SLC22A16 AND NOT-MSMO1	AML	0.8439	0.8969	0.7968
FMNL1 AND NOT-BTN3A1	AML	0.8352	0.788	0.8884
FMNL1 AND NOT-BTN3A3	AML	0.8318	0.7914	0.8765
EMP3 AND NOT-S1PR1	AML	0.8247	0.8247	0.8247
FMNL1 AND NOT-S1PR1	AML	0.818	0.8038	0.8327
FMNL1 AND NOT-LAT	AML	0.8272	0.8068	0.8486
CD44 AND NOT-KCNK6	AML	0.8624	0.7993	0.9363
FMNL1 AND NOT-STEAP4	AML	0.8007	0.7399	0.8725
FMNL1 AND NOT-KCNK6	AML	0.7985	0.7571	0.8446
FMNL1 AND NOT-IFNAR2	AML	0.7954	0.7715	0.8207
FMNL1 AND NOT-SEMA4D	AML	0.7851	0.7083	0.8805
TTYH3 AND NOT-SLC38A1	Liposarcoma	0.6364	0.7	0.5833
STAB1 AND NOT-NFASC	Liposarcoma	0.7302	0.8519	0.6389
STAB1 AND NOT-ATP13A5	Liposarcoma	0.6129	0.7308	0.5278
STAB1 AND NOT-CD1A	Liposarcoma	0.6765	0.7188	0.6389
STAB1 AND NOT-ATP8A1	Liposarcoma	0.6333	0.7917	0.5278
STAB1 AND NOT-SEMA4D	Liposarcoma	0.6667	0.9048	0.5278
STAB1 AND NOT-ADAM29	Liposarcoma	0.6849	0.6757	0.6944
STAB1 AND NOT-DPP10	Liposarcoma	0.6494	0.6098	0.6944
STAB1 AND NOT-ASTN1	Liposarcoma	0.6494	0.6098	0.6944
STAB1 AND NOT-SLC23A2	Liposarcoma	0.6575	0.6486	0.6667
STAB1 AND NOT-HTR1E	Liposarcoma	0.6269	0.6774	0.5833
STAB1 AND NOT-CACNG1	Liposarcoma	0.6027	0.5946	0.6111
STAB1 AND NOT-MSMO1	Liposarcoma	0.641	0.5952	0.6944
STAB1 AND NOT-CALHM1	Liposarcoma	0.6567	0.7097	0.6111
STAB1 AND NOT-SEMA4B	Liposarcoma	0.6102	0.7826	0.5

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
CD70 AND NOT-EDNRB	Mantle-Cell Lymphoma	0.8986	1	0.8158
CD70 AND NOT-SDC1	Mantle-Cell Lymphoma	0.8986	1	0.8158
CD70 AND NOT-EGFR	Mantle-Cell Lymphoma	0.8986	1	0.8158
CD70 AND NOT-CD34	Mantle-Cell Lymphoma	0.8986	1	0.8158
CD70 AND NOT-TPBG	Mantle-Cell Lymphoma	0.8986	1	0.8158
CD70 AND NOT-AXL	Mantle-Cell Lymphoma	0.8986	1	0.8158
CD70 AND NOT-IL11RA	Mantle-Cell Lymphoma	0.8824	1	0.7895
CD70 AND NOT-PTK7	Mantle-Cell Lymphoma	0.8986	1	0.8158
CD70 AND NOT-CLDN1	Mantle-Cell Lymphoma	0.8986	1	0.8158
CD70 AND NOT-ITGAV	Mantle-Cell Lymphoma	0.8857	0.9688	0.8158
CD70 AND NOT-MET	Mantle-Cell Lymphoma	0.8986	1	0.8158
CD70 AND NOT-ERBB2	Mantle-Cell Lymphoma	0.8986	1	0.8158
CD70 AND NOT-CD276	Mantle-Cell Lymphoma	0.8986	1	0.8158
CD70 AND NOT-CLDN11	Mantle-Cell Lymphoma	0.8986	1	0.8158
CD70 AND NOT-FAP	Mantle-Cell Lymphoma	0.8986	1	0.8158
CD70 AND NOT-STEAP2	Mantle-Cell Lymphoma	0.8657	1	0.7632
CD70 AND NOT-STEAP1	Mantle-Cell Lymphoma	0.8657	1	0.7632
FCRL1 AND NOT-ITGB3	Mantle-Cell Lymphoma	0.7826	0.871	0.7105
CXCR5 AND NOT-CSPG4	Mantle-Cell Lymphoma	0.7576	0.8929	0.6579
CD70 AND NOT-MUC4	Mantle-Cell Lymphoma	0.8824	1	0.7895
CD70 AND NOT-LGR5	Mantle-Cell Lymphoma	0.8657	1	0.7632
CD70 AND NOT-SSTR5	Mantle-Cell Lymphoma	0.8824	1	0.7895
CD70 AND NOT-ERBB3	Mantle-Cell Lymphoma	0.8986	1	0.8158
CD52 AND NOT-ITGB3	Mantle-Cell Lymphoma	0.9114	0.878	0.9474
TNFRSF13C AND NOT-NCAM1	Mantle-Cell Lymphoma	0.8736	0.7755	1
EDNRB AND NOT-ITGB6	Melanoma	0.9	1	0.8182
EDNRB AND NOT-CLDN18	Melanoma	0.9	1	0.8182
EDNRB AND NOT-TNFRSF13C	Melanoma	0.9	1	0.8182
EDNRB AND NOT-CLDN6	Melanoma	0.9	1	0.8182
EDNRB AND NOT-MSLN	Melanoma	0.9	1	0.8182
EDNRB AND NOT-CLDN5	Melanoma	0.9	1	0.8182
EDNRB AND NOT-GUCY2C	Melanoma	0.9	1	0.8182
EDNRB AND NOT-SSTR1	Melanoma	0.9	1	0.8182
EDNRB AND NOT-CLDN7	Melanoma	0.9	1	0.8182
EDNRB AND NOT-FOLR1	Melanoma	0.9	1	0.8182
EDNRB AND NOT-GPA33	Melanoma	0.9	1	0.8182
EDNRB AND NOT-EPCAM	Melanoma	0.9	1	0.8182
EDNRB AND NOT-SLC34A2	Melanoma	0.9	1	0.8182
EDNRB AND NOT-ENG	Melanoma	0.9	1	0.8182
CLDN12 AND NOT-ULBP1	Melanoma	0.7778	1	0.6364
CLDN12 AND NOT-VTCN1	Melanoma	0.7778	1	0.6364
CLDN12 AND NOT-SSTR1	Melanoma	0.7778	1	0.6364
CLDN12 AND NOT-FOLH1	Melanoma	0.7778	1	0.6364
CLDN12 AND NOT-SSTR3	Melanoma	0.7778	1	0.6364
CLDN12 AND NOT-SSTR4	Melanoma	0.7778	1	0.6364
CLDN12 AND NOT-SSTR5	Melanoma	0.7778	1	0.6364
CLDN12 AND NOT-CLDN7	Melanoma	0.7778	1	0.6364
CLDN12 AND NOT-EPCAM	Melanoma	0.7778	1	0.6364
CLDN12 AND NOT-ERBB4	Melanoma	0.7778	1	0.6364
CLDN12 AND NOT-ERBB2	Melanoma	0.7778	1	0.6364
CLDN12 AND NOT-MUC16	Melanoma	0.7778	1	0.6364
CLDN12 AND NOT-SLC34A2	Melanoma	0.7778	1	0.6364
CLDN12 AND NOT-CLDN9	Melanoma	0.7778	1	0.6364
CLDN12 AND NOT-CLDN1	Melanoma	0.7778	1	0.6364
CLDN12 AND NOT-CLDN2	Melanoma	0.7778	1	0.6364
CLDN12 AND NOT-CLDN6	Melanoma	0.7778	1	0.6364
CLDN12 AND NOT-CLDN8	Melanoma	0.7778	1	0.6364
CLDN12 AND NOT-IL20RA	Melanoma	0.7778	1	0.6364
CLDN12 AND NOT-ENPP3	Melanoma	0.7778	1	0.6364
CLDN12 AND NOT-MUC4	Melanoma	0.7778	1	0.6364
CLDN12 AND NOT-STEAP1	Melanoma	0.7778	1	0.6364
CLDN12 AND NOT-LGR5	Melanoma	0.7778	1	0.6364
CLDN12 AND NOT-EPHA3	Melanoma	0.7778	1	0.6364
CLDN12 AND NOT-CLDN23	Melanoma	0.7778	1	0.6364
CLDN12 AND NOT-GUCY2C	Melanoma	0.7778	1	0.6364
CLDN12 AND NOT-MUC13	Melanoma	0.7778	1	0.6364
CLDN12 AND NOT-GPA33	Melanoma	0.7778	1	0.6364
ALK AND NOT-CLDN11	Neuroblastoma	0.8625	0.9079	0.8214
ALK AND NOT-ROR1	Neuroblastoma	0.8625	0.9079	0.8214
ALK AND NOT-SLC34A2	Neuroblastoma	0.8535	0.9178	0.7976
ALK AND NOT-MOK	Neuroblastoma	0.8625	0.9079	0.8214
ALK AND NOT-AXL	Neuroblastoma	0.8625	0.9079	0.8214
ALK AND NOT-CSPG4	Neuroblastoma	0.8625	0.9079	0.8214
ALK AND NOT-TNC	Neuroblastoma	0.8535	0.9178	0.7976
ALK AND NOT-EPHB2	Neuroblastoma	0.8553	0.9067	0.8095
ALK AND NOT-SLC7A5	Neuroblastoma	0.8481	0.9054	0.7976
ALK AND NOT-TPBG	Neuroblastoma	0.8466	0.8734	0.8214

FIG. 12 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
STAB1 AND NOT-GABRB2	Liposarcoma	0.6567	0.7097	0.6111
KCNJ10 AND NOT-NFASC	B-Cell Diffuse	0.7826	0.8438	0.7297
KCNJ10 AND NOT-OPCML	B-Cell Diffuse	0.7532	0.725	0.7838
KCNJ10 AND NOT-CDH10	B-Cell Diffuse	0.7324	0.7647	0.7027
KCNJ10 AND NOT-DTNA	B-Cell Diffuse	0.7297	0.7297	0.7297
KCNJ10 AND NOT-ASTN1	B-Cell Diffuse	0.7632	0.7436	0.7838
KCNJ10 AND NOT-SLCO1C1	B-Cell Diffuse	0.7059	0.625	0.8108
KCNJ10 AND NOT-FXYD7	B-Cell Diffuse	0.7013	0.675	0.7297
KCNJ10 AND NOT-OMG	B-Cell Diffuse	0.7013	0.675	0.7297
KCNJ10 AND NOT-ATP2B2	B-Cell Diffuse	0.6897	0.6	0.8108
KCNJ10 AND NOT-KIRREL3	B-Cell Diffuse	0.6897	0.6	0.8108
KCNJ10 AND NOT-CSMD3	B-Cell Diffuse	0.6977	0.6122	0.8108
KCNJ10 AND NOT-KCNA2	B-Cell Diffuse	0.6824	0.6042	0.7838
KCNJ10 AND NOT-GRM3	B-Cell Diffuse	0.6824	0.6042	0.7838
KCNJ10 AND NOT-DISP2	B-Cell Diffuse	0.7324	0.7647	0.7027
KCNJ10 AND NOT-SYT4	B-Cell Diffuse	0.6667	0.566	0.8108
KCNJ10 AND NOT-MEGF10	B-Cell Diffuse	0.6667	0.6136	0.7297
KCNJ10 AND NOT-TRPM3	B-Cell Diffuse	0.6667	0.5957	0.7568
KCNJ10 AND NOT-GPR158	B-Cell Diffuse	0.6512	0.5714	0.7568
KCNJ10 AND NOT-SLC30A10	B-Cell Diffuse	0.6452	0.5357	0.8108
KCNJ10 AND NOT-SLC24A2	B-Cell Diffuse	0.6383	0.5263	0.8108
KCNJ10 AND NOT-KCNK10	B-Cell Diffuse	0.6353	0.5625	0.7297
KCNJ10 AND NOT-CNTNAP2	B-Cell Diffuse	0.6301	0.6389	0.6216
KCNJ10 AND NOT-CACNG3	B-Cell Diffuse	0.6279	0.551	0.7297
KCNJ10 AND NOT-ANO4	B-Cell Diffuse	0.6279	0.551	0.7297
KCNJ10 AND NOT-NRG3	B-Cell Diffuse	0.6105	0.5	0.7838
KCNJ10 AND NOT-GABRG1	B-Cell Diffuse	0.6024	0.5435	0.6757
GPR19 AND NOT-GABRB2	Anaplastic Lymphoma	0.6	0.6	0.6
LAPTM5 AND NOT-ANPEP	Mantle-Cell Lymphoma	0.7789	0.6491	0.9737
LAPTM5 AND NOT-CD93	Mantle-Cell Lymphoma	0.7677	0.623	1
LAPTM5 AND NOT-SLC31A1	Mantle-Cell Lymphoma	0.7037	0.5429	1
LAPTM5 AND NOT-STEAP4	Mantle-Cell Lymphoma	0.7835	0.6441	1
LAPTM5 AND NOT-CXCL16	Mantle-Cell Lymphoma	0.6966	0.6078	0.8158
LAPTM5 AND NOT-DYSF	Mantle-Cell Lymphoma	0.6972	0.5352	1
LAPTM5 AND NOT-CD163	Mantle-Cell Lymphoma	0.6786	0.5135	1
SLC38A1 AND NOT-MLC1	Mantle-Cell Lymphoma	0.6761	0.7273	0.6316
LAPTM5 AND NOT-CD36	Mantle-Cell Lymphoma	0.6607	0.5	0.9737
SLC38A1 AND NOT-SLC31A1	Mantle-Cell Lymphoma	0.6588	0.5957	0.7368
CLDN15 AND NOT-CD36	Mantle-Cell Lymphoma	0.65	0.619	0.6842
CLDN15 AND NOT-SLC16A5	Mantle-Cell Lymphoma	0.6462	0.7778	0.5526
SLC38A1 AND NOT-CD93	Mantle-Cell Lymphoma	0.6437	0.5714	0.7368
CLDN15 AND NOT-PROCR	Mantle-Cell Lymphoma	0.6506	0.6	0.7105
SLC38A1 AND NOT-MPL	Mantle-Cell Lymphoma	0.6588	0.5957	0.7368
SLC38A1 AND NOT-KCNC2	Mantle-Cell Lymphoma	0.7105	0.7105	0.7105
CLDN15 AND NOT-SLC22A18	Mantle-Cell Lymphoma	0.6506	0.6	0.7105
SLC38A1 AND NOT-SLC22A13	Mantle-Cell Lymphoma	0.6301	0.6571	0.6053
SLC38A1 AND NOT-CALHM3	Mantle-Cell Lymphoma	0.7013	0.6923	0.7105
SLC38A1 AND NOT-GRIN1	Mantle-Cell Lymphoma	0.6667	0.6279	0.7105
SLC38A1 AND NOT-CALY	Mantle-Cell Lymphoma	0.7027	0.7222	0.6842
SLC38A1 AND NOT-CD36	Mantle-Cell Lymphoma	0.6207	0.551	0.7105
SLC38A1 AND NOT-ANPEP	Mantle-Cell Lymphoma	0.6207	0.551	0.7105
SLC38A1 AND NOT-DYSF	Mantle-Cell Lymphoma	0.7	0.6667	0.7368
LAPTM5 AND NOT-TGFBI	Mantle-Cell Lymphoma	0.6154	0.4848	0.8421
SLC38A1 AND NOT-KCNK4	Mantle-Cell Lymphoma	0.6571	0.7188	0.6053
BTN3A1 AND NOT-DYSF	Mantle-Cell Lymphoma	0.7294	0.6596	0.8158
SLC38A1 AND NOT-SYT6	Mantle-Cell Lymphoma	0.7164	0.8276	0.6316
SLC38A1 AND NOT-STEAP4	Mantle-Cell Lymphoma	0.6437	0.5714	0.7368
CLDN15 AND NOT-CXCL16	Mantle-Cell Lymphoma	0.6053	0.6053	0.6053
SLC38A1 AND NOT-GRIK1	Mantle-Cell Lymphoma	0.6027	0.6286	0.5789
BTN3A1 AND NOT-MLC1	Mantle-Cell Lymphoma	0.6022	0.5091	0.7368
SLC38A1 AND NOT-AJAP1	Mantle-Cell Lymphoma	0.6761	0.7273	0.6316
SLC38A1 AND NOT-PTGER4	Mantle-Cell Lymphoma	0.6265	0.5778	0.6842
SLC38A1 AND NOT-CD163	Mantle-Cell Lymphoma	0.6292	0.549	0.7368
SLC38A1 AND NOT-GPR78	Mantle-Cell Lymphoma	0.6265	0.5778	0.6842
CLDN15 AND NOT-BEST2	Mantle-Cell Lymphoma	0.642	0.6047	0.6842
SLC38A1 AND NOT-ATP6V0A4	Mantle-Cell Lymphoma	0.6341	0.5909	0.6842
SLC38A1 AND NOT-LPPR3	Mantle-Cell Lymphoma	0.6316	0.6316	0.6316
SLC38A1 AND NOT-CHRNA4	Mantle-Cell Lymphoma	0.6849	0.7143	0.6579
CLDN15 AND NOT-CD93	Mantle-Cell Lymphoma	0.6585	0.6136	0.7105
SLC38A1 AND NOT-SLC28A1	Mantle-Cell Lymphoma	0.6842	0.6842	0.6842
LAPTM5 AND NOT-SORL1	Mantle-Cell Lymphoma	0.7551	0.6167	0.9737
SLC38A1 AND NOT-GALR2	Mantle-Cell Lymphoma	0.6585	0.6136	0.7105
SLC38A1 AND NOT-SLC39A2	Mantle-Cell Lymphoma	0.6506	0.6	0.7105
SLC38A1 AND NOT-SLC5A8	Mantle-Cell Lymphoma	0.6316	0.6316	0.6316
SLC38A1 AND NOT-OR3A2	Mantle-Cell Lymphoma	0.6154	0.6	0.6316
SLC38A1 AND NOT-OXTR	Mantle-Cell Lymphoma	0.6486	0.6667	0.6316
SLC38A1 AND NOT-KCNA4	Mantle-Cell Lymphoma	0.619	0.5652	0.6842

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
ALK AND NOT-FAP	Neuroblastoma	0.8364	0.8519	0.8214
L1CAM AND NOT-BCAN	Neuroblastoma	0.8222	0.7708	0.881
L1CAM AND NOT-SSTR1	Neuroblastoma	0.8	0.7327	0.881
L1CAM AND NOT-CLDN11	Neuroblastoma	0.7957	0.7255	0.881
L1CAM AND NOT-ITGB3	Neuroblastoma	0.759	0.6667	0.881
L1CAM AND NOT-CLDN18	Neuroblastoma	0.7513	0.6549	0.881
L1CAM AND NOT-FOLH1	Neuroblastoma	0.7419	0.6765	0.8214
BMPR1B AND NOT-RNF43	Neuroblastoma	0.7413	0.8983	0.631
L1CAM AND NOT-EGFR	Neuroblastoma	0.7389	0.6303	0.8929
BMPR1B AND NOT-ENPP3	Neuroblastoma	0.7361	0.8833	0.631
BMPR1B AND NOT-CLDN4	Neuroblastoma	0.7361	0.8833	0.631
BMPR1B AND NOT-CEACAM5	Neuroblastoma	0.7347	0.8571	0.6429
BMPR1B AND NOT-ITGB6	Neuroblastoma	0.7338	0.9273	0.6071
L1CAM AND NOT-TYR	Neuroblastoma	0.7317	0.6198	0.8929
BMPR1B AND NOT-ERBB2	Neuroblastoma	0.731	0.8689	0.631
BMPR1B AND NOT-CLDN8	Neuroblastoma	0.7297	0.8438	0.6429
BMPR1B AND NOT-RAET1E	Neuroblastoma	0.7297	0.8438	0.6429
BMPR1B AND NOT-CLDN3	Neuroblastoma	0.7297	0.8438	0.6429
BMPR1B AND NOT-CLDN7	Neuroblastoma	0.7273	0.8814	0.619
BMPR1B AND NOT-PSCA	Neuroblastoma	0.7248	0.8308	0.6429
L1CAM AND NOT-ITGB6	Neuroblastoma	0.7192	0.6134	0.869
BMPR1B AND NOT-HHLA2	Neuroblastoma	0.7152	0.806	0.6429
BMPR1B AND NOT-CLDN1	Neuroblastoma	0.7143	0.8929	0.5952
BMPR1B AND NOT-FOLR1	Neuroblastoma	0.7123	0.8387	0.619
EDNRB AND NOT-SDC1	Oligodendroglioma	0.6154	0.7273	0.5333
CSPG4 AND NOT-TPBG	Oligodendroglioma	0.6667	0.8889	0.5333
EDNRB AND NOT-TPBG	Oligodendroglioma	0.6667	0.8889	0.5333
EDNRB AND NOT-ENG	Oligodendroglioma	0.64	0.8	0.5333
EDNRB AND NOT-KDR	Oligodendroglioma	0.6087	0.875	0.4667
EDNRB AND NOT-CLDN1	Oligodendroglioma	0.64	0.8	0.5333
EDNRB AND NOT-ERBB2	Oligodendroglioma	0.6154	0.7273	0.5333
MUC13 AND NOT-TRPM4	Oligodendroglioma	0.6	0.6	0.6
MUC16 AND NOT-PROM1	Ovarian	0.7273	0.8	0.6667
MUC16 AND NOT-SDC1	Ovarian	0.6957	0.7273	0.6667
MUC16 AND NOT-MUC4	Ovarian	0.6667	0.6667	0.6667
MUC16 AND NOT-TPBG	Ovarian	0.6667	0.7778	0.5833
MUC16 AND NOT-PTK7	Ovarian	0.6667	0.6667	0.6667
MUC16 AND NOT-CLDN8	Ovarian	0.6364	0.7	0.5833
MUC16 AND NOT-CD52	Ovarian	0.6154	0.5714	0.6667
MUC16 AND NOT-MUC13	Ovarian	0.6154	0.5714	0.6667
STEAP2 AND NOT-B4GALNT1	Prostate	0.8571	1	0.75
STEAP2 AND NOT-NCAM1	Prostate	0.8889	0.8	1
STEAP2 AND NOT-EDNRB	Prostate	0.75	0.75	0.75
STEAP2 AND NOT-IL3RA	Prostate	0.8571	1	0.75
STEAP2 AND NOT-DPEP1	Prostate	0.8571	1	0.75
STEAP2 AND NOT-KDR	Prostate	0.8	0.6667	1
STEAP2 AND NOT-IL11RA	Prostate	0.7273	0.5714	1
STEAP2 AND NOT-TNC	Prostate	0.7273	0.5714	1
STEAP2 AND NOT-IGF1R	Prostate	0.7273	0.5714	1
STEAP2 AND NOT-DKK1	Prostate	0.7273	0.5714	1
STEAP2 AND NOT-CSPG4	Prostate	0.7273	0.5714	1
STEAP2 AND NOT-HSPA5	Prostate	0.7273	0.5714	1
STEAP2 AND NOT-AXL	Prostate	0.7273	0.5714	1
STEAP2 AND NOT-VTCN1	Prostate	0.7273	0.5714	1
STEAP2 AND NOT-L1CAM	Prostate	0.7273	0.5714	1
STEAP2 AND NOT-PSCA	Prostate	1	1	1
STEAP1 AND NOT-CD33	Prostate	0.75	0.75	0.75
STEAP1 AND NOT-VCAM1	Prostate	0.75	0.75	0.75
STEAP1 AND NOT-FOLR2	Prostate	0.75	0.75	0.75
STEAP1 AND NOT-AXL	Prostate	0.75	0.75	0.75
CLDN12 AND NOT-PROM1	Prostate	0.6667	1	0.5
STEAP2 AND NOT-CLDN5	Prostate	0.6667	1	0.5
STEAP1 AND NOT-FCRL5	Prostate	0.6667	1	0.5
STEAP2 AND NOT-SLC39A6	Prostate	0.6667	1	0.5
STEAP1 AND NOT-B4GALNT1	Prostate	0.6667	1	0.5
STEAP1 AND NOT-CSPG4	Prostate	0.6667	0.6	0.75
STEAP1 AND NOT-ST8SIA1	Prostate	0.6667	0.6	0.75
STEAP1 AND NOT-DKK1	Prostate	0.6667	0.6	0.75
STEAP1 AND NOT-CLDN1	Prostate	0.6667	0.6	0.75
STEAP2 AND NOT-ALDH1A1	Prostate	0.6667	0.6	0.75
STEAP1 AND NOT-CLDN5	Prostate	0.6667	0.6	0.75
STEAP2 AND NOT-CD38	Prostate	0.6667	0.6	0.75
FAP AND NOT-CLDN8	Sarcoma	0.8276	1	0.7059
FAP AND NOT-CLDN6	Sarcoma	0.8667	1	0.7647
FAP AND NOT-IL20RA	Sarcoma	0.8667	1	0.7647
FAP AND NOT-SSTR1	Sarcoma	0.7857	1	0.6471
FAP AND NOT-RNF43	Sarcoma	0.7407	1	0.5882

FIG. 12 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
SLC38A1 AND NOT-SLC22A6	Mantle-Cell Lymphoma	0.6857	0.75	0.6316
SLC38A1 AND NOT-OPCML	Mantle-Cell Lymphoma	0.6024	0.5556	0.6579
FAT1 AND NOT-ATP8B1	Melanoma	0.8421	1	0.7273
FAT1 AND NOT-NOX1	Melanoma	0.8421	1	0.7273
GPR19 AND NOT-GABRB2	Melanoma	0.9	1	0.8182
FAT1 AND NOT-OR1C1	Melanoma	0.8421	1	0.7273
FAT1 AND NOT-SLC17A3	Melanoma	0.8421	1	0.7273
GPR137B AND NOT-CD36	Melanoma	0.8	0.8889	0.7273
FAT1 AND NOT-GPR6	Melanoma	0.8421	1	0.7273
GPR19 AND NOT-CDH18	Melanoma	0.9524	1	0.9091
FAT1 AND NOT-BEST3	Melanoma	0.8421	1	0.7273
FAT1 AND NOT-LCT	Melanoma	0.7778	1	0.6364
GPR19 AND NOT-GABRB1	Melanoma	0.7778	1	0.6364
FAT1 AND NOT-PCDHGC4	Melanoma	0.8421	1	0.7273
FAT1 AND NOT-AOC3	Melanoma	0.8421	1	0.7273
FAT1 AND NOT-TMEM235	Melanoma	0.7778	1	0.6364
CACNA1B AND NOT-KCNA2	Neuroblastoma	0.9434	1	0.8929
CACNA1B AND NOT-KCNA1	Neuroblastoma	0.939	0.9625	0.9167
CACNA1B AND NOT-GABRB2	Neuroblastoma	0.939	0.9625	0.9167
CACNA1B AND NOT-ATP2B2	Neuroblastoma	0.9383	0.9744	0.9048
KCNQ2 AND NOT-KCNA1	Neuroblastoma	0.9333	0.9506	0.9167
KCNQ2 AND NOT-SLC12A5	Neuroblastoma	0.9317	0.974	0.8929
CACNA1B AND NOT-GRM3	Neuroblastoma	0.9317	0.974	0.8929
KCNQ2 AND NOT-GRM3	Neuroblastoma	0.9317	0.974	0.8929
KCNQ2 AND NOT-KCNA2	Neuroblastoma	0.9299	1	0.869
KCNQ2 AND NOT-GABRB2	Neuroblastoma	0.9277	0.939	0.9167
CACNA1B AND NOT-OMG	Neuroblastoma	0.925	0.9737	0.881
KCNQ2 AND NOT-ATP2B2	Neuroblastoma	0.9222	0.9277	0.9167
KCNQ2 AND NOT-MLC1	Neuroblastoma	0.9222	0.9277	0.9167
CACNA1B AND NOT-OPCML	Neuroblastoma	0.9202	0.9494	0.8929
KCNQ2 AND NOT-CSMD3	Neuroblastoma	0.9182	0.9733	0.869
NKAIN1 AND NOT-SLC13A5	Neuroblastoma	0.9193	0.961	0.881
NKAIN1 AND NOT-TRPC3	Neuroblastoma	0.9172	0.9863	0.8571
CACNA1B AND NOT-GABRA1	Neuroblastoma	0.9157	0.9268	0.9048
GPR19 AND NOT-GRM3	Neuroblastoma	0.9143	0.8791	0.9524
NKAIN1 AND NOT-GRM3	Neuroblastoma	0.9114	0.973	0.8571
NKAIN1 AND NOT-GABRB2	Neuroblastoma	0.9103	0.9861	0.8452
KCNQ2 AND NOT-OMG	Neuroblastoma	0.9091	1	0.8333
SCN2A AND NOT-GRM3	Neuroblastoma	0.9068	0.9481	0.869
SLC6A15 AND NOT-KCNA1	Neuroblastoma	0.9045	0.9726	0.8452
CACNA1B AND NOT-GABBR2	Neuroblastoma	0.9036	0.9146	0.8929
KCNQ2 AND NOT-SCN1A	Neuroblastoma	0.9024	0.925	0.881
CACNA1B AND NOT-MLC1	Neuroblastoma	0.9006	0.8851	0.9167
CACNA1B AND NOT-GRIN2A	Neuroblastoma	0.9006	0.8851	0.9167
KCNQ2 AND NOT-OPALIN	Neuroblastoma	0.9006	0.8851	0.9167
NKAIN1 AND NOT-CACNG8	Neuroblastoma	0.9	0.9474	0.8571
SCN2A AND NOT-CSMD3	Neuroblastoma	0.9	0.9474	0.8571
GPR19 AND NOT-GABRB2	Neuroblastoma	0.9	0.9474	0.8571
CACNA1B AND NOT-SLC12A5	Neuroblastoma	0.8994	0.8941	0.9048
KCNQ2 AND NOT-SLCO1C1	Neuroblastoma	0.8982	0.9036	0.8929
NKAIN1 AND NOT-MLC1	Neuroblastoma	0.9202	0.9494	0.8929
KCNQ2 AND NOT-NKAIN2	Neuroblastoma	0.8961	0.9857	0.8214
NKAIN1 AND NOT-NKAIN2	Neuroblastoma	0.8961	0.9857	0.8214
NKAIN1 AND NOT-MOG	Neuroblastoma	0.8929	0.8929	0.8929
NKAIN1 AND NOT-SLC9B1	Neuroblastoma	0.8982	0.9036	0.8929
SLC6A15 AND NOT-KCNA2	Neuroblastoma	0.8903	0.9718	0.8214
CACNA1B AND NOT-GABRD	Neuroblastoma	0.8889	0.8736	0.9048
NKAIN1 AND NOT-GABRD	Neuroblastoma	0.8929	0.8929	0.8929
NKAIN1 AND NOT-ATP2B2	Neuroblastoma	0.9146	0.9375	0.8929
NKAIN1 AND NOT-GRM1	Neuroblastoma	0.8929	0.8929	0.8929
ASTN1 AND NOT-NKAIN2	Neuroblastoma	0.8931	0.9467	0.8452
SCN2A AND NOT-KCNA1	Neuroblastoma	0.8862	0.8916	0.881
SCN2A AND NOT-MLC1	Neuroblastoma	0.8862	0.8916	0.881
NKAIN1 AND NOT-OPCML	Neuroblastoma	0.8862	0.8916	0.881
NKAIN1 AND NOT-OPRK1	Neuroblastoma	0.8848	0.9012	0.869
NKAIN1 AND NOT-GRIN2C	Neuroblastoma	0.8846	0.9583	0.8214
NKAIN1 AND NOT-SLC1A6	Neuroblastoma	0.8846	0.9583	0.8214
NKAIN1 AND NOT-HTR5A	Neuroblastoma	0.8876	0.8824	0.8929
KCNQ2 AND NOT-GABRD	Neuroblastoma	0.8837	0.8636	0.9048
SLC6A15 AND NOT-CSMD3	Neuroblastoma	0.8831	0.9714	0.8095
NKAIN1 AND NOT-SCN1A	Neuroblastoma	0.8824	0.8721	0.8929
NKAIN1 AND NOT-GGTLC1	Neuroblastoma	0.8824	0.8721	0.8929
NKAIN1 AND NOT-ACSL6	Neuroblastoma	0.8824	0.8721	0.8929
NKAIN1 AND NOT-CALHM1	Neuroblastoma	0.882	0.9221	0.8452
GPR85 AND NOT-GABRB2	Neuroblastoma	0.8816	0.9853	0.7976
NKAIN1 AND NOT-OR1Q1	Neuroblastoma	0.881	0.881	0.881
NKAIN1 AND NOT-NPHS1	Neuroblastoma	0.881	0.881	0.881

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
FAP AND NOT-ERBB3	Sarcoma	0.7407	1	0.5882
FAP AND NOT-CD22	Sarcoma	0.7857	1	0.6471
FAP AND NOT-MSLN	Sarcoma	0.7857	1	0.6471
FAP AND NOT-PMEL	Sarcoma	0.8276	1	0.7059
FAP AND NOT-EPCAM	Sarcoma	0.8667	1	0.7647
FAP AND NOT-PROM1	Sarcoma	0.8667	1	0.7647
FAP AND NOT-MST1R	Sarcoma	0.7857	1	0.6471
FAP AND NOT-ENPP3	Sarcoma	0.7407	1	0.5882
FAP AND NOT-CR2	Sarcoma	0.7857	1	0.6471
FAP AND NOT-ERBB4	Sarcoma	0.8276	1	0.7059
FAP AND NOT-CLDN11	Sarcoma	0.7407	1	0.5882
FAP AND NOT-ITGB6	Sarcoma	0.7407	1	0.5882
FAP AND NOT-MUC4	Sarcoma	0.6923	1	0.5294
FAP AND NOT-FCRL2	Sarcoma	0.6923	1	0.5294
FAP AND NOT-RAET1E	Sarcoma	0.8276	1	0.7059
FAP AND NOT-SSTR2	Sarcoma	0.7857	1	0.6471
FAP AND NOT-TNFRSF8	Sarcoma	0.6923	1	0.5294
FAP AND NOT-NCAM1	Sarcoma	0.8667	1	0.7647
ROR1 AND NOT-CLDN6	Sarcoma	0.8667	1	0.7647
FAP AND NOT-MUC1	Sarcoma	0.7857	1	0.6471
FAP AND NOT-CLDN2	Sarcoma	0.8276	1	0.7059
FAP AND NOT-SSTR5	Sarcoma	0.8667	1	0.7647
ROR1 AND NOT-ERBB3	Sarcoma	0.7857	1	0.6471
FAP AND NOT-EDNRB	Sarcoma	0.7407	1	0.5882
FAP AND NOT-CXCR5	Sarcoma	0.8667	1	0.7647
FAP AND NOT-SLC34A2	Sarcoma	0.8667	1	0.7647
FAP AND NOT-MUC13	Sarcoma	0.6923	1	0.5294
FAP AND NOT-VTCN1	Sarcoma	0.8276	1	0.7059
FAP AND NOT-CLDN18	Sarcoma	0.6923	1	0.5294
FAP AND NOT-MUC16	Sarcoma	0.8667	1	0.7647
FAP AND NOT-TNFRSF17	Sarcoma	0.6923	1	0.5294
FAP AND NOT-KDR	Sarcoma	0.8276	1	0.7059
ROR1 AND NOT-MSLN	Sarcoma	0.8276	1	0.7059
ROR1 AND NOT-EPCAM	Sarcoma	0.8387	0.9286	0.7647
FAP AND NOT-GUCY2C	Sarcoma	0.8667	1	0.7647
ROR1 AND NOT-IL20RA	Sarcoma	0.8387	0.9286	0.7647
FAP AND NOT-GPA33	Sarcoma	0.8276	1	0.7059
FAP AND NOT-FOLR1	Sarcoma	0.8276	1	0.7059
ROR1 AND NOT-CLDN1	Sarcoma	0.7143	0.9091	0.5882
FAP AND NOT-CLDN23	Sarcoma	0.8667	1	0.7647
ROR1 AND NOT-RNF43	Sarcoma	0.8276	1	0.7059
ROR1 AND NOT-CLDN8	Sarcoma	0.8667	1	0.7647
FAP AND NOT-TNFRSF13C	Sarcoma	0.7407	1	0.5882
EPHB2 AND NOT-RNF43	Sarcoma	0.64	1	0.4706
EPHB2 AND NOT-PROM1	Sarcoma	0.64	1	0.4706
FAP AND NOT-CLDN1	Sarcoma	0.64	1	0.4706
EPHB2 AND NOT-GUCY2C	Sarcoma	0.64	1	0.4706
EPHB2 AND NOT-EPCAM	Sarcoma	0.64	1	0.4706
ROR1 AND NOT-SLC7A5	Sarcoma	0.8387	0.9286	0.7647
EPHB2 AND NOT-ENPP3	Sarcoma	0.64	1	0.4706
EPHB2 AND NOT-GPA33	Sarcoma	0.64	1	0.4706
ROR1 AND NOT-MST1R	Sarcoma	0.8276	1	0.7059
EPHB2 AND NOT-ITGB6	Sarcoma	0.64	1	0.4706
FAP AND NOT-MS4A1	Sarcoma	0.7857	1	0.6471
FAP AND NOT-CLDN7	Sarcoma	0.8667	1	0.7647
FAP AND NOT-CLDN12	Sarcoma	0.8667	1	0.7647
ROR1 AND NOT-FCRL2	Sarcoma	0.8667	1	0.7647
ROR1 AND NOT-ERBB4	Sarcoma	0.8276	1	0.7059
ROR1 AND NOT-CR2	Sarcoma	0.8667	1	0.7647
EPHB2 AND NOT-MUC13	Sarcoma	0.64	1	0.4706
FAP AND NOT-SDC1	Sarcoma	0.8276	1	0.7059
ROR1 AND NOT-SSTR1	Sarcoma	0.8276	1	0.7059
EPHB2 AND NOT-MUC4	Sarcoma	0.64	1	0.4706
FAP AND NOT-SSTR4	Sarcoma	0.8276	1	0.7059
CD276 AND NOT-RNF43	Sarcoma	0.6923	1	0.5294
ROR1 AND NOT-GUCY2C	Sarcoma	0.8667	1	0.7647
FAP AND NOT-FCRL1	Sarcoma	0.8276	1	0.7059
ROR1 AND NOT-SDC1	Sarcoma	0.8387	0.9286	0.7647
ROR1 AND NOT-ENPP3	Sarcoma	0.7407	1	0.5882
EPHB2 AND NOT-CLDN23	Sarcoma	0.64	1	0.4706
ROR1 AND NOT-SSTR5	Sarcoma	0.8667	1	0.7647
ROR1 AND NOT-ITGB6	Sarcoma	0.7857	1	0.6471
EPHB2 AND NOT-SSTR1	Sarcoma	0.64	1	0.4706
ROR1 AND NOT-PMEL	Sarcoma	0.8276	1	0.7059
ROR1 AND NOT-MS4A1	Sarcoma	0.8667	1	0.7647
ROR1 AND NOT-CLDN11	Sarcoma	0.7143	0.9091	0.5882
ROR1 AND NOT-CXCR5	Sarcoma	0.8667	1	0.7647

FIG. 12 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
NKAIN1 AND NOT-SLCO1C1	Neuroblastoma	0.8916	0.9024	0.881
CACNA1B AND NOT-SLCO1C1	Neuroblastoma	0.8786	0.8539	0.9048
JPH3 AND NOT-CSMD3	Neuroblastoma	0.8861	0.9459	0.8333
SCN2A AND NOT-KCNA2	Neuroblastoma	0.878	0.9	0.8571
NKAIN1 AND NOT-KCNA2	Neuroblastoma	0.8774	0.9577	0.8095
GPR85 AND NOT-KCNA1	Neuroblastoma	0.8774	0.9577	0.8095
NKAIN1 AND NOT-ROS1	Neuroblastoma	0.8772	0.8621	0.8929
NKAIN1 AND NOT-GPRC6A	Neuroblastoma	0.8772	0.8621	0.8929
KCNQ2 AND NOT-OPCML	Neuroblastoma	0.8772	0.8621	0.8929
NKAIN1 AND NOT-GUCY2F	Neuroblastoma	0.8772	0.8621	0.8929
NKAIN1 AND NOT-ADAM7	Neuroblastoma	0.8929	0.8929	0.8929
NKAIN1 AND NOT-CSMD3	Neuroblastoma	0.8758	0.971	0.7976
NKAIN1 AND NOT-BEST3	Neuroblastoma	0.8757	0.8706	0.881
NKAIN1 AND NOT-SLC22A11	Neuroblastoma	0.8757	0.8706	0.881
NKAIN1 AND NOT-GSG1L	Neuroblastoma	0.8757	0.8706	0.881
NKAIN1 AND NOT-KCNA7	Neuroblastoma	0.8757	0.8706	0.881
NKAIN1 AND NOT-SLC5A11	Neuroblastoma	0.8743	0.8795	0.869
NKAIN1 AND NOT-KCNJ9	Neuroblastoma	0.8743	0.8795	0.869
SCN2A AND NOT-NKAIN2	Neuroblastoma	0.8742	0.9851	0.7857
CACNA1B AND NOT-CSMD3	Neuroblastoma	0.8736	0.8444	0.9048
SLC6A15 AND NOT-GABBR2	Neuroblastoma	0.8734	0.9324	0.8214
SYT11 AND NOT-MLC1	Neuroblastoma	0.8729	0.8144	0.9405
NKAIN1 AND NOT-UPK3A	Neuroblastoma	0.8721	0.8523	0.8929
NKAIN1 AND NOT-TRPM1	Neuroblastoma	0.8721	0.8523	0.8929
NKAIN1 AND NOT-KCNA10	Neuroblastoma	0.8721	0.8523	0.8929
NKAIN1 AND NOT-TAS1R1	Neuroblastoma	0.8721	0.8523	0.8929
SCN2A AND NOT-OPCML	Neuroblastoma	0.8718	0.9444	0.8095
GPR85 AND NOT-ATP2B2	Neuroblastoma	0.8718	0.9444	0.8095
NKAIN1 AND NOT-TSPAN16	Neuroblastoma	0.8712	0.8987	0.8452
SLC6A15 AND NOT-MLC1	Neuroblastoma	0.8712	0.8987	0.8452
UNC5A AND NOT-GRM3	Neuroblastoma	0.8712	0.8987	0.8452
SLC6A15 AND NOT-GABRB2	Neuroblastoma	0.8712	0.8987	0.8452
SLC6A15 AND NOT-GRM3	Neuroblastoma	0.8712	0.8987	0.8452
SYT11 AND NOT-NKAIN2	Neuroblastoma	0.8706	0.8605	0.881
NKAIN1 AND NOT-MIP	Neuroblastoma	0.8706	0.8605	0.881
SCN2A AND NOT-OMG	Neuroblastoma	0.8701	0.9571	0.7976
KCNQ2 AND NOT-GABRA2	Neuroblastoma	0.869	0.869	0.869
NKAIN4 AND NOT-CDH20	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-KCNJ9	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-IL13	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-AQP6	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-KCNA1	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-KCNC2	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-KCNJ6	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-SLC6A18	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-OR52D1	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-MIP	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-TRPM1	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-CLDN20	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-OPRK1	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-HTR5A	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-GRM4	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-GRIA4	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-GRIK1	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-GRIN2B	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-GRM1	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-GRM3	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-PCDHB1	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-HTR2C	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-SLC39A2	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-OPRM1	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-OR3A1	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-SLC22A18	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-GJD2	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-PTGIR	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-PTPRR	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-CACNG8	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-CACNG7	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-OXTR	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-PORCN	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-SLC1A6	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-PCDH15	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-NMUR2	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-ADCY10	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-P2RX3	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-P2RY4	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-SLC22A9	Oligodendroglioma	0.8889	1	0.8

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
ROR1 AND NOT-IL3RA	Sarcoma	0.8276	1	0.7059
ROR1 AND NOT-CLDN12	Sarcoma	0.8387	0.9286	0.7647
ROR1 AND NOT-CLDN23	Sarcoma	0.8387	0.9286	0.7647
ROR1 AND NOT-TNFRSF10A	Sarcoma	0.8276	1	0.7059
ROR1 AND NOT-GPA33	Sarcoma	0.6923	1	0.5294
TPBG AND NOT-EPCAM	Sarcoma	0.6154	0.8889	0.4706
MUC13 AND NOT-ENPP3	Stomach	0.9512	1	0.907
MUC13 AND NOT-CLDN8	Stomach	0.9524	0.9756	0.9302
CLDN18 AND NOT-SLC34A2	Stomach	0.7912	0.75	0.8372
MUC13 AND NOT-HHLA2	Stomach	0.7714	1	0.6279
CLDN18 AND NOT-CLDN5	Stomach	0.7778	0.7447	0.814
CLDN18 AND NOT-CD52	Stomach	0.7556	0.7234	0.7907
MST1R AND NOT-CLDN8	Stomach	0.8989	0.8696	0.9302
CLDN18 AND NOT-FOLR1	Stomach	0.7356	0.7273	0.7442
CLDN18 AND NOT-CD33	Stomach	0.766	0.7059	0.8372
EPHB2 AND NOT-ENPP3	Stomach	0.7255	0.6271	0.8605
CLDN18 AND NOT-GPC3	Stomach	0.7416	0.7174	0.7674
CLDN18 AND NOT-CLEC14A	Stomach	0.7	0.7568	0.6512
CLDN18 AND NOT-ERBB4	Stomach	0.6923	0.7714	0.6279
CEACAM5 AND NOT-CLDN8	Stomach	0.8	0.8095	0.7907
MUC13 AND NOT-CLDN3	Stomach	0.7174	0.6735	0.7674
MUC13 AND NOT-MUC4	Stomach	0.7778	0.7447	0.814
CLDN18 AND NOT-IL11RA	Stomach	0.6829	0.7179	0.6512
CLDN18 AND NOT-CD34	Stomach	0.6824	0.6905	0.6744
MUC13 AND NOT-ABCA5	Stomach	0.68	0.5965	0.7907
MUC13 AND NOT-GPA33	Stomach	0.9512	1	0.907
EPCAM AND NOT-CLDN8	Stomach	0.7857	0.8049	0.7674
CLDN18 AND NOT-CD37	Stomach	0.6667	0.6591	0.6744
CLDN18 AND NOT-EDNRB	Stomach	0.6667	0.6591	0.6744
CLDN18 AND NOT-IL3RA	Stomach	0.6667	0.6591	0.6744
CEACAM5 AND NOT-ENPP3	Stomach	0.8046	0.7955	0.814
EPCAM AND NOT-ENPP3	Stomach	0.8155	0.7	0.9767
MUC13 AND NOT-GUCY2C	Stomach	0.7416	0.7174	0.7674
CLDN18 AND NOT-CD160	Stomach	0.7294	0.7381	0.7209
CLDN18 AND NOT-ENG	Stomach	0.6506	0.675	0.6279
CLDN18 AND NOT-KDR	Stomach	0.7865	0.7609	0.814
MUC13 AND NOT-CLDN7	Stomach	0.6835	0.75	0.6279
MST1R AND NOT-ABCA5	Stomach	0.6602	0.5667	0.7907
CLDN7 AND NOT-CLDN8	Stomach	0.8478	0.7959	0.907
CLDN18 AND NOT-GPNMB	Stomach	0.6966	0.6739	0.7209
CLDN18 AND NOT-ALK	Stomach	0.8312	0.9412	0.7442
MST1R AND NOT-ENPP3	Stomach	0.8966	0.8864	0.907
CLDN18 AND NOT-AXL	Stomach	0.6966	0.6739	0.7209
PROM1 AND NOT-ENPP3	Stomach	0.7111	0.6809	0.7442
MUC13 AND NOT-CD160	Stomach	0.8409	0.8222	0.8605
NOT-TPBG AND CLDN7	Liver	0.7143	0.625	0.8333
NOT-IL20RA AND RNF43	Liver	0.6667	1	0.5
NOT-TPBG AND RNF43	Liver	0.6667	1	0.5
NOT-TPBG AND MUC13	Liver	0.6	0.75	0.5
NOT-ITGB6 AND MUC13	Liver	0.6667	1	0.5
NOT-IL11RA AND AXL	Renal	0.8	1	0.6667
NOT-CLDN11 AND AXL	Renal	0.7273	0.8	0.6667
NOT-ENPP3 AND CLDN3	Esophagus	0.6	1	0.4286
NOT-ENPP3 AND CLDN2	Esophagus	0.6	1	0.4286
NOT-ENPP3 AND CEACAM5	Esophagus	0.6667	0.8	0.5714
NOT-EPCAM AND SLC39A6	Glioma	0.8254	0.9123	0.7536
NOT-EPCAM AND CLDN12	Glioma	0.7848	0.6966	0.8986
NOT-RNF43 AND CLDN12	Glioma	0.7967	0.9074	0.7101
NOT-TRPM4 AND CD276	Glioma	0.7333	0.679	0.7971
NOT-EPCAM AND CD276	Glioma	0.7799	0.6889	0.8986
NOT-TRPM4 AND CLDN12	Glioma	0.7639	0.7333	0.7971
NOT-SLAMF7 AND EPHB2	Glioma	0.6971	0.5755	0.8841
NOT-RNF43 AND CD276	Glioma	0.7643	0.6818	0.8696
NOT-ITGB6 AND PROM1	Glioma	0.7353	0.7463	0.7246
NOT-ERBB3 AND PTK7	Leiomyosarcoma	0.6667	0.7273	0.6154
NOT-ERBB3 AND CD276	Leiomyosarcoma	0.7826	0.9	0.6923
NOT-RNF43 AND AXL	Leiomyosarcoma	0.6667	0.7273	0.6154
NOT-EPCAM AND AXL	Leiomyosarcoma	0.64	0.6667	0.6154
NOT-ERBB3 AND TPBG	Leiomyosarcoma	0.75	0.8182	0.6923
NOT-ERBB3 AND AXL	Leiomyosarcoma	0.6667	0.875	0.5385
NOT-ERBB3 AND GPNMB	Leiomyosarcoma	0.6207	0.5625	0.6923
NOT-EPCAM AND TPBG	Leiomyosarcoma	0.6667	0.7273	0.6154
NOT-ITGAV AND CD70	AML	0.7957	0.7285	0.8765
NOT-ITGAV AND SLC7A5	AML	0.792	0.7252	0.8725
NOT-EDNRB AND SLC7A5	AML	0.7709	0.6387	0.9721
NOT-CLDN12 AND CD37	AML	0.7708	0.6716	0.9044
NOT-ITGAV AND SLC39A6	AML	0.769	0.6899	0.8685

FIG. 12 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
NKAIN4 AND NOT-CALY	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-RXFP3	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-KCNK9	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-ATP8A2	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-FXYD7	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-KCNK10	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-GPR85	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-BEST2	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-SLC6A15	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-SLC30A10	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-GPR22	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-DIO1	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-LRRN4	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-KCNV2	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-HTR3C	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-LRRC55	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-DSC1	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-DSCAM	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-DTNA	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-EFNB2	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-EPHA2	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-EMP3	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-PAQR7	Oligodendroglioma	0.8889	1	0.8
EPHA10 AND NOT-EMP3	Ovarian	0.7273	0.8	0.6667
EPHA10 AND NOT-GABRA2	Ovarian	0.72	0.6923	0.75
EPHA10 AND NOT-ADAM2	Ovarian	0.7	0.875	0.5833
EPHA10 AND NOT-MRGPRF	Ovarian	0.6957	0.7273	0.6667
EPHA10 AND NOT-GGTLC1	Ovarian	0.6957	0.7273	0.6667
EPHA10 AND NOT-ANO4	Ovarian	0.6957	0.7273	0.6667
EPHA10 AND NOT-MTNR1B	Ovarian	0.6957	0.7273	0.6667
EPHA10 AND NOT-SCARA5	Ovarian	0.6923	0.6429	0.75
EPHA10 AND NOT-CDH9	Ovarian	0.6923	0.6429	0.75
EPHA10 AND NOT-CD36	Ovarian	0.72	0.6923	0.75
EPHA10 AND NOT-XCR1	Ovarian	0.6957	0.7273	0.6667
EPHA10 AND NOT-SLC22A13	Ovarian	0.7	0.875	0.5833
EPHA10 AND NOT-TAS2R1	Ovarian	0.75	0.75	0.75
CSPG5 AND NOT-PAQR8	Ovarian	0.8	1	0.6667
EPHA10 AND NOT-CHRNB4	Ovarian	0.6667	0.6667	0.6667
EPHA10 AND NOT-MLNR	Ovarian	0.6667	0.6667	0.6667
EPHA10 AND NOT-KCNA4	Ovarian	0.6667	0.6667	0.6667
EPHA10 AND NOT-GSG1L	Ovarian	0.6667	0.6667	0.6667
EPHA10 AND NOT-PCDH15	Ovarian	0.6667	0.6667	0.6667
EPHA10 AND NOT-SLC1A6	Ovarian	0.6667	0.6667	0.6667
EPHA10 AND NOT-CNIH2	Ovarian	0.6667	0.6667	0.6667
EPHA10 AND NOT-PCDHA6	Ovarian	0.6667	0.6667	0.6667
EPHA10 AND NOT-ATP1B4	Ovarian	0.6667	0.6667	0.6667
EPHA10 AND NOT-NMBR	Ovarian	0.6667	0.6667	0.6667
EPHA10 AND NOT-LRFN2	Ovarian	0.6667	0.6667	0.6667
EPHA10 AND NOT-GABRD	Ovarian	0.6667	0.6667	0.6667
EPHA10 AND NOT-BEST3	Ovarian	0.6667	0.6667	0.6667
EPHA10 AND NOT-KCNK6	Ovarian	0.6667	0.6667	0.6667
EPHA10 AND NOT-SLC6A18	Ovarian	0.6667	0.6	0.75
EPHA10 AND NOT-AOC3	Ovarian	0.6667	0.6	0.75
EPHA10 AND NOT-MLC1	Ovarian	0.8	1	0.6667
EPHA10 AND NOT-OR4N4	Ovarian	0.6667	0.6667	0.6667
EPHA10 AND NOT-SLC23A2	Ovarian	0.6667	0.6667	0.6667
EPHA10 AND NOT-ABCA12	Ovarian	0.7	0.875	0.5833
EPHA10 AND NOT-HCN1	Ovarian	0.6667	0.6667	0.6667
EPHA10 AND NOT-ABCG4	Ovarian	0.7368	1	0.5833
EPHA10 AND NOT-USH2A	Ovarian	0.6667	0.6667	0.6667
EPHA10 AND NOT-NOX1	Ovarian	0.6429	0.5625	0.75
EPHA10 AND NOT-UMODL1	Ovarian	0.64	0.6154	0.6667
EPHA10 AND NOT-ASTN1	Ovarian	0.64	0.6154	0.6667
EPHA10 AND NOT-OR2L2	Ovarian	0.64	0.6154	0.6667
EPHA10 AND NOT-SLC9B1	Ovarian	0.64	0.6154	0.6667
EPHA10 AND NOT-ADAM20	Ovarian	0.64	0.6154	0.6667
EPHA10 AND NOT-LRRN4	Ovarian	0.64	0.6154	0.6667
EPHA10 AND NOT-GABRG1	Ovarian	0.6667	0.6667	0.6667
EPHA10 AND NOT-OR3A1	Ovarian	0.64	0.6154	0.6667
EPHA10 AND NOT-ATP6V0A4	Ovarian	0.64	0.6154	0.6667
EPHA10 AND NOT-LPPR3	Ovarian	0.64	0.6154	0.6667
EPHA10 AND NOT-GABRG3	Ovarian	0.64	0.6154	0.6667
EPHA10 AND NOT-TRPM1	Ovarian	0.64	0.6154	0.6667
EPHA10 AND NOT-SLC5A11	Ovarian	0.64	0.6154	0.6667
EPHA10 AND NOT-ERVW-1	Ovarian	0.64	0.6154	0.6667
EPHA10 AND NOT-KCNJ10	Ovarian	0.64	0.6154	0.6667
EPHA10 AND NOT-GRID2	Ovarian	0.64	0.6154	0.6667

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
NOT-CLDN5 AND SLC7A5	AML	0.7519	0.6131	0.9721
NOT-ITGAV AND ENG	AML	0.7406	0.7362	0.745
NOT-CLDN12 AND SLC7A5	AML	0.7255	0.615	0.8845
NOT-EDNRB AND ENG	AML	0.7244	0.6508	0.8167
NOT-EDNRB AND MUC1	AML	0.7185	0.76	0.6813
NOT-CLDN12 AND CD70	AML	0.7161	0.5927	0.9044
NOT-ERBB3 AND MUC1	AML	0.7152	0.7731	0.6653
NOT-ITGAV AND PROM1	AML	0.7149	0.7671	0.6693
NOT-STEAP2 AND MUC1	AML	0.7134	0.7636	0.6693
NOT-ITGAV AND CD38	AML	0.7122	0.6492	0.7888
NOT-AXL AND CD70	AML	0.6892	0.5982	0.8127
NOT-CLDN5 AND CD70	AML	0.6838	0.5988	0.7968
NOT-ITGAV AND MUC1	AML	0.6818	0.7937	0.5976
NOT-AXL AND SLC7A5	AML	0.6731	0.5148	0.9721
NOT-AXL AND CD38	AML	0.6711	0.5734	0.8088
NOT-AXL AND MUC1	AML	0.6667	0.6849	0.6494
NOT-GPNMB AND SLC7A5	AML	0.6627	0.5271	0.8924
NOT-GPNMB AND MUC1	AML	0.6611	0.7009	0.6255
NOT-ITGAV AND CD79A	AML	0.6581	0.6109	0.7131
NOT-CLDN12 AND MUC1	AML	0.651	0.7037	0.6056
NOT-KDR AND ENG	AML	0.6471	0.5291	0.8327
NOT-AXL AND ENG	AML	0.6467	0.5352	0.8167
NOT-GPNMB AND ENG	AML	0.6388	0.5504	0.761
NOT-EGFR AND VCAM1	B-Cell Diffuse	0.6765	0.7419	0.6216
NOT-EGFR AND GPNMB	B-Cell Diffuse	0.6761	0.7059	0.6486
NOT-STEAP2 AND GPNMB	B-Cell Diffuse	0.6118	0.5417	0.7027
NOT-TRPM4 AND ROR1	Mantle-Cell Lymphoma	0.8182	0.9643	0.7105
NOT-SLC34A2 AND ENG	Melanoma	1	1	1
NOT-VTCN1 AND ENG	Melanoma	0.9	1	0.8182
NOT-SLC34A2 AND FAP	Melanoma	0.9	1	0.8182
NOT-SLC34A2 AND GPNMB	Melanoma	1	1	1
NOT-VTCN1 AND GPNMB	Melanoma	0.9	1	0.8182
NOT-VTCN1 AND FAP	Melanoma	0.8421	1	0.7273
NOT-VTCN1 AND ERBB2	Melanoma	0.8421	1	0.7273
NOT-CLDN7 AND ERBB2	Melanoma	0.9524	1	0.9091
NOT-VTCN1 AND STEAP1	Melanoma	0.8421	1	0.7273
NOT-CLDN7 AND TRPM4	Melanoma	0.9524	1	0.9091
NOT-VTCN1 AND CD276	Melanoma	0.8421	1	0.7273
NOT-CLDN7 AND PTK7	Melanoma	0.9524	1	0.9091
NOT-SLC34A2 AND STEAP1	Melanoma	0.9524	1	0.9091
NOT-CLDN7 AND STEAP1	Melanoma	0.9524	1	0.9091
NOT-SLC34A2 AND ERBB2	Melanoma	0.9524	1	0.9091
NOT-VTCN1 AND VCAM1	Melanoma	0.9	1	0.8182
NOT-CLDN7 AND SDC1	Melanoma	0.9524	1	0.9091
NOT-SLC34A2 AND PTK7	Melanoma	0.9091	0.9091	0.9091
NOT-CLDN7 AND GPNMB	Melanoma	1	1	1
NOT-SLC34A2 AND CD276	Melanoma	0.9524	1	0.9091
NOT-CLDN7 AND CD276	Melanoma	0.9	1	0.8182
NOT-SLC34A2 AND ERBB3	Melanoma	0.7778	1	0.6364
NOT-CLDN7 AND ERBB3	Melanoma	0.7778	1	0.6364
NOT-VTCN1 AND ERBB3	Melanoma	0.9	1	0.8182
NOT-CLDN7 AND ENG	Melanoma	1	1	1
NOT-VTCN1 AND SDC1	Melanoma	0.9	1	0.8182
NOT-SLC34A2 AND VCAM1	Melanoma	1	1	1
NOT-EGFR AND PTK7	Neuroblastoma	0.9294	0.9186	0.9405
NOT-EGFR AND CD276	Neuroblastoma	0.904	0.8602	0.9524
NOT-EGFR AND BMPR1B	Neuroblastoma	0.878	0.9	0.8571
NOT-TRPM4 AND CD276	Neuroblastoma	0.7771	0.7473	0.8095
NOT-EGFR AND CLDN12	Neuroblastoma	0.7418	0.6124	0.9405
NOT-EGFR AND GPNMB	Neuroblastoma	0.7317	0.6198	0.8929
NOT-CLDN7 AND PTK7	Neuroblastoma	0.73	0.6293	0.869
NOT-ERBB2 AND PTK7	Neuroblastoma	0.7263	0.6842	0.7738
NOT-IL20RA AND CD276	Neuroblastoma	0.7255	0.6167	0.881
NOT-PCYT1A AND CD276	Neuroblastoma	0.7882	0.7791	0.7976
NOT-RNF43 AND PTK7	Neuroblastoma	0.7192	0.6134	0.869
NOT-ERBB3 AND GPNMB	Sarcoma	0.625	0.6667	0.5882
NOT-IL20RA AND GPNMB	Sarcoma	0.6154	0.5455	0.7059
PTPRZ1 AND NOT-GRM3	Astrocytoma	0.881	0.9737	0.8043
PTPRZ1 AND NOT-GABRA2	Astrocytoma	0.8817	0.8723	0.8913
PTPRZ1 AND NOT-KCNJ6	Astrocytoma	0.8276	0.878	0.7826
NRCAM AND NOT-CDH18	Astrocytoma	0.8235	0.8974	0.7609
PTPRZ1 AND NOT-KIAA0319	Astrocytoma	0.8235	0.8974	0.7609
NKAIN4 AND NOT-CDH7	Astrocytoma	0.8205	1	0.6957
PTPRZ1 AND NOT-GPR83	Astrocytoma	0.8095	0.8947	0.7391
NKAIN4 AND NOT-TMEM130	Astrocytoma	0.8095	0.8947	0.7391
NKAIN4 AND NOT-DIO2	Astrocytoma	0.8095	0.8947	0.7391
NKAIN4 AND NOT-ADRA2C	Astrocytoma	0.8052	1	0.6739

FIG. 12 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
EPHA10 AND NOT-SLC17A2	Ovarian	0.6364	0.7	0.5833
EPHA10 AND NOT-GALR1	Ovarian	0.6364	0.7	0.5833
EPHA10 AND NOT-CALY	Ovarian	0.6364	0.7	0.5833
EPHA10 AND NOT-KCNQ2	Ovarian	0.64	0.6154	0.6667
EPHA10 AND NOT-SLC22A25	Ovarian	0.6957	0.7273	0.6667
EPHA10 AND NOT-ADCY8	Ovarian	0.6667	0.6667	0.6667
EPHA10 AND NOT-CD58	Ovarian	0.6207	0.5294	0.75
EPHA10 AND NOT-TTYH3	Ovarian	0.6207	0.5294	0.75
EPHA10 AND NOT-ANTXR2	Ovarian	0.6207	0.5294	0.75
EPHA10 AND NOT-CD74	Ovarian	0.6207	0.5294	0.75
EPHA10 AND NOT-GYPC	Ovarian	0.6207	0.5294	0.75
EPHA10 AND NOT-IL17RA	Ovarian	0.6207	0.5294	0.75
EPHA10 AND NOT-F2R	Ovarian	0.72	0.6923	0.75
EPHA10 AND NOT-GNRHR	Ovarian	0.6667	0.6667	0.6667
EPHA10 AND NOT-CRB2	Ovarian	0.6667	0.6667	0.6667
EPHA10 AND NOT-GLRA3	Ovarian	0.72	0.6923	0.75
EPHA10 AND NOT-ZDHHC2	Ovarian	0.6667	0.7778	0.5833
EPHA10 AND NOT-GHSR	Ovarian	0.6154	0.5714	0.6667
EPHA10 AND NOT-KCNA10	Ovarian	0.6154	0.5714	0.6667
EPHA10 AND NOT-KCNF1	Ovarian	0.6154	0.5714	0.6667
EPHA10 AND NOT-NFASC	Ovarian	0.6154	0.5714	0.6667
EPHA10 AND NOT-GRIN1	Ovarian	0.6154	0.5714	0.6667
EPHA10 AND NOT-MUC17	Ovarian	0.6154	0.5714	0.6667
EPHA10 AND NOT-CLCN1	Ovarian	0.6154	0.5714	0.6667
EPHA10 AND NOT-PTPRR	Ovarian	0.6154	0.5714	0.6667
EPHA10 AND NOT-ATP8B2	Ovarian	0.6154	0.5714	0.6667
EPHA10 AND NOT-NTSR2	Ovarian	0.6154	0.5714	0.6667
EPHA10 AND NOT-HTR1D	Ovarian	0.6154	0.5714	0.6667
EPHA10 AND NOT-HCN2	Ovarian	0.6154	0.5714	0.6667
EPHA10 AND NOT-KCNK12	Ovarian	0.6154	0.5714	0.6667
EPHA10 AND NOT-KCNK10	Ovarian	0.6154	0.5714	0.6667
EPHA10 AND NOT-GRIK4	Ovarian	0.6154	0.5714	0.6667
EPHA10 AND NOT-SLC12A9	Ovarian	0.6154	0.5714	0.6667
EPHA10 AND NOT-SYT4	Ovarian	0.6154	0.5714	0.6667
EPHA10 AND NOT-MEGF11	Ovarian	0.6154	0.5714	0.6667
KCNU1 AND NOT-ADAM2	Ovarian	0.8571	1	0.75
TRPM8 AND NOT-SLC16A5	Prostate	0.8571	1	0.75
TRPM8 AND NOT-SLC39A2	Prostate	1	1	1
GHR AND NOT-CD36	Prostate	0.8571	1	0.75
TRPM8 AND NOT-SLC4A8	Prostate	0.8889	0.8	1
TRPM8 AND NOT-NRG3	Prostate	0.8571	1	0.75
TRPM8 AND NOT-SYT4	Prostate	0.8	0.6667	1
TRPM8 AND NOT-GRIN2C	Prostate	0.8889	0.8	1
TRPM8 AND NOT-FMNL1	Prostate	1	1	1
TRPM8 AND NOT-SLC9A1	Prostate	0.75	0.75	0.75
TRPM8 AND NOT-CALN1	Prostate	0.75	0.75	0.75
TRPM8 AND NOT-SLC17A2	Prostate	0.75	0.75	0.75
TRPM8 AND NOT-SLC22A18	Prostate	0.7273	0.5714	1
TRPM8 AND NOT-GUCY2D	Prostate	0.7273	0.5714	1
TRPM8 AND NOT-CD36	Prostate	0.7273	0.5714	1
TRPM8 AND NOT-TTYH2	Prostate	0.7273	0.5714	1
TRPM8 AND NOT-PTGER4	Prostate	0.7273	0.5714	1
TRPM8 AND NOT-CHRNA2	Prostate	0.7273	0.5714	1
TRPM8 AND NOT-PIK3IP1	Prostate	0.7273	0.5714	1
TRPM8 AND NOT-PAQR8	Prostate	0.7273	0.5714	1
TRPM8 AND NOT-LAT	Prostate	0.7273	0.5714	1
TRPM8 AND NOT-DYSF	Prostate	0.7273	0.5714	1
TRPM8 AND NOT-CDH10	Prostate	0.7273	0.5714	1
TRPM8 AND NOT-NFASC	Prostate	0.7273	0.5714	1
TRPM8 AND NOT-GYPC	Prostate	0.7273	0.5714	1
TRPM8 AND NOT-PAQR7	Prostate	0.7273	0.5714	1
TRPM8 AND NOT-STAB1	Prostate	0.7273	0.5714	1
TRPM8 AND NOT-EMP3	Prostate	0.7273	0.5714	1
TRPM8 AND NOT-SEMA4D	Prostate	0.7273	0.5714	1
TRPM8 AND NOT-EPHA2	Prostate	0.7273	0.5714	1
TRPM8 AND NOT-IL17RA	Prostate	0.7273	0.5714	1
TRPM8 AND NOT-AGER	Prostate	0.7273	0.5714	1
TRPM8 AND NOT-BST2	Prostate	0.7273	0.5714	1
TRPM8 AND NOT-GABRA1	Prostate	0.7273	0.5714	1
TRPM8 AND NOT-TNFRSF12A	Prostate	0.7273	0.5714	1
TRPM8 AND NOT-SLC5A2	Prostate	0.7273	0.5714	1
UPK3A AND NOT-SCARA5	Prostate	0.75	0.75	0.75
ADAM2 AND NOT-MC2R	Prostate	0.8571	1	0.75
UPK3A AND NOT-CACNG6	Prostate	0.75	0.75	0.75
UPK3A AND NOT-NTSR2	Prostate	0.75	0.75	0.75
TRPM8 AND NOT-GALR1	Prostate	0.6667	1	0.5
TRPM8 AND NOT-SLC23A2	Prostate	0.6667	1	0.5

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
PTPRZ1 AND NOT-CDH18	Astrocytoma	0.8043	0.8043	0.8043
PTPRZ1 AND NOT-DGKE	Astrocytoma	0.8043	0.8043	0.8043
PTPRZ1 AND NOT-CDH8	Astrocytoma	0.7957	0.7872	0.8043
VANGL2 AND NOT-STEAP4	Astrocytoma	0.8041	0.7647	0.8478
NKAIN4 AND NOT-KCNJ12	Astrocytoma	0.7952	0.8919	0.7174
NKAIN4 AND NOT-SLC22A24	Astrocytoma	0.7952	0.8919	0.7174
NKAIN4 AND NOT-PLXNA2	Astrocytoma	0.7901	0.9143	0.6957
NKAIN4 AND NOT-CDH12	Astrocytoma	0.7901	0.9143	0.6957
PTPRZ1 AND NOT-GABRA1	Astrocytoma	0.7857	0.8684	0.7174
NRCAM AND NOT-CDH8	Astrocytoma	0.7857	0.8684	0.7174
NRCAM AND NOT-GABRA1	Astrocytoma	0.7848	0.9394	0.6739
PTPRZ1 AND NOT-DYSF	Astrocytoma	0.781	0.6949	0.8913
NKAIN4 AND NOT-KCNH3	Astrocytoma	0.7805	0.8889	0.6957
NKAIN4 AND NOT-CHRM3	Astrocytoma	0.7805	0.8889	0.6957
NKAIN4 AND NOT-JAG2	Astrocytoma	0.7805	0.8889	0.6957
NKAIN4 AND NOT-CACNA1A	Astrocytoma	0.7805	0.8889	0.6957
NKAIN4 AND NOT-LRRC8B	Astrocytoma	0.7805	0.8889	0.6957
NKAIN4 AND NOT-SEMA4F	Astrocytoma	0.7805	0.8889	0.6957
VANGL2 AND NOT-LRIG3	Astrocytoma	0.7879	0.7358	0.8478
NKAIN4 AND NOT-NPY1R	Astrocytoma	0.7792	0.9677	0.6522
NKAIN4 AND NOT-EPHA7	Astrocytoma	0.7733	1	0.6304
NRCAM AND NOT-DGKE	Astrocytoma	0.7727	0.8095	0.7391
NKAIN4 AND NOT-ANO3	Astrocytoma	0.7654	0.8857	0.6739
NKAIN4 AND NOT-DNAJC5	Astrocytoma	0.7654	0.8857	0.6739
NKAIN4 AND NOT-VIPR1	Astrocytoma	0.7654	0.8857	0.6739
NKAIN4 AND NOT-EFNB3	Astrocytoma	0.7654	0.8857	0.6739
NKAIN4 AND NOT-ADCY1	Astrocytoma	0.7654	0.8857	0.6739
PTPRZ1 AND NOT-SLC12A5	Astrocytoma	0.7654	0.8857	0.6739
NKAIN4 AND NOT-GABRB3	Astrocytoma	0.7654	0.8857	0.6739
VANGL2 AND NOT-EPHA2	Astrocytoma	0.7647	0.6964	0.8478
ADAM12 AND NOT-PROCR	Breast	0.8411	0.8491	0.8333
ADAM12 AND NOT-SCARA5	Breast	0.8381	0.8627	0.8148
ADAM12 AND NOT-SLC2A1	Breast	0.8257	0.8182	0.8333
ADAM12 AND NOT-SLC22A11	Breast	0.8182	0.8036	0.8333
ADAM12 AND NOT-AOC3	Breast	0.8182	0.8036	0.8333
ADAM12 AND NOT-MRGPRF	Breast	0.8073	0.8	0.8148
ADAM12 AND NOT-ERVW-1	Breast	0.8073	0.8	0.8148
ADAM12 AND NOT-ANTXR2	Breast	0.8037	0.8113	0.7963
ADAM12 AND NOT-MRAP	Breast	0.7963	0.7963	0.7963
ADAM12 AND NOT-THBD	Breast	0.7921	0.8511	0.7407
ADAM12 AND NOT-S1PR1	Breast	0.7885	0.82	0.7593
ADAM12 AND NOT-PHLDB2	Breast	0.785	0.7925	0.7778
SLC5A6 AND NOT-PAQR7	Breast	0.7778	0.9722	0.6481
PPAPDC1A AND NOT-DPP6	Breast	0.7917	0.9048	0.7037
PPAPDC1A AND NOT-SLC4A4	Breast	0.8085	0.95	0.7037
PPAPDC1A AND NOT-LRRTM2	Breast	0.7835	0.8837	0.7037
PPAPDC1A AND NOT-PCDH9	Breast	0.7917	0.9048	0.7037
PPAPDC1A AND NOT-ADCYAP1R1	Breast	0.7677	0.8444	0.7037
LRRC8E AND NOT-GJB5	Breast	0.7477	0.7547	0.7407
PPAPDC1A AND NOT-CADM2	Breast	0.7755	0.8636	0.7037
LRRC8E AND NOT-DNER	Breast	0.78	0.8478	0.7222
VANGL1 AND NOT-CLCA4	Breast	0.7451	0.7917	0.7037
PPAPDC1A AND NOT-IGSF11	Breast	0.7451	0.7917	0.7037
PRLR AND NOT-XCR1	Breast	0.7447	0.875	0.6481
PPAPDC1A AND NOT-ATP1A2	Breast	0.7423	0.8372	0.6667
PPAPDC1A AND NOT-FLRT1	Breast	0.7826	0.9474	0.6667
PPAPDC1A AND NOT-PLP1	Breast	0.7551	0.8409	0.6852
PPAPDC1A AND NOT-ATP1B2	Breast	0.7835	0.8837	0.7037
PPAPDC1A AND NOT-MAL	Breast	0.7447	0.875	0.6481
SLC5A6 AND NOT-SLC9B1	Breast	0.7778	0.9722	0.6481
PPAPDC1A AND NOT-DNER	Breast	0.76	0.8261	0.7037
PPAPDC1A AND NOT-SLC6A1	Breast	0.7379	0.7755	0.7037
SLC5A6 AND NOT-MPL	Breast	0.7527	0.8974	0.6481
PRLR AND NOT-CACNG3	Breast	0.7368	0.8537	0.6481
PRLR AND NOT-NPHS1	Breast	0.7368	0.8537	0.6481
PRLR AND NOT-SLC22A6	Breast	0.7368	0.8537	0.6481
LRRC8E AND NOT-CLCA4	Breast	0.7358	0.75	0.7222
SLC5A6 AND NOT-GPR22	Breast	0.7447	0.875	0.6481
PPAPDC1A AND NOT-GPM6B	Breast	0.7551	0.8409	0.6852
PPAPDC1A AND NOT-GPM6A	Breast	0.7333	0.9167	0.6111
PRLR AND NOT-CDH18	Breast	0.7447	0.875	0.6481
SLC5A6 AND NOT-EMP3	Breast	0.7609	0.9211	0.6481
PPAPDC1A AND NOT-SLC7A14	Breast	0.7525	0.8085	0.7037
PPAPDC1A AND NOT-AQP4	Breast	0.7708	0.881	0.6852
SLC5A6 AND NOT-DGKE	Breast	0.764	0.9714	0.6296
SLC5A6 AND NOT-SEMA4D	Breast	0.7778	0.9722	0.6481
PPAPDC1A AND NOT-KCNH8	Breast	0.7308	0.76	0.7037

FIG. 12 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
TRPM8 AND NOT-CLSTN3	Prostate	0.6667	1	0.5
UPK3A AND NOT-CACNG1	Prostate	0.6667	1	0.5
TRPM8 AND NOT-MEGF10	Prostate	0.6667	1	0.5
UPK3A AND NOT-HTR5A	Prostate	0.6667	1	0.5
TRPM8 AND NOT-TRPC3	Prostate	0.6667	1	0.5
STEAP4 AND NOT-CD58	Prostate	0.6667	1	0.5
TRPM8 AND NOT-OPRK1	Prostate	0.6667	1	0.5
TRPM8 AND NOT-SEZ6	Prostate	0.6667	1	0.5
UPK3A AND NOT-VN1R2	Prostate	0.6667	1	0.5
UPK3A AND NOT-KCNA7	Prostate	0.6667	1	0.5
UPK3A AND NOT-ATP1B4	Prostate	0.6667	0.6	0.75
TRPM8 AND NOT-PLVAP	Prostate	0.6667	0.6	0.75
TRPM8 AND NOT-CALHM3	Prostate	0.6667	0.6	0.75
UPK3A AND NOT-GYPC	Prostate	0.6667	0.6	0.75
UPK3A AND NOT-DYSF	Prostate	0.6667	0.6	0.75
UPK3A AND NOT-BEST3	Prostate	0.6667	0.6	0.75
UPK3A AND NOT-CACNA1S	Prostate	0.6667	0.6	0.75
UPK3A AND NOT-CD36	Prostate	0.6667	0.6	0.75
TRPM8 AND NOT-DTNA	Prostate	0.6667	0.6	0.75
UPK3A AND NOT-TTYH2	Prostate	0.6667	0.6	0.75
UPK3A AND NOT-CHRNA1	Prostate	0.6667	0.6	0.75
ADAM2 AND NOT-OR1Q1	Prostate	0.8571	1	0.75
ADAM2 AND NOT-MRGPRX2	Prostate	0.6667	0.6	0.75
ADAM2 AND NOT-DRD1	Prostate	0.6667	1	0.5
ADAM2 AND NOT-CACNG1	Prostate	0.8571	1	0.75
ADAM2 AND NOT-LRRC52	Prostate	0.8571	1	0.75
ADAM2 AND NOT-SLC22A9	Prostate	0.8571	1	0.75
ADAM2 AND NOT-KCNA4	Prostate	0.8571	1	0.75
ADAM2 AND NOT-CD82	Prostate	0.8571	1	0.75
ADAM2 AND NOT-CLDN19	Prostate	0.8571	1	0.75
ADAM2 AND NOT-OMG	Prostate	0.8571	1	0.75
TRPM8 AND NOT-CD82	Prostate	0.6667	1	0.5
ADAM2 AND NOT-CACNG6	Prostate	0.6667	1	0.5
ADAM2 AND NOT-SLC22A14	Prostate	0.6667	1	0.5
ADAM2 AND NOT-FMNL1	Prostate	0.6667	1	0.5
ADAM2 AND NOT-CD1A	Prostate	0.6667	0.6	0.75
ADAM2 AND NOT-RGR	Prostate	0.6667	1	0.5
UPK3A AND NOT-CLDN17	Prostate	0.6667	1	0.5
UPK3A AND NOT-MC2R	Prostate	0.6667	1	0.5
UPK3A AND NOT-MEGF11	Prostate	0.6667	1	0.5
UPK3A AND NOT-PTGER4	Prostate	0.6667	0.6	0.75
GGTLC1 AND NOT-DYSF	Prostate	0.8571	1	0.75
ADAM2 AND NOT-PORCN	Prostate	0.6667	0.6	0.75
ADAM2 AND NOT-FUT3	Prostate	0.6667	0.6	0.75
STAB1 AND NOT-NFASC	Sarcoma	0.7586	0.9167	0.6471
STAB1 AND NOT-ATP13A5	Sarcoma	0.7097	0.7857	0.6471
STAB1 AND NOT-CDH17	Sarcoma	0.7333	0.8462	0.6471
STAB1 AND NOT-CD1A	Sarcoma	0.7333	0.8462	0.6471
STAB1 AND NOT-CLDN19	Sarcoma	0.7407	1	0.5882
STAB1 AND NOT-NRG3	Sarcoma	0.7407	1	0.5882
STAB1 AND NOT-CLDN10	Sarcoma	0.6875	0.7333	0.6471
STAB1 AND NOT-ABCG8	Sarcoma	0.6875	0.7333	0.6471
STAB1 AND NOT-ATP8A2	Sarcoma	0.7273	0.75	0.7059
STAB1 AND NOT-ATP8A1	Sarcoma	0.7407	1	0.5882
STAB1 AND NOT-KCNJ10	Sarcoma	0.7857	1	0.6471
STAB1 AND NOT-HRH3	Sarcoma	0.7097	0.7857	0.6471
STAB1 AND NOT-FXYD6	Sarcoma	0.6923	1	0.5294
STAB1 AND NOT-GGTLC1	Sarcoma	0.7333	0.8462	0.6471
STAB1 AND NOT-HTR1D	Sarcoma	0.7059	0.7059	0.7059
STAB1 AND NOT-GUCY2F	Sarcoma	0.6875	0.7333	0.6471
STAB1 AND NOT-SEMA4D	Sarcoma	0.8276	1	0.7059
STAB1 AND NOT-KCNQ2	Sarcoma	0.6667	0.9	0.5294
STAB1 AND NOT-SLC22A6	Sarcoma	0.6667	0.7692	0.5882
STAB1 AND NOT-ADAM30	Sarcoma	0.6667	0.6875	0.6471
STAB1 AND NOT-OR1Q1	Sarcoma	0.6667	0.9	0.5294
STAB1 AND NOT-XCR1	Sarcoma	0.6667	0.7692	0.5882
STAB1 AND NOT-WNT7A	Sarcoma	0.6667	0.9	0.5294
STAB1 AND NOT-PVRL1	Sarcoma	0.6857	0.6667	0.7059
STAB1 AND NOT-LHFPL5	Sarcoma	0.6667	0.6875	0.6471
STAB1 AND NOT-TSPAN16	Sarcoma	0.7742	0.8571	0.7059
STAB1 AND NOT-CDH9	Sarcoma	0.6667	0.6316	0.7059
STAB1 AND NOT-GRM3	Sarcoma	0.7586	0.9167	0.6471
STAB1 AND NOT-SLC1A2	Sarcoma	0.6667	0.7692	0.5882
STAB1 AND NOT-GALR1	Sarcoma	0.6897	0.8333	0.5882
STAB1 AND NOT-SLC13A5	Sarcoma	0.7586	0.9167	0.6471
STAB1 AND NOT-SLC28A1	Sarcoma	0.7097	0.7857	0.6471
STAB1 AND NOT-HTR1E	Sarcoma	0.7097	0.7857	0.6471

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
PRLR AND NOT-SLC22A11	Breast	0.7292	0.8333	0.6481
PRLR AND NOT-C8B	Breast	0.7292	0.8333	0.6481
PRLR AND NOT-TRPM3	Breast	0.7292	0.8333	0.6481
PRLR AND NOT-MTNR1B	Breast	0.7292	0.8333	0.6481
PPAPDC1A AND NOT-NRXN1	Breast	0.7273	0.8	0.6667
PPAPDC1A AND NOT-GRIK2	Breast	0.7273	0.8	0.6667
PPAPDC1A AND NOT-ATP10B	Breast	0.7629	0.8605	0.6852
PPAPDC1A AND NOT-NTRK3	Breast	0.7255	0.7708	0.6852
SLCO1B1 AND NOT-CLEC1B	Liver	0.9091	1	0.8333
SLCO1B1 AND NOT-KCNN2	Liver	1	1	1
SLCO1B1 AND NOT-MARCO	Liver	0.8	0.6667	1
SLC17A2 AND NOT-CLEC1B	Liver	0.8	1	0.6667
SLC43A1 AND NOT-NPY1R	Liver	0.8	1	0.6667
SLCO1B1 AND NOT-STAB2	Liver	0.8	1	0.6667
SLCO1B1 AND NOT-GPR182	Liver	0.8	1	0.6667
SLCO1B1 AND NOT-CLEC4G	Liver	0.8	1	0.6667
SLCO1B1 AND NOT-PTH1R	Liver	0.8	1	0.6667
SLC10A1 AND NOT-CLEC4G	Liver	0.8	1	0.6667
SLCO1B1 AND NOT-CLEC4M	Liver	0.8	1	0.6667
SLC4A2 AND NOT-SLC9A1	Liver	0.8	1	0.6667
ABCG8 AND NOT-CDHR2	Liver	0.8	1	0.6667
SLC10A1 AND NOT-CLEC1B	Liver	0.8	1	0.6667
SLC4A2 AND NOT-THBD	Liver	0.8	1	0.6667
SLC2A2 AND NOT-CLEC1B	Liver	0.9091	1	0.8333
SLC4A2 AND NOT-PAQR7	Liver	0.8	1	0.6667
SLC17A2 AND NOT-KCNN2	Liver	0.9091	1	0.8333
SLC30A8 AND NOT-FUT1	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-ANO5	Pancreas	0.7059	1	0.5455
SLC30A8 AND NOT-LAG3	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-LIFR	Pancreas	0.625	1	0.4545
SLC30A8 AND NOT-PRLR	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-CHRM3	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-OR1J4	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-TRHDE	Pancreas	0.7778	1	0.6364
NOX4 AND NOT-PAQR7	Pancreas	0.6087	0.5833	0.6364
SLC30A8 AND NOT-NRCAM	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-OR3A3	Pancreas	0.7059	1	0.5455
SLC30A8 AND NOT-SIGLEC11	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-CLCNKB	Pancreas	0.7059	1	0.5455
SLC30A8 AND NOT-LPAR3	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-KCNJ13	Pancreas	0.7059	1	0.5455
SLC30A8 AND NOT-AQP11	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-STX1B	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-SLC38A4	Pancreas	0.625	1	0.4545
SLC30A8 AND NOT-KCNJ5	Pancreas	0.7059	1	0.5455
SLC30A8 AND NOT-TMEM8A	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-NRG4	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-SLC22A7	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-CD300C	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-SLC16A10	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-JPH1	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-INSR	Pancreas	0.7059	1	0.5455
SLC30A8 AND NOT-EPOR	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-SLC23A1	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-KLB	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-SLC26A6	Pancreas	0.7059	1	0.5455
SLC30A8 AND NOT-CNGB3	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-RYR2	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-CHRNA3	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-ABCB8	Pancreas	0.7059	1	0.5455
SLC30A8 AND NOT-PRPH2	Pancreas	0.7059	1	0.5455
SLC30A8 AND NOT-SDK1	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-PLB1	Pancreas	0.625	1	0.4545
SLC30A8 AND NOT-NUCB2	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-KCNN2	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-CA4	Pancreas	0.7059	1	0.5455
SLC30A8 AND NOT-AQP10	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-ADRB1	Pancreas	0.7059	1	0.5455
SLC30A8 AND NOT-FGF10	Pancreas	0.625	1	0.4545
CDH6 AND NOT-SLC9B1	Renal	0.8571	1	0.75
CDH6 AND NOT-PPAPDC1A	Renal	0.8571	1	0.75
CDH6 AND NOT-PHLDB2	Renal	0.8571	1	0.75
CDH6 AND NOT-KCNJ10	Renal	0.8571	1	0.75
CDH6 AND NOT-CDH11	Renal	0.8571	1	0.75
CDH6 AND NOT-OXTR	Renal	0.8571	1	0.75
CDH6 AND NOT-SCN2A	Renal	0.8571	1	0.75
CDH6 AND NOT-SYT11	Renal	0.8571	1	0.75

FIG. 12 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
STAB1 AND NOT-MC2R	Sarcoma	0.7586	0.9167	0.6471
STAB1 AND NOT-TAS2R1	Sarcoma	0.7143	0.9091	0.5882
STAB1 AND NOT-IGDCC3	Sarcoma	0.6897	0.8333	0.5882
STAB1 AND NOT-CACNG3	Sarcoma	0.6667	0.9	0.5294
STAB1 AND NOT-SLC6A13	Sarcoma	0.6667	0.6875	0.6471
STAB1 AND NOT-ACSL6	Sarcoma	0.7742	0.8571	0.7059
STAB1 AND NOT-PCDHB1	Sarcoma	0.6875	0.7333	0.6471
STAB1 AND NOT-HTR3C	Sarcoma	0.6897	0.8333	0.5882
STAB1 AND NOT-SMPD2	Sarcoma	0.6667	0.7692	0.5882
STAB1 AND NOT-SEZ6	Sarcoma	0.6667	0.9	0.5294
STAB1 AND NOT-MTNR1B	Sarcoma	0.7333	0.8462	0.6471
STAB1 AND NOT-C8B	Sarcoma	0.7407	1	0.5882
STAB1 AND NOT-SLC5A11	Sarcoma	0.6471	0.6471	0.6471
STAB1 AND NOT-GRM5	Sarcoma	0.6471	0.6471	0.6471
STAB1 AND NOT-RGSL1	Sarcoma	0.6471	0.6471	0.6471
STAB1 AND NOT-FGF6	Sarcoma	0.6471	0.6471	0.6471
STAB1 AND NOT-RGR	Sarcoma	0.6471	0.6471	0.6471
STAB1 AND NOT-CACNG6	Sarcoma	0.6471	0.6471	0.6471
STAB1 AND NOT-DSCAM	Sarcoma	0.6471	0.6471	0.6471
STAB1 AND NOT-OTOF	Sarcoma	0.6667	0.7692	0.5882
STAB1 AND NOT-AMHR2	Sarcoma	0.6471	0.6471	0.6471
STAB1 AND NOT-CACNG4	Sarcoma	0.6471	0.6471	0.6471
STAB1 AND NOT-OR8D1	Sarcoma	0.6897	0.8333	0.5882
SLC6A7 AND NOT-CLDN10	Sarcoma	0.6667	0.7692	0.5882
STAB1 AND NOT-MAG	Sarcoma	0.6452	0.7143	0.5882
STAB1 AND NOT-KREMEN2	Sarcoma	0.6452	0.7143	0.5882
STAB1 AND NOT-UPK3A	Sarcoma	0.6452	0.7143	0.5882
STAB1 AND NOT-SYT4	Sarcoma	0.6452	0.7143	0.5882
STAB1 AND NOT-BEST3	Sarcoma	0.6429	0.8182	0.5294
STAB1 AND NOT-ATP2B2	Sarcoma	0.6429	0.8182	0.5294
STAB1 AND NOT-ZDHHC2	Sarcoma	0.6471	0.6471	0.6471
STAB1 AND NOT-CABP7	Sarcoma	0.6486	0.6	0.7059
STAB1 AND NOT-EPHA8	Sarcoma	0.7857	1	0.6471
STAB1 AND NOT-NPHS1	Sarcoma	0.6857	0.6667	0.7059
STAB1 AND NOT-CLRN1	Sarcoma	0.6897	0.8333	0.5882
TTYH3 AND NOT-GPR37L1	Sarcoma	0.64	1	0.4706
TTYH3 AND NOT-NRG3	Sarcoma	0.64	1	0.4706
TTYH3 AND NOT-SLC22A13	Sarcoma	0.64	1	0.4706
TTYH3 AND NOT-TTYH2	Sarcoma	0.64	1	0.4706
TTYH3 AND NOT-UMOD	Sarcoma	0.64	1	0.4706
TTYH3 AND NOT-GPR12	Sarcoma	0.64	1	0.4706
TTYH3 AND NOT-CD207	Sarcoma	0.64	1	0.4706
TTYH3 AND NOT-CACNG7	Sarcoma	0.64	1	0.4706
TTYH3 AND NOT-GABRD	Sarcoma	0.64	1	0.4706
TTYH3 AND NOT-SLC12A3	Sarcoma	0.64	1	0.4706
TTYH3 AND NOT-EPHA8	Sarcoma	0.64	1	0.4706
TTYH3 AND NOT-GABRG1	Sarcoma	0.64	1	0.4706
TTYH3 AND NOT-CALY	Sarcoma	0.64	1	0.4706
STAB1 AND NOT-SLC22A9	Sarcoma	0.64	1	0.4706
TTYH3 AND NOT-PCDHGC4	Sarcoma	0.64	1	0.4706
TTYH3 AND NOT-PCDHAC2	Sarcoma	0.64	1	0.4706
TTYH3 AND NOT-LRRC55	Sarcoma	0.64	1	0.4706
TTYH3 AND NOT-RGR	Sarcoma	0.64	1	0.4706
TTYH3 AND NOT-SLC22A11	Sarcoma	0.64	1	0.4706
TTYH3 AND NOT-CCKBR	Sarcoma	0.64	1	0.4706
TTYH3 AND NOT-CACNG3	Sarcoma	0.64	1	0.4706
TTYH3 AND NOT-CD1B	Sarcoma	0.64	1	0.4706
TTYH3 AND NOT-MC2R	Sarcoma	0.64	1	0.4706
TTYH3 AND NOT-KCNK10	Sarcoma	0.64	1	0.4706
TTYH3 AND NOT-HCN4	Sarcoma	0.64	1	0.4706
TTYH3 AND NOT-OR1Q1	Sarcoma	0.64	1	0.4706
TTYH3 AND NOT-GRM3	Sarcoma	0.64	1	0.4706
TTYH3 AND NOT-ATP8A1	Sarcoma	0.64	1	0.4706
TTYH3 AND NOT-OR3A1	Sarcoma	0.64	1	0.4706
CDH17 AND NOT-SLC30A10	Stomach	0.7664	0.6406	0.9535
MUC17 AND NOT-SLC30A10	Stomach	0.8533	1	0.7442
CDH17 AND NOT-BEST2	Stomach	0.6724	0.5342	0.907
MUC17 AND NOT-SLC13A2	Stomach	0.697	1	0.5349
MUC17 AND NOT-SLC22A5	Stomach	0.8378	1	0.7209
MUC17 AND NOT-SLC5A11	Stomach	0.7945	0.9667	0.6744
MUC17 AND NOT-PTPRR	Stomach	0.7945	0.9667	0.6744
CDH17 AND NOT-SLC22A5	Stomach	0.65	0.5065	0.907
MUC17 AND NOT-ABCG8	Stomach	0.8312	0.9412	0.7442
MUC17 AND NOT-SLC31A1	Stomach	0.7671	0.9333	0.6512
MUC17 AND NOT-ABCG5	Stomach	0.7324	0.9286	0.6047
MUC17 AND NOT-FLVCR1	Stomach	0.6364	0.913	0.4884
CDH17 AND NOT-NOX1	Stomach	0.6341	0.4875	0.907

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
CDH6 AND NOT-MSMO1	Renal	0.8571	1	0.75
CDH6 AND NOT-CCKAR	Renal	0.8571	1	0.75
CDH6 AND NOT-ZDHHC2	Renal	0.8571	1	0.75
CDH6 AND NOT-KCNK10	Renal	0.8571	1	0.75
CDH6 AND NOT-PAQR7	Renal	0.8571	1	0.75
CDH6 AND NOT-DSCAM	Renal	0.8571	1	0.75
CDH6 AND NOT-GPR22	Renal	0.8571	1	0.75
CDH6 AND NOT-CLDN16	Renal	0.8571	1	0.75
CDH6 AND NOT-MEGF10	Renal	0.8571	1	0.75
CDH6 AND NOT-CSPG5	Renal	0.8571	1	0.75
CDH6 AND NOT-NKAIN2	Renal	0.8571	1	0.75
CDH6 AND NOT-ZACN	Renal	0.8571	1	0.75
CDH6 AND NOT-HCN1	Renal	0.8571	1	0.75
CDH6 AND NOT-CHRNB4	Renal	0.8571	1	0.75
CDH6 AND NOT-DRD5	Renal	0.8571	1	0.75
CDH6 AND NOT-CDH10	Renal	0.8571	1	0.75
CDH6 AND NOT-CDH8	Renal	0.8571	1	0.75
CDH6 AND NOT-SLC12A5	Renal	0.8571	1	0.75
CDH6 AND NOT-PAQR8	Renal	0.8571	1	0.75
CDH6 AND NOT-MLC1	Renal	0.8571	1	0.75
CDH6 AND NOT-ANTXR2	Renal	0.8571	1	0.75
CDH6 AND NOT-GNRHR	Renal	0.8571	1	0.75
CDH6 AND NOT-ENPP1	Renal	0.8571	1	0.75
CDH6 AND NOT-GABRA2	Renal	0.8571	1	0.75
CDH6 AND NOT-GABRA3	Renal	0.8571	1	0.75
CDH6 AND NOT-AQP6	Renal	0.8571	1	0.75
CDH6 AND NOT-CDH20	Renal	0.8571	1	0.75
CDH6 AND NOT-GABRA5	Renal	0.8571	1	0.75
CDH6 AND NOT-CNTNAP2	Renal	0.8571	1	0.75
CDH6 AND NOT-KIAA0319	Renal	0.8571	1	0.75
CDH6 AND NOT-GPR26	Renal	0.8571	1	0.75
CDH6 AND NOT-SLC4A8	Renal	0.8571	1	0.75
CDH6 AND NOT-FLVCR1	Renal	0.8571	1	0.75
CDH6 AND NOT-CASR	Renal	0.8571	1	0.75
CDH6 AND NOT-NPHS2	Renal	0.8571	1	0.75
CDH6 AND NOT-NPHS1	Renal	0.8571	1	0.75
CDH6 AND NOT-CLDN20	Renal	0.8571	1	0.75
CDH6 AND NOT-SLC6A7	Renal	0.8571	1	0.75
CDH6 AND NOT-SLC6A6	Renal	0.8571	1	0.75
CDH6 AND NOT-SLC30A1	Renal	0.8571	1	0.75
CDH6 AND NOT-ATP8A1	Renal	0.8571	1	0.75
CDH6 AND NOT-CD207	Renal	0.8571	1	0.75
CDH6 AND NOT-SLC12A3	Renal	0.8571	1	0.75
CDH6 AND NOT-SCN1A	Renal	0.8571	1	0.75
CDH6 AND NOT-CACNG8	Renal	0.8571	1	0.75
CDH6 AND NOT-ZP4	Renal	0.8571	1	0.75
CDH6 AND NOT-DCSTAMP	Renal	0.8571	1	0.75
CDH6 AND NOT-SLC32A1	Renal	0.8571	1	0.75
CDH6 AND NOT-SLC30A10	Renal	0.8571	1	0.75
CDH6 AND NOT-ASTN1	Renal	0.8571	1	0.75
CDH6 AND NOT-DTNA	Renal	0.8571	1	0.75
CDH6 AND NOT-DISP2	Renal	0.8571	1	0.75
CDH6 AND NOT-GPR83	Renal	0.8571	1	0.75
CDH6 AND NOT-MTNR1B	Renal	0.8571	1	0.75
CDH6 AND NOT-AOC3	Renal	0.8571	1	0.75
CDH6 AND NOT-SCN10A	Renal	0.8571	1	0.75
CDH6 AND NOT-PCDHAC2	Renal	0.8571	1	0.75
CDH6 AND NOT-KCNJ9	Renal	0.8571	1	0.75
CDH6 AND NOT-CD1B	Renal	0.8571	1	0.75
CDH6 AND NOT-CACNG7	Renal	0.8571	1	0.75
CDH6 AND NOT-GPR37L1	Renal	0.8571	1	0.75
CDH6 AND NOT-GRM1	Renal	0.8571	1	0.75
CDH6 AND NOT-OPALIN	Renal	0.8571	1	0.75
CDH6 AND NOT-KCNK6	Renal	0.8571	1	0.75
CDH6 AND NOT-TRHR	Renal	0.8571	1	0.75
CDH6 AND NOT-OR52D1	Renal	0.8571	1	0.75
CDH6 AND NOT-OR1C1	Renal	0.8571	1	0.75
CDH6 AND NOT-GHRHR	Renal	0.8571	1	0.75
CDH6 AND NOT-OPCML	Renal	0.8571	1	0.75
CDH6 AND NOT-SLC10A1	Renal	0.8571	1	0.75
CDH6 AND NOT-HTR5A	Renal	0.8571	1	0.75
CDH6 AND NOT-EBP	Renal	0.8571	1	0.75
CDH6 AND NOT-SLC5A11	Renal	0.8571	1	0.75
CDH6 AND NOT-ANO4	Renal	0.8571	1	0.75
CDH6 AND NOT-CNGA4	Renal	0.8571	1	0.75
CDH6 AND NOT-CLDN14	Renal	0.8571	1	0.75
CDH6 AND NOT-MUC17	Renal	0.8571	1	0.75

FIG. 12 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
MUC17 AND NOT-P2RX4	Stomach	0.7945	0.9667	0.6744
MUC17 AND NOT-GABRA2	Stomach	0.6957	0.9231	0.5581
CDH17 AND NOT-MUC12	Stomach	0.6271	0.4933	0.8605
MUC17 AND NOT-BEST2	Stomach	0.8219	1	0.6977
MUC17 AND NOT-LCT	Stomach	0.8158	0.9394	0.7209
MUC17 AND NOT-CCR9	Stomach	0.8158	0.9394	0.7209
MUC17 AND NOT-FUT3	Stomach	0.697	1	0.5349
MUC17 AND NOT-CD36	Stomach	0.8	0.9375	0.6977
MUC17 AND NOT-GPR22	Stomach	0.8312	0.9412	0.7442
MUC17 AND NOT-KCNK10	Stomach	0.7324	0.9286	0.6047
MUC17 AND NOT-EBP	Stomach	0.7429	0.963	0.6047
MUC17 AND NOT-CACNG4	Stomach	0.7324	0.9286	0.6047
MUC17 AND NOT-ENPP1	Stomach	0.8	0.9375	0.6977
MUC17 AND NOT-SLC22A18	Stomach	0.6129	1	0.4419
MUC17 AND NOT-NKAIN1	Stomach	0.6154	0.9091	0.4651
MUC17 AND NOT-NOX1	Stomach	0.8219	1	0.6977
MUC17 AND NOT-PTPRT	Stomach	0.75	0.931	0.6279
MUC17 AND NOT-MUC12	Stomach	0.8056	1	0.6744
MUC17 AND NOT-SLC17A2	Stomach	0.7246	0.9615	0.5814
MUC17 AND NOT-KIAA0319	Stomach	0.7143	0.9259	0.5814
MUC17 AND NOT-PCDHA9	Stomach	0.8158	0.9394	0.7209
MUC17 AND NOT-PAQR7	Stomach	0.6957	0.9231	0.5581
MUC17 AND NOT-PVRL1	Stomach	0.6567	0.9167	0.5116
MUC17 AND NOT-IL17RA	Stomach	0.8312	0.9412	0.7442
MUC17 AND NOT-AGER	Stomach	0.6567	0.9167	0.5116
MUC17 AND NOT-SLC39A8	Stomach	0.7429	0.963	0.6047
MUC17 AND NOT-AMHR2	Stomach	0.7778	0.9655	0.6512
MUC17 AND NOT-KCNV1	Stomach	0.8	0.9375	0.6977
MUC17 AND NOT-ADAM2	Stomach	0.6667	0.9565	0.5116
MUC17 AND NOT-CD58	Stomach	0.6129	1	0.4419
MUC17 AND NOT-KCNK16	Stomach	0.6032	0.95	0.4419
MUC17 AND NOT-SLCO1B1	Stomach	0.6032	0.95	0.4419
MUC17 AND NOT-RGR	Stomach	0.8312	0.9412	0.7442
MUC17 AND NOT-EFNB2	Stomach	0.7838	0.9355	0.6744
MUC17 AND NOT-ANTXR2	Stomach	0.6957	0.9231	0.5581
MUC17 AND NOT-CEACAM7	Stomach	0.8421	0.9697	0.7442
MUC17 AND NOT-SLC9A1	Stomach	0.7838	0.9355	0.6744
MUC17 AND NOT-CHRNA1	Stomach	0.8312	0.9412	0.7442
MUC17 AND NOT-KCNJ9	Stomach	0.6462	0.9545	0.4884
MUC17 AND NOT-GALR1	Stomach	0.7143	0.9259	0.5814
MUC17 AND NOT-CDH9	Stomach	0.7945	0.9667	0.6744
MUC17 AND NOT-NPFFR1	Stomach	0.8158	0.9394	0.7209
MUC17 AND NOT-CLDN10	Stomach	0.8158	0.9394	0.7209
MUC17 AND NOT-SCN8A	Stomach	0.6154	0.9091	0.4651
MUC17 AND NOT-GRM7	Stomach	0.6765	0.92	0.5349
MUC17 AND NOT-EPHA10	Stomach	0.8267	0.9688	0.7209
MUC17 AND NOT-SORL1	Stomach	0.6765	0.92	0.5349
IL15RA AND NOT-CD36	Stomach	0.6522	0.6122	0.6977
NOT-CLDN10 AND CDH11	Colon	0.75	1	0.6
NOT-ATP2B2 AND HCN1	Colon	0.75	1	0.6
NOT-CLDN10 AND EPHA2	Colon	0.6667	0.75	0.6
NOT-MEGF10 AND HCN1	Colon	0.75	1	0.6
NOT-CLDN10 AND SLC2A1	Colon	0.6667	0.75	0.6
NOT-SLC6A6 AND LRFN2	Ependymoma	1	1	1
NOT-SLC6A6 AND UMODL1	Ependymoma	0.6667	0.5	1
NOT-SLC6A6 AND SLC4A5	Ependymoma	0.6667	0.5	1
NOT-SLC6A6 AND ATP8A1	Ependymoma	0.6667	0.5	1
NOT-BTN3A1 AND TNFRSF10B	Ependymoma	0.6667	0.5	1
NOT-SLC6A6 AND CLDN16	Ependymoma	0.6667	0.5	1
NOT-CD36 AND IL15RA	Esophagus	0.6667	0.625	0.7143
NOT-ATP8A1 AND SLC6A18	Leiomyosarcoma	0.6957	0.8	0.6154
NOT-SLC16A5 AND VANGL1	Leiomyosarcoma	0.7619	1	0.6154
NOT-ATP8A1 AND ATP8B2	Leiomyosarcoma	0.7826	0.9	0.6923
NOT-ATP8A1 AND EMP3	Leiomyosarcoma	0.6957	0.8	0.6154
NOT-ATP8A1 AND SYT11	Leiomyosarcoma	0.64	0.6667	0.6154
NOT-ATP8A1 AND CHRND	Leiomyosarcoma	0.6207	0.5625	0.6923
NOT-ATP8A1 AND PTGIR	Leiomyosarcoma	0.6957	0.8	0.6154
NOT-ATP8A1 AND CHRNA2	Leiomyosarcoma	0.6452	0.5556	0.7692
NOT-ATP8A1 AND FLVCR1	Leiomyosarcoma	0.6316	1	0.4615
NOT-ATP8A1 AND GABRA3	Leiomyosarcoma	0.75	0.8182	0.6923
NOT-ATP8A1 AND NPFFR1	Leiomyosarcoma	0.8696	1	0.7692
NOT-ATP8A1 AND GABRA5	Leiomyosarcoma	0.6429	0.6	0.6923
NOT-ATP8A1 AND GYPC	Leiomyosarcoma	0.7619	1	0.6154
NOT-ATP8A1 AND IL13	Leiomyosarcoma	0.6957	0.8	0.6154
NOT-ATP8A1 AND SLC6A7	Leiomyosarcoma	0.6364	0.7778	0.5385
NOT-CLDN10 AND SLC6A7	Leiomyosarcoma	0.6316	1	0.4615
NOT-ATP8A1 AND CLDN15	Leiomyosarcoma	0.6	0.8571	0.4615

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
CDH6 AND NOT-DIO1	Renal	0.8571	1	0.75
CDH6 AND NOT-KCNA2	Renal	0.8571	1	0.75
CDH6 AND NOT-UNC5A	Renal	0.8571	1	0.75
CDH6 AND NOT-SLC22A8	Renal	0.8571	1	0.75
CDH6 AND NOT-BTN3A1	Renal	0.8571	1	0.75
CDH6 AND NOT-SLC22A16	Renal	0.8571	1	0.75
NOX1 AND NOT-BEST4	Colon	1	1	1
NOX1 AND NOT-FLVCR2	Colon	0.9091	0.8333	1
NOX1 AND NOT-SLC22A23	Colon	0.9091	0.8333	1
NOX1 AND NOT-TSPAN7	Colon	1	1	1
CDH17 AND NOT-FLVCR2	Colon	0.9091	0.8333	1
B3GNT3 AND NOT-SLC9A1	Colon	0.8889	1	0.8
PMEPA1 AND NOT-STEAP4	Colon	0.8889	1	0.8
FAT1 AND NOT-LPAR1	Colon	0.8889	1	0.8
NOX1 AND NOT-SEMA4G	Colon	0.8889	1	0.8
CDH17 AND NOT-SEMA4G	Colon	0.8889	1	0.8
CD320 AND NOT-SLC9A1	Colon	0.8889	1	0.8
HEPH AND NOT-SLC9A1	Colon	1	1	1
IFI6 AND NOT-NAALAD2	Colon	0.9091	0.8333	1
PMEPA1 AND NOT-GHR	Colon	0.8889	1	0.8
NOX1 AND NOT-SEMA6D	Colon	0.8889	1	0.8
NOX1 AND NOT-CA12	Colon	0.9091	0.8333	1
PSENEN AND NOT-SLC16A5	Colon	0.8889	1	0.8
IFI6 AND NOT-SCN11A	Colon	0.8889	1	0.8
MEP1A AND NOT-SLC9A1	Colon	1	1	1
NOX1 AND NOT-IL1R2	Colon	0.8333	0.7143	1
NOX1 AND NOT-PTGDR	Colon	1	1	1
PMEPA1 AND NOT-SPAM1	Colon	0.8889	1	0.8
NOX1 AND NOT-SLC4A4	Colon	0.8889	1	0.8
MEP1A AND NOT-SLC30A10	Colon	0.8889	1	0.8
CDH17 AND NOT-SLC4A4	Colon	0.8889	1	0.8
IFI6 AND NOT-ATP8B4	Colon	0.8333	0.7143	1
IFI6 AND NOT-CLEC12A	Colon	1	1	1
NOX1 AND NOT-PAQR5	Colon	1	1	1
CDH17 AND NOT-SLC22A23	Colon	0.9091	0.8333	1
IFI6 AND NOT-CD274	Colon	0.8333	0.7143	1
FZD3 AND NOT-SLC24A2	Ependymoma	1	1	1
FZD3 AND NOT-ABCG4	Ependymoma	1	1	1
FZD3 AND NOT-GABRB2	Ependymoma	1	1	1
SLC1A3 AND NOT-LAPTM5	Ependymoma	1	1	1
SLC1A3 AND NOT-SLC31A1	Ependymoma	1	1	1
FZD3 AND NOT-CDH8	Ependymoma	1	1	1
FZD3 AND NOT-CACNA1B	Ependymoma	1	1	1
FZD3 AND NOT-CACNA1E	Ependymoma	1	1	1
NRG3 AND NOT-SLC2A3	Ependymoma	1	1	1
SLC1A3 AND NOT-PHLDB2	Ependymoma	1	1	1
FZD3 AND NOT-NETO1	Ependymoma	1	1	1
FZD3 AND NOT-SCN8A	Ependymoma	1	1	1
FZD3 AND NOT-DGKE	Ependymoma	1	1	1
MEGF10 AND NOT-KCNJ2	Ependymoma	1	1	1
FZD3 AND NOT-SCN2A	Ependymoma	1	1	1
FZD3 AND NOT-SYT4	Ependymoma	1	1	1
SLC1A3 AND NOT-SLC30A1	Ependymoma	1	1	1
SLC1A3 AND NOT-PROCR	Ependymoma	1	1	1
FZD3 AND NOT-GRM5	Ependymoma	1	1	1
FZD3 AND NOT-GRM3	Ependymoma	1	1	1
SLC7A11 AND NOT-SLC6A6	Ependymoma	1	1	1
FZD3 AND NOT-KCNA1	Ependymoma	1	1	1
FZD3 AND NOT-SLC6A6	Ependymoma	1	1	1
SLC1A3 AND NOT-CD36	Ependymoma	1	1	1
SLC1A3 AND NOT-IL15RA	Ependymoma	1	1	1
FZD3 AND NOT-ATP2B2	Ependymoma	1	1	1
ZDHHC17 AND NOT-SLC6A6	Ependymoma	1	1	1
FZD3 AND NOT-GABRA1	Ependymoma	1	1	1
FZD3 AND NOT-GABRA2	Ependymoma	1	1	1
SLC16A2 AND NOT-PHLDB2	Ependymoma	1	1	1
FZD3 AND NOT-OPRK1	Ependymoma	1	1	1
SLC1A3 AND NOT-STAB1	Ependymoma	1	1	1
FZD3 AND NOT-UNC5A	Ependymoma	1	1	1
FZD3 AND NOT-ATP8A2	Ependymoma	1	1	1
FZD3 AND NOT-CACNG8	Ependymoma	1	1	1
SLC1A3 AND NOT-DYSF	Ependymoma	1	1	1
SLC1A3 AND NOT-CXCL16	Ependymoma	1	1	1
FZD3 AND NOT-SLC32A1	Ependymoma	1	1	1
FZD3 AND NOT-PTPRR	Ependymoma	1	1	1
FZD3 AND NOT-SLC12A5	Ependymoma	1	1	1
SLC1A3 AND NOT-SLC38A1	Ependymoma	1	1	1

FIG. 12 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
NOT-ATP8A1 AND GPR135	Leiomyosarcoma	0.6667	0.5882	0.7692
NOT-SMPD2 AND VANGL1	Leiomyosarcoma	0.8696	1	0.7692
NOT-ATP8A1 AND CHRNB4	Leiomyosarcoma	0.7143	0.6667	0.7692
NOT-ATP8A1 AND CCKAR	Leiomyosarcoma	0.7273	0.8889	0.6154
NOT-ATP8A1 AND TTYH3	Leiomyosarcoma	0.7826	0.9	0.6923
NOT-ATP8A1 AND JPH3	Leiomyosarcoma	0.6429	0.6	0.6923
NOT-ATP8A1 AND BTN3A3	Leiomyosarcoma	0.7619	1	0.6154
NOT-ATP8A1 AND MDGA2	Leiomyosarcoma	0.6667	0.5882	0.7692
NOT-ATP8A1 AND MIP	Leiomyosarcoma	0.72	0.75	0.6923
NOT-ATP8A1 AND MPL	Leiomyosarcoma	0.6957	0.8	0.6154
NOT-ATP8A1 AND GABRG3	Leiomyosarcoma	0.6207	0.5625	0.6923
NOT-ATP8A1 AND SLC26A8	Leiomyosarcoma	0.6207	0.5625	0.6923
NOT-ATP8A1 AND TNFRSF12A	Leiomyosarcoma	0.6667	0.7273	0.6154
NOT-ATP8A1 AND OPRM1	Leiomyosarcoma	0.6061	0.5	0.7692
NOT-ATP8A1 AND AGER	Leiomyosarcoma	0.6957	0.8	0.6154
NOT-CLDN10 AND FAM26D	Leiomyosarcoma	0.6957	0.8	0.6154
NOT-ATP8A1 AND BEST3	Leiomyosarcoma	0.6061	0.5	0.7692
NOT-ATP8A1 AND PPAPDC1A	Leiomyosarcoma	0.6316	1	0.4615
NOT-ATP8A1 AND ATP7A	Leiomyosarcoma	0.7	1	0.5385
NOT-ATP8A1 AND FAM26D	Leiomyosarcoma	0.6087	0.7	0.5385
NOT-PHLDB2 AND EMP3	AML	0.9377	0.9163	0.9602
NOT-PHLDB2 AND CD44	AML	0.9367	0.9037	0.9721
NOT-PHLDB2 AND SLC39A8	AML	0.9275	0.9655	0.8924
NOT-STEAP4 AND EMP3	AML	0.8919	0.8652	0.9203
NOT-STEAP4 AND CD44	AML	0.9126	0.8902	0.9363
NOT-PHLDB2 AND CD36	AML	0.8784	0.8649	0.8924
NOT-PHLDB2 AND GYPC	AML	0.8773	0.802	0.9681
NOT-PHLDB2 AND LAPTM5	AML	0.9118	0.8464	0.988
NOT-PHLDB2 AND P2RX4	AML	0.9295	0.8905	0.9721
NOT-GPR137B AND SLC39A8	AML	0.8518	0.8947	0.8127
NOT-ATP8B1 AND SLC39A8	AML	0.8515	0.8465	0.8566
NOT-GPR137B AND P2RX4	AML	0.8499	0.9054	0.8008
NOT-PHLDB2 AND ANPEP	AML	0.8505	0.7763	0.9402
NOT-GPR137B AND SLC30A1	AML	0.849	0.8703	0.8287
NOT-PHLDB2 AND SLCO1B3	AML	0.8909	0.831	0.9602
NOT-PHLDB2 AND SLC43A1	AML	0.8455	0.9163	0.7849
NOT-PHLDB2 AND SLC30A1	AML	0.8671	0.7726	0.988
NOT-PHLDB2 AND FMNL1	AML	0.8436	0.7759	0.9243
NOT-PHLDB2 AND VAMP8	AML	0.9168	0.8552	0.988
NOT-LRIG3 AND SLC39A8	AML	0.8436	0.8386	0.8486
NOT-GPR137B AND CD44	AML	0.8412	0.8718	0.8127
NOT-GPR137B AND LAPTM5	AML	0.8404	0.8525	0.8287
NOT-FAT1 AND SLC39A8	AML	0.8306	0.8408	0.8207
NOT-GPR137B AND EMP3	AML	0.8204	0.841	0.8008
NOT-PHLDB2 AND SLC16A5	AML	0.813	0.8947	0.745
NOT-GPR137B AND FMNL1	AML	0.8126	0.8616	0.7689
NOT-LRIG3 AND SLC43A1	AML	0.81	0.8509	0.7729
NOT-PHLDB2 AND IL15RA	AML	0.8097	0.9104	0.7291
NOT-GHR AND SLC43A1	AML	0.8077	0.871	0.753
NOT-AOC3 AND SLC43A1	AML	0.8075	0.8502	0.7689
NOT-ATP8B1 AND SLC43A1	AML	0.8075	0.8502	0.7689
NOT-STEAP4 AND SLC43A1	AML	0.7922	0.8673	0.7291
NOT-PHLDB2 AND SLC5A2	AML	0.7919	0.7363	0.8566
NOT-SEMA4B AND P2RX4	AML	0.7919	0.6882	0.9323
NOT-ATP8B1 AND P2RX4	AML	0.7907	0.6781	0.9482
NOT-PHLDB2 AND STAB1	AML	0.7905	0.8632	0.7291
NOT-CDH11 AND SLC43A1	AML	0.7902	0.8083	0.7729
NOT-PHLDB2 AND BST2	AML	0.7902	0.7518	0.8327
NOT-ATP8B1 AND VAMP8	AML	0.7899	0.6545	0.996
NOT-SCARA5 AND SLC43A1	AML	0.7884	0.8225	0.757
NOT-ATP8B1 AND SLC30A1	AML	0.7862	0.6494	0.996
NOT-ATP8B1 AND CD44	AML	0.7847	0.6543	0.9801
NOT-TGFBI AND EMP3	AML	0.7826	0.8612	0.7171
NOT-AOC3 AND CD44	AML	0.7783	0.6489	0.9721
NOT-SORL1 AND IFNAR2	Liposarcoma	0.6129	0.7308	0.5278
NOT-SORL1 AND PTGER4	Liposarcoma	0.6087	0.6364	0.5833
NOT-ANPEP AND CLDN15	B-Cell Diffuse	0.6	0.5581	0.6486
NOT-VANGL1 AND CLDN15	Mantle-Cell Lymphoma	0.775	0.7381	0.8158
NOT-SLC31A1 AND CLDN15	Mantle-Cell Lymphoma	0.7692	0.75	0.7895
NOT-ANPEP AND HLA-DRB1	Mantle-Cell Lymphoma	0.8471	0.766	0.9474
NOT-VANGL1 AND SLC38A1	Mantle-Cell Lymphoma	0.7416	0.6471	0.8684
NOT-STEAP4 AND EMP3	Mantle-Cell Lymphoma	0.7333	0.6346	0.8684
NOT-VANGL1 AND HLA-DRB1	Mantle-Cell Lymphoma	0.7048	0.5522	0.9737
NOT-ANPEP AND CLDN15	Mantle-Cell Lymphoma	0.7027	0.7222	0.6842
NOT-VANGL1 AND SLC50A1	Mantle-Cell Lymphoma	0.7021	0.5893	0.8684
NOT-STEAP4 AND CD44	Mantle-Cell Lymphoma	0.72	0.5806	0.9474
NOT-SLC31A1 AND HLA-DRB1	Mantle-Cell Lymphoma	0.6981	0.5441	0.9737

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
PTPRA AND NOT-SLC6A6	Ependymoma	1	1	1
SLC52A2 AND NOT-DPP10	Esophagus	0.7273	1	0.5714
SLC52A2 AND NOT-SCARA5	Esophagus	0.7273	1	0.5714
CDH17 AND NOT-SEMA6D	Esophagus	0.7692	0.8333	0.7143
SLC52A2 AND NOT-STAB1	Esophagus	0.7273	1	0.5714
SLC52A2 AND NOT-SLC13A1	Esophagus	0.6667	1	0.5
SLC52A2 AND NOT-SLC22A5	Esophagus	0.6667	1	0.5
CXCL16 AND NOT-ADCY7	Esophagus	0.7273	1	0.5714
SLC52A2 AND NOT-TMEFF2	Esophagus	0.6667	1	0.5
SLC52A2 AND NOT-HCN1	Esophagus	0.6667	1	0.5
CXCL16 AND NOT-ADRB2	Esophagus	0.64	0.7273	0.5714
SLC52A2 AND NOT-CD36	Esophagus	0.7273	1	0.5714
CXCL16 AND NOT-TLR1	Esophagus	0.72	0.8182	0.6429
SLC52A2 AND NOT-MDGA2	Esophagus	0.6667	1	0.5
SLC52A2 AND NOT-TAS2R10	Esophagus	0.7826	1	0.6429
SLC52A2 AND NOT-NFASC	Esophagus	0.7826	1	0.6429
SLC52A2 AND NOT-TAS1R1	Esophagus	0.7273	1	0.5714
SLC52A2 AND NOT-MLANA	Esophagus	0.7273	1	0.5714
SLC52A2 AND NOT-ATP8A2	Esophagus	0.8333	1	0.7143
SLC52A2 AND NOT-BEST2	Esophagus	0.7273	1	0.5714
SLC52A2 AND NOT-MMP24	Esophagus	0.8333	1	0.7143
SLC52A2 AND NOT-DSC1	Esophagus	0.8333	1	0.7143
SLC52A2 AND NOT-FMNL1	Esophagus	0.7826	1	0.6429
SLC52A2 AND NOT-AGER	Esophagus	0.6667	1	0.5
CXCL16 AND NOT-PILRA	Esophagus	0.6667	1	0.5
SLC52A2 AND NOT-CATSPER3	Esophagus	0.7826	1	0.6429
SLC52A2 AND NOT-SLC1A2	Esophagus	0.8333	1	0.7143
SLC52A2 AND NOT-SCN1A	Esophagus	0.6667	1	0.5
SLC52A2 AND NOT-TSHR	Esophagus	0.6667	1	0.5
SLC52A2 AND NOT-PCDH15	Esophagus	0.7826	1	0.6429
SLC52A2 AND NOT-ATP13A5	Esophagus	0.7826	1	0.6429
SLC52A2 AND NOT-GRIA4	Esophagus	0.6667	1	0.5
SLC52A2 AND NOT-HTR1F	Esophagus	0.7273	1	0.5714
SLC52A2 AND NOT-DTNA	Esophagus	0.7273	1	0.5714
SLC52A2 AND NOT-ABHD3	Esophagus	0.7273	1	0.5714
SLC52A2 AND NOT-SLC30A10	Esophagus	0.8333	1	0.7143
CXCL16 AND NOT-FCER1A	Esophagus	0.6154	0.6667	0.5714
SLC52A2 AND NOT-UMOD	Esophagus	0.6667	1	0.5
SLC52A2 AND NOT-TSPAN16	Esophagus	0.7826	1	0.6429
SLC52A2 AND NOT-ATP8B2	Esophagus	0.8333	1	0.7143
SLC52A2 AND NOT-EPHA10	Esophagus	0.8333	1	0.7143
SLC52A2 AND NOT-GABRA5	Esophagus	0.7273	1	0.5714
SLC52A2 AND NOT-OPALIN	Esophagus	0.6667	1	0.5
SLC52A2 AND NOT-FXYD7	Esophagus	0.7273	1	0.5714
SLC52A2 AND NOT-GABRA3	Esophagus	0.7273	1	0.5714
SLC52A2 AND NOT-CNGA4	Esophagus	0.7826	1	0.6429
SLC52A2 AND NOT-GUCY2D	Esophagus	0.8333	1	0.7143
SLC52A2 AND NOT-KCNV1	Esophagus	0.6667	1	0.5
SLC52A2 AND NOT-OR3A1	Esophagus	0.6667	1	0.5
SLC52A2 AND NOT-NTSR2	Esophagus	0.6	1	0.4286
SLC52A2 AND NOT-CSMD3	Esophagus	0.6	1	0.4286
CDH17 AND NOT-CA4	Esophagus	0.6	1	0.4286
SLC52A2 AND NOT-CHRNA9	Esophagus	0.6667	1	0.5
SLC52A2 AND NOT-GABRD	Esophagus	0.6	1	0.4286
SLC52A2 AND NOT-CSMD2	Esophagus	0.6	1	0.4286
SLC52A2 AND NOT-GABRA4	Esophagus	0.6	1	0.4286
SLC52A2 AND NOT-DRD5	Esophagus	0.6	1	0.4286
SLC52A2 AND NOT-CHRNA1	Esophagus	0.6	1	0.4286
SLC52A2 AND NOT-AGTR2	Esophagus	0.6	1	0.4286
SLC52A2 AND NOT-GABRG1	Esophagus	0.6	1	0.4286
SLC52A2 AND NOT-OR2L13	Esophagus	0.6	1	0.4286
SLC52A2 AND NOT-OR8B2	Esophagus	0.7273	1	0.5714
SLC52A2 AND NOT-CACNG3	Esophagus	0.6667	1	0.5
SLC52A2 AND NOT-TRPM3	Esophagus	0.6	1	0.4286
SLC52A2 AND NOT-FAM26D	Esophagus	0.7273	1	0.5714
SLC52A2 AND NOT-NOX1	Esophagus	0.7273	1	0.5714
SLC52A2 AND NOT-GPR12	Esophagus	0.6667	1	0.5
SLC52A2 AND NOT-CNTNAP4	Esophagus	0.6	1	0.4286
SLC52A2 AND NOT-SLC26A8	Esophagus	0.6	1	0.4286
SLC52A2 AND NOT-CEACAM7	Esophagus	0.7273	1	0.5714
SLC52A2 AND NOT-CALN1	Esophagus	0.6667	1	0.5
SLC52A2 AND NOT-SLC23A2	Esophagus	0.7273	1	0.5714
SLC52A2 AND NOT-HTR6	Esophagus	0.7273	1	0.5714
SLC52A2 AND NOT-UMODL1	Esophagus	0.7273	1	0.5714
SLC52A2 AND NOT-CACNG7	Esophagus	0.6667	1	0.5
SLC52A2 AND NOT-SLC4A5	Esophagus	0.6	1	0.4286
SLC52A2 AND NOT-CNIH2	Esophagus	0.8333	1	0.7143

FIG. 12 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
NOT-CXCL16 AND HLA-DRB1	Mantle-Cell Lymphoma	0.6835	0.6585	0.7105
NOT-CXCL16 AND BTN3A1	Mantle-Cell Lymphoma	0.6857	0.75	0.6316
NOT-CD93 AND BTN3A1	Mantle-Cell Lymphoma	0.7561	0.7045	0.8158
NOT-ANPEP AND BTN3A1	Mantle-Cell Lymphoma	0.7561	0.7045	0.8158
NOT-VANGL1 AND SEMA4B	Mantle-Cell Lymphoma	0.6517	0.5686	0.7632
NOT-STEAP4 AND BTN3A1	Mantle-Cell Lymphoma	0.8158	0.8158	0.8158
NOT-VANGL1 AND LAPTM5	Mantle-Cell Lymphoma	0.6441	0.475	1
NOT-CD36 AND BTN3A1	Mantle-Cell Lymphoma	0.7126	0.6327	0.8158
NOT-CHRNG AND SLC38A1	Mantle-Cell Lymphoma	0.6377	0.7097	0.5789
NOT-ANPEP AND SLC50A1	Mantle-Cell Lymphoma	0.6279	0.5625	0.7105
NOT-VANGL1 AND BTN3A1	Mantle-Cell Lymphoma	0.6239	0.4789	0.8947
NOT-VANGL1 AND CD44	Mantle-Cell Lymphoma	0.6154	0.4557	0.9474
NOT-ANPEP AND IFNAR2	Mantle-Cell Lymphoma	0.6596	0.5536	0.8158
NOT-STEAP4 AND GYPC	Mantle-Cell Lymphoma	0.6076	0.5854	0.6316
NOT-ANPEP AND SEMA4B	Mantle-Cell Lymphoma	0.6067	0.5294	0.7105
NOT-STEAP4 AND HLA-DRB1	Mantle-Cell Lymphoma	0.7957	0.6727	0.9737
NOT-CD93 AND EMP3	Mantle-Cell Lymphoma	0.6022	0.5091	0.7368
NOT-VANGL1 AND IFNAR2	Mantle-Cell Lymphoma	0.6018	0.4533	0.8947
NOT-VANGL1 AND EMP3	Mantle-Cell Lymphoma	0.6	0.4583	0.8684
NOT-ANPEP AND CD44	Mantle-Cell Lymphoma	0.6034	0.4487	0.9211
NOT-CD36 AND HLA-DRB1	Mantle-Cell Lymphoma	0.7826	0.6667	0.9474
NOT-CHRNA2 AND ICAM1	Melanoma	0.9	1	0.8182
NOT-NMUR2 AND ICAM1	Melanoma	0.9	1	0.8182
NOT-CLDN19 AND ICAM1	Melanoma	0.9	1	0.8182
NOT-GLRA2 AND ICAM1	Melanoma	0.9	1	0.8182
NOT-SLC22A9 AND FAT1	Melanoma	0.8421	1	0.7273
NOT-CHRNA2 AND DCSTAMP	Melanoma	0.8421	1	0.7273
NOT-ADAM29 AND FAT1	Melanoma	0.9524	1	0.9091
NOT-SLC10A2 AND ICAM1	Melanoma	0.8571	0.9	0.8182
NOT-CHRNA2 AND IFI6	Melanoma	0.8571	0.9	0.8182
NOT-CHRNA2 AND FAT1	Melanoma	1	1	1
NOT-ADAM29 AND VANGL1	Melanoma	0.9091	0.9091	0.9091
NOT-C8B AND ICAM1	Melanoma	0.9	1	0.8182
NOT-SLC6A13 AND ICAM1	Melanoma	0.8421	1	0.7273
NOT-CACNG6 AND ICAM1	Melanoma	0.9	1	0.8182
NOT-CLDN19 AND IFI6	Melanoma	0.8182	0.8182	0.8182
NOT-SLC22A8 AND ICAM1	Melanoma	0.8182	0.8182	0.8182
NOT-CHRNA2 AND GPR19	Melanoma	1	1	1
NOT-ADAM29 AND SLC23A2	Melanoma	0.8421	1	0.7273
NOT-ADAM29 AND LRIG3	Melanoma	1	1	1
NOT-NMUR2 AND FAT1	Melanoma	1	1	1
NOT-NPFFR1 AND GPR19	Melanoma	0.9524	1	0.9091
NOT-SCN10A AND ICAM1	Melanoma	0.8421	1	0.7273
NOT-SLC8A3 AND VANGL1	Melanoma	0.8421	1	0.7273
NOT-ADAM29 AND PLVAP	Melanoma	1	1	1
NOT-SLC22A9 AND LRIG3	Melanoma	0.8421	1	0.7273
NOT-CHRNA2 AND CDH11	Melanoma	0.9524	1	0.9091
NOT-CHRNA2 AND TNFRSF12A	Melanoma	0.9524	1	0.9091
NOT-NMUR2 AND DCSTAMP	Melanoma	0.8421	1	0.7273
NOT-TMEM235 AND VANGL1	Melanoma	0.8571	0.9	0.8182
NOT-SLC10A2 AND TTYH2	Melanoma	0.8	0.7143	0.9091
NOT-NMUR2 AND TTYH2	Melanoma	0.8	0.7143	0.9091
NOT-ADAM29 AND TTYH2	Melanoma	0.8	0.7143	0.9091
NOT-CLDN19 AND TTYH2	Melanoma	0.8	0.7143	0.9091
NOT-CLDN19 AND VANGL1	Melanoma	0.9524	1	0.9091
NOT-CACNG6 AND TTYH2	Melanoma	0.8	0.7143	0.9091
NOT-NMUR2 AND IFI6	Melanoma	0.8182	0.8182	0.8182
NOT-CHRNA2 AND LRIG3	Melanoma	1	1	1
NOT-CLDN19 AND FAT1	Melanoma	0.9524	1	0.9091
NOT-CHRNA2 AND ENPP1	Melanoma	0.8696	0.8333	0.9091
NOT-NMUR2 AND EPHA2	Melanoma	0.8571	0.9	0.8182
NOT-HTR1E AND VANGL1	Melanoma	0.9091	0.9091	0.9091
NOT-CLDN19 AND GPR19	Melanoma	1	1	1
NOT-CHRNA2 AND GJA3	Melanoma	0.9	1	0.8182
NOT-CLDN19 AND TNFRSF12A	Melanoma	0.9524	1	0.9091
NOT-CDH10 AND GPR158	Melanoma	0.8421	1	0.7273
NOT-CHRNA2 AND PLVAP	Melanoma	1	1	1
NOT-ADAM29 AND ENPP1	Melanoma	0.9524	1	0.9091
NOT-C8B AND TTYH2	Melanoma	0.8182	0.8182	0.8182
NOT-SLC17A2 AND VANGL1	Melanoma	0.8571	0.9	0.8182
NOT-NTSR2 AND GPR19	Melanoma	1	1	1
NOT-TMEM235 AND ICAM1	Melanoma	0.8421	1	0.7273
NOT-SLC22A8 AND IFI6	Melanoma	0.8182	0.8182	0.8182
NOT-SLC10A2 AND IFI6	Melanoma	0.8182	0.8182	0.8182
NOT-GLRA2 AND IFI6	Melanoma	0.8182	0.8182	0.8182
NOT-CACNG6 AND IFI6	Melanoma	0.8182	0.8182	0.8182
NOT-CLDN19 AND LRIG3	Melanoma	1	1	1

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
SLC52A2 AND NOT-ZP4	Esophagus	0.8333	1	0.7143
SLC52A2 AND NOT-GRM3	Esophagus	0.6667	1	0.5
SLC52A2 AND NOT-NRG3	Esophagus	0.8333	1	0.7143
SLC52A2 AND NOT-MEGF10	Esophagus	0.7826	1	0.6429
SLC52A2 AND NOT-TRPM1	Esophagus	0.7826	1	0.6429
TM4SF5 AND NOT-SLC30A10	Esophagus	0.7778	0.6364	1
SLC52A2 AND NOT-GRIN2A	Esophagus	0.7826	1	0.6429
SLC52A2 AND NOT-CD207	Esophagus	0.6667	1	0.5
SLC52A2 AND NOT-DRD1	Esophagus	0.6667	1	0.5
SLC52A2 AND NOT-SLC1A6	Esophagus	0.8333	1	0.7143
SLC52A2 AND NOT-CHRNB4	Esophagus	0.7273	1	0.5714
SLC52A2 AND NOT-GHR	Esophagus	0.7273	1	0.5714
SLC52A2 AND NOT-CD1A	Esophagus	0.7826	1	0.6429
SLC52A2 AND NOT-SELP	Esophagus	0.6667	1	0.5
SLC52A2 AND NOT-SLC2A2	Esophagus	0.6	1	0.4286
NRCAM AND NOT-CNTNAP2	Glioblastoma	0.875	0.9333	0.8235
PTPRZ1 AND NOT-CLDN10	Glioblastoma	0.8667	1	0.7647
PTPRZ1 AND NOT-KIAA0319	Glioblastoma	0.875	0.9333	0.8235
PTPRZ1 AND NOT-GRM3	Glioblastoma	0.9032	1	0.8235
PTPRZ1 AND NOT-GABRA2	Glioblastoma	0.8667	1	0.7647
PTPRZ1 AND NOT-SLC6A15	Glioblastoma	0.875	0.9333	0.8235
NRCAM AND NOT-GRM5	Glioblastoma	0.875	0.9333	0.8235
PTPRZ1 AND NOT-CDH22	Glioblastoma	0.9091	0.9375	0.8824
PTPRZ1 AND NOT-SLC1A6	Glioblastoma	0.9091	0.9375	0.8824
PTPRZ1 AND NOT-ATP8A2	Glioblastoma	0.875	0.9333	0.8235
PTPRZ1 AND NOT-CACNA1B	Glioblastoma	0.875	0.9333	0.8235
PTPRZ1 AND NOT-KCNJ6	Glioblastoma	0.9091	0.9375	0.8824
PTPRZ1 AND NOT-SCN8A	Glioblastoma	0.875	0.9333	0.8235
PTPRZ1 AND NOT-GPR85	Glioblastoma	0.8387	0.9286	0.7647
PTPRZ1 AND NOT-GABRG1	Glioblastoma	0.9032	1	0.8235
NRCAM AND NOT-SCN8A	Glioblastoma	0.8485	0.875	0.8235
PTPRZ1 AND NOT-SCN2A	Glioblastoma	0.9032	1	0.8235
NRCAM AND NOT-KIAA0319	Glioblastoma	0.8387	0.9286	0.7647
PTPRZ1 AND NOT-GPR83	Glioblastoma	0.9091	0.9375	0.8824
PTPRZ1 AND NOT-SLC32A1	Glioblastoma	0.8387	0.9286	0.7647
PTPRZ1 AND NOT-KCNA1	Glioblastoma	0.875	0.9333	0.8235
PTPRZ1 AND NOT-HCN1	Glioblastoma	0.875	0.9333	0.8235
PTPRZ1 AND NOT-CNTNAP2	Glioblastoma	0.875	0.9333	0.8235
PTPRZ1 AND NOT-CDH18	Glioblastoma	0.875	0.9333	0.8235
PTPRZ1 AND NOT-CALN1	Glioblastoma	0.8387	0.9286	0.7647
PTPRZ1 AND NOT-CDH8	Glioblastoma	0.875	0.9333	0.8235
PTPRZ1 AND NOT-GABRA1	Glioblastoma	0.8387	0.9286	0.7647
NRCAM AND NOT-ATP2B2	Glioblastoma	0.9091	0.9375	0.8824
PTPRZ1 AND NOT-SLC12A5	Glioblastoma	0.875	0.9333	0.8235
PTPRZ1 AND NOT-PTPRR	Glioblastoma	0.9091	0.9375	0.8824
PTPRZ1 AND NOT-GPR26	Glioblastoma	0.875	0.9333	0.8235
PTPRZ1 AND NOT-GRM5	Glioblastoma	0.875	0.9333	0.8235
PTPRZ1 AND NOT-ATP2B2	Glioblastoma	0.875	0.9333	0.8235
PTPRZ1 AND NOT-KCNC1	Glioblastoma	0.9091	0.9375	0.8824
PTPRZ1 AND NOT-DGKE	Glioblastoma	0.9091	0.9375	0.8824
PTPRZ1 AND NOT-CDH10	Glioblastoma	0.8387	0.9286	0.7647
PTPRZ1 AND NOT-GRM7	Glioblastoma	0.9091	0.9375	0.8824
PTPRZ1 AND NOT-GPR22	Glioblastoma	0.875	0.9333	0.8235
NRCAM AND NOT-SCN2A	Glioblastoma	0.8387	0.9286	0.7647
PTPRZ1 AND NOT-SYT4	Glioblastoma	0.875	0.9333	0.8235
PTPRZ1 AND NOT-OPCML	Glioblastoma	0.8	0.9231	0.7059
PTPRZ1 AND NOT-NRG3	Glioblastoma	0.8	0.9231	0.7059
PTPRZ1 AND NOT-GRM1	Glioblastoma	0.8	0.9231	0.7059
PTPRZ1 AND NOT-CACNA1E	Glioblastoma	0.9032	1	0.8235
PTPRZ1 AND NOT-ANO4	Glioblastoma	0.8667	1	0.7647
PTPRZ1 AND NOT-GABRA4	Glioblastoma	0.875	0.9333	0.8235
PTPRZ1 AND NOT-GHR	Glioblastoma	0.8	0.9231	0.7059
NRCAM AND NOT-GABRA4	Glioblastoma	0.8235	0.8235	0.8235
PTPRZ1 AND NOT-DPP10	Glioblastoma	0.8387	0.9286	0.7647
PTPRZ1 AND NOT-PIK3IP1	Glioblastoma	0.9091	0.9375	0.8824
PTPRZ1 AND NOT-CLSTN3	Glioblastoma	0.9091	0.9375	0.8824
PTPRZ1 AND NOT-CCKBR	Glioblastoma	0.9091	0.9375	0.8824
PTPRZ1 AND NOT-OR2L13	Glioblastoma	0.875	0.9333	0.8235
PTPRZ1 AND NOT-SLC6A12	Glioblastoma	0.875	0.9333	0.8235
PTPRZ1 AND NOT-PTPRT	Glioblastoma	0.875	0.9333	0.8235
PTPRZ1 AND NOT-SLC16A5	Glioma	0.9855	0.9855	0.9855
PTPRZ1 AND NOT-SLC6A15	Glioma	0.9781	0.9853	0.971
NRCAM AND NOT-KCNA2	Glioma	0.9385	1	0.8841
NRCAM AND NOT-KCNJ9	Glioma	0.9385	1	0.8841
PTPRZ1 AND NOT-NKAIN2	Glioma	0.9855	0.9855	0.9855
PTPRZ1 AND NOT-ABCG4	Glioma	0.9781	0.9853	0.971
PTPRZ1 AND NOT-TMPRSS11D	Glioma	0.9781	0.9853	0.971

FIG. 12 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
NOT-NMUR2 AND LRIG3	Melanoma	1	1	1
NOT-NOX1 AND VANGL1	Melanoma	0.9	1	0.8182
NOT-AMHR2 AND ICAM1	Melanoma	0.8182	0.8182	0.8182
NOT-ADAM29 AND GJA3	Melanoma	0.9	1	0.8182
NOT-NMUR2 AND CDH11	Melanoma	0.9524	1	0.9091
NOT-CLDN19 AND PLVAP	Melanoma	0.9524	1	0.9091
NOT-SLC6A13 AND VANGL1	Melanoma	0.9	1	0.8182
NOT-CHRNA2 AND SLC23A2	Melanoma	0.7778	1	0.6364
NOT-CLDN19 AND SLC22A18	Melanoma	0.7778	1	0.6364
NOT-GJD2 AND ICAM1	Melanoma	0.9	1	0.8182
NOT-NMUR2 AND SLC23A2	Melanoma	0.7778	1	0.6364
NOT-ADAM29 AND LRRC8E	Melanoma	0.9524	1	0.9091
NOT-CALHM1 AND SLC22A18	Melanoma	0.8182	0.8182	0.8182
NOT-HTR1E AND SLC23A2	Melanoma	0.7778	1	0.6364
NOT-ANO4 AND DCC	Ovarian	0.7826	0.8182	0.75
NOT-EMP3 AND GPRC6A	Ovarian	0.7619	0.8889	0.6667
NOT-KCNK12 AND GPRC6A	Ovarian	0.7	0.875	0.5833
NOT-GRID2 AND SLC4A5	Ovarian	0.64	0.6154	0.6667
NOT-KCNK12 AND DCC	Ovarian	0.6364	0.7	0.5833
NOT-EMP3 AND NMUR2	Ovarian	0.8	1	0.6667
NOT-KCNK12 AND KCNV1	Ovarian	0.6364	0.7	0.5833
NOT-KCNK12 AND LRRC8E	Ovarian	0.6667	0.7778	0.5833
NOT-EMP3 AND LRRC8E	Ovarian	0.8	1	0.6667
NOT-MTNR1B AND KCNU1	Ovarian	0.7619	0.8889	0.6667
NKAIN4 AND NOT-P2RX5	Astrocytoma	0.7952	0.8919	0.7174
NKAIN4 AND NOT-L1CAM	Astrocytoma	0.75	0.8824	0.6522
NKAIN4 AND NOT-SLC34A2	Astrocytoma	0.6667	0.8621	0.5435
CRB1 AND NOT-B4GALNT1	Astrocytoma	0.6606	0.5714	0.7826
EDNRB AND NOT-AOC3	Astrocytoma	0.6538	0.5862	0.7391
EDNRB AND NOT-PHLDB2	Astrocytoma	0.6531	0.6154	0.6957
CRB1 AND NOT-L1CAM	Astrocytoma	0.6316	0.8	0.5217
CRB1 AND NOT-EPCAM	Astrocytoma	0.623	0.5	0.8261
EDNRB AND NOT-CD36	Astrocytoma	0.6168	0.541	0.7174
NKAIN4 AND NOT-ITGB3	Astrocytoma	0.6111	0.8462	0.4783
BEST3 AND NOT-CD34	Astrocytoma	0.6098	0.6944	0.5435
CSPG5 AND NOT-P2RX5	Astrocytoma	0.6047	0.65	0.5652
CSPG5 AND NOT-EPCAM	Astrocytoma	0.6	0.6136	0.587
FAP AND NOT-SCARA5	Breast	0.8431	0.8958	0.7963
FAP AND NOT-EMP3	Breast	0.835	0.8776	0.7963
FAP AND NOT-MRGPRF	Breast	0.84	0.913	0.7778
FAP AND NOT-CLDN10	Breast	0.8081	0.8889	0.7407
FAP AND NOT-MLANA	Breast	0.8462	0.88	0.8148
FAP AND NOT-SLC39A2	Breast	0.8462	0.88	0.8148
FAP AND NOT-LRIG3	Breast	0.8155	0.8571	0.7778
FAP AND NOT-DSC1	Breast	0.8627	0.9167	0.8148
FAP AND NOT-CD207	Breast	0.8381	0.8627	0.8148
FAP AND NOT-CD1A	Breast	0.8269	0.86	0.7963
FAP AND NOT-EPHA2	Breast	0.7551	0.8409	0.6852
PPAPDC1A AND NOT-NCAM1	Breast	0.7551	0.8409	0.6852
FAP AND NOT-TMPRSS11E	Breast	0.82	0.8913	0.7593
FAP AND NOT-ANTXR2	Breast	0.7416	0.9429	0.6111
FAP AND NOT-SLC1A6	Breast	0.8119	0.8723	0.7593
FAP AND NOT-AOC3	Breast	0.8431	0.8958	0.7963
FAP AND NOT-PHLDB2	Breast	0.7391	0.8947	0.6296
FAP AND NOT-MSMO1	Breast	0.7292	0.8333	0.6481
FAP AND NOT-CDH22	Breast	0.8155	0.8571	0.7778
ENPP1 AND NOT-ALDH1A1	Breast	0.7551	0.8409	0.6852
FAP AND NOT-FXYD6	Breast	0.7917	0.9048	0.7037
FAP AND NOT-PTGIS	Breast	0.8283	0.9111	0.7593
FAP AND NOT-ADCY8	Breast	0.8462	0.88	0.8148
FAP AND NOT-SELP	Breast	0.7742	0.9231	0.6667
FAP AND NOT-SLC2A1	Breast	0.8235	0.875	0.7778
FAP AND NOT-TRPM1	Breast	0.8381	0.8627	0.8148
FAP AND NOT-ERVW-1	Breast	0.8269	0.86	0.7963
FAP AND NOT-PVRL1	Breast	0.8381	0.8627	0.8148
FAP AND NOT-MRAP	Breast	0.8039	0.8542	0.7593
FAP AND NOT-SLC23A2	Breast	0.8039	0.8542	0.7593
FAP AND NOT-GRIK4	Breast	0.8544	0.898	0.8148
FAP AND NOT-CD44	Breast	0.8381	0.8627	0.8148
FAP AND NOT-GABRA4	Breast	0.8462	0.88	0.8148
FAP AND NOT-TSPAN16	Breast	0.8431	0.8958	0.7963
FAP AND NOT-AGTR2	Breast	0.8627	0.9167	0.8148
FAP AND NOT-GJA3	Breast	0.8713	0.9362	0.8148
FAP AND NOT-DRD5	Breast	0.835	0.8776	0.7963
FAP AND NOT-PROCR	Breast	0.7045	0.9118	0.5741
FAP AND NOT-SLC6A11	Breast	0.8381	0.8627	0.8148
FAP AND NOT-CHRNG	Breast	0.835	0.8776	0.7963

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
PTPRZ1 AND NOT-KIAA0319	Glioma	0.9855	0.9855	0.9855
PTPRZ1 AND NOT-CD1A	Glioma	0.9927	1	0.9855
PTPRZ1 AND NOT-HCN1	Glioma	0.9781	0.9853	0.971
PTPRZ1 AND NOT-ATP8A2	Glioma	0.9855	0.9855	0.9855
PTPRZ1 AND NOT-SCN8A	Glioma	0.9781	0.9853	0.971
PTPRZ1 AND NOT-SMPD2	Glioma	0.9706	0.9851	0.9565
PTPRZ1 AND NOT-TMPRSS11E	Glioma	0.9855	0.9855	0.9855
NRCAM AND NOT-HTR1E	Glioma	0.9385	1	0.8841
NRCAM AND NOT-CDH8	Glioma	0.9313	0.9839	0.8841
NLGN1 AND NOT-CACNA1B	Glioma	0.9385	1	0.8841
NLGN1 AND NOT-GPR83	Glioma	0.9385	1	0.8841
PTPRZ1 AND NOT-GPR6	Glioma	0.9706	0.9851	0.9565
PTPRZ1 AND NOT-CDH8	Glioma	0.9855	0.9855	0.9855
PTPRZ1 AND NOT-DGKE	Glioma	0.9855	0.9855	0.9855
NLGN1 AND NOT-HCN1	Glioma	0.9302	1	0.8696
NRCAM AND NOT-SCN8A	Glioma	0.9302	1	0.8696
NLGN1 AND NOT-SCN8A	Glioma	0.9302	1	0.8696
PTPRZ1 AND NOT-CACNA1B	Glioma	0.9855	0.9855	0.9855
PTPRZ1 AND NOT-HTR5A	Glioma	0.9781	0.9853	0.971
NRCAM AND NOT-HCN1	Glioma	0.9313	0.9839	0.8841
NLGN1 AND NOT-GPR6	Glioma	0.9385	1	0.8841
NRCAM AND NOT-ATP2B2	Glioma	0.9313	0.9839	0.8841
NLGN1 AND NOT-GABRA2	Glioma	0.9385	1	0.8841
PTPRZ1 AND NOT-GRM3	Glioma	0.9927	1	0.9855
NLGN1 AND NOT-KCNA1	Glioma	0.9302	1	0.8696
PTPRZ1 AND NOT-SLC32A1	Glioma	0.9706	0.9851	0.9565
PTPRZ1 AND NOT-SLC26A8	Glioma	0.9855	0.9855	0.9855
PTPRZ1 AND NOT-CD207	Glioma	0.9855	0.9855	0.9855
PTPRZ1 AND NOT-GABRA2	Glioma	0.9855	0.9855	0.9855
NRCAM AND NOT-GABRB2	Glioma	0.9385	1	0.8841
SYT11 AND NOT-SYT2	Glioma	0.9853	1	0.971
PTPRZ1 AND NOT-GRM1	Glioma	0.9706	0.9851	0.9565
PTPRZ1 AND NOT-KCNK9	Glioma	0.9781	0.9853	0.971
NLGN1 AND NOT-KIAA0319	Glioma	0.9385	1	0.8841
PTPRZ1 AND NOT-SLC13A2	Glioma	0.9855	0.9855	0.9855
PTPRZ1 AND NOT-GPR83	Glioma	0.9781	0.9853	0.971
PTPRZ1 AND NOT-GGTLC1	Glioma	0.9927	1	0.9855
PTPRZ1 AND NOT-NPHS1	Glioma	0.963	0.9848	0.942
PTPRZ1 AND NOT-KCNA7	Glioma	0.9781	0.9853	0.971
PTPRZ1 AND NOT-MTNR1B	Glioma	0.9778	1	0.9565
PTPRZ1 AND NOT-ABCA12	Glioma	0.9781	0.9853	0.971
NRCAM AND NOT-CNTNAP2	Glioma	0.9302	1	0.8696
NLGN1 AND NOT-KCNK9	Glioma	0.9302	1	0.8696
PTPRZ1 AND NOT-EPHA10	Glioma	0.9855	0.9855	0.9855
NRCAM AND NOT-KCNV1	Glioma	0.9242	0.9683	0.8841
NRCAM AND NOT-NKAIN2	Glioma	0.9242	0.9683	0.8841
NRCAM AND NOT-KCNA1	Glioma	0.9242	0.9683	0.8841
NRCAM AND NOT-CCKBR	Glioma	0.9242	0.9683	0.8841
NRCAM AND NOT-SLC24A2	Glioma	0.9242	0.9683	0.8841
PTPRZ1 AND NOT-PVRL1	Glioma	0.9781	0.9853	0.971
PTPRZ1 AND NOT-HCN4	Glioma	0.9855	0.9855	0.9855
NLGN1 AND NOT-SLC32A1	Glioma	0.9302	1	0.8696
NRCAM AND NOT-TMEM235	Glioma	0.9242	0.9683	0.8841
NLGN1 AND NOT-DGKE	Glioma	0.9385	1	0.8841
PTPRZ1 AND NOT-MRAP	Glioma	0.9855	0.9855	0.9855
NLGN1 AND NOT-ATP8A2	Glioma	0.9302	1	0.8696
PTPRZ1 AND NOT-SLC22A9	Glioma	0.9778	1	0.9565
NLGN1 AND NOT-GPR26	Glioma	0.9385	1	0.8841
PTPRZ1 AND NOT-KCNJ6	Glioma	0.9855	0.9855	0.9855
NRCAM AND NOT-CACNA1E	Glioma	0.9231	0.9836	0.8696
NRCAM AND NOT-GPR26	Glioma	0.9231	0.9836	0.8696
NLGN1 AND NOT-GRM1	Glioma	0.9385	1	0.8841
PTPRZ1 AND NOT-KCNA1	Glioma	0.9781	0.9853	0.971
NRCAM AND NOT-CACNG8	Glioma	0.9242	0.9683	0.8841
NRCAM AND NOT-CLSTN3	Glioma	0.9313	0.9839	0.8841
NRCAM AND NOT-CACNG3	Glioma	0.9385	1	0.8841
NRCAM AND NOT-CACNA1B	Glioma	0.9385	1	0.8841
NRCAM AND NOT-GRM1	Glioma	0.9313	0.9839	0.8841
PTPRZ1 AND NOT-OR8D1	Glioma	0.9855	0.9855	0.9855
PTPRZ1 AND NOT-HTR1E	Glioma	0.9855	0.9855	0.9855
PTPRZ1 AND NOT-CACNA1S	Glioma	0.9781	0.9853	0.971
PTPRZ1 AND NOT-FGF6	Glioma	0.9855	0.9855	0.9855
PTPRZ1 AND NOT-GUCY2F	Glioma	0.9855	0.9855	0.9855
PTPRZ1 AND NOT-LRRC8E	Glioma	0.9706	0.9851	0.9565
PTPRZ1 AND NOT-CLSTN3	Glioma	0.9855	0.9855	0.9855
ATP8B2 AND NOT-FCER1A	Leiomyosarcoma	0.7826	0.9	0.6923
ATP8B2 AND NOT-ABCB1	Leiomyosarcoma	0.8333	0.9091	0.7692

FIG. 12 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
FAP AND NOT-CHRNB4	Breast	0.8544	0.898	0.8148
FAP AND NOT-SLC9B1	Breast	0.8381	0.8627	0.8148
FAP AND NOT-MMP24	Breast	0.8269	0.86	0.7963
FAP AND NOT-ESAM	Breast	0.8381	0.8627	0.8148
FAP AND NOT-ABCA12	Breast	0.7921	0.8511	0.7407
FAP AND NOT-ATP6V0A4	Breast	0.7677	0.8444	0.7037
FAP AND NOT-SLC30A1	Breast	0.7447	0.875	0.6481
FAP AND NOT-GRIN2A	Breast	0.8269	0.86	0.7963
FAP AND NOT-PTPRR	Breast	0.8544	0.898	0.8148
FAP AND NOT-MUC17	Breast	0.8381	0.8627	0.8148
FAP AND NOT-SLC22A11	Breast	0.8269	0.86	0.7963
FAP AND NOT-OR7C1	Breast	0.8269	0.86	0.7963
FAP AND NOT-GPR22	Breast	0.8627	0.9167	0.8148
FAP AND NOT-GYPC	Breast	0.8544	0.898	0.8148
FAP AND NOT-CLDN16	Breast	0.8155	0.8571	0.7778
FAP AND NOT-TSHR	Breast	0.8269	0.86	0.7963
FAP AND NOT-CCKAR	Breast	0.8381	0.8627	0.8148
FAP AND NOT-MPL	Breast	0.8381	0.8627	0.8148
FAP AND NOT-HCN1	Breast	0.8381	0.8627	0.8148
PPAPDC1A AND NOT-BCAN	Breast	0.6972	0.6909	0.7037
FAP AND NOT-CALN1	Breast	0.8381	0.8627	0.8148
FAP AND NOT-PCDHAC2	Breast	0.8544	0.898	0.8148
FAP AND NOT-NRG3	Breast	0.7755	0.8636	0.7037
FAP AND NOT-GALR2	Breast	0.8381	0.8627	0.8148
ABCG8 AND NOT-FOLH1	Liver	0.8	1	0.6667
FAT1 AND NOT-TPBG	Liver	0.8	1	0.6667
ALCAM AND NOT-TPBG	Liver	0.7692	0.7143	0.8333
CLDN15 AND NOT-TPBG	Liver	0.9091	1	0.8333
ALCAM AND NOT-IGF1R	Liver	0.9091	1	0.8333
ALCAM AND NOT-ERBB4	Liver	0.9091	1	0.8333
ABCG8 AND NOT-TPBG	Liver	0.8	1	0.6667
ABCG8 AND NOT-GPA33	Liver	0.8	1	0.6667
ABCG8 AND NOT-ENPP3	Liver	0.8	1	0.6667
ALCAM AND NOT-IL20RA	Liver	0.7143	0.625	0.8333
ABCG8 AND NOT-MUC13	Liver	0.8	1	0.6667
ABCG8 AND NOT-GUCY2C	Liver	0.8	1	0.6667
ABCG8 AND NOT-CLDN7	Liver	0.8	1	0.6667
ABCG8 AND NOT-MST1R	Liver	0.8	1	0.6667
ABCG8 AND NOT-SLC34A2	Liver	0.8	1	0.6667
ALCAM AND NOT-FOLR1	Liver	0.7143	0.625	0.8333
SLC30A10 AND NOT-CLDN7	Liver	0.6667	1	0.5
ABCG8 AND NOT-PROM1	Liver	0.6667	1	0.5
SLC30A10 AND NOT-GUCY2C	Liver	0.6667	1	0.5
SLC30A10 AND NOT-IL20RA	Liver	0.6667	1	0.5
ABCG8 AND NOT-FOLR1	Liver	0.8	1	0.6667
SLC30A10 AND NOT-MUC4	Liver	0.6667	1	0.5
SLC30A10 AND NOT-MUC13	Liver	0.6667	1	0.5
SLC30A10 AND NOT-ITGB6	Liver	0.6667	1	0.5
SLC30A10 AND NOT-TRPM4	Liver	0.6667	1	0.5
ABCG8 AND NOT-ITGB6	Liver	0.8	1	0.6667
SLC30A10 AND NOT-MST1R	Liver	0.6667	1	0.5
SLC30A10 AND NOT-GPA33	Liver	0.6667	1	0.5
SLC30A10 AND NOT-PROM1	Liver	0.6667	1	0.5
FAP AND NOT-PAQR7	Pancreas	0.8421	1	0.7273
FAP AND NOT-MEGF10	Pancreas	0.7059	1	0.5455
FAP AND NOT-ENPP1	Pancreas	0.8421	1	0.7273
FAP AND NOT-MRGPRF	Pancreas	0.7778	1	0.6364
FAP AND NOT-AOC3	Pancreas	0.8421	1	0.7273
FAP AND NOT-DGKE	Pancreas	0.8421	1	0.7273
FAP AND NOT-FXYD6	Pancreas	0.7778	1	0.6364
FAP AND NOT-SLC22A14	Pancreas	0.7778	1	0.6364
FAP AND NOT-CDH9	Pancreas	0.8421	1	0.7273
SLC30A8 AND NOT-CD70	Pancreas	0.7778	1	0.6364
FAP AND NOT-NFASC	Pancreas	0.7778	1	0.6364
FAP AND NOT-EMP3	Pancreas	0.7778	1	0.6364
FAP AND NOT-AGTR2	Pancreas	0.8421	1	0.7273
FAP AND NOT-ATP8B2	Pancreas	0.8421	1	0.7273
FAP AND NOT-ATP13A5	Pancreas	0.625	1	0.4545
SLC30A8 AND NOT-ERBB4	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-SSTR2	Pancreas	0.7778	1	0.6364
FAP AND NOT-SELP	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-L1CAM	Pancreas	0.7778	1	0.6364
SLC30A8 AND NOT-CLDN8	Pancreas	0.7778	1	0.6364
FAP AND NOT-GRM7	Pancreas	0.8421	1	0.7273
SLC30A8 AND NOT-SSTR4	Pancreas	0.7778	1	0.6364
FAP AND NOT-CACNG1	Pancreas	0.625	1	0.4545
FAP AND NOT-SLC22A16	Pancreas	0.8421	1	0.7273

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
ATP8B2 AND NOT-IL27RA	Leiomyosarcoma	0.8333	0.9091	0.7692
ATP8B2 AND NOT-MAL	Leiomyosarcoma	0.8333	0.9091	0.7692
FGFR1 AND NOT-SLC6A11	Leiomyosarcoma	0.7619	1	0.6154
ATP8B2 AND NOT-AQP3	Leiomyosarcoma	0.8333	0.9091	0.7692
ATP8B2 AND NOT-CD1C	Leiomyosarcoma	0.7826	0.9	0.6923
FGFR1 AND NOT-SLC5A2	Leiomyosarcoma	0.8182	1	0.6923
DDR2 AND NOT-ATP8A1	Leiomyosarcoma	0.7619	1	0.6154
FGFR1 AND NOT-SLC30A1	Leiomyosarcoma	0.7826	0.9	0.6923
FGFR1 AND NOT-KIAA0319	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-CD44	Leiomyosarcoma	0.7273	0.8889	0.6154
ATP8B2 AND NOT-CCR6	Leiomyosarcoma	0.8696	1	0.7692
FGFR1 AND NOT-CD207	Leiomyosarcoma	0.7619	1	0.6154
FGFR1 AND NOT-OR10J1	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-DSCAM	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-FGF6	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-RGSL1	Leiomyosarcoma	0.8182	1	0.6923
TGFB3 AND NOT-CLDN10	Leiomyosarcoma	0.6667	0.875	0.5385
ROR2 AND NOT-CLDN10	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-CLDN10	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-HCN2	Leiomyosarcoma	0.8182	1	0.6923
ATP8B2 AND NOT-PTGDR2	Leiomyosarcoma	0.6667	0.6429	0.6923
GPR161 AND NOT-CLDN10	Leiomyosarcoma	0.8182	1	0.6923
ATP8B2 AND NOT-MFAP3L	Leiomyosarcoma	0.6667	0.7273	0.6154
ATP8B2 AND NOT-P2RY14	Leiomyosarcoma	0.6667	0.875	0.5385
ATP8B2 AND NOT-SLC18A2	Leiomyosarcoma	0.7692	0.7692	0.7692
FGFR1 AND NOT-FSHR	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-GPR12	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-CHRNA4	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-ATP6V0A4	Leiomyosarcoma	0.8182	1	0.6923
ROR2 AND NOT-GGTLC1	Leiomyosarcoma	0.7619	1	0.6154
ATP8B2 AND NOT-IL12RB1	Leiomyosarcoma	0.7	1	0.5385
FGFR1 AND NOT-DSC1	Leiomyosarcoma	0.7619	1	0.6154
FGFR1 AND NOT-GPR37L1	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-SLC34A1	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-LRIG3	Leiomyosarcoma	0.6667	0.875	0.5385
ATP8B2 AND NOT-CD101	Leiomyosarcoma	0.7826	0.9	0.6923
EMP3 AND NOT-P2RY14	Leiomyosarcoma	0.6667	0.7273	0.6154
FGFR1 AND NOT-ATP13A5	Leiomyosarcoma	0.7619	1	0.6154
FGFR1 AND NOT-GPR6	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-HTR1D	Leiomyosarcoma	0.7619	1	0.6154
FGFR1 AND NOT-CLDN19	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-PROCR	Leiomyosarcoma	0.7826	0.9	0.6923
FGFR1 AND NOT-SLC23A2	Leiomyosarcoma	0.6667	0.875	0.5385
FGFR1 AND NOT-TMPRSS11E	Leiomyosarcoma	0.7619	1	0.6154
FGFR1 AND NOT-KCNK4	Leiomyosarcoma	0.8182	1	0.6923
ATP8B2 AND NOT-IL18R1	Leiomyosarcoma	0.6667	0.875	0.5385
FGFR1 AND NOT-HRH3	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-MLANA	Leiomyosarcoma	0.7826	0.9	0.6923
FGFR1 AND NOT-CD1A	Leiomyosarcoma	0.7826	0.9	0.6923
FGFR1 AND NOT-CACNG6	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-CACNG7	Leiomyosarcoma	0.8182	1	0.6923
ATP8B2 AND NOT-ATP2A3	Leiomyosarcoma	0.7826	0.9	0.6923
FGFR1 AND NOT-HCN4	Leiomyosarcoma	0.7619	1	0.6154
FGFR1 AND NOT-KCNH4	Leiomyosarcoma	0.7619	1	0.6154
FGFR1 AND NOT-ABCA12	Leiomyosarcoma	0.7826	0.9	0.6923
FGFR1 AND NOT-SLC13A2	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-PVRL1	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-CDH22	Leiomyosarcoma	0.8182	1	0.6923
ROR2 AND NOT-NKAIN1	Leiomyosarcoma	0.7619	1	0.6154
FGFR1 AND NOT-SELP	Leiomyosarcoma	0.7826	0.9	0.6923
ROR2 AND NOT-SELP	Leiomyosarcoma	0.75	0.8182	0.6923
FGFR1 AND NOT-LRRC8E	Leiomyosarcoma	0.7273	0.8889	0.6154
FGFR1 AND NOT-CHRNG	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-CD1B	Leiomyosarcoma	0.7	1	0.5385
TGFBI AND NOT-F11R	Leiomyosarcoma	0.75	0.8182	0.6923
FGFR1 AND NOT-SMPD2	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-KREMEN2	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-NPBWR1	Leiomyosarcoma	0.7	1	0.5385
DDR2 AND NOT-STEAP4	Leiomyosarcoma	0.8182	1	0.6923
PCDH18 AND NOT-SELP	Leiomyosarcoma	0.7273	0.8889	0.6154
FGFR1 AND NOT-MAG	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-AQP6	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-SLC22A11	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-NPHS2	Leiomyosarcoma	0.7	1	0.5385
FGFR1 AND NOT-ABCG5	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-CEACAM7	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-ADAM29	Leiomyosarcoma	0.8182	1	0.6923

FIG. 12 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
FAP AND NOT-ESAM	Pancreas	0.8421	1	0.7273
SLC22A2 AND NOT-CLDN8	Renal	0.7368	1	0.5833
SLC22A2 AND NOT-L1CAM	Renal	0.7	0.875	0.5833
CLDN2 AND NOT-MUC17	Renal	0.6957	0.7273	0.6667
VCAM1 AND NOT-SCARA5	Renal	0.8571	1	0.75
TNFRSF12A AND NOT-IL11RA	Renal	0.8	1	0.6667
VCAM1 AND NOT-MRGPRF	Renal	0.8571	1	0.75
SLC22A2 AND NOT-LGR5	Renal	0.6667	0.7778	0.5833
CLDN2 AND NOT-SLC13A5	Renal	0.6667	0.7778	0.5833
CLDN2 AND NOT-SLC22A25	Renal	0.6667	0.7778	0.5833
CLDN2 AND NOT-ENPP1	Renal	0.6667	0.7778	0.5833
CLDN2 AND NOT-DIO1	Renal	0.6957	0.7273	0.6667
GGTLC1 AND NOT-CLDN11	Renal	0.6667	0.7778	0.5833
CLDN2 AND NOT-SLCO1A2	Renal	0.7273	0.8	0.6667
CLDN2 AND NOT-DISP2	Renal	0.7273	0.8	0.6667
CLDN2 AND NOT-SLC16A5	Renal	0.6667	0.6667	0.6667
VCAM1 AND NOT-SLC43A1	Renal	0.7368	1	0.5833
CLDN2 AND NOT-TMPRSS11E	Renal	0.7	0.875	0.5833
VCAM1 AND NOT-SLC16A5	Renal	0.7368	1	0.5833
SLC22A2 AND NOT-SSTR4	Renal	0.6364	0.7	0.5833
CLDN2 AND NOT-CALHM1	Renal	0.6364	0.7	0.5833
VCAM1 AND NOT-KCNK6	Renal	0.8571	1	0.75
SLC6A13 AND NOT-CLDN8	Renal	0.6316	0.8571	0.5
SLC17A1 AND NOT-CLDN11	Renal	0.6316	0.8571	0.5
SLC6A13 AND NOT-L1CAM	Renal	0.6316	0.8571	0.5
VCAM1 AND NOT-FMNL1	Renal	0.7368	1	0.5833
GGTLC1 AND NOT-IL11RA	Renal	0.6207	0.5294	0.75
CLDN2 AND NOT-LRRC52	Renal	0.8	1	0.6667
CLDN2 AND NOT-GHRHR	Renal	0.7368	1	0.5833
CLDN2 AND NOT-CCR9	Renal	0.6154	0.5714	0.6667
CLDN2 AND NOT-SLC5A11	Renal	0.6154	0.5714	0.6667
CLDN2 AND NOT-KCNJ10	Renal	0.6154	0.5714	0.6667
CLDN2 AND NOT-PTPRT	Renal	0.6667	0.7778	0.5833
CLDN2 AND NOT-GABRA4	Renal	0.6667	1	0.5
CLDN2 AND NOT-GABRA2	Renal	0.6087	0.6364	0.5833
CLDN2 AND NOT-GJB4	Renal	0.6364	0.7	0.5833
VCAM1 AND NOT-PTGER4	Renal	0.8	1	0.6667
CLDN2 AND NOT-LAMP5	Renal	0.6667	0.6667	0.6667
SLC17A1 AND NOT-LGR5	Renal	0.6	0.75	0.5
CLDN2 AND NOT-USH2A	Renal	0.6	0.75	0.5
VCAM1 AND NOT-SLC6A7	Renal	0.7368	1	0.5833
SLC6A13 AND NOT-LGR5	Renal	0.6	0.75	0.5
SLC13A1 AND NOT-CR2	Renal	0.6	0.75	0.5
SLC6A13 AND NOT-CR2	Renal	0.6	0.75	0.5
CLDN2 AND NOT-CNTNAP2	Renal	0.6	0.75	0.5
SLC13A1 AND NOT-LGR5	Renal	0.6	0.75	0.5
CLDN2 AND NOT-OPALIN	Renal	0.6364	0.7	0.5833
VCAM1 AND NOT-LAMP5	Renal	0.8	1	0.6667
TNFRSF12A AND NOT-CLDN11	Renal	0.6087	0.6364	0.5833
CLDN2 AND NOT-CSMD3	Renal	0.6667	1	0.5
CLDN2 AND NOT-KCNK10	Renal	0.6154	0.5714	0.6667
CLDN2 AND NOT-SLC12A3	Renal	0.6087	0.6364	0.5833
CLDN2 AND NOT-TRPC3	Renal	0.6364	0.7	0.5833
DPEP1 AND NOT-IL13	Colon	1	1	1
DPEP1 AND NOT-SLC28A1	Colon	1	1	1
DPEP1 AND NOT-SLC13A1	Colon	1	1	1
DPEP1 AND NOT-CD1B	Colon	1	1	1
DPEP1 AND NOT-GPR26	Colon	1	1	1
DPEP1 AND NOT-EPHA8	Colon	1	1	1
DPEP1 AND NOT-ANPEP	Colon	1	1	1
DPEP1 AND NOT-SPAM1	Colon	1	1	1
DPEP1 AND NOT-ENPP1	Colon	1	1	1
DPEP1 AND NOT-OXTR	Colon	1	1	1
DPEP1 AND NOT-HTR3C	Colon	1	1	1
GUCY2C AND NOT-SLC9A1	Colon	1	1	1
DPEP1 AND NOT-KCNV2	Colon	1	1	1
CLDN2 AND NOT-SLC28A1	Colon	0.8889	1	0.8
DPEP1 AND NOT-FXYD7	Colon	0.8889	1	0.8
DPEP1 AND NOT-CALHM1	Colon	0.8889	1	0.8
NOX1 AND NOT-GPA33	Colon	0.8889	1	0.8
EPHB2 AND NOT-EPHA10	Colon	0.8889	1	0.8
CLDN2 AND NOT-SLC13A1	Colon	0.8889	1	0.8
EPHB2 AND NOT-SLC39A8	Colon	0.8889	1	0.8
EPHB2 AND NOT-SLC9A1	Colon	0.8889	1	0.8
DPEP1 AND NOT-PCDH11Y	Colon	0.8889	1	0.8
CLDN2 AND NOT-SLC2A2	Colon	0.8889	1	0.8
DPEP1 AND NOT-OR1C1	Colon	1	1	1

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
FGFR1 AND NOT-CLDN16	Leiomyosarcoma	0.7619	1	0.6154
FGFR1 AND NOT-MRGPRF	Leiomyosarcoma	0.7273	0.8889	0.6154
FGFR1 AND NOT-PIK3IP1	Leiomyosarcoma	0.7273	0.8889	0.6154
FGFR1 AND NOT-GABRA2	Leiomyosarcoma	0.7619	1	0.6154
FGFR1 AND NOT-MLNR	Leiomyosarcoma	0.7826	0.9	0.6923
FGFR1 AND NOT-MLC1	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-SLC22A18	Leiomyosarcoma	0.7826	0.9	0.6923
FGFR1 AND NOT-WNT7A	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-OR3A1	Leiomyosarcoma	0.7	1	0.5385
FGFR1 AND NOT-SLC13A5	Leiomyosarcoma	0.8182	1	0.6923
FGFR1 AND NOT-SLC39A2	Leiomyosarcoma	0.8182	1	0.6923
FLT3 AND NOT-BTN3A1	AML	0.9526	0.9872	0.9203
FLT3 AND NOT-S1PR1	AML	0.9524	0.9914	0.9163
FLT3 AND NOT-ATP8B1	AML	0.9506	0.983	0.9203
FLT3 AND NOT-GHR	AML	0.9506	0.983	0.9203
FLT3 AND NOT-BTN3A3	AML	0.9504	0.9871	0.9163
FLT3 AND NOT-IFNAR2	AML	0.9485	0.9829	0.9163
FLT3 AND NOT-PHLDB2	AML	0.9482	0.9871	0.9124
FLT3 AND NOT-EFNB2	AML	0.948	0.9913	0.9084
FLT3 AND NOT-CSPG5	AML	0.9467	0.9747	0.9203
FLT3 AND NOT-GPR19	AML	0.9467	0.9747	0.9203
FLT3 AND NOT-GRM5	AML	0.9467	0.9747	0.9203
FLT3 AND NOT-SLC31A1	AML	0.9465	0.9787	0.9163
FLT3 AND NOT-SLC22A5	AML	0.9448	0.9706	0.9203
FLT3 AND NOT-KCNK6	AML	0.9448	0.9706	0.9203
FLT3 AND NOT-SLC8A3	AML	0.9448	0.9706	0.9203
FLT3 AND NOT-SEMA4D	AML	0.9448	0.9706	0.9203
FLT3 AND NOT-NKAIN2	AML	0.9446	0.9746	0.9163
FLT3 AND NOT-PROCR	AML	0.9441	0.9828	0.9084
FLT3 AND NOT-FXYD6	AML	0.9436	0.9912	0.9004
FLT3 AND NOT-CD93	AML	0.9429	0.9665	0.9203
FLT3 AND NOT-SLC9A1	AML	0.9429	0.9665	0.9203
FLT3 AND NOT-GPR37L1	AML	0.9429	0.9665	0.9203
FLT3 AND NOT-KCNV1	AML	0.9429	0.9665	0.9203
FLT3 AND NOT-ZACN	AML	0.9429	0.9665	0.9203
FLT3 AND NOT-SLC5A2	AML	0.9429	0.9665	0.9203
FLT3 AND NOT-PAQR7	AML	0.9429	0.9665	0.9203
FLT3 AND NOT-ADAM29	AML	0.9429	0.9665	0.9203
FLT3 AND NOT-CALN1	AML	0.9424	0.9745	0.9124
SLC22A16 AND NOT-PERP	AML	0.9412	0.9587	0.9243
FLT3 AND NOT-SLC43A1	AML	0.9407	0.9664	0.9163
FLT3 AND NOT-SELP	AML	0.9407	0.9664	0.9163
FLT3 AND NOT-SMPD2	AML	0.9407	0.9664	0.9163
FLT3 AND NOT-SLC2A1	AML	0.9407	0.9664	0.9163
FLT3 AND NOT-SLC38A1	AML	0.9407	0.9664	0.9163
FLT3 AND NOT-IL15RA	AML	0.9405	0.9703	0.9124
FLT3 AND NOT-LAT	AML	0.9402	0.9744	0.9084
FLT3 AND NOT-PAQR8	AML	0.9402	0.9744	0.9084
SLC22A16 AND NOT-PROS1	AML	0.9416	0.9512	0.9323
SLC22A16 AND NOT-PODXL	AML	0.9398	0.9474	0.9323
FLT3 AND NOT-SLC16A5	AML	0.9363	0.9661	0.9084
SLC22A16 AND NOT-TMEM47	AML	0.936	0.9398	0.9323
FLT3 AND NOT-SYT11	AML	0.9347	0.9911	0.8845
SLC22A16 AND NOT-TUSC3	AML	0.9341	0.936	0.9323
SLC22A16 AND NOT-TRPC1	AML	0.9352	0.9506	0.9203
FLT3 AND NOT-ZDHHC2	AML	0.932	0.9658	0.9004
P2RX1 AND NOT-PHLDB2	AML	0.9377	0.9163	0.9602
FLT3 AND NOT-FLVCR1	AML	0.9314	0.9739	0.8924
SLC22A16 AND NOT-ANTXR1	AML	0.9304	0.9286	0.9323
SLC22A16 AND NOT-NTN4	AML	0.9304	0.9286	0.9323
SLC22A16 AND NOT-SLC7A2	AML	0.9301	0.932	0.9283
SLC22A16 AND NOT-LIFR	AML	0.9301	0.932	0.9283
SLC22A16 AND NOT-LAMP3	AML	0.9286	0.9249	0.9323
SLC22A16 AND NOT-EMP2	AML	0.9286	0.9249	0.9323
SLC22A16 AND NOT-LPAR1	AML	0.9339	0.9395	0.9283
CSF3R AND NOT-PHLDB2	AML	0.9283	0.9283	0.9283
SLC22A16 AND NOT-MTUS1	AML	0.9283	0.9283	0.9283
SLC22A16 AND NOT-ATP9A	AML	0.9393	0.9547	0.9243
SLC22A16 AND NOT-OSMR	AML	0.9304	0.9286	0.9323
SLC22A16 AND NOT-ENPP5	AML	0.9264	0.9246	0.9283
SLC22A16 AND NOT-SEMA5A	AML	0.9264	0.9246	0.9283
SLC22A16 AND NOT-TSPAN6	AML	0.9264	0.9246	0.9283
SLC22A16 AND NOT-TMEM150C	AML	0.9249	0.9176	0.9323
SLC22A16 AND NOT-SLC25A4	AML	0.9249	0.9176	0.9323
SLC22A16 AND NOT-BMPR1A	AML	0.932	0.9357	0.9283
SLC22A16 AND NOT-SSPN	AML	0.9246	0.9458	0.9044
SLC22A16 AND NOT-ACVR2A	AML	0.924	0.9277	0.9203

FIG. 12 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
DPEP1 AND NOT-DSC1	Colon	1	1	1
DPEP1 AND NOT-CHRND	Colon	1	1	1
DPEP1 AND NOT-CSMD3	Colon	1	1	1
DPEP1 AND NOT-FGF6	Colon	0.8889	1	0.8
DPEP1 AND NOT-LRRC52	Colon	1	1	1
DPEP1 AND NOT-OR1Q1	Colon	1	1	1
CLDN2 AND NOT-KCNV2	Colon	0.8889	1	0.8
DPEP1 AND NOT-KCNK10	Colon	1	1	1
CLDN2 AND NOT-ENPP1	Colon	0.8889	1	0.8
GPA33 AND NOT-SLC9A1	Colon	1	1	1
NOX1 AND NOT-MUC4	Colon	0.8889	1	0.8
DPEP1 AND NOT-GPR22	Colon	1	1	1
DPEP1 AND NOT-ABCG8	Colon	0.8889	1	0.8
DPEP1 AND NOT-ABCA12	Colon	0.8889	1	0.8
DPEP1 AND NOT-SLC10A2	Colon	1	1	1
DPEP1 AND NOT-SLC22A8	Colon	0.8889	1	0.8
DPEP1 AND NOT-CDH8	Colon	1	1	1
MUC13 AND NOT-SLC9A1	Colon	1	1	1
DPEP1 AND NOT-SLC5A11	Colon	0.8889	1	0.8
DPEP1 AND NOT-KCNF1	Colon	0.8889	1	0.8
DPEP1 AND NOT-TRPC7	Colon	1	1	1
DPEP1 AND NOT-UMOD	Colon	1	1	1
DPEP1 AND NOT-SLC39A12	Colon	1	1	1
DPEP1 AND NOT-GJB4	Colon	1	1	1
DPEP1 AND NOT-SLCO6A1	Colon	1	1	1
DPEP1 AND NOT-TMPRSS11B	Colon	1	1	1
DPEP1 AND NOT-OTOF	Colon	1	1	1
DPEP1 AND NOT-CLDN20	Colon	0.8333	0.7143	1
DPEP1 AND NOT-SLC1A6	Colon	1	1	1
DPEP1 AND NOT-PIRT	Colon	1	1	1
DPEP1 AND NOT-OR2C3	Colon	0.9091	0.8333	1
EPHB2 AND NOT-CHRND	Colon	0.8889	1	0.8
CLDN2 AND NOT-GPR26	Colon	0.8889	1	0.8
CLDN2 AND NOT-SPAM1	Colon	0.8889	1	0.8
CLDN2 AND NOT-GRM4	Colon	0.8889	1	0.8
GUCY2C AND NOT-SLC30A10	Colon	0.8889	1	0.8
DPEP1 AND NOT-SLCO1C1	Colon	0.8889	1	0.8
DPEP1 AND NOT-CALHM3	Colon	0.8889	1	0.8
GRIN2A AND NOT-STEAP2	Ependymoma	1	1	1
SSTR1 AND NOT-EBP	Ependymoma	1	1	1
SSTR1 AND NOT-SLC6A6	Ependymoma	1	1	1
GRIN2A AND NOT-FOLH1	Ependymoma	1	1	1
GRIN2A AND NOT-ERBB4	Ependymoma	1	1	1
CRB2 AND NOT-EPCAM	Ependymoma	0.6667	0.5	1
GRIN2A AND NOT-B4GALNT1	Ependymoma	0.6667	0.5	1
SSTR1 AND NOT-PTPRR	Ependymoma	1	1	1
SSTR1 AND NOT-SLC31A1	Ependymoma	1	1	1
TNFRSF10B AND NOT-TNFRSF10A	Ependymoma	0.6667	0.5	1
CDH17 AND NOT-ENPP3	Esophagus	0.625	0.5556	0.7143
CBX3 AND NOT-CD36	Esophagus	0.64	0.7273	0.5714
SLCO1B3 AND NOT-IL11RA	Esophagus	0.6	1	0.4286
MUC17 AND NOT-HHLA2	Esophagus	0.6	1	0.4286
SLCO1B3 AND NOT-AFP	Esophagus	0.6	1	0.4286
CBX3 AND NOT-ATP8B2	Esophagus	0.6957	0.8889	0.5714
MUC17 AND NOT-ENPP3	Esophagus	0.6667	1	0.5
EPHB2 AND NOT-BEST2	Esophagus	0.64	0.7273	0.5714
EPHB2 AND NOT-SLC22A5	Esophagus	0.625	0.5556	0.7143
CXCL16 AND NOT-CD33	Esophagus	0.64	0.7273	0.5714
MUC17 AND NOT-CD160	Esophagus	0.7826	1	0.6429
LYPD1 AND NOT-TPBG	Glioblastoma	0.6842	0.619	0.7647
LYPD1 AND NOT-FOLH1	Glioblastoma	0.6667	0.6316	0.7059
LYPD1 AND NOT-EPCAM	Glioblastoma	0.6667	0.7692	0.5882
LYPD1 AND NOT-STEAP2	Glioblastoma	0.6667	0.5909	0.7647
LYPD1 AND NOT-CLDN11	Glioblastoma	0.6667	0.56	0.8235
LYPD1 AND NOT-RNF43	Glioblastoma	0.6667	0.56	0.8235
LYPD1 AND NOT-IL20RA	Glioblastoma	0.6667	0.56	0.8235
LYPD1 AND NOT-TRPM4	Glioblastoma	0.6667	0.56	0.8235
LYPD1 AND NOT-CD22	Glioblastoma	0.6667	0.56	0.8235
LYPD1 AND NOT-P2RX5	Glioblastoma	0.6061	0.625	0.5882
LYPD1 AND NOT-ERBB3	Glioblastoma	0.6667	0.7692	0.5882
SYT11 AND NOT-EPCAM	Glioma	0.9714	0.9577	0.9855
SYT11 AND NOT-P2RX5	Glioma	0.9571	0.9437	0.971
SYT11 AND NOT-B4GALNT1	Glioma	0.9496	0.9429	0.9565
SYT11 AND NOT-CLDN6	Glioma	0.964	0.9571	0.971
SYT11 AND NOT-STEAP2	Glioma	0.9645	0.9444	0.9855
SYT11 AND NOT-CD22	Glioma	0.964	0.9571	0.971
SYT11 AND NOT-CLDN11	Glioma	0.9496	0.9429	0.9565

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
SLC22A16 AND NOT-SGCB	AML	0.924	0.9277	0.9203
SLC22A16 AND NOT-DAG1	AML	0.9339	0.9395	0.9283
SLC22A16 AND NOT-WNT5A	AML	0.9231	0.9141	0.9323
SLC22A16 AND NOT-SLC2A12	AML	0.9286	0.9249	0.9323
SLC22A16 AND NOT-SGCE	AML	0.9317	0.9393	0.9243
SLC22A16 AND NOT-ITGB5	AML	0.9215	0.9309	0.9124
SLC22A16 AND NOT-PTPRM	AML	0.9209	0.9137	0.9283
SLC22A16 AND NOT-DCBLD2	AML	0.9286	0.9249	0.9323
SLC22A16 AND NOT-KITLG	AML	0.9203	0.9203	0.9203
SLC22A16 AND NOT-ABHD6	AML	0.9194	0.907	0.9323
SLC22A16 AND NOT-AMIGO2	AML	0.9182	0.92	0.9163
ADAM12 AND NOT-SLC2A1	Liposarcoma	0.7407	0.6667	0.8333
ADAM12 AND NOT-SELP	Liposarcoma	0.7059	0.6122	0.8333
ADAM12 AND NOT-ERVW-1	Liposarcoma	0.7381	0.6458	0.8611
ADAM12 AND NOT-SLC1A2	Liposarcoma	0.6667	0.5439	0.8611
ADAM12 AND NOT-MSMO1	Liposarcoma	0.6444	0.537	0.8056
FAT4 AND NOT-NFASC	Liposarcoma	0.6667	0.9048	0.5278
SLC2A10 AND NOT-SORL1	Liposarcoma	0.6364	0.7	0.5833
ADAM12 AND NOT-ADAM29	Liposarcoma	0.6316	0.5085	0.8333
CNTNAP1 AND NOT-NFASC	Liposarcoma	0.6316	0.8571	0.5
FAT4 AND NOT-SORL1	Liposarcoma	0.6774	0.8077	0.5833
EMP3 AND NOT-CXCR2	Liposarcoma	0.6286	0.6471	0.6111
TTYH3 AND NOT-CD46	Liposarcoma	0.6197	0.6286	0.6111
TTYH3 AND NOT-KCNA3	Liposarcoma	0.6197	0.6286	0.6111
CNTNAP1 AND NOT-OMG	Liposarcoma	0.6567	0.7097	0.6111
SLC50A1 AND NOT-ABCB1	Lung Adenocarcinoma	0.6491	0.8409	0.5286
SLC50A1 AND NOT-VIPR1	Lung Adenocarcinoma	0.6207	0.7826	0.5143
OSMR AND NOT-GHR	Lung Adenocarcinoma	0.6154	0.9412	0.4571
SLC50A1 AND NOT-ZFYVE27	Lung Adenocarcinoma	0.6116	0.7255	0.5286
PON2 AND NOT-SLC30A10	Lung Adenocarcinoma	0.614	0.7955	0.5
ADAM12 AND NOT-ENPP1	Lung Carcinoma	0.6265	0.5098	0.8125
KCNJ10 AND NOT-MFAP3L	B-Cell Diffuse	0.8451	0.8824	0.8108
KCNJ10 AND NOT-SLC4A4	B-Cell Diffuse	0.8056	0.8286	0.7838
KCNJ10 AND NOT-ATP1B2	B-Cell Diffuse	0.8108	0.8108	0.8108
KCNJ10 AND NOT-LRP4	B-Cell Diffuse	0.8286	0.8788	0.7838
CXCL9 AND NOT-PROCR	B-Cell Diffuse	0.7869	1	0.6486
KCNJ10 AND NOT-NTRK2	B-Cell Diffuse	0.7895	0.7692	0.8108
KCNJ10 AND NOT-PLP1	B-Cell Diffuse	0.8	0.7895	0.8108
KCNJ10 AND NOT-NDRG4	B-Cell Diffuse	0.7692	0.8929	0.6757
KCNJ10 AND NOT-SEMA6D	B-Cell Diffuse	0.7671	0.7778	0.7568
CXCL9 AND NOT-GJA3	B-Cell Diffuse	0.7647	0.8387	0.7027
KCNJ10 AND NOT-SLC6A8	B-Cell Diffuse	0.7761	0.8667	0.7027
KCNJ10 AND NOT-DPP6	B-Cell Diffuse	0.8333	0.8571	0.8108
CXCL9 AND NOT-PCDH11Y	B-Cell Diffuse	0.7576	0.8621	0.6757
KCNJ10 AND NOT-TYRO3	B-Cell Diffuse	0.7536	0.8125	0.7027
CXCL9 AND NOT-SLC22A16	B-Cell Diffuse	0.7536	0.8125	0.7027
KCNJ10 AND NOT-PTPRD	B-Cell Diffuse	0.75	0.7714	0.7297
KCNJ10 AND NOT-CNTN1	B-Cell Diffuse	0.75	0.7714	0.7297
KCNJ10 AND NOT-ADCYAP1R1	B-Cell Diffuse	0.75	0.7714	0.7297
MCOLN2 AND NOT-SLC30A10	B-Cell Diffuse	0.7826	0.8438	0.7297
KCNJ10 AND NOT-FGFR3	B-Cell Diffuse	0.7436	0.7073	0.7838
CXCL9 AND NOT-TMPRSS11E	B-Cell Diffuse	0.7429	0.7879	0.7027
CD80 AND NOT-GJA3	B-Cell Diffuse	0.7429	0.7879	0.7027
KCNJ10 AND NOT-PCDH19	B-Cell Diffuse	0.7407	0.6818	0.8108
KCNJ10 AND NOT-NEO1	B-Cell Diffuse	0.7397	0.75	0.7297
TNFRSF9 AND NOT-PCDH11Y	B-Cell Diffuse	0.7419	0.92	0.6216
KCNJ10 AND NOT-BACE1	B-Cell Diffuse	0.7353	0.8065	0.6757
CXCL9 AND NOT-SLC39A2	B-Cell Diffuse	0.7324	0.7647	0.7027
CXCL9 AND NOT-CLDN10	B-Cell Diffuse	0.7324	0.7647	0.7027
CXCL9 AND NOT-DSC1	B-Cell Diffuse	0.7324	0.7647	0.7027
CXCL9 AND NOT-AMHR2	B-Cell Diffuse	0.7324	0.7647	0.7027
CXCL9 AND NOT-GHR	B-Cell Diffuse	0.7273	0.8276	0.6486
CXCL9 AND NOT-STEAP4	B-Cell Diffuse	0.7273	0.8276	0.6486
CD80 AND NOT-KCNC2	B-Cell Diffuse	0.7246	0.7813	0.6757
KCNJ10 AND NOT-TSPAN7	B-Cell Diffuse	0.7246	0.7813	0.6757
CXCL9 AND NOT-C8B	B-Cell Diffuse	0.7246	0.7813	0.6757
CXCL9 AND NOT-GPR37L1	B-Cell Diffuse	0.7222	0.7429	0.7027
CXCL9 AND NOT-EPHA8	B-Cell Diffuse	0.7222	0.7429	0.7027
CXCL9 AND NOT-CACNA1S	B-Cell Diffuse	0.7222	0.7429	0.7027
CXCL9 AND NOT-NMUR2	B-Cell Diffuse	0.7222	0.7429	0.7027
CXCL9 AND NOT-VN1R2	B-Cell Diffuse	0.7222	0.7429	0.7027
CXCL9 AND NOT-GPR26	B-Cell Diffuse	0.7222	0.7429	0.7027
CXCL9 AND NOT-GABRA2	B-Cell Diffuse	0.7222	0.7429	0.7027
CXCL9 AND NOT-KCNF1	B-Cell Diffuse	0.7222	0.7429	0.7027
CXCL9 AND NOT-CASR	B-Cell Diffuse	0.7222	0.7429	0.7027
CXCL9 AND NOT-SLC10A1	B-Cell Diffuse	0.7222	0.7429	0.7027
CXCL9 AND NOT-MLANA	B-Cell Diffuse	0.7222	0.7429	0.7027

FIG. 12 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
SYT11 AND NOT-CLDN7	Glioma	0.964	0.9571	0.971
SYT11 AND NOT-FOLH1	Glioma	0.9496	0.9429	0.9565
SYT11 AND NOT-ITGB3	Glioma	0.9481	0.9697	0.9275
LYPD1 AND NOT-EPCAM	Glioma	0.9048	1	0.8261
ITGAV AND NOT-ATP8B1	Glioma	0.9197	0.9265	0.913
ITGAV AND NOT-SLC16A5	Glioma	0.9343	0.9412	0.9275
ITGAV AND NOT-STEAP4	Glioma	0.9323	0.9688	0.8986
SYT11 AND NOT-ULBP2	Glioma	0.8504	0.931	0.7826
SLC39A6 AND NOT-ATP8B1	Glioma	0.8462	0.9016	0.7971
ATP8B2 AND NOT-CD79B	Leiomyosarcoma	0.8333	0.9091	0.7692
ATP8B2 AND NOT-TNFRSF10A	Leiomyosarcoma	0.7826	0.9	0.6923
FAP AND NOT-CLDN10	Leiomyosarcoma	0.8182	1	0.6923
CD276 AND NOT-SLC16A5	Leiomyosarcoma	0.6	0.8571	0.4615
FAP AND NOT-SELP	Leiomyosarcoma	0.7619	1	0.6154
CD276 AND NOT-SMPD2	Leiomyosarcoma	0.6957	0.8	0.6154
FAP AND NOT-PAQR7	Leiomyosarcoma	0.6316	1	0.4615
ATP8B2 AND NOT-CD79A	Leiomyosarcoma	0.8333	0.9091	0.7692
FAP AND NOT-GRIN1	Leiomyosarcoma	0.8182	1	0.6923
SLC22A16 AND NOT-STEAP2	AML	0.9231	0.9141	0.9323
SLC22A16 AND NOT-MET	AML	0.9225	0.9206	0.9243
SLC22A16 AND NOT-BMPR1B	AML	0.9191	0.9102	0.9283
SLC22A16 AND NOT-ERBB3	AML	0.9141	0.8966	0.9323
SLC22A16 AND NOT-ERBB2	AML	0.9141	0.8966	0.9323
SLC22A16 AND NOT-GPNMB	AML	0.9136	0.9447	0.8845
SLC22A16 AND NOT-STEAP1	AML	0.913	0.9059	0.9203
SLC22A16 AND NOT-CLDN23	AML	0.9155	0.9031	0.9283
SLC22A16 AND NOT-CLDN12	AML	0.9128	0.9298	0.8964
SLC22A16 AND NOT-CLDN1	AML	0.9105	0.8897	0.9323
SLC22A16 AND NOT-RNF43	AML	0.9105	0.8897	0.9323
SLC22A16 AND NOT-ERBB4	AML	0.9231	0.9141	0.9323
SLC22A16 AND NOT-SLC34A2	AML	0.9066	0.8859	0.9283
SLC22A16 AND NOT-ITGAV	AML	0.9057	0.9325	0.8805
CD37 AND NOT-PHLDB2	AML	0.9049	0.8826	0.9283
SLC22A16 AND NOT-ENPP3	AML	0.9035	0.8764	0.9323
SLC22A16 AND NOT-MUC16	AML	0.9035	0.8764	0.9323
SLC22A16 AND NOT-IL20RA	AML	0.9017	0.8731	0.9323
SLC22A16 AND NOT-VTCN1	AML	0.9017	0.8731	0.9323
SLC22A16 AND NOT-CLDN8	AML	0.9	0.8699	0.9323
SLC22A16 AND NOT-SDC1	AML	0.9	0.8699	0.9323
SLC22A16 AND NOT-MUC13	AML	0.9052	0.8797	0.9323
SLC22A16 AND NOT-AXL	AML	0.9256	0.935	0.9163
SLC22A16 AND NOT-FOLH1	AML	0.8971	0.875	0.9203
SLC22A16 AND NOT-FOLR1	AML	0.8966	0.8635	0.9323
SLC22A16 AND NOT-MUC4	AML	0.8966	0.8635	0.9323
SLC22A16 AND NOT-CLDN7	AML	0.8966	0.8635	0.9323
SLC22A16 AND NOT-EPCAM	AML	0.8923	0.8625	0.9243
CD37 AND NOT-BTN3A3	AML	0.8919	0.8652	0.9203
CD37 AND NOT-BTN3A1	AML	0.8835	0.8363	0.9363
SLC22A16 AND NOT-MUC1	AML	0.8811	0.8626	0.9004
CD37 AND NOT-S1PR1	AML	0.8672	0.8506	0.8845
CD37 AND NOT-LAT	AML	0.8582	0.7979	0.9283
CD37 AND NOT-GPR137B	AML	0.8537	0.8714	0.8367
SLC22A16 AND NOT-TPBG	AML	0.8525	0.8776	0.8287
CD37 AND NOT-STEAP4	AML	0.8514	0.7891	0.9243
CD37 AND NOT-IFNAR2	AML	0.8417	0.8165	0.8685
CD37 AND NOT-TGFBI	AML	0.835	0.8144	0.8566
CD37 AND NOT-SEMA4D	AML	0.8269	0.7429	0.9323
FMNL1 AND NOT-FCRL5	AML	0.8209	0.7719	0.8765
FMNL1 AND NOT-MS4A1	AML	0.8349	0.7971	0.8765
SLC22A16 AND NOT-PROM1	AML	0.8148	0.8426	0.7888
FMNL1 AND NOT-SLAMF7	AML	0.8103	0.7534	0.8765
FMNL1 AND NOT-TNFRSF17	AML	0.8081	0.7526	0.8725
FMNL1 AND NOT-FCRL1	AML	0.8073	0.7483	0.8765
FMNL1 AND NOT-CD22	AML	0.8102	0.7475	0.8845
FMNL1 AND NOT-CR2	AML	0.8147	0.7551	0.8845
CD37 AND NOT-PIK3IP1	AML	0.8056	0.8024	0.8088
FMNL1 AND NOT-CD72	AML	0.7971	0.7309	0.8765
P2RX4 AND NOT-CLDN23	AML	0.7893	0.6801	0.9402
CD44 AND NOT-IL11RA	AML	0.7821	0.7343	0.8367
FMNL1 AND NOT-CD79A	AML	0.7807	0.7065	0.8725
EMP3 AND NOT-IL11RA	AML	0.7797	0.7393	0.8247
FMNL1 AND NOT-IL11RA	AML	0.7794	0.7235	0.8446
FMNL1 AND NOT-CD19	AML	0.8152	0.781	0.8526
CD37 AND NOT-MSMO1	AML	0.7678	0.7244	0.8167
FMNL1 AND NOT-CD79B	AML	0.7674	0.7472	0.7888
CD276 AND NOT-CD1A	Liposarcoma	0.6914	0.6222	0.7778
CD276 AND NOT-SEMA4B	Liposarcoma	0.6667	0.8333	0.5556

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
CXCL9 AND NOT-KIRREL3	B-Cell Diffuse	0.7222	0.7429	0.7027
CXCL9 AND NOT-CD207	B-Cell Diffuse	0.7222	0.7429	0.7027
CXCL9 AND NOT-ADAM29	B-Cell Diffuse	0.7222	0.7429	0.7027
CD80 AND NOT-CACNG1	B-Cell Diffuse	0.7222	0.7429	0.7027
CXCL9 AND NOT-HTR6	B-Cell Diffuse	0.7222	0.7429	0.7027
CXCL9 AND NOT-CABP7	B-Cell Diffuse	0.7222	0.7429	0.7027
CXCL9 AND NOT-ANPEP	B-Cell Diffuse	0.7213	0.9167	0.5946
MCOLN2 AND NOT-NOX1	B-Cell Diffuse	0.72	0.7105	0.7297
CXCL9 AND NOT-STAB1	B-Cell Diffuse	0.7188	0.8519	0.6216
CXCL9 AND NOT-SLC6A12	B-Cell Diffuse	0.7164	0.8	0.6486
CD80 AND NOT-FXYD7	B-Cell Diffuse	0.7164	0.8	0.6486
CXCL9 AND NOT-FAT1	B-Cell Diffuse	0.7164	0.8	0.6486
KCNJ10 AND NOT-ATP1A2	B-Cell Diffuse	0.716	0.6591	0.7838
CD40 AND NOT-CCR9	B-Cell Diffuse	0.7879	0.8966	0.7027
KCNJ10 AND NOT-GLDN	B-Cell Diffuse	0.7143	0.7576	0.6757
CXCL9 AND NOT-LRRC8E	B-Cell Diffuse	0.7143	0.7576	0.6757
CXCL9 AND NOT-CD1A	B-Cell Diffuse	0.7143	0.7576	0.6757
CXCL9 AND NOT-GRIN2B	B-Cell Diffuse	0.7143	0.7576	0.6757
CD40 AND NOT-LRRN4	B-Cell Diffuse	0.7143	0.7576	0.6757
CXCL9 AND NOT-SLC17A1	B-Cell Diffuse	0.7123	0.7222	0.7027
CXCL9 AND NOT-CYP4A11	B-Cell Diffuse	0.7123	0.7222	0.7027
CXCL9 AND NOT-SLC22A25	B-Cell Diffuse	0.7123	0.7222	0.7027
CXCL9 AND NOT-SLC6A13	B-Cell Diffuse	0.7123	0.7222	0.7027
CXCL9 AND NOT-CLRN1	B-Cell Diffuse	0.7123	0.7222	0.7027
CXCL9 AND NOT-ATP6V0A4	B-Cell Diffuse	0.7123	0.7222	0.7027
CXCL9 AND NOT-ATP2B2	B-Cell Diffuse	0.7123	0.7222	0.7027
CXCL9 AND NOT-KCNH4	B-Cell Diffuse	0.7123	0.7222	0.7027
CXCL9 AND NOT-SLC1A6	B-Cell Diffuse	0.7123	0.7222	0.7027
CD80 AND NOT-SLC6A13	B-Cell Diffuse	0.7123	0.7222	0.7027
CXCL9 AND NOT-SLC13A5	B-Cell Diffuse	0.7123	0.7222	0.7027
CD80 AND NOT-HTR6	B-Cell Diffuse	0.7123	0.7222	0.7027
CXCL9 AND NOT-IGDCC3	B-Cell Diffuse	0.7123	0.7222	0.7027
CXCL9 AND NOT-SLC22A8	B-Cell Diffuse	0.7123	0.7222	0.7027
CXCL9 AND NOT-DIO1	B-Cell Diffuse	0.7123	0.7222	0.7027
CXCL9 AND NOT-OR1C1	B-Cell Diffuse	0.7123	0.7222	0.7027
CXCL9 AND NOT-NOX1	B-Cell Diffuse	0.7123	0.7222	0.7027
CXCL9 AND NOT-CSMD3	B-Cell Diffuse	0.7123	0.7222	0.7027
CXCL9 AND NOT-DRD5	B-Cell Diffuse	0.7123	0.7222	0.7027
CXCL9 AND NOT-MSMO1	B-Cell Diffuse	0.7123	0.7222	0.7027
CXCL9 AND NOT-BEST2	B-Cell Diffuse	0.7123	0.7222	0.7027
CXCL9 AND NOT-PIRT	B-Cell Diffuse	0.7123	0.7222	0.7027
CXCL9 AND NOT-ABCG4	B-Cell Diffuse	0.7123	0.7222	0.7027
CXCL9 AND NOT-SLCO1B3	Anaplastic Lymphoma	0.6207	0.6429	0.6
CXCL9 AND NOT-MMP24	Anaplastic Lymphoma	0.6207	0.6429	0.6
CXCL9 AND NOT-EPHA8	Anaplastic Lymphoma	0.6	0.6	0.6
CXCL9 AND NOT-MRAP	Anaplastic Lymphoma	0.6923	0.8182	0.6
CXCL9 AND NOT-DISP2	Anaplastic Lymphoma	0.6207	0.6429	0.6
CXCL9 AND NOT-GABRB2	Anaplastic Lymphoma	0.6429	0.6923	0.6
CXCL9 AND NOT-ERVW-1	Anaplastic Lymphoma	0.6	0.6	0.6
CXCL9 AND NOT-MEGF11	Anaplastic Lymphoma	0.6207	0.6429	0.6
CXCL9 AND NOT-KCNJ6	Anaplastic Lymphoma	0.6	0.6	0.6
CXCL9 AND NOT-ATP7A	Anaplastic Lymphoma	0.6	0.6	0.6
CXCL9 AND NOT-CHRNG	Anaplastic Lymphoma	0.6207	0.6429	0.6
LAG3 AND NOT-GABRB2	Anaplastic Lymphoma	0.6061	0.5556	0.6667
CXCL9 AND NOT-CSMD3	Anaplastic Lymphoma	0.6429	0.6923	0.6
CXCL9 AND NOT-DCSTAMP	Anaplastic Lymphoma	0.6429	0.6923	0.6
CLECL1 AND NOT-CHRNG	Mantle-Cell Lymphoma	0.9444	1	0.8947
CLECL1 AND NOT-MPL	Mantle-Cell Lymphoma	0.962	0.9268	1
CLECL1 AND NOT-TSPAN16	Mantle-Cell Lymphoma	0.9474	0.9474	0.9474
CLECL1 AND NOT-CD36	Mantle-Cell Lymphoma	0.961	0.9487	0.9737
CLECL1 AND NOT-PAQR7	Mantle-Cell Lymphoma	0.9351	0.9231	0.9474
CLECL1 AND NOT-MLC1	Mantle-Cell Lymphoma	0.9315	0.9714	0.8947
CLECL1 AND NOT-TAS1R1	Mantle-Cell Lymphoma	0.9474	0.9474	0.9474
CLECL1 AND NOT-ANPEP	Mantle-Cell Lymphoma	0.9737	0.9737	0.9737
CLECL1 AND NOT-ACSL6	Mantle-Cell Lymphoma	0.987	0.9744	1
CLECL1 AND NOT-SLC9B1	Mantle-Cell Lymphoma	0.9383	0.8837	1
CLECL1 AND NOT-SLC6A6	Mantle-Cell Lymphoma	0.9315	0.9714	0.8947
CLECL1 AND NOT-STEAP4	Mantle-Cell Lymphoma	0.9744	0.95	1
CLECL1 AND NOT-MUC17	Mantle-Cell Lymphoma	0.9383	0.8837	1
CLECL1 AND NOT-KCNK6	Mantle-Cell Lymphoma	0.9744	0.95	1
CLECL1 AND NOT-CD93	Mantle-Cell Lymphoma	0.9744	0.95	1
CLECL1 AND NOT-CD163	Mantle-Cell Lymphoma	0.9744	0.95	1
CLECL1 AND NOT-GALR2	Mantle-Cell Lymphoma	0.9048	0.8261	1
CLECL1 AND NOT-PTGER4	Mantle-Cell Lymphoma	0.9048	0.8261	1
CLECL1 AND NOT-CHRNB4	Mantle-Cell Lymphoma	0.9091	0.8974	0.9211
CLECL1 AND NOT-VANGL1	Mantle-Cell Lymphoma	0.9048	0.8261	1
CLECL1 AND NOT-GUCY2F	Mantle-Cell Lymphoma	0.9024	0.8409	0.9737

FIG. 12 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
ROR1 AND NOT-ALCAM	Liposarcoma	0.6429	0.9	0.5
ROR1 AND NOT-SLC10A1	Liposarcoma	0.6316	0.8571	0.5
ROR1 AND NOT-ADAM29	Liposarcoma	0.6316	0.8571	0.5
ROR1 AND NOT-NFASC	Liposarcoma	0.6316	0.8571	0.5
ROR1 AND NOT-SCN10A	Liposarcoma	0.6316	0.8571	0.5
ROR1 AND NOT-SLC38A1	Liposarcoma	0.6207	0.8182	0.5
ROR1 AND NOT-SORL1	Liposarcoma	0.6207	0.8182	0.5
ROR1 AND NOT-SLC2A1	Liposarcoma	0.6207	0.8182	0.5
CD276 AND NOT-SLC1A6	Liposarcoma	0.6176	0.6563	0.5833
EPHB2 AND NOT-SLC30A10	Liposarcoma	0.8108	0.7895	0.8333
ROR1 AND NOT-PIK3IP1	Liposarcoma	0.6102	0.7826	0.5
ROR1 AND NOT-ANO4	Liposarcoma	0.6102	0.7826	0.5
ROR1 AND NOT-EFNB2	Liposarcoma	0.6102	0.7826	0.5
ROR1 AND NOT-SLC16A5	Liposarcoma	0.6071	0.85	0.4722
ROR1 AND NOT-CD1B	Liposarcoma	0.6071	0.85	0.4722
EPHB2 AND NOT-SLC22A18	Liposarcoma	0.6429	0.5625	0.75
STAB1 AND NOT-EPCAM	Liposarcoma	0.6757	0.6579	0.6944
CD276 AND NOT-MSMO1	Liposarcoma	0.75	0.6346	0.9167
STAB1 AND NOT-CLDN1	Liposarcoma	0.6133	0.5897	0.6389
STAB1 AND NOT-SSTR2	Liposarcoma	0.6557	0.8	0.5556
CD276 AND NOT-SLC22A18	Liposarcoma	0.6567	0.7097	0.6111
CD276 AND NOT-DSC1	Liposarcoma	0.7105	0.675	0.75
CD276 AND NOT-CD207	Liposarcoma	0.6197	0.6286	0.6111
SLC34A2 AND NOT-SLC16A5	Lung Adenocarcinoma	0.6716	0.7031	0.6429
SLC34A2 AND NOT-CATSPERD	Lung Adenocarcinoma	0.6483	0.6267	0.6714
SLC34A2 AND NOT-LPPR3	Lung Adenocarcinoma	0.6456	0.5795	0.7286
SLC34A2 AND NOT-GRM7	Lung Adenocarcinoma	0.6456	0.5795	0.7286
SLC34A2 AND NOT-CNGA4	Lung Adenocarcinoma	0.6323	0.5765	0.7
SLC34A2 AND NOT-CNIH2	Lung Adenocarcinoma	0.6115	0.5517	0.6857
SLC34A2 AND NOT-NFASC	Lung Adenocarcinoma	0.6071	0.8095	0.4857
SLC34A2 AND NOT-CRB1	Lung Adenocarcinoma	0.6323	0.5765	0.7
SLC34A2 AND NOT-GALR1	Lung Adenocarcinoma	0.6111	0.5946	0.6286
SLC34A2 AND NOT-MSMO1	Lung Adenocarcinoma	0.6438	0.6184	0.6714
MAGEA11 AND NOT-ENPP1	Lung Carcinoma	0.6316	0.9677	0.4688
MAGEA11 AND NOT-SLC16A5	Lung Carcinoma	0.6237	1	0.4531
MAGEA11 AND NOT-SLC22A5	Lung Carcinoma	0.6383	1	0.4688
MAGEA11 AND NOT-PAQR7	Lung Carcinoma	0.6383	1	0.4688
MAGEA11 AND NOT-ESAM	Lung Carcinoma	0.6237	1	0.4531
MAGEA11 AND NOT-SLC23A2	Lung Carcinoma	0.617	0.9667	0.4531
MAGEA11 AND NOT-SLC22A11	Lung Carcinoma	0.6122	0.8824	0.4688
MAGEA11 AND NOT-MRAP	Lung Carcinoma	0.6237	1	0.4531
MAGEA11 AND NOT-AQP6	Lung Carcinoma	0.625	0.9375	0.4688
KCNJ10 AND NOT-BMPR1B	B-Cell Diffuse	0.8	0.7895	0.8108
KCNJ10 AND NOT-NCAM1	B-Cell Diffuse	0.7368	0.7179	0.7568
KCNJ10 AND NOT-CLDN11	B-Cell Diffuse	0.7273	0.7	0.7568
KCNJ10 AND NOT-FOLH1	B-Cell Diffuse	0.7123	0.7222	0.7027
CD180 AND NOT-CD36	B-Cell Diffuse	0.6875	0.8148	0.5946
CD79B AND NOT-ACSL6	B-Cell Diffuse	0.6857	0.7273	0.6486
CD79B AND NOT-SLC30A1	B-Cell Diffuse	0.6849	0.6944	0.6757
CD79B AND NOT-DRD5	B-Cell Diffuse	0.6667	0.7188	0.6216
CD180 AND NOT-ACSL6	B-Cell Diffuse	0.6667	0.6341	0.7027
KCNJ10 AND NOT-ERBB4	B-Cell Diffuse	0.6667	0.6857	0.6486
CD79B AND NOT-ATP8B2	B-Cell Diffuse	0.6667	0.6857	0.6486
FCRL5 AND NOT-PPAPDC1B	B-Cell Diffuse	0.6667	0.7188	0.6216
CD79B AND NOT-CD93	B-Cell Diffuse	0.6667	0.6579	0.6757
CD180 AND NOT-MLC1	B-Cell Diffuse	0.6579	0.641	0.6757
CD72 AND NOT-ACSL6	B-Cell Diffuse	0.6571	0.697	0.6216
CD72 AND NOT-MLC1	B-Cell Diffuse	0.6571	0.697	0.6216
CD79B AND NOT-CD36	B-Cell Diffuse	0.6567	0.7333	0.5946
CD72 AND NOT-CD36	B-Cell Diffuse	0.6479	0.6765	0.6216
P2RX5 AND NOT-ATP8B2	B-Cell Diffuse	0.6506	0.587	0.7297
P2RX5 AND NOT-CD93	B-Cell Diffuse	0.642	0.5909	0.7027
CD79B AND NOT-LAT	B-Cell Diffuse	0.6389	0.6571	0.6216
CD72 AND NOT-ATP8B2	B-Cell Diffuse	0.6389	0.6571	0.6216
CD79A AND NOT-MLC1	B-Cell Diffuse	0.6389	0.6571	0.6216
CD72 AND NOT-CD93	B-Cell Diffuse	0.6389	0.6571	0.6216
CD180 AND NOT-CD93	B-Cell Diffuse	0.6389	0.6571	0.6216
FCRL5 AND NOT-ATP8B2	B-Cell Diffuse	0.6849	0.6944	0.6757
P2RX5 AND NOT-ACSL6	B-Cell Diffuse	0.6582	0.619	0.7027
P2RX5 AND NOT-CD36	B-Cell Diffuse	0.641	0.6098	0.6757
CD79A AND NOT-ACSL6	B-Cell Diffuse	0.6364	0.7241	0.5676
CD79B AND NOT-BTN3A1	B-Cell Diffuse	0.6364	0.7241	0.5676
FCRL5 AND NOT-S1PR1	B-Cell Diffuse	0.6377	0.6875	0.5946
FCRL5 AND NOT-MPL	B-Cell Diffuse	0.6753	0.65	0.7027
FCRL5 AND NOT-CD93	B-Cell Diffuse	0.6842	0.6667	0.7027
FCRL5 AND NOT-CNTNAP2	B-Cell Diffuse	0.6316	0.6154	0.6486
P2RX5 AND NOT-MLC1	B-Cell Diffuse	0.6585	0.6	0.7297

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
CLECL1 AND NOT-MMP24	Mantle-Cell Lymphoma	0.8941	0.8085	1
IL10RA AND NOT-ANPEP	Mantle-Cell Lymphoma	0.9048	0.8261	1
IL10RA AND NOT-CD36	Mantle-Cell Lymphoma	0.9136	0.8605	0.9737
CLECL1 AND NOT-SLC12A9	Mantle-Cell Lymphoma	0.8889	0.9412	0.8421
CLECL1 AND NOT-DRD5	Mantle-Cell Lymphoma	0.8889	0.9412	0.8421
CLECL1 AND NOT-SLC16A5	Mantle-Cell Lymphoma	0.8861	0.8537	0.9211
CLECL1 AND NOT-DYSF	Mantle-Cell Lymphoma	0.95	0.9048	1
CLECL1 AND NOT-CXCL16	Mantle-Cell Lymphoma	0.8857	0.9688	0.8158
CLECL1 AND NOT-LAMP5	Mantle-Cell Lymphoma	0.9091	0.8974	0.9211
CLECL1 AND NOT-CALHM3	Mantle-Cell Lymphoma	0.881	0.8043	0.9737
CLECL1 AND NOT-CNTNAP2	Mantle-Cell Lymphoma	0.88	0.8919	0.8684
QSOX2 AND NOT-KCNC2	Mantle-Cell Lymphoma	0.8889	0.8372	0.9474
QSOX2 AND NOT-SLC5A11	Mantle-Cell Lymphoma	0.8831	0.8718	0.8947
IL10RA AND NOT-CD93	Mantle-Cell Lymphoma	0.8941	0.8085	1
IL10RA AND NOT-CD163	Mantle-Cell Lymphoma	0.9383	0.8837	1
QSOX2 AND NOT-MEGF11	Mantle-Cell Lymphoma	0.9024	0.8409	0.9737
QSOX2 AND NOT-CHRNG	Mantle-Cell Lymphoma	0.9333	0.9459	0.9211
QSOX2 AND NOT-GPR78	Mantle-Cell Lymphoma	0.9	0.8571	0.9474
CLECL1 AND NOT-SLC4A8	Mantle-Cell Lymphoma	0.8837	0.7917	1
KCNN4 AND NOT-ANPEP	Mantle-Cell Lymphoma	0.9114	0.878	0.9474
CLECL1 AND NOT-THBD	Mantle-Cell Lymphoma	0.9383	0.8837	1
CELSR1 AND NOT-LRRC8E	Mantle-Cell Lymphoma	0.9231	0.9	0.9474
CLECL1 AND NOT-ABHD3	Mantle-Cell Lymphoma	0.9157	0.8444	1
KCNN4 AND NOT-SLC31A1	Mantle-Cell Lymphoma	0.925	0.881	0.9737
QSOX2 AND NOT-SLC39A2	Mantle-Cell Lymphoma	0.8889	0.8372	0.9474
QSOX2 AND NOT-GABRD	Mantle-Cell Lymphoma	0.8889	0.8372	0.9474
QSOX2 AND NOT-SCN8A	Mantle-Cell Lymphoma	0.8947	0.8947	0.8947
QSOX2 AND NOT-ZACN	Mantle-Cell Lymphoma	0.9189	0.9444	0.8947
QSOX2 AND NOT-OR3A2	Mantle-Cell Lymphoma	0.875	0.8333	0.9211
QSOX2 AND NOT-CRB2	Mantle-Cell Lymphoma	0.8889	0.8372	0.9474
QSOX2 AND NOT-ANPEP	Mantle-Cell Lymphoma	0.961	0.9487	0.9737
QSOX2 AND NOT-KCNK4	Mantle-Cell Lymphoma	0.9024	0.8409	0.9737
QSOX2 AND NOT-ATP6V0A4	Mantle-Cell Lymphoma	0.8889	0.8372	0.9474
QSOX2 AND NOT-CALY	Mantle-Cell Lymphoma	0.8974	0.875	0.9211
QSOX2 AND NOT-DRD5	Mantle-Cell Lymphoma	0.8857	0.9688	0.8158
CLECL1 AND NOT-GPR78	Mantle-Cell Lymphoma	0.8605	0.7708	0.9737
CLECL1 AND NOT-STAB1	Mantle-Cell Lymphoma	0.8837	0.7917	1
QSOX2 AND NOT-SLC28A1	Mantle-Cell Lymphoma	0.9024	0.8409	0.9737
QSOX2 AND NOT-SLC5A8	Mantle-Cell Lymphoma	0.875	0.8333	0.9211
QSOX2 AND NOT-MRGPRX2	Mantle-Cell Lymphoma	0.8608	0.8293	0.8947
QSOX2 AND NOT-LPPR3	Mantle-Cell Lymphoma	0.8889	0.8372	0.9474
QSOX2 AND NOT-TSPAN16	Mantle-Cell Lymphoma	0.9333	0.9459	0.9211
QSOX2 AND NOT-SYT6	Mantle-Cell Lymphoma	0.8889	0.8372	0.9474
QSOX2 AND NOT-GABRA4	Mantle-Cell Lymphoma	0.9024	0.8409	0.9737
CLECL1 AND NOT-TGFBI	Mantle-Cell Lymphoma	0.8889	0.9412	0.8421
QSOX2 AND NOT-SLC22A6	Mantle-Cell Lymphoma	0.8889	0.8372	0.9474
QSOX2 AND NOT-CLCN1	Mantle-Cell Lymphoma	0.875	0.8333	0.9211
QSOX2 AND NOT-GALR2	Mantle-Cell Lymphoma	0.9367	0.9024	0.9737
QSOX2 AND NOT-SEZ6	Mantle-Cell Lymphoma	0.9024	0.8409	0.9737
QSOX2 AND NOT-MAG	Mantle-Cell Lymphoma	0.9024	0.8409	0.9737
QSOX2 AND NOT-OR52D1	Mantle-Cell Lymphoma	0.875	0.8333	0.9211
QSOX2 AND NOT-CD207	Mantle-Cell Lymphoma	0.9024	0.8409	0.9737
QSOX2 AND NOT-SLC6A6	Mantle-Cell Lymphoma	0.9189	0.9444	0.8947
QSOX2 AND NOT-MMP24	Mantle-Cell Lymphoma	0.9367	0.9024	0.9737
QSOX2 AND NOT-SLC6A11	Mantle-Cell Lymphoma	0.8608	0.8293	0.8947
QSOX2 AND NOT-BEST2	Mantle-Cell Lymphoma	0.8889	0.8372	0.9474
QSOX2 AND NOT-GUCY2D	Mantle-Cell Lymphoma	0.9	0.8571	0.9474
QSOX2 AND NOT-PVRL1	Mantle-Cell Lymphoma	0.9024	0.8409	0.9737
CLECL1 AND NOT-FMNL1	Mantle-Cell Lymphoma	0.8837	0.7917	1
CLECL1 AND NOT-ATP8B2	Mantle-Cell Lymphoma	0.9157	0.8444	1
CLECL1 AND NOT-TTYH2	Mantle-Cell Lymphoma	0.9268	0.8636	1
QSOX2 AND NOT-OPCML	Mantle-Cell Lymphoma	0.875	0.8333	0.9211
QSOX2 AND NOT-MLC1	Mantle-Cell Lymphoma	0.9315	0.9714	0.8947
QSOX2 AND NOT-GRM3	Mantle-Cell Lymphoma	0.9367	0.9024	0.9737
QSOX2 AND NOT-CALN1	Mantle-Cell Lymphoma	0.875	0.8333	0.9211
QSOX2 AND NOT-TSHR	Mantle-Cell Lymphoma	0.9024	0.8409	0.9737
QSOX2 AND NOT-SLC1A6	Mantle-Cell Lymphoma	0.8608	0.8293	0.8947
QSOX2 AND NOT-GJB4	Mantle-Cell Lymphoma	0.8889	0.8372	0.9474
QSOX2 AND NOT-CLDN16	Mantle-Cell Lymphoma	0.9024	0.8409	0.9737
CLECL1 AND NOT-GRM3	Mantle-Cell Lymphoma	0.8539	0.7451	1
QSOX2 AND NOT-GGTLC1	Mantle-Cell Lymphoma	0.8533	0.8649	0.8421
CLECL1 AND NOT-SLC39A8	Mantle-Cell Lymphoma	0.8837	0.7917	1
CXCL9 AND NOT-SLC30A1	T-Cell, Peripheral	0.7917	0.95	0.6786
CXCL9 AND NOT-ATP8B1	T-Cell, Peripheral	0.7586	0.7333	0.7857
CXCL9 AND NOT-FAT1	T-Cell, Peripheral	0.7458	0.7097	0.7857
CXCL9 AND NOT-SLC22A18	T-Cell, Peripheral	0.7458	0.7097	0.7857
CXCL9 AND NOT-IFI6	T-Cell, Peripheral	0.8	0.9091	0.7143

FIG. 12 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
IL2RA AND NOT-FXYD7	B-Cell Diffuse	0.6452	0.8	0.5405
FCRL5 AND NOT-MLC1	B-Cell Diffuse	0.6757	0.6757	0.6757
CD79B AND NOT-MPL	B-Cell Diffuse	0.6269	0.7	0.5676
CD79B AND NOT-BTN3A3	B-Cell Diffuse	0.6269	0.7	0.5676
CD79A AND NOT-CD36	B-Cell Diffuse	0.6269	0.7	0.5676
CD79B AND NOT-ANPEP	B-Cell Diffuse	0.6269	0.7	0.5676
CD180 AND NOT-ATP8B2	B-Cell Diffuse	0.6269	0.7	0.5676
FCRL5 AND NOT-CCR9	B-Cell Diffuse	0.625	0.5814	0.6757
FCRL5 AND NOT-DRD5	B-Cell Diffuse	0.6486	0.6486	0.6486
FCRL5 AND NOT-GPR22	B-Cell Diffuse	0.6842	0.6667	0.7027
P2RX5 AND NOT-DGKE	B-Cell Diffuse	0.6216	0.6216	0.6216
CD79A AND NOT-PAQR8	B-Cell Diffuse	0.6216	0.6216	0.6216
CD180 AND NOT-S1PR1	B-Cell Diffuse	0.6197	0.6471	0.5946
CD72 AND NOT-CNTNAP2	B-Cell Diffuse	0.6197	0.6471	0.5946
IL2RA AND NOT-SLC6A6	B-Cell Diffuse	0.6364	0.7241	0.5676
FCRL5 AND NOT-ACSL6	B-Cell Diffuse	0.6757	0.6757	0.6757
FCRL5 AND NOT-PAQR7	B-Cell Diffuse	0.6575	0.6667	0.6486
CD72 AND NOT-MPL	B-Cell Diffuse	0.6176	0.6774	0.5676
CD79B AND NOT-SLC16A5	B-Cell Diffuse	0.6154	0.7143	0.5405
CD79B AND NOT-ANTXR2	B-Cell Diffuse	0.6154	0.7143	0.5405
FCRL5 AND NOT-ANPEP	B-Cell Diffuse	0.6154	0.5854	0.6486
P2RX5 AND NOT-PAQR8	B-Cell Diffuse	0.6136	0.5294	0.7297
FCRL5 AND NOT-ATP8A2	B-Cell Diffuse	0.6133	0.6053	0.6216
GPNMB AND NOT-MRGPRF	B-Cell Diffuse	0.6133	0.6053	0.6216
FCRL5 AND NOT-CHRNB4	B-Cell Diffuse	0.6118	0.5417	0.7027
FCRL5 AND NOT-SLC9B1	B-Cell Diffuse	0.6111	0.6286	0.5946
IL2RA AND NOT-CCR9	B-Cell Diffuse	0.6389	0.6571	0.6216
CD79A AND NOT-CD93	B-Cell Diffuse	0.6087	0.6563	0.5676
IL2RA AND NOT-TRPM3	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-MRGPRX2	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-ZP4	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-CACNA1S	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-FXYD7	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-LRRN4	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-KCNA10	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-AGTR2	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-MEGF11	Anaplastic Lymphoma	0.72	0.9	0.6
IL2RA AND NOT-OR5P3	Anaplastic Lymphoma	0.6923	0.8182	0.6
IL2RA AND NOT-MOG	Anaplastic Lymphoma	0.6923	0.8182	0.6
IL2RA AND NOT-CNTNAP4	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-KCNK10	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-KIAA0319	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-KCNH4	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-SLCO1B3	Anaplastic Lymphoma	0.6364	1	0.4667
IL2RA AND NOT-CNTNAP2	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-SLC6A15	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-MUC17	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-CNGA4	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-CALY	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-OR52D1	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-TSPAN16	Anaplastic Lymphoma	0.6154	0.7273	0.5333
IL2RA AND NOT-SLC9B1	Anaplastic Lymphoma	0.6154	0.7273	0.5333
IL2RA AND NOT-SLC22A13	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-GRM3	Anaplastic Lymphoma	0.6154	0.7273	0.5333
IL2RA AND NOT-SLC5A11	Anaplastic Lymphoma	0.6154	0.7273	0.5333
IL2RA AND NOT-GABRB2	Anaplastic Lymphoma	0.6154	0.7273	0.5333
IL2RA AND NOT-DRD5	Anaplastic Lymphoma	0.6154	0.7273	0.5333
IL2RA AND NOT-CDH18	Anaplastic Lymphoma	0.6154	0.7273	0.5333
IL2RA AND NOT-CALHM1	Anaplastic Lymphoma	0.6429	0.6923	0.6
IL2RA AND NOT-MMP24	Anaplastic Lymphoma	0.72	0.9	0.6
IL2RA AND NOT-ASTN1	Anaplastic Lymphoma	0.6154	0.7273	0.5333
IL2RA AND NOT-MTNR1B	Anaplastic Lymphoma	0.6154	0.7273	0.5333
IL2RA AND NOT-SV2C	Anaplastic Lymphoma	0.6154	0.7273	0.5333
IL2RA AND NOT-TRPM1	Anaplastic Lymphoma	0.6154	0.7273	0.5333
IL2RA AND NOT-EPHA8	Anaplastic Lymphoma	0.6154	0.7273	0.5333
IL2RA AND NOT-RXFP3	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-ADAM30	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-KCNA4	Anaplastic Lymphoma	0.6154	0.7273	0.5333
IL2RA AND NOT-DSC1	Anaplastic Lymphoma	0.6154	0.7273	0.5333
IL2RA AND NOT-LCT	Anaplastic Lymphoma	0.6154	0.7273	0.5333
IL2RA AND NOT-GABRA2	Anaplastic Lymphoma	0.6429	0.6923	0.6
IL2RA AND NOT-GRIA4	Anaplastic Lymphoma	0.6154	0.7273	0.5333
IL2RA AND NOT-KCNA1	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-LPPR3	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-SLC12A3	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-SEZ6	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-CCKAR	Anaplastic Lymphoma	0.6667	0.75	0.6

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
CXCL9 AND NOT-SEMA4B	T-Cell, Peripheral	0.7458	0.7097	0.7857
CXCL9 AND NOT-PVRL1	T-Cell, Peripheral	0.7333	0.6875	0.7857
CXCL9 AND NOT-CXCL16	T-Cell, Peripheral	0.75	0.9	0.6429
CXCL9 AND NOT-PTGIS	T-Cell, Peripheral	0.7333	0.6875	0.7857
CXCL9 AND NOT-DSC1	T-Cell, Peripheral	0.7586	0.7333	0.7857
CXCL9 AND NOT-SLC2A1	T-Cell, Peripheral	0.7458	0.7097	0.7857
CXCL9 AND NOT-SLC16A5	T-Cell, Peripheral	0.7273	0.7407	0.7143
CXCL9 AND NOT-ABCG4	T-Cell, Peripheral	0.7333	0.6875	0.7857
CXCL9 AND NOT-SMPD2	T-Cell, Peripheral	0.7458	0.7097	0.7857
CXCL9 AND NOT-GHR	T-Cell, Peripheral	0.717	0.76	0.6786
CXCL9 AND NOT-THBD	T-Cell, Peripheral	0.7586	0.7333	0.7857
CXCL9 AND NOT-FXYD6	T-Cell, Peripheral	0.7458	0.7097	0.7857
CXCL9 AND NOT-PHLDB2	T-Cell, Peripheral	0.7458	0.7097	0.7857
CXCL9 AND NOT-CD207	T-Cell, Peripheral	0.7143	0.7143	0.7143
CXCL9 AND NOT-EPHA2	T-Cell, Peripheral	0.7143	0.7143	0.7143
CXCL9 AND NOT-ZACN	T-Cell, Peripheral	0.7143	0.7143	0.7143
CXCL9 AND NOT-SCARA5	T-Cell, Peripheral	0.7368	0.7241	0.75
CXCL9 AND NOT-LRIG3	T-Cell, Peripheral	0.7119	0.6774	0.75
CXCL9 AND NOT-PAQR7	T-Cell, Peripheral	0.7119	0.6774	0.75
CXCL9 AND NOT-MSMO1	T-Cell, Peripheral	0.7119	0.6774	0.75
CXCL9 AND NOT-SLC6A18	T-Cell, Peripheral	0.7333	0.6875	0.7857
CXCL9 AND NOT-SLC22A16	T-Cell, Peripheral	0.7857	0.7857	0.7857
CXCL9 AND NOT-KCNA7	T-Cell, Peripheral	0.7119	0.6774	0.75
CXCL9 AND NOT-SLC39A8	T-Cell, Peripheral	0.7308	0.7917	0.6786
CXCL9 AND NOT-STEAP4	T-Cell, Peripheral	0.7059	0.7826	0.6429
CXCL9 AND NOT-ANPEP	T-Cell, Peripheral	0.8235	0.913	0.75
CXCL9 AND NOT-CHRNA9	T-Cell, Peripheral	0.7119	0.6774	0.75
CXCL9 AND NOT-CHRNA1	T-Cell, Peripheral	0.7119	0.6774	0.75
CXCL9 AND NOT-CHRNB2	T-Cell, Peripheral	0.7018	0.6897	0.7143
CXCL9 AND NOT-SLC5A2	T-Cell, Peripheral	0.7018	0.6897	0.7143
CXCL9 AND NOT-SLC17A2	T-Cell, Peripheral	0.7333	0.6875	0.7857
CXCL9 AND NOT-LHFPL5	T-Cell, Peripheral	0.7119	0.6774	0.75
CXCL9 AND NOT-SLC17A1	T-Cell, Peripheral	0.7119	0.6774	0.75
CXCL9 AND NOT-ABCG8	T-Cell, Peripheral	0.7333	0.6875	0.7857
CXCL9 AND NOT-SLC2A2	T-Cell, Peripheral	0.7333	0.6875	0.7857
CXCL9 AND NOT-ATP2B2	T-Cell, Peripheral	0.7333	0.6875	0.7857
CXCL9 AND NOT-PTPRT	T-Cell, Peripheral	0.7333	0.6875	0.7857
CXCL9 AND NOT-TAS2R1	T-Cell, Peripheral	0.7018	0.6897	0.7143
CXCL9 AND NOT-SLC43A1	T-Cell, Peripheral	0.7333	0.6875	0.7857
CXCL9 AND NOT-ALCAM	T-Cell, Peripheral	0.7719	0.7586	0.7857
CXCL9 AND NOT-SLC13A2	T-Cell, Peripheral	0.7333	0.6875	0.7857
CXCL9 AND NOT-GABBR2	T-Cell, Peripheral	0.7333	0.6875	0.7857
CXCL9 AND NOT-KREMEN2	T-Cell, Peripheral	0.6909	0.7037	0.6786
CXCL9 AND NOT-DYSF	T-Cell, Peripheral	0.6909	0.7037	0.6786
CXCL9 AND NOT-PROCR	T-Cell, Peripheral	0.7826	1	0.6429
CXCL9 AND NOT-LRRC55	T-Cell, Peripheral	0.6897	0.6667	0.7143
CXCL9 AND NOT-TTYH3	T-Cell, Peripheral	0.6897	0.6667	0.7143
CXCL9 AND NOT-DSCAM	T-Cell, Peripheral	0.6897	0.6667	0.7143
CXCL9 AND NOT-CEACAM7	T-Cell, Peripheral	0.6897	0.6667	0.7143
CXCL9 AND NOT-SLC9A1	T-Cell, Peripheral	0.6897	0.6667	0.7143
CXCL9 AND NOT-GGTLC1	T-Cell, Peripheral	0.7018	0.6897	0.7143
CXCL9 AND NOT-UNC5A	T-Cell, Peripheral	0.7333	0.6875	0.7857
CXCL9 AND NOT-EBP	T-Cell, Peripheral	0.7333	0.6875	0.7857
CXCL9 AND NOT-CD1B	T-Cell, Peripheral	0.6897	0.6667	0.7143
CXCL9 AND NOT-SLC12A5	T-Cell, Peripheral	0.7333	0.6875	0.7857
CXCL9 AND NOT-CHRND	T-Cell, Peripheral	0.7333	0.6875	0.7857
CXCL9 AND NOT-SLC6A6	T-Cell, Peripheral	0.6809	0.8421	0.5714
CXCL9 AND NOT-CD74	T-Cell, Peripheral	0.6786	0.6786	0.6786
CXCL9 AND NOT-IL13	T-Cell, Peripheral	0.6786	0.6786	0.6786
CXCL9 AND NOT-KCNJ6	T-Cell, Peripheral	0.6786	0.6786	0.6786
CXCL9 AND NOT-CD36	T-Cell, Peripheral	0.6667	0.8824	0.5357
CXCL9 AND NOT-SLCO1A2	T-Cell, Peripheral	0.6667	0.6552	0.6786
CXCL9 AND NOT-MUC12	T-Cell, Peripheral	0.6667	0.6552	0.6786
CXCL9 AND NOT-SLC6A12	T-Cell, Peripheral	0.6667	0.7391	0.6071
CXCL9 AND NOT-MEGF10	T-Cell, Peripheral	0.6667	0.6552	0.6786
CXCL9 AND NOT-CD1A	T-Cell, Peripheral	0.6667	0.6923	0.6429
CXCL9 AND NOT-SLC10A1	T-Cell, Peripheral	0.6667	0.6552	0.6786
CXCL9 AND NOT-TGFBI	T-Cell, Peripheral	0.8235	0.913	0.75
CXCL9 AND NOT-RGR	T-Cell, Peripheral	0.6667	0.6552	0.6786
CXCL9 AND NOT-VANGL1	T-Cell, Peripheral	0.7333	0.6875	0.7857
CXCL9 AND NOT-GJA3	T-Cell, Peripheral	0.6923	0.75	0.6429
CXCL9 AND NOT-RGSL1	T-Cell, Peripheral	0.6545	0.6667	0.6429
CXCL9 AND NOT-OR2C3	T-Cell, Peripheral	0.6545	0.6667	0.6429
CXCL9 AND NOT-TNFRSF10B	T-Cell, Peripheral	0.6545	0.6667	0.6429
CXCL9 AND NOT-SLC23A2	T-Cell, Peripheral	0.6531	0.7619	0.5714
CXCL9 AND NOT-SLCO1B3	T-Cell, Peripheral	0.6512	0.9333	0.5
CXCL10 AND NOT-ATP8B1	T-Cell, Peripheral	0.6441	0.6129	0.6786

FIG. 12 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
IL2RA AND NOT-OR3A2	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-TRPC7	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-LYPD1	Anaplastic Lymphoma	0.6154	0.7273	0.5333
IL2RA AND NOT-GABRD	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-MPL	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-GUCY2F	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-SLC39A12	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-OR2L2	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-SLC22A16	Anaplastic Lymphoma	0.6923	0.8182	0.6
IL2RA AND NOT-GABRA4	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-PCDHB1	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-CACNG2	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-SLC28A1	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-HTR6	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-MAG	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-ADAM7	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-P2RY4	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-GALR2	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-SCN1A	Anaplastic Lymphoma	0.6429	0.6923	0.6
IL2RA AND NOT-KCNC1	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-GABRA5	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-OR4N4	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-KCNQ2	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-GPR83	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-GNRHR	Anaplastic Lymphoma	0.6429	0.6923	0.6
IL2RA AND NOT-HCN1	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-PCDHGC4	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-DCC	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-LRRTM4	Anaplastic Lymphoma	0.6154	0.7273	0.5333
IL2RA AND NOT-GRM1	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-OR1Q1	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-GLRA3	Anaplastic Lymphoma	0.6429	0.6923	0.6
IL2RA AND NOT-CACNA1E	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-SLC13A5	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-P2RX3	Anaplastic Lymphoma	0.6923	0.8182	0.6
IL2RA AND NOT-GPR6	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-CALN1	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-CACNG7	Anaplastic Lymphoma	0.6154	0.7273	0.5333
IL2RA AND NOT-CD207	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-AJAP1	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-SLC1A6	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-GRM7	Anaplastic Lymphoma	0.6154	0.7273	0.5333
IL2RA AND NOT-ANO4	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-GRIK4	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-UMODL1	Anaplastic Lymphoma	0.6429	0.6923	0.6
IL2RA AND NOT-GPR12	Anaplastic Lymphoma	0.6429	0.6923	0.6
IL2RA AND NOT-HCN2	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-CASR	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-KREMEN2	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-OPRM1	Anaplastic Lymphoma	0.6667	0.75	0.6
IL2RA AND NOT-DTNA	Anaplastic Lymphoma	0.6667	0.75	0.6
MS4A1 AND NOT-SORL1	Mantle-Cell Lymphoma	0.9136	0.8605	0.9737
CD52 AND NOT-CD36	Mantle-Cell Lymphoma	0.8642	0.814	0.9211
CD52 AND NOT-ANPEP	Mantle-Cell Lymphoma	0.8642	0.814	0.9211
CD52 AND NOT-CD93	Mantle-Cell Lymphoma	0.8889	0.8372	0.9474
MS4A1 AND NOT-CALHM3	Mantle-Cell Lymphoma	0.8506	0.7551	0.9737
MS4A1 AND NOT-MLC1	Mantle-Cell Lymphoma	0.85	0.8095	0.8947
MS4A1 AND NOT-PTGER4	Mantle-Cell Lymphoma	0.8444	0.7308	1
MS4A1 AND NOT-CCR9	Mantle-Cell Lymphoma	0.8434	0.7778	0.9211
P2RX5 AND NOT-CHRNG	Mantle-Cell Lymphoma	0.8421	0.8421	0.8421
P2RX5 AND NOT-MLC1	Mantle-Cell Lymphoma	0.8421	0.8421	0.8421
CD52 AND NOT-MLC1	Mantle-Cell Lymphoma	0.8649	0.8889	0.8421
CD52 AND NOT-CD163	Mantle-Cell Lymphoma	0.8675	0.8	0.9474
MS4A1 AND NOT-XCR1	Mantle-Cell Lymphoma	0.8352	0.717	1
MS4A1 AND NOT-KCNA4	Mantle-Cell Lymphoma	0.8315	0.7255	0.9737
MS4A1 AND NOT-DRD5	Mantle-Cell Lymphoma	0.8312	0.8205	0.8421
MS4A1 AND NOT-ADAM2	Mantle-Cell Lymphoma	0.8312	0.8205	0.8421
P2RX5 AND NOT-CD163	Mantle-Cell Lymphoma	0.9	0.8571	0.9474
MS4A1 AND NOT-CACNG4	Mantle-Cell Lymphoma	0.8261	0.7037	1
P2RX5 AND NOT-SLC12A9	Mantle-Cell Lymphoma	0.8267	0.8378	0.8158
MS4A1 AND NOT-TSPAN16	Mantle-Cell Lymphoma	0.8182	0.72	0.9474
MS4A1 AND NOT-CLDN10	Mantle-Cell Lymphoma	0.8172	0.6909	1
MS4A1 AND NOT-ADCY8	Mantle-Cell Lymphoma	0.8172	0.6909	1
MS4A1 AND NOT-KCNK12	Mantle-Cell Lymphoma	0.814	0.7292	0.9211
MS4A1 AND NOT-GALR1	Mantle-Cell Lymphoma	0.8132	0.6981	0.9737
CD79B AND NOT-SORL1	Mantle-Cell Lymphoma	0.8116	0.9032	0.7368
TNFSF11 AND NOT-STAB1	T-Cell, Peripheral	0.6047	0.8667	0.4643

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
CXCL9 AND NOT-PIRT	T-Cell, Peripheral	0.6429	0.6429	0.6429
CXCL9 AND NOT-SCN10A	T-Cell, Peripheral	0.6429	0.6429	0.6429
CXCL9 AND NOT-NPHS2	T-Cell, Peripheral	0.6429	0.6429	0.6429
CXCL9 AND NOT-SLC39A12	T-Cell, Peripheral	0.6429	0.6429	0.6429
CXCL9 AND NOT-KCNJ9	T-Cell, Peripheral	0.6429	0.6429	0.6429
CXCL9 AND NOT-TNFRSF12A	T-Cell, Peripheral	0.7241	0.7	0.75
CXCL9 AND NOT-MRGPRF	T-Cell, Peripheral	0.7273	0.7407	0.7143
C11orf24 AND NOT-ANPEP	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-LHFPL5	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-SLC10A1	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-OR10J1	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-NMUR2	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-NPFFR1	Melanoma	0.9	1	0.8182
TNFSF9 AND NOT-TMEM235	Melanoma	0.9524	1	0.9091
C11orf24 AND NOT-TMEM235	Melanoma	0.9	1	0.8182
SLC5A6 AND NOT-ATP13A5	Melanoma	0.9524	1	0.9091
TNFSF9 AND NOT-CDH20	Melanoma	0.9524	1	0.9091
C11orf24 AND NOT-OR1Q1	Melanoma	0.9	1	0.8182
MC1R AND NOT-OR5P3	Melanoma	0.9	1	0.8182
MC1R AND NOT-NPFFR1	Melanoma	0.9	1	0.8182
MC1R AND NOT-KCNJ6	Melanoma	0.9	1	0.8182
MC1R AND NOT-TMEM235	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-ATP13A5	Melanoma	0.9	1	0.8182
MC1R AND NOT-OR10J1	Melanoma	0.9	1	0.8182
MC1R AND NOT-NMUR2	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-NOX1	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-OR3A1	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-HCN4	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-CACNG4	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-OR1C1	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-CD82	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-SLC22A8	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-FSHR	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-CCR9	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-AMHR2	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-CLDN19	Melanoma	0.9	1	0.8182
TNFSF9 AND NOT-KCNJ6	Melanoma	0.9524	1	0.9091
C11orf24 AND NOT-MC2R	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-CHRNA2	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-RGSL1	Melanoma	0.9	1	0.8182
TNFSF9 AND NOT-KCNQ2	Melanoma	0.9524	1	0.9091
C11orf24 AND NOT-GJD2	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-C8B	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-BEST2	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-ABCG8	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-SLC2A2	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-SLC6A13	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-SCN10A	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-SLC22A9	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-CALHM1	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-LRFN2	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-SLC6A18	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-HTR1E	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-GLRA2	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-CACNG6	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-SLC10A2	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-SLC13A1	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-PCDHGC4	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-OTOF	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-HRH3	Melanoma	0.9	1	0.8182
TNFSF9 AND NOT-NMUR2	Melanoma	0.9524	1	0.9091
C11orf24 AND NOT-ADAM29	Melanoma	0.9	1	0.8182
TNFSF9 AND NOT-SYT4	Melanoma	0.9524	1	0.9091
C11orf24 AND NOT-KCNJ6	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-GRIN2B	Melanoma	0.9	1	0.8182
MC1R AND NOT-PCDHB1	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-PCDH11Y	Melanoma	0.9	1	0.8182
MC1R AND NOT-CHRNA2	Melanoma	0.9	1	0.8182
MC1R AND NOT-SLC6A13	Melanoma	0.9	1	0.8182
MC1R AND NOT-KCNQ2	Melanoma	0.9	1	0.8182
MC1R AND NOT-SLC34A1	Melanoma	0.9	1	0.8182
MC1R AND NOT-OTOF	Melanoma	0.9	1	0.8182
MC1R AND NOT-PCDH11X	Melanoma	0.9	1	0.8182
MC1R AND NOT-OR1C1	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-IL13	Melanoma	0.9	1	0.8182
MC1R AND NOT-CALHM1	Melanoma	0.9	1	0.8182
MC1R AND NOT-GRIN2B	Melanoma	0.9	1	0.8182

FIG. 12 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
EDNRB AND NOT-VN1R2	Melanoma	0.9	1	0.8182
EDNRB AND NOT-ERVW-1	Melanoma	0.9	1	0.8182
EDNRB AND NOT-PCDHB1	Melanoma	0.9	1	0.8182
EDNRB AND NOT-GRM8	Melanoma	0.9	1	0.8182
EDNRB AND NOT-GRM6	Melanoma	0.9	1	0.8182
EDNRB AND NOT-GRM4	Melanoma	0.9	1	0.8182
EDNRB AND NOT-GRM3	Melanoma	0.9	1	0.8182
EDNRB AND NOT-GRIN2B	Melanoma	0.9	1	0.8182
EDNRB AND NOT-GRID2	Melanoma	0.9	1	0.8182
EDNRB AND NOT-MLNR	Melanoma	0.9	1	0.8182
EDNRB AND NOT-GPR26	Melanoma	0.9	1	0.8182
EDNRB AND NOT-TMEM235	Melanoma	0.9	1	0.8182
EDNRB AND NOT-CDH20	Melanoma	0.9	1	0.8182
EDNRB AND NOT-HTR1E	Melanoma	0.9	1	0.8182
EDNRB AND NOT-HTR5A	Melanoma	0.9	1	0.8182
EDNRB AND NOT-SLCO1B3	Melanoma	0.9	1	0.8182
EDNRB AND NOT-ATP13A5	Melanoma	0.9	1	0.8182
EDNRB AND NOT-MIP	Melanoma	0.9	1	0.8182
EDNRB AND NOT-MC2R	Melanoma	0.9	1	0.8182
EDNRB AND NOT-CDH17	Melanoma	0.9	1	0.8182
EDNRB AND NOT-SLC22A25	Melanoma	0.9	1	0.8182
EDNRB AND NOT-KCNJ6	Melanoma	0.9	1	0.8182
EDNRB AND NOT-KCNH1	Melanoma	0.9	1	0.8182
EDNRB AND NOT-KCND2	Melanoma	0.9	1	0.8182
EDNRB AND NOT-KCNA4	Melanoma	0.9	1	0.8182
EDNRB AND NOT-KCNA1	Melanoma	0.9	1	0.8182
EDNRB AND NOT-CD82	Melanoma	0.9	1	0.8182
EDNRB AND NOT-GPR6	Melanoma	0.9	1	0.8182
EDNRB AND NOT-GLRA2	Melanoma	0.9	1	0.8182
EDNRB AND NOT-OR3A1	Melanoma	0.9	1	0.8182
EDNRB AND NOT-SV2C	Melanoma	0.9	1	0.8182
EDNRB AND NOT-LHFPL5	Melanoma	0.9	1	0.8182
EDNRB AND NOT-GPRC6A	Melanoma	0.9	1	0.8182
EDNRB AND NOT-SLC39A12	Melanoma	0.9	1	0.8182
EDNRB AND NOT-EPHA8	Melanoma	0.9	1	0.8182
EDNRB AND NOT-CHRNB2	Melanoma	0.9	1	0.8182
EDNRB AND NOT-AGTR2	Melanoma	0.9	1	0.8182
EDNRB AND NOT-CHRNA4	Melanoma	0.9	1	0.8182
EDNRB AND NOT-SLC17A2	Melanoma	0.9	1	0.8182
EDNRB AND NOT-PIK3IP1	Melanoma	0.9	1	0.8182
EDNRB AND NOT-DSCAM	Melanoma	0.9	1	0.8182
EDNRB AND NOT-AGER	Melanoma	0.9	1	0.8182
EDNRB AND NOT-NTSR2	Melanoma	0.9	1	0.8182
EDNRB AND NOT-PCDH11X	Melanoma	0.9	1	0.8182
EDNRB AND NOT-ADAM2	Melanoma	0.9	1	0.8182
EDNRB AND NOT-CACNG4	Melanoma	0.9	1	0.8182
EDNRB AND NOT-NOX1	Melanoma	0.9	1	0.8182
EDNRB AND NOT-AMHR2	Melanoma	0.9	1	0.8182
EDNRB AND NOT-GHSR	Melanoma	0.9	1	0.8182
EDNRB AND NOT-OR10J1	Melanoma	0.9	1	0.8182
EDNRB AND NOT-OR2L2	Melanoma	0.9	1	0.8182
EDNRB AND NOT-OR1C1	Melanoma	0.9	1	0.8182
EDNRB AND NOT-CATSPERD	Melanoma	0.9	1	0.8182
EDNRB AND NOT-GABRB2	Melanoma	0.9	1	0.8182
EDNRB AND NOT-SLC22A9	Melanoma	0.9	1	0.8182
EDNRB AND NOT-CALHM1	Melanoma	0.9	1	0.8182
NKAIN1 AND NOT-EGFR	Neuroblastoma	0.908	0.9367	0.881
NKAIN1 AND NOT-SSTR1	Neuroblastoma	0.8929	0.8929	0.8929
NKAIN1 AND NOT-BCAN	Neuroblastoma	0.8929	0.8929	0.8929
NKAIN1 AND NOT-GAGE1	Neuroblastoma	0.8862	0.8916	0.881
L1CAM AND NOT-NKAIN2	Neuroblastoma	0.8846	0.9583	0.8214
L1CAM AND NOT-GRM3	Neuroblastoma	0.8848	0.9012	0.869
NKAIN1 AND NOT-GPA33	Neuroblastoma	0.881	0.881	0.881
L1CAM AND NOT-KCNA1	Neuroblastoma	0.9375	0.9868	0.8929
NKAIN1 AND NOT-DPEP1	Neuroblastoma	0.8721	0.8523	0.8929
NKAIN1 AND NOT-WT1	Neuroblastoma	0.8712	0.8987	0.8452
NKAIN1 AND NOT-RNF43	Neuroblastoma	0.8706	0.8605	0.881
L1CAM AND NOT-SLCO1C1	Neuroblastoma	0.9182	0.9733	0.869
NKAIN1 AND NOT-ULBP3	Neuroblastoma	0.8671	0.8427	0.8929
L1CAM AND NOT-TRPC3	Neuroblastoma	0.8671	0.8427	0.8929
NKAIN1 AND NOT-TNFRSF10A	Neuroblastoma	0.8671	0.8427	0.8929
NKAIN1 AND NOT-HHLA2	Neuroblastoma	0.8655	0.8506	0.881
NKAIN1 AND NOT-CLDN2	Neuroblastoma	0.8655	0.8506	0.881
NKAIN1 AND NOT-TYR	Neuroblastoma	0.8639	0.8588	0.869
L1CAM AND NOT-CSMD3	Neuroblastoma	0.882	0.9221	0.8452
L1CAM AND NOT-MLC1	Neuroblastoma	0.8621	0.8333	0.8929
NKAIN1 AND NOT-CLDN18	Neuroblastoma	0.8621	0.8333	0.8929

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
MC1R AND NOT-FSHR	Melanoma	0.9	1	0.8182
MC1R AND NOT-SLC17A2	Melanoma	0.9	1	0.8182
MC1R AND NOT-PCDHGC4	Melanoma	0.9	1	0.8182
MC1R AND NOT-CDH20	Melanoma	0.9	1	0.8182
MC1R AND NOT-VN1R2	Melanoma	0.9	1	0.8182
MC1R AND NOT-MIP	Melanoma	0.9	1	0.8182
MC1R AND NOT-MC2R	Melanoma	0.9	1	0.8182
MC1R AND NOT-HTR5A	Melanoma	0.9	1	0.8182
MC1R AND NOT-OR1Q1	Melanoma	0.9	1	0.8182
MC1R AND NOT-AOC3	Melanoma	0.9	1	0.8182
MC1R AND NOT-GPRC6A	Melanoma	0.9	1	0.8182
MC1R AND NOT-DSCAM	Melanoma	0.9	1	0.8182
C11orf24 AND NOT-TAS2R1	Melanoma	0.9	1	0.8182
MC1R AND NOT-GJD2	Melanoma	0.9	1	0.8182
SLC10A4 AND NOT-KCNJ10	Neuroblastoma	0.994	1	0.9881
SLC10A4 AND NOT-NKAIN2	Neuroblastoma	0.994	1	0.9881
SLC10A4 AND NOT-OPALIN	Neuroblastoma	0.9881	0.9881	0.9881
SLC10A4 AND NOT-TMEM235	Neuroblastoma	0.9881	0.9881	0.9881
SLC10A4 AND NOT-OMG	Neuroblastoma	0.988	1	0.9762
SLC10A4 AND NOT-KCNA1	Neuroblastoma	0.9822	0.9765	0.9881
SLC10A4 AND NOT-MAG	Neuroblastoma	0.9822	0.9765	0.9881
SLC10A4 AND NOT-MOG	Neuroblastoma	0.9822	0.9765	0.9881
SLC10A4 AND NOT-SLC5A11	Neuroblastoma	0.9822	0.9765	0.9881
SLC10A4 AND NOT-HCN2	Neuroblastoma	0.9822	0.9765	0.9881
SLC10A4 AND NOT-GABRB2	Neuroblastoma	0.9822	0.9765	0.9881
SLC10A4 AND NOT-SLC24A2	Neuroblastoma	0.9822	0.9765	0.9881
SLC10A4 AND NOT-OR1Q1	Neuroblastoma	0.9822	0.9765	0.9881
SLC10A4 AND NOT-MLC1	Neuroblastoma	0.9822	0.9765	0.9881
SLC10A4 AND NOT-TTYH2	Neuroblastoma	0.9822	0.9765	0.9881
SLC10A4 AND NOT-KCNA2	Neuroblastoma	0.9818	1	0.9643
SLC10A4 AND NOT-HTR1F	Neuroblastoma	0.994	1	0.9881
SLC10A4 AND NOT-TRPC3	Neuroblastoma	0.982	0.988	0.9762
SLC10A4 AND NOT-SLC13A1	Neuroblastoma	0.9822	0.9765	0.9881
SLC10A4 AND NOT-SLC9B1	Neuroblastoma	0.9765	0.9651	0.9881
SLC10A4 AND NOT-GNRHR	Neuroblastoma	0.9765	0.9651	0.9881
SLC10A4 AND NOT-ADAM7	Neuroblastoma	0.9765	0.9651	0.9881
SLC10A4 AND NOT-MRGPRX2	Neuroblastoma	0.9765	0.9651	0.9881
SLC10A4 AND NOT-XCR1	Neuroblastoma	0.9765	0.9651	0.9881
SLC10A4 AND NOT-GABRA1	Neuroblastoma	0.9762	0.9762	0.9762
SLC10A4 AND NOT-GPRC6A	Neuroblastoma	0.9762	0.9762	0.9762
SLC10A4 AND NOT-KCNU1	Neuroblastoma	0.9762	0.9762	0.9762
SLC10A4 AND NOT-GRIN2B	Neuroblastoma	0.9881	0.9881	0.9881
SLC10A4 AND NOT-OR5P3	Neuroblastoma	0.9822	0.9765	0.9881
SLC10A4 AND NOT-TMPRSS11B	Neuroblastoma	0.9822	0.9765	0.9881
SLC10A4 AND NOT-GRM6	Neuroblastoma	0.9822	0.9765	0.9881
SLC10A4 AND NOT-OR8B2	Neuroblastoma	0.9765	0.9651	0.9881
SLC10A4 AND NOT-LCT	Neuroblastoma	0.9765	0.9651	0.9881
SLC10A4 AND NOT-GLRA3	Neuroblastoma	0.994	1	0.9881
SLC10A4 AND NOT-CACNG6	Neuroblastoma	0.9762	0.9762	0.9762
SLC10A4 AND NOT-ADAM30	Neuroblastoma	0.9708	0.954	0.9881
SLC10A4 AND NOT-ACSL6	Neuroblastoma	0.9708	0.954	0.9881
SLC10A4 AND NOT-SLCO1A2	Neuroblastoma	0.9708	0.954	0.9881
SLC10A4 AND NOT-GPR156	Neuroblastoma	0.9708	0.954	0.9881
SLC10A4 AND NOT-LRRC8E	Neuroblastoma	0.9708	0.954	0.9881
SLC10A4 AND NOT-GRM1	Neuroblastoma	0.9708	0.954	0.9881
SLC10A4 AND NOT-PCDHB1	Neuroblastoma	0.9708	0.954	0.9881
SLC10A4 AND NOT-SLC22A6	Neuroblastoma	0.9708	0.954	0.9881
SLC10A4 AND NOT-CALHM1	Neuroblastoma	0.9708	0.954	0.9881
SLC10A4 AND NOT-SCN10A	Neuroblastoma	0.9708	0.954	0.9881
SLC10A4 AND NOT-SCN1A	Neuroblastoma	0.9708	0.954	0.9881
SLC10A4 AND NOT-GRM3	Neuroblastoma	0.9708	0.954	0.9881
SLC10A4 AND NOT-CD82	Neuroblastoma	0.9704	0.9647	0.9762
SLC10A4 AND NOT-GRID2	Neuroblastoma	0.9704	0.9647	0.9762
SLC10A4 AND NOT-SLC22A16	Neuroblastoma	0.9704	0.9647	0.9762
SLC10A4 AND NOT-CD207	Neuroblastoma	0.9704	0.9647	0.9762
SLC10A4 AND NOT-GABBR2	Neuroblastoma	0.9701	0.9759	0.9643
SLC10A4 AND NOT-PCDH11X	Neuroblastoma	0.9701	0.9759	0.9643
SLC10A4 AND NOT-CLDN16	Neuroblastoma	0.9701	0.9759	0.9643
SLC10A4 AND NOT-HTR3C	Neuroblastoma	0.9701	0.9759	0.9643
SLC10A4 AND NOT-GSG1L	Neuroblastoma	0.9697	0.9877	0.9524
SLC10A4 AND NOT-CSMD3	Neuroblastoma	0.9693	1	0.9405
SLC10A4 AND NOT-HTR5A	Neuroblastoma	0.9765	0.9651	0.9881
SLC10A4 AND NOT-SLC22A11	Neuroblastoma	0.9762	0.9762	0.9762
SLC10A4 AND NOT-MEGF10	Neuroblastoma	0.9651	0.9432	0.9881
SLC10A4 AND NOT-DTNA	Neuroblastoma	0.9651	0.9432	0.9881
SLC10A4 AND NOT-CACNG8	Neuroblastoma	0.9651	0.9432	0.9881
SLC10A4 AND NOT-ROS1	Neuroblastoma	0.9651	0.9432	0.9881

FIG. 12 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
NKAIN1 AND NOT-ITGB3	Neuroblastoma	0.8605	0.8409	0.881
NKAIN1 AND NOT-CLDN7	Neuroblastoma	0.8588	0.8488	0.869
NKAIN1 AND NOT-DNAJB8	Neuroblastoma	0.8555	0.8315	0.881
L1CAM AND NOT-KCNA2	Neuroblastoma	0.8553	0.9559	0.7738
NKAIN1 AND NOT-SSTR5	Neuroblastoma	0.8538	0.8391	0.869
NKAIN1 AND NOT-MUC13	Neuroblastoma	0.8506	0.8222	0.881
NKAIN1 AND NOT-CLDN11	Neuroblastoma	0.85	0.8947	0.8095
ALK AND NOT-PAQR7	Neuroblastoma	0.8553	0.9067	0.8095
ALK AND NOT-SLC2A1	Neuroblastoma	0.8625	0.9079	0.8214
ALK AND NOT-CHRND	Neuroblastoma	0.8679	0.92	0.8214
ALK AND NOT-CLDN17	Neuroblastoma	0.8662	0.9315	0.8095
ALK AND NOT-DCSTAMP	Neuroblastoma	0.859	0.9306	0.7976
ALK AND NOT-OPCML	Neuroblastoma	0.879	0.9452	0.8214
ALK AND NOT-LRRC8E	Neuroblastoma	0.8734	0.9324	0.8214
ALK AND NOT-GABBR2	Neuroblastoma	0.8701	0.9571	0.7976
ALK AND NOT-MLC1	Neuroblastoma	0.8846	0.9583	0.8214
ALK AND NOT-SLC6A13	Neuroblastoma	0.8774	0.9577	0.8095
ALK AND NOT-GPR137B	Neuroblastoma	0.8608	0.9189	0.8095
ALK AND NOT-ADAM29	Neuroblastoma	0.879	0.9452	0.8214
ALK AND NOT-DYSF	Neuroblastoma	0.8625	0.9079	0.8214
ALK AND NOT-OR1C1	Neuroblastoma	0.8625	0.9079	0.8214
ALK AND NOT-GPR37L1	Neuroblastoma	0.8625	0.9079	0.8214
ALK AND NOT-TNFRSF12A	Neuroblastoma	0.8553	0.9067	0.8095
ALK AND NOT-ATP13A5	Neuroblastoma	0.8625	0.9079	0.8214
ALK AND NOT-VN1R2	Neuroblastoma	0.8684	0.9706	0.7857
ALK AND NOT-LYPD1	Neuroblastoma	0.8625	0.9079	0.8214
ALK AND NOT-FAT1	Neuroblastoma	0.8519	0.8846	0.8214
ALK AND NOT-ZDHHC2	Neuroblastoma	0.8553	0.9067	0.8095
ALK AND NOT-SCN10A	Neuroblastoma	0.8679	0.92	0.8214
ALK AND NOT-PHLDB2	Neuroblastoma	0.8625	0.9079	0.8214
ALK AND NOT-PCDH11X	Neuroblastoma	0.8734	0.9324	0.8214
ALK AND NOT-OXTR	Neuroblastoma	0.8625	0.9079	0.8214
ALK AND NOT-EMP3	Neuroblastoma	0.8625	0.9079	0.8214
L1CAM AND NOT-GABRB2	Neuroblastoma	0.8488	0.8295	0.869
NKAIN1 AND NOT-MUC1	Neuroblastoma	0.8488	0.8295	0.869
ALK AND NOT-PPAPDC1B	Neuroblastoma	0.8481	0.9054	0.7976
NKAIN1 AND NOT-RAET1E	Neuroblastoma	0.8475	0.8065	0.8929
L1CAM AND NOT-SLC39A12	Neuroblastoma	0.8475	0.8065	0.8929
L1CAM AND NOT-OMG	Neuroblastoma	0.8462	0.9167	0.7857
NKAIN1 AND NOT-IL2RA	Neuroblastoma	0.8452	0.8452	0.8452
NKAIN1 AND NOT-FCRL2	Neuroblastoma	0.8503	0.8554	0.8452
NKAIN1 AND NOT-PSCA	Neuroblastoma	0.8427	0.7979	0.8929
L1CAM AND NOT-MOG	Neuroblastoma	0.8427	0.7979	0.8929
NKAIN1 AND NOT-CLDN6	Neuroblastoma	0.8421	0.8276	0.8571
FLVCR1 AND NOT-PCYT1A	Neuroblastoma	0.8395	0.8718	0.8095
NKAIN1 AND NOT-TRPM4	Neuroblastoma	0.8395	0.8718	0.8095
NKAIN1 AND NOT-ITGB6	Neuroblastoma	0.838	0.7895	0.8929
NKAIN4 AND NOT-GUCY2C	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-SSTR5	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-ROR1	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-P2RX5	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-RNF43	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-PTK7	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-SLAMF7	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-SDC1	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-SSTR1	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-SSTR2	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-CD79A	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-MUC16	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-CD19	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-CLDN6	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-SSTR4	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-PROM1	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-VTCN1	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-MSLN	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-MET	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-FOLR1	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-SLC34A2	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-CD52	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-RAET1E	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-CLDN7	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-CR2	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-ENG	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-EPHB2	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-ERBB2	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-TPBG	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-SLC39A6	Oligodendroglioma	0.8889	1	0.8

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
SLC10A4 AND NOT-SLC39A12	Neuroblastoma	0.9651	0.9432	0.9881
SLC10A4 AND NOT-SLC28A1	Neuroblastoma	0.9651	0.9432	0.9881
SLC10A4 AND NOT-HTR6	Neuroblastoma	0.9651	0.9432	0.9881
SLC10A4 AND NOT-TRPM3	Neuroblastoma	0.9765	0.9651	0.9881
SLC10A4 AND NOT-CNGA4	Neuroblastoma	0.9651	0.9432	0.9881
SLC10A4 AND NOT-ATP2B2	Neuroblastoma	0.9647	0.9535	0.9762
SLC10A4 AND NOT-GJA10	Neuroblastoma	0.9647	0.9535	0.9762
SLC10A4 AND NOT-SLC12A5	Neuroblastoma	0.9647	0.9535	0.9762
SLC10A4 AND NOT-CLRN1	Neuroblastoma	0.9643	0.9643	0.9643
SLC10A4 AND NOT-HCN4	Neuroblastoma	0.9643	0.9643	0.9643
SLC10A4 AND NOT-GRIN2C	Neuroblastoma	0.9639	0.9756	0.9524
SLC10A4 AND NOT-SLC6A12	Neuroblastoma	0.9639	0.9756	0.9524
SLC10A4 AND NOT-MC2R	Neuroblastoma	0.9639	0.9756	0.9524
SLC10A4 AND NOT-OPRK1	Neuroblastoma	0.9634	0.9875	0.9405
SLC10A4 AND NOT-KCNJ6	Neuroblastoma	0.963	1	0.9286
SLC10A4 AND NOT-SLC12A3	Neuroblastoma	0.9595	0.9326	0.9881
SLC10A4 AND NOT-SEMA4D	Neuroblastoma	0.9595	0.9326	0.9881
SLC10A4 AND NOT-OR2L2	Neuroblastoma	0.9595	0.9326	0.9881
SLC10A4 AND NOT-CLDN17	Neuroblastoma	0.9591	0.9425	0.9762
NKAIN4 AND NOT-AQP11	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-ACE2	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-TRPV4	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-RXFP1	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-PRPH2	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-RHAG	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-EDA2R	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-CDH26	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-SLC5A7	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-ROBO2	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-CEACAM1	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-OPN1SW	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-RYR1	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-SORT1	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-SLC22A23	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-SCN5A	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-ABCD4	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-PTPRS	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-NLGN2	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-PCDH10	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-KCNT1	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-ISLR2	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-STIM2	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-HCN3	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-SLC7A14	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-PTPN2	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-WDR19	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-CADM3	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-PTPRC	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-PTPRM	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-PTPRN	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-PTPRO	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-CX3CL1	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-PCDH19	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-SLC16A2	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-SLC9A5	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-SLC12A1	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-SLC12A4	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-SLC15A2	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-SLC16A1	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-SLC18A3	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-SRR	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-SLCO2A1	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-SLC22A3	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-CYP4F12	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-SMPD1	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-SIGLEC1	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-DST	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-SPINT1	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-BPI	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-BRCA1	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-SLC5A3	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-SLC4A2	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-SDC2	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-ANO3	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-SELE	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-CDH23	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-C10orf54	Oligodendroglioma	0.8889	1	0.8

FIG. 12 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
NKAIN4 AND NOT-EPCAM	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-CD180	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-IL2RA	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-IGF1R	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-STEAP2	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-B4GALNT1	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-EPHA3	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-FAP	Oligodendroglioma	0.8462	1	0.7333
NKAIN4 AND NOT-CD38	Oligodendroglioma	0.8462	1	0.7333
NKAIN4 AND NOT-ERBB4	Oligodendroglioma	0.8462	1	0.7333
KCNQ2 AND NOT-B4GALNT1	Oligodendroglioma	0.8387	0.8125	0.8667
MUC16 AND NOT-STEAP4	Ovarian	0.8	1	0.6667
MUC16 AND NOT-CD74	Ovarian	0.7619	0.8889	0.6667
MUC16 AND NOT-ANTXR2	Ovarian	0.7273	0.8	0.6667
EPHA10 AND NOT-P2RX5	Ovarian	0.7273	0.8	0.6667
EPHA10 AND NOT-CLDN5	Ovarian	0.7826	0.8182	0.75
MUC16 AND NOT-CD44	Ovarian	0.8	1	0.6667
MUC16 AND NOT-PCDH15	Ovarian	0.7	0.875	0.5833
MUC16 AND NOT-PPAPDC1B	Ovarian	0.6957	0.7273	0.6667
MUC16 AND NOT-MRGPRF	Ovarian	0.6667	1	0.5
EPHA10 AND NOT-ENG	Ovarian	0.6667	0.6667	0.6667
MUC16 AND NOT-ALCAM	Ovarian	0.6667	0.6667	0.6667
MUC16 AND NOT-ADAM2	Ovarian	0.6667	1	0.5
EPHA10 AND NOT-AXL	Ovarian	0.6429	0.5625	0.75
EPHA10 AND NOT-SSTR1	Ovarian	0.64	0.6154	0.6667
EPHA10 AND NOT-FCRL2	Ovarian	0.64	0.6154	0.6667
EPHA10 AND NOT-CXCR5	Ovarian	0.6364	0.7	0.5833
MUC16 AND NOT-USH2A	Ovarian	0.6316	0.8571	0.5
EPHA10 AND NOT-RAET1E	Ovarian	0.6429	0.5625	0.75
EPHA10 AND NOT-PTK7	Ovarian	0.6667	0.6	0.75
MUC16 AND NOT-LRIG3	Ovarian	0.6154	0.5714	0.6667
EPHA10 AND NOT-CLDN8	Ovarian	0.6154	0.5714	0.6667
MUC16 AND NOT-CDH9	Ovarian	0.6154	0.5714	0.6667
EPHA10 AND NOT-CSPG4	Ovarian	0.6154	0.5714	0.6667
EPHA10 AND NOT-ULBP1	Ovarian	0.6087	0.6364	0.5833
MUC16 AND NOT-CLDN10	Ovarian	0.6087	0.6364	0.5833
MUC16 AND NOT-SLC6A6	Ovarian	0.6087	0.6364	0.5833
MUC16 AND NOT-ANPEP	Ovarian	0.6316	0.8571	0.5
MUC16 AND NOT-GYPC	Ovarian	0.7273	0.8	0.6667
EPHA10 AND NOT-CD276	Ovarian	0.6207	0.5294	0.75
EPHA10 AND NOT-CD19	Ovarian	0.6923	0.6429	0.75
MUC16 AND NOT-TAS2R1	Ovarian	0.6	0.75	0.5
EPHA10 AND NOT-SSTR2	Ovarian	0.6923	0.6429	0.75
ERBB4 AND NOT-ANO4	Ovarian	0.6087	0.6364	0.5833
EPHA10 AND NOT-CD34	Ovarian	0.6207	0.5294	0.75
KCNU1 AND NOT-CXCR5	Ovarian	0.7273	0.8	0.6667
STEAP2 AND NOT-SLC16A5	Prostate	0.8571	1	0.75
STEAP2 AND NOT-SLC39A2	Prostate	1	1	1
TRPM8 AND NOT-B4GALNT1	Prostate	0.8571	1	0.75
STEAP2 AND NOT-SLC4A8	Prostate	0.8889	0.8	1
TRPM8 AND NOT-NCAM1	Prostate	0.8889	0.8	1
STEAP2 AND NOT-NRG3	Prostate	0.8571	1	0.75
STEAP2 AND NOT-SYT4	Prostate	0.8	0.6667	1
TRPM8 AND NOT-LGR5	Prostate	0.8889	0.8	1
TRPM8 AND NOT-IL3RA	Prostate	0.8571	1	0.75
STEAP2 AND NOT-GRIN2C	Prostate	0.8889	0.8	1
TRPM8 AND NOT-DPEP1	Prostate	0.8571	1	0.75
STEAP1 AND NOT-SLC22A18	Prostate	0.8571	1	0.75
TRPM8 AND NOT-FOLR1	Prostate	0.8571	1	0.75
TRPM8 AND NOT-KDR	Prostate	0.8	0.6667	1
TRPM8 AND NOT-PSCA	Prostate	1	1	1
STEAP1 AND NOT-DYSF	Prostate	0.75	0.75	0.75
STEAP2 AND NOT-SLC17A2	Prostate	0.75	0.75	0.75
STEAP1 AND NOT-CD36	Prostate	0.75	0.75	0.75
STEAP2 AND NOT-SLC9A1	Prostate	0.75	0.75	0.75
STEAP1 AND NOT-PTGER4	Prostate	0.75	0.75	0.75
STEAP2 AND NOT-CALN1	Prostate	0.75	0.75	0.75
TRPM8 AND NOT-EDNRB	Prostate	0.75	0.75	0.75
STEAP2 AND NOT-MLC1	Prostate	0.75	0.75	0.75
TRPM8 AND NOT-MUC16	Prostate	0.75	0.75	0.75
STEAP2 AND NOT-FMNL1	Prostate	1	1	1
STEAP2 AND NOT-GRIN2A	Prostate	0.8571	1	0.75
STEAP1 AND NOT-IL17RA	Prostate	0.75	0.75	0.75
GGTLC1 AND NOT-FOLR1	Prostate	0.8571	1	0.75
STEAP2 AND NOT-GYPC	Prostate	0.7273	0.5714	1
STEAP2 AND NOT-SLC22A18	Prostate	0.7273	0.5714	1
STEAP2 AND NOT-ASTN1	Prostate	0.7273	0.5714	1

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
NKAIN4 AND NOT-DNAJC1	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-SFRP1	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-SGCG	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-CLEC7A	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-SHH	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-SLC26A6	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-SLC1A1	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-SLC4A3	Oligodendroglioma	0.8889	1	0.8
NKAIN4 AND NOT-SLC1A4	Oligodendroglioma	0.8889	1	0.8
KCNK15 AND NOT-THBD	Ovarian	1	1	1
KCNK15 AND NOT-GHR	Ovarian	0.9231	0.8571	1
KCNK15 AND NOT-PROCR	Ovarian	0.96	0.9231	1
KCNK15 AND NOT-KCNK6	Ovarian	1	1	1
KCNK15 AND NOT-ANTXR2	Ovarian	1	1	1
KCNK15 AND NOT-PTGER4	Ovarian	0.8889	0.8	1
KCNK15 AND NOT-GYPC	Ovarian	0.9231	0.8571	1
KCNK15 AND NOT-GPR137B	Ovarian	0.9231	0.8571	1
KCNK15 AND NOT-PHLDB2	Ovarian	0.8571	0.75	1
KCNK15 AND NOT-TTYH2	Ovarian	0.9167	0.9167	0.9167
KCNK15 AND NOT-KCNA4	Ovarian	0.9167	0.9167	0.9167
KCNK15 AND NOT-CD163	Ovarian	0.96	0.9231	1
KCNK15 AND NOT-ADAM2	Ovarian	0.9091	1	0.8333
KCNK15 AND NOT-PTGIS	Ovarian	0.8571	0.75	1
KCNK15 AND NOT-EMP3	Ovarian	0.9091	1	0.8333
KCNK15 AND NOT-DTNA	Ovarian	0.8571	0.75	1
KCNK15 AND NOT-SLC39A8	Ovarian	0.8696	0.9091	0.8333
KCNK15 AND NOT-SCARA5	Ovarian	1	1	1
KCNK15 AND NOT-CD44	Ovarian	0.96	0.9231	1
KCNK15 AND NOT-MTNR1B	Ovarian	0.8571	1	0.75
KCNK15 AND NOT-IL17RA	Ovarian	0.9231	0.8571	1
KCNK15 AND NOT-ATP8B2	Ovarian	0.9231	0.8571	1
KCNK15 AND NOT-BTN3A1	Ovarian	0.8571	0.75	1
KCNK15 AND NOT-LAT	Ovarian	0.9231	0.8571	1
KCNK15 AND NOT-MLNR	Ovarian	0.8182	0.9	0.75
KCNK15 AND NOT-KCNK12	Ovarian	0.8182	0.9	0.75
KCNK15 AND NOT-PCDHA6	Ovarian	0.8182	0.9	0.75
KCNK15 AND NOT-GRID2	Ovarian	0.8182	0.9	0.75
KCNK15 AND NOT-ATP1B4	Ovarian	0.8696	0.9091	0.8333
KCNK15 AND NOT-PLVAP	Ovarian	0.8571	0.75	1
KCNK15 AND NOT-AOC3	Ovarian	0.8571	0.75	1
KCNK15 AND NOT-CD93	Ovarian	0.96	0.9231	1
KCNK15 AND NOT-SELP	Ovarian	1	1	1
KCNK15 AND NOT-CDH11	Ovarian	0.8276	0.7059	1
KCNK15 AND NOT-PCDH11Y	Ovarian	0.8333	0.8333	0.8333
KCNK15 AND NOT-STAB1	Ovarian	0.8	0.6667	1
KCNK15 AND NOT-SLC23A2	Ovarian	0.8	0.6667	1
KCNK15 AND NOT-MRGPRF	Ovarian	0.8	0.6667	1
KCNK15 AND NOT-S1PR1	Ovarian	0.8	0.6667	1
KCNK15 AND NOT-PCDH15	Ovarian	0.8	0.6667	1
KCNK15 AND NOT-PIK3IP1	Ovarian	0.8	0.6667	1
KCNK15 AND NOT-ESAM	Ovarian	0.8	0.6667	1
KCNK15 AND NOT-ASTN1	Ovarian	0.8	0.6667	1
KCNK15 AND NOT-ABCA12	Ovarian	0.9167	0.9167	0.9167
KCNK15 AND NOT-IL15RA	Ovarian	0.9231	0.8571	1
KCNK15 AND NOT-CD74	Ovarian	0.8	0.6667	1
KCNK15 AND NOT-CACNG4	Ovarian	0.7857	0.6875	0.9167
KCNK15 AND NOT-NMBR	Ovarian	0.7857	0.6875	0.9167
KCNK15 AND NOT-LPPR3	Ovarian	0.7826	0.8182	0.75
EPHA10 AND NOT-ITM2A	Ovarian	0.8571	1	0.75
KCNK15 AND NOT-FXYD6	Ovarian	0.8571	0.75	1
KCNK15 AND NOT-CD58	Ovarian	0.7742	0.6316	1
KCNK15 AND NOT-SLC9A1	Ovarian	0.7742	0.6316	1
KCNK15 AND NOT-ATP8A1	Ovarian	0.7742	0.6316	1
KCNK15 AND NOT-TMEM235	Ovarian	0.7742	0.6316	1
KCNK15 AND NOT-SYT4	Ovarian	0.7742	0.6316	1
KCNK15 AND NOT-FAT1	Ovarian	0.7742	0.6316	1
KCNK15 AND NOT-MEGF11	Ovarian	0.7742	0.6316	1
KCNK15 AND NOT-LRRN4	Ovarian	0.7742	0.6316	1
KCNK15 AND NOT-PPAPDC1A	Ovarian	0.7742	0.6316	1
KCNK15 AND NOT-MMP24	Ovarian	0.7742	0.6316	1
KCNK15 AND NOT-ATP7A	Ovarian	0.7742	0.6316	1
KCNK15 AND NOT-BTN3A3	Ovarian	0.7742	0.6316	1
KCNK15 AND NOT-SLC43A1	Ovarian	0.7742	0.6316	1
KCNK15 AND NOT-PAQR7	Ovarian	0.7742	0.6316	1
KCNK15 AND NOT-MSMO1	Ovarian	0.7742	0.6316	1
KCNK15 AND NOT-CNTNAP4	Ovarian	0.7742	0.6316	1
KCNK15 AND NOT-SLC16A5	Ovarian	0.7742	0.6316	1

FIG. 12 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
STEAP2 AND NOT-EPHA2	Prostate	0.7273	0.5714	1
STEAP2 AND NOT-CLDN10	Prostate	0.7273	0.5714	1
STEAP2 AND NOT-CSMD3	Prostate	0.7273	0.5714	1
TRPM8 AND NOT-SDC1	Prostate	0.7273	0.5714	1
TRPM8 AND NOT-TNFRSF17	Prostate	0.7273	0.5714	1
TRPM8 AND NOT-L1CAM	Prostate	0.7273	0.5714	1
STEAP2 AND NOT-GUCY2D	Prostate	0.7273	0.5714	1
STEAP2 AND NOT-PAQR8	Prostate	0.7273	0.5714	1
TRPM8 AND NOT-CD72	Prostate	0.7273	0.5714	1
STEAP2 AND NOT-LAT	Prostate	0.7273	0.5714	1
STEAP2 AND NOT-PAQR7	Prostate	0.7273	0.5714	1
STEAP2 AND NOT-DCC	Prostate	0.7273	0.5714	1
STEAP2 AND NOT-TTYH2	Prostate	0.7273	0.5714	1
STEAP2 AND NOT-CD36	Prostate	0.7273	0.5714	1
STEAP2 AND NOT-SEMA4D	Prostate	0.7273	0.5714	1
FAP AND NOT-NFASC	Sarcoma	0.8667	1	0.7647
FAP AND NOT-AOC3	Sarcoma	0.8276	1	0.7059
FAP AND NOT-CLDN10	Sarcoma	0.8276	1	0.7059
FAP AND NOT-PAQR7	Sarcoma	0.8276	1	0.7059
FAP AND NOT-ATP13A5	Sarcoma	0.8276	1	0.7059
FAP AND NOT-NRG3	Sarcoma	0.8667	1	0.7647
FAP AND NOT-NKAIN1	Sarcoma	0.7857	1	0.6471
FAP AND NOT-GPR6	Sarcoma	0.7857	1	0.6471
FAP AND NOT-SELP	Sarcoma	0.7407	1	0.5882
FAP AND NOT-KCNA2	Sarcoma	0.7407	1	0.5882
FAP AND NOT-GRIN1	Sarcoma	0.7407	1	0.5882
FAP AND NOT-DGKE	Sarcoma	0.7857	1	0.6471
FAP AND NOT-SLC1A6	Sarcoma	0.8276	1	0.7059
FAP AND NOT-SLC16A5	Sarcoma	0.7407	1	0.5882
FAP AND NOT-AGTR2	Sarcoma	0.8667	1	0.7647
FAP AND NOT-GUCY2F	Sarcoma	0.8276	1	0.7059
FAP AND NOT-GGTLC1	Sarcoma	0.7857	1	0.6471
FAP AND NOT-RXFP3	Sarcoma	0.7857	1	0.6471
FAP AND NOT-CD1A	Sarcoma	0.7407	1	0.5882
FAP AND NOT-KCNK6	Sarcoma	0.7857	1	0.6471
FAP AND NOT-AQP6	Sarcoma	0.7407	1	0.5882
FAP AND NOT-LHFPL5	Sarcoma	0.8667	1	0.7647
FAP AND NOT-SLC22A9	Sarcoma	0.7407	1	0.5882
FAP AND NOT-LPPR3	Sarcoma	0.8667	1	0.7647
FAP AND NOT-MMP24	Sarcoma	0.7407	1	0.5882
FAP AND NOT-MTNR1B	Sarcoma	0.8667	1	0.7647
FAP AND NOT-CACNG4	Sarcoma	0.8276	1	0.7059
FAP AND NOT-MRGPRF	Sarcoma	0.7857	1	0.6471
STAB1 AND NOT-CLDN6	Sarcoma	0.7273	0.75	0.7059
FAP AND NOT-XCR1	Sarcoma	0.8276	1	0.7059
FAP AND NOT-GPR135	Sarcoma	0.7407	1	0.5882
FAP AND NOT-ADAM30	Sarcoma	0.8276	1	0.7059
FAP AND NOT-GALR1	Sarcoma	0.7857	1	0.6471
FAP AND NOT-OR8D1	Sarcoma	0.8667	1	0.7647
FAP AND NOT-CLDN19	Sarcoma	0.7857	1	0.6471
FAP AND NOT-CDH9	Sarcoma	0.8667	1	0.7647
FAP AND NOT-ABCG8	Sarcoma	0.8276	1	0.7059
FAP AND NOT-OR3A1	Sarcoma	0.8667	1	0.7647
FAP AND NOT-KIRREL2	Sarcoma	0.7857	1	0.6471
FAP AND NOT-KCNQ2	Sarcoma	0.7407	1	0.5882
FAP AND NOT-ATP2B2	Sarcoma	0.7407	1	0.5882
FAP AND NOT-SLC17A2	Sarcoma	0.7857	1	0.6471
FAP AND NOT-KCNJ10	Sarcoma	0.7857	1	0.6471
FAP AND NOT-TSPAN16	Sarcoma	0.8667	1	0.7647
FAP AND NOT-SLC6A11	Sarcoma	0.8276	1	0.7059
FAP AND NOT-SLC5A11	Sarcoma	0.8276	1	0.7059
FAP AND NOT-KCNA7	Sarcoma	0.7407	1	0.5882
FAP AND NOT-CLCN1	Sarcoma	0.7407	1	0.5882
FAP AND NOT-CATSPER3	Sarcoma	0.7407	1	0.5882
FAP AND NOT-VN1R2	Sarcoma	0.7407	1	0.5882
FAP AND NOT-CDH17	Sarcoma	0.8276	1	0.7059
FAP AND NOT-KREMEN2	Sarcoma	0.7857	1	0.6471
FAP AND NOT-SLC22A6	Sarcoma	0.7857	1	0.6471
FAP AND NOT-GRM3	Sarcoma	0.7407	1	0.5882
FAP AND NOT-UMOD	Sarcoma	0.7857	1	0.6471
FAP AND NOT-HRH3	Sarcoma	0.7407	1	0.5882
FAP AND NOT-RGSL1	Sarcoma	0.8667	1	0.7647
FAP AND NOT-PVRL1	Sarcoma	0.8667	1	0.7647
FAP AND NOT-IGDCC3	Sarcoma	0.8276	1	0.7059
FAP AND NOT-CD207	Sarcoma	0.8276	1	0.7059
FAP AND NOT-MLC1	Sarcoma	0.8667	1	0.7647
FAP AND NOT-OR5P3	Sarcoma	0.7857	1	0.6471

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
KCNK15 AND NOT-CLDN15	Ovarian	0.7742	0.6316	1
KCNK15 AND NOT-GABRG1	Ovarian	0.7742	0.6316	1
KCNK15 AND NOT-SMPD2	Ovarian	0.7742	0.6316	1
KCNK15 AND NOT-ATP8A2	Ovarian	0.7742	0.6316	1
KCNK15 AND NOT-DPP10	Ovarian	0.7742	0.6316	1
KCNK15 AND NOT-GABRB2	Ovarian	0.7742	0.6316	1
KCNK15 AND NOT-NKAIN2	Ovarian	0.7742	0.6316	1
KCNK15 AND NOT-GABRA2	Ovarian	0.7742	0.6316	1
KCNK15 AND NOT-ALCAM	Ovarian	0.7742	0.6316	1
KCNK15 AND NOT-LRRC8E	Ovarian	0.7742	0.6316	1
KCNK15 AND NOT-NPHS2	Ovarian	0.7742	0.6316	1
KCNK15 AND NOT-OR10J1	Ovarian	0.7742	0.6316	1
KCNK15 AND NOT-CD36	Ovarian	0.7742	0.6316	1
KCNK15 AND NOT-SLC22A18	Ovarian	0.8333	0.8333	0.8333
KCNK15 AND NOT-ERVW-1	Ovarian	0.7692	0.7143	0.8333
KCNK15 AND NOT-TRPM1	Ovarian	0.7692	0.7143	0.8333
KCNK15 AND NOT-LRFN2	Ovarian	0.7692	0.7143	0.8333
CSPG5 AND NOT-TSPAN5	Ovarian	0.8	1	0.6667
CSPG5 AND NOT-SLC6A1	Ovarian	0.7619	0.8889	0.6667
KCNK15 AND NOT-XCR1	Ovarian	0.7586	0.6471	0.9167
KCNK15 AND NOT-CLSTN3	Ovarian	0.7586	0.6471	0.9167
KCNK15 AND NOT-TAS1R1	Ovarian	0.7586	0.6471	0.9167
KCNK15 AND NOT-USH2A	Ovarian	0.7586	0.6471	0.9167
KCNK15 AND NOT-PAQR8	Ovarian	0.8696	0.9091	0.8333
KCNK15 AND NOT-NFASC	Ovarian	0.75	0.75	0.75
KCNK15 AND NOT-ANO4	Ovarian	0.75	0.75	0.75
KCNK15 AND NOT-GGTLC1	Ovarian	0.75	0.75	0.75
EPHA10 AND NOT-STIM1	Ovarian	0.8571	1	0.75
TRPM8 AND NOT-SLC2A5	Prostate	1	1	1
OR51E2 AND NOT-IL17RA	Prostate	0.8889	0.8	1
OR51E2 AND NOT-EMP3	Prostate	0.8889	0.8	1
OR51E2 AND NOT-CD36	Prostate	0.8889	0.8	1
TRPM8 AND NOT-FADS2	Prostate	0.8889	0.8	1
OR51E2 AND NOT-CHRNA2	Prostate	0.8889	0.8	1
OR51E2 AND NOT-SLC16A5	Prostate	0.8889	0.8	1
OR51E2 AND NOT-GABRA1	Prostate	0.8889	0.8	1
OR51E2 AND NOT-SEMA4D	Prostate	0.8889	0.8	1
OR51E2 AND NOT-GUCY2D	Prostate	0.8889	0.8	1
OR51E2 AND NOT-SLC22A18	Prostate	0.8889	0.8	1
OR51E2 AND NOT-SYT4	Prostate	0.8889	0.8	1
OR51E2 AND NOT-SLC5A2	Prostate	0.8889	0.8	1
OR51E2 AND NOT-PTGER4	Prostate	0.8889	0.8	1
OR51E2 AND NOT-TNFRSF10B	Prostate	0.8889	0.8	1
OR51E2 AND NOT-CDH10	Prostate	0.8889	0.8	1
OR51E2 AND NOT-GYPC	Prostate	0.8889	0.8	1
OR51E2 AND NOT-SHISA9	Prostate	0.8889	0.8	1
TRPM8 AND NOT-FZD1	Prostate	0.8889	0.8	1
OR51E2 AND NOT-STAB1	Prostate	0.8889	0.8	1
OR51E2 AND NOT-EPHA2	Prostate	0.8889	0.8	1
TRPM8 AND NOT-SGCD	Prostate	1	1	1
OR51E2 AND NOT-DYSF	Prostate	0.8889	0.8	1
TRPM8 AND NOT-SLC52A3	Prostate	0.8889	0.8	1
OR51E2 AND NOT-PAQR8	Prostate	0.8889	0.8	1
OR51E2 AND NOT-NFASC	Prostate	0.8889	0.8	1
OR51E2 AND NOT-BEST3	Prostate	0.8889	0.8	1
OR51E2 AND NOT-CD207	Prostate	0.8889	0.8	1
OR51E2 AND NOT-CNTNAP2	Prostate	0.8889	0.8	1
OR51E2 AND NOT-SLC4A8	Prostate	0.8889	0.8	1
OR51E2 AND NOT-LAT	Prostate	0.8889	0.8	1
OR51E2 AND NOT-TNFRSF12A	Prostate	0.8889	0.8	1
OR51E2 AND NOT-NOX1	Prostate	0.8889	0.8	1
OR51E2 AND NOT-CLDN10	Prostate	0.8889	0.8	1
OR51E2 AND NOT-KCNJ10	Prostate	0.8889	0.8	1
OR51E2 AND NOT-BTN3A1	Prostate	0.8889	0.8	1
OR51E2 AND NOT-SLC6A6	Prostate	0.8889	0.8	1
OR51E2 AND NOT-FMNL1	Prostate	1	1	1
TRPM8 AND NOT-LSAMP	Prostate	1	1	1
OR51E2 AND NOT-SLC39A2	Prostate	1	1	1
OR51E2 AND NOT-SCN8A	Prostate	0.8889	0.8	1
OR51E2 AND NOT-PCDH8	Prostate	0.8889	0.8	1
OR51E2 AND NOT-BST2	Prostate	0.8889	0.8	1
OR51E2 AND NOT-TTYH2	Prostate	0.8889	0.8	1
OR51E2 AND NOT-CEACAM7	Prostate	0.8889	0.8	1
OR51E2 AND NOT-PIK3IP1	Prostate	0.8889	0.8	1
TRPM8 AND NOT-IL6ST	Prostate	0.8571	1	0.75
TRPM8 AND NOT-ANTXR1	Prostate	0.8571	1	0.75
TRPM8 AND NOT-IL6R	Prostate	0.8571	1	0.75

FIG. 12 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
FAP AND NOT-GRM7	Sarcoma	0.6923	1	0.5294
FAP AND NOT-SEZ6	Sarcoma	0.6923	1	0.5294
FAP AND NOT-UPK3A	Sarcoma	0.6923	1	0.5294
FAP AND NOT-KCNK4	Sarcoma	0.6923	1	0.5294
FAP AND NOT-HCN4	Sarcoma	0.6923	1	0.5294
FAP AND NOT-FXYD6	Sarcoma	0.6923	1	0.5294
FAP AND NOT-LAMP5	Sarcoma	0.6923	1	0.5294
FAP AND NOT-CACNG6	Sarcoma	0.8276	1	0.7059
FAP AND NOT-KCNA10	Sarcoma	0.6923	1	0.5294
FAP AND NOT-GABBR2	Sarcoma	0.6923	1	0.5294
FAP AND NOT-GRM5	Sarcoma	0.8276	1	0.7059
FAP AND NOT-GRM1	Sarcoma	0.6923	1	0.5294
FAP AND NOT-PCDHB1	Sarcoma	0.8276	1	0.7059
FAP AND NOT-NPHS1	Sarcoma	0.8276	1	0.7059
FAP AND NOT-CACNG7	Sarcoma	0.7857	1	0.6471
FAP AND NOT-KCNV1	Sarcoma	0.6923	1	0.5294
FAP AND NOT-KCNK16	Sarcoma	0.7407	1	0.5882
FAP AND NOT-ADAM2	Sarcoma	0.8276	1	0.7059
FAP AND NOT-EPHA8	Sarcoma	0.8276	1	0.7059
FAP AND NOT-ATP6V0A4	Sarcoma	0.8276	1	0.7059
FAP AND NOT-SLC4A5	Sarcoma	0.8276	1	0.7059
FAP AND NOT-SLC23A2	Sarcoma	0.7857	1	0.6471
FAP AND NOT-KCNH1	Sarcoma	0.7857	1	0.6471
FAP AND NOT-TMPRSS11E	Sarcoma	0.7857	1	0.6471
FAP AND NOT-GPR37L1	Sarcoma	0.8276	1	0.7059
FAP AND NOT-PCDH11Y	Sarcoma	0.6923	1	0.5294
FAP AND NOT-BEST2	Sarcoma	0.7407	1	0.5882
FAP AND NOT-SLC34A1	Sarcoma	0.8276	1	0.7059
FAP AND NOT-KIAA0319	Sarcoma	0.8276	1	0.7059
CLDN18 AND NOT-AGER	Stomach	0.75	0.7333	0.7674
CLDN18 AND NOT-EMP3	Stomach	0.7609	0.7143	0.814
CLDN18 AND NOT-ROS1	Stomach	0.7416	0.7174	0.7674
CLDN18 AND NOT-THBD	Stomach	0.7356	0.7273	0.7442
CLDN18 AND NOT-CD36	Stomach	0.7294	0.7381	0.7209
CLDN18 AND NOT-ICAM1	Stomach	0.75	0.7333	0.7674
MUC13 AND NOT-SLC22A5	Stomach	0.8889	0.9474	0.8372
CLDN18 AND NOT-S1PR1	Stomach	0.7692	0.7292	0.814
CLDN18 AND NOT-AOC3	Stomach	0.7711	0.8	0.7442
CLDN18 AND NOT-FXYD6	Stomach	0.7143	0.7317	0.6977
CDH17 AND NOT-CLDN8	Stomach	0.6964	0.5652	0.907
CLDN18 AND NOT-LAPTM5	Stomach	0.7391	0.6939	0.7907
CLDN18 AND NOT-ESAM	Stomach	0.6905	0.7073	0.6744
CLDN18 AND NOT-SELP	Stomach	0.7253	0.6875	0.7674
CLDN18 AND NOT-GYPC	Stomach	0.6966	0.6739	0.7209
MUC17 AND NOT-ENPP3	Stomach	0.8378	1	0.7209
CLDN18 AND NOT-SLC17A3	Stomach	0.6829	0.7179	0.6512
CLDN18 AND NOT-STEAP4	Stomach	0.6744	0.6744	0.6744
CLDN18 AND NOT-CD163	Stomach	0.766	0.7059	0.8372
CLDN18 AND NOT-DYSF	Stomach	0.7527	0.7	0.814
CDH17 AND NOT-ENPP3	Stomach	0.6726	0.5429	0.8837
MUC17 AND NOT-CD160	Stomach	0.7714	1	0.6279
MUC13 AND NOT-BEST2	Stomach	0.9762	1	0.9535
CLDN18 AND NOT-SLC39A8	Stomach	0.6667	0.6591	0.6744
MUC13 AND NOT-SORL1	Stomach	0.675	0.7297	0.6279
MUC13 AND NOT-SLC30A10	Stomach	0.9438	0.913	0.9767
MUC17 AND NOT-GUCY2C	Stomach	0.7164	1	0.5581
MUC13 AND NOT-SLC22A18	Stomach	0.7436	0.8286	0.6744
CLDN18 AND NOT-CD93	Stomach	0.6585	0.6923	0.6279
MUC17 AND NOT-ABCA5	Stomach	0.7143	0.9259	0.5814
CLDN18 AND NOT-CATSPERD	Stomach	0.766	0.7059	0.8372
NOT-ATP8B1 AND ERBB2	Ependymoma	0.6667	0.5	1
NOT-PHLDB2 AND ROR1	Ependymoma	1	1	1
NOT-SLC6A6 AND SLC39A6	Ependymoma	0.6667	0.5	1
NOT-SLC6A6 AND SSTR3	Ependymoma	0.6667	0.5	1
NOT-SLC6A6 AND SSTR5	Ependymoma	1	1	1
NOT-CD36 AND CLDN18	Esophagus	0.6	1	0.4286
NOT-IL11RA AND PROCR	Esophagus	0.7333	0.6875	0.7857
NOT-IL11RA AND IL15RA	Esophagus	0.75	0.9	0.6429
NOT-CD52 AND LAPTM5	Glioblastoma	0.7429	0.7222	0.7647
NOT-STEAP4 AND ITGAV	Glioblastoma	0.619	0.52	0.7647
NOT-RNF43 AND VANGL1	Leiomyosarcoma	0.8889	0.8571	0.9231
NOT-EPCAM AND VANGL1	Leiomyosarcoma	0.8966	0.8125	1
NOT-ERBB3 AND FAT1	Leiomyosarcoma	0.7826	0.9	0.6923
NOT-ERBB3 AND CDH11	Leiomyosarcoma	0.6957	0.8	0.6154
NOT-ERBB3 AND VANGL1	Leiomyosarcoma	0.8462	0.8462	0.8462
NOT-IL20RA AND VANGL1	Leiomyosarcoma	0.64	0.6667	0.6154
NOT-CLDN10 AND CD276	Leiomyosarcoma	0.8	0.8333	0.7692

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
OR51E1 AND NOT-CD36	Prostate	0.8571	1	0.75
OR51E2 AND NOT-UPK3A	Prostate	0.8889	0.8	1
OR51E2 AND NOT-CSPG5	Prostate	0.8889	0.8	1
OR51E2 AND NOT-UMOD	Prostate	0.8571	1	0.75
TRPM8 AND NOT-TGOLN2	Prostate	0.8571	1	0.75
OR51E2 AND NOT-GRIN2A	Prostate	0.8571	1	0.75
TRPM8 AND NOT-KL	Prostate	0.8571	1	0.75
TRPM8 AND NOT-PCDH9	Prostate	0.8571	1	0.75
OR51E2 AND NOT-NRG3	Prostate	0.8571	1	0.75
OR51E1 AND NOT-DYSF	Prostate	0.8571	1	0.75
OR51E2 AND NOT-OR2C3	Prostate	0.8571	1	0.75
TRPM8 AND NOT-KCNK3	Prostate	0.8571	1	0.75
TRPM8 AND NOT-PTPRF	Prostate	0.8571	1	0.75
TRPM8 AND NOT-PRRT2	Prostate	0.8571	1	0.75
UPK3A AND NOT-SGCD	Prostate	0.8571	1	0.75
OR51E1 AND NOT-IL17RA	Prostate	0.8571	1	0.75
OR51E2 AND NOT-GRM1	Prostate	0.8889	0.8	1
OR51E2 AND NOT-HTR1E	Prostate	0.8889	0.8	1
OR51E2 AND NOT-HTR1D	Prostate	0.8889	0.8	1
OR51E2 AND NOT-VN1R2	Prostate	0.8889	0.8	1
OR51E2 AND NOT-CLDN16	Prostate	0.8889	0.8	1
OR51E2 AND NOT-MDGA2	Prostate	0.8889	0.8	1
OR51E2 AND NOT-SLC5A8	Prostate	0.8889	0.8	1
OR51E2 AND NOT-GRM3	Prostate	0.8889	0.8	1
OR51E2 AND NOT-OR1Q1	Prostate	0.8889	0.8	1
OR51E2 AND NOT-GRM5	Prostate	0.8889	0.8	1
OR51E2 AND NOT-GRM4	Prostate	0.8889	0.8	1
OR51E2 AND NOT-PCDHB1	Prostate	0.8889	0.8	1
OR51E2 AND NOT-GRM6	Prostate	0.8889	0.8	1
OR51E2 AND NOT-CHRND	Prostate	0.8889	0.8	1
OR51E2 AND NOT-GRM7	Prostate	0.8889	0.8	1
OR51E2 AND NOT-ERVW-1	Prostate	0.8889	0.8	1
OR51E2 AND NOT-CYP4A11	Prostate	0.8889	0.8	1
OR51E2 AND NOT-KCNU1	Prostate	0.8889	0.8	1
OR51E2 AND NOT-GRM8	Prostate	0.8889	0.8	1
OR51E2 AND NOT-GUCY2F	Prostate	0.8889	0.8	1
OR51E2 AND NOT-CDH8	Prostate	0.8889	0.8	1
OR51E2 AND NOT-CHRNB4	Prostate	0.8889	0.8	1
OR51E2 AND NOT-HTR2C	Prostate	0.8889	0.8	1
OR51E2 AND NOT-KCND2	Prostate	0.8889	0.8	1
OR51E2 AND NOT-HTR5A	Prostate	0.8889	0.8	1
OR51E2 AND NOT-KCNC1	Prostate	0.8889	0.8	1
OR51E2 AND NOT-SLC10A2	Prostate	0.8889	0.8	1
OR51E2 AND NOT-SLC10A1	Prostate	0.8889	0.8	1
OR51E2 AND NOT-KCNH1	Prostate	0.8889	0.8	1
OR51E2 AND NOT-KCNJ6	Prostate	0.8889	0.8	1
OR51E2 AND NOT-KCNJ9	Prostate	0.8889	0.8	1
OR51E2 AND NOT-GLRA3	Prostate	0.8889	0.8	1
OR51E2 AND NOT-TRPM3	Prostate	0.8889	0.8	1
OR51E2 AND NOT-SLC8A3	Prostate	0.8889	0.8	1
OR51E2 AND NOT-KCNQ2	Prostate	0.8889	0.8	1
TGFBI AND NOT-F11R	Sarcoma	0.9375	1	0.8824
TGFBI AND NOT-AQP3	Sarcoma	0.7857	1	0.6471
CNTNAP1 AND NOT-NFASC	Sarcoma	0.7879	0.8125	0.7647
DDR2 AND NOT-AOC3	Sarcoma	0.7586	0.9167	0.6471
ATP8B2 AND NOT-ABCB1	Sarcoma	0.8	0.9231	0.7059
OSMR AND NOT-STEAP4	Sarcoma	0.8	0.9231	0.7059
OSMR AND NOT-NFASC	Sarcoma	0.8276	1	0.7059
ADAM12 AND NOT-SELP	Sarcoma	0.7333	0.8462	0.6471
ADAM12 AND NOT-NFASC	Sarcoma	0.7692	0.6818	0.8824
DDR2 AND NOT-KCNA2	Sarcoma	0.7407	1	0.5882
OSMR AND NOT-AOC3	Sarcoma	0.8485	0.875	0.8235
OSMR AND NOT-SLC16A5	Sarcoma	0.8	0.9231	0.7059
STAB1 AND NOT-GDPD2	Sarcoma	0.7742	0.8571	0.7059
ADAM12 AND NOT-SLC16A5	Sarcoma	0.7222	0.6842	0.7647
STAB1 AND NOT-ABCB1	Sarcoma	0.7742	0.8571	0.7059
ADAM12 AND NOT-PAQR7	Sarcoma	0.7805	0.6667	0.9412
DDR2 AND NOT-MRAP	Sarcoma	0.8276	1	0.7059
STAB1 AND NOT-BPI	Sarcoma	0.8276	1	0.7059
OSMR AND NOT-SELP	Sarcoma	0.7143	0.9091	0.5882
STAB1 AND NOT-CLEC1B	Sarcoma	0.7407	1	0.5882
OSMR AND NOT-CLDN10	Sarcoma	0.7143	0.9091	0.5882
STAB1 AND NOT-ART4	Sarcoma	0.8276	1	0.7059
STAB1 AND NOT-KCNA3	Sarcoma	0.8276	1	0.7059
MMP14 AND NOT-PAQR7	Sarcoma	0.7407	1	0.5882
MMP14 AND NOT-CLDN19	Sarcoma	0.7407	1	0.5882
ADAM12 AND NOT-CLDN10	Sarcoma	0.8125	0.8667	0.7647

FIG. 12 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
NOT-ERBB3 AND LRIG3	Leiomyosarcoma	0.7619	1	0.6154
NOT-ERBB3 AND FCAMR	Leiomyosarcoma	0.6154	0.6154	0.6154
NOT-ERBB3 AND PRND	Leiomyosarcoma	0.7143	0.6667	0.7692
NOT-ERBB3 AND MOG	Leiomyosarcoma	0.6061	0.5	0.7692
NOT-CLDN10 AND AXL	Leiomyosarcoma	0.6957	0.8	0.6154
NOT-ERBB3 AND SLC39A8	AML	0.834	0.809	0.8606
NOT-STEAP1 AND SLC39A8	AML	0.8208	0.7948	0.8486
NOT-MET AND SLC39A8	AML	0.8131	0.7873	0.8406
NOT-ITGAV AND P2RX4	AML	0.8127	0.7668	0.8645
NOT-CLDN12 AND P2RX4	AML	0.8112	0.7641	0.8645
NOT-EDNRB AND SLC43A1	AML	0.8085	0.8676	0.757
NOT-KDR AND SLC43A1	AML	0.8075	0.8502	0.7689
NOT-ITGAV AND SLC39A8	AML	0.8043	0.863	0.753
NOT-GPR137B AND SLC7A5	AML	0.8031	0.7863	0.8207
NOT-ITGAV AND CD44	AML	0.8015	0.744	0.8685
NOT-ITGAV AND PTGER4	AML	0.8015	0.7383	0.8765
NOT-CLDN12 AND SLC39A8	AML	0.7958	0.8341	0.761
NOT-ITGAV AND EMP3	AML	0.7928	0.7237	0.8765
NOT-ITGAV AND SLC30A1	AML	0.7893	0.7152	0.8805
NOT-AXL AND SLC39A8	AML	0.7885	0.8818	0.7131
NOT-CLDN12 AND SLC30A1	AML	0.7883	0.6939	0.9124
NOT-GPNMB AND SLC43A1	AML	0.7867	0.8894	0.7052
NOT-AXL AND SLC43A1	AML	0.7804	0.8394	0.7291
NOT-ITGAV AND FMNL1	AML	0.7794	0.7042	0.8725
NOT-EDNRB AND P2RX4	AML	0.7745	0.6565	0.9442
NOT-ITGAV AND SLCO1B3	AML	0.774	0.7086	0.8526
NOT-EDNRB AND SLC39A8	AML	0.772	0.8906	0.6813
NOT-CDH11 AND SLC7A5	AML	0.7709	0.6387	0.9721
NOT-ITGAV AND VAMP8	AML	0.77	0.6842	0.8805
NOT-AXL AND SLC30A1	AML	0.77	0.634	0.9801
NOT-ITGAV AND ZDHHC2	AML	0.769	0.703	0.8486
NOT-ITGAV AND SLC43A1	AML	0.7682	0.8326	0.7131
NOT-EPCAM AND VANGL1	Liposarcoma	0.6296	0.9444	0.4722
NOT-SORL1 AND EPHB2	Liposarcoma	0.6452	0.7692	0.5556
NOT-FAT1 AND GPNMB	B-Cell Diffuse	0.6757	0.6757	0.6757
NOT-ATP8B1 AND GPNMB	B-Cell Diffuse	0.6667	0.7188	0.6216
NOT-LRIG3 AND GPNMB	B-Cell Diffuse	0.64	0.6316	0.6486
NOT-EMP3 AND IL2RA	Ovarian	0.6667	1	0.5
NOT-KCNK12 AND CLDN9	Ovarian	0.6	0.75	0.5
NOT-GRID2 AND RNF43	Ovarian	0.6667	1	0.5
NOT-EMP3 AND RNF43	Ovarian	0.7368	1	0.5833
NOT-RNF43 AND VANGL1	Sarcoma	0.7742	0.8571	0.7059
NOT-EPCAM AND VANGL1	Sarcoma	0.7895	0.7143	0.8824
NOT-IL20RA AND VANGL1	Sarcoma	0.7059	0.7059	0.7059
VANGL2 AND NOT-SDC1	Astrocytoma	0.8298	0.8125	0.8478
VANGL2 AND NOT-TPBG	Astrocytoma	0.809	0.8372	0.7826
PTPRZ1 AND NOT-STEAP2	Astrocytoma	0.8039	0.7321	0.8913
PTPRZ1 AND NOT-RNF43	Astrocytoma	0.7921	0.7273	0.8696
VANGL2 AND NOT-EPCAM	Astrocytoma	0.8211	0.7959	0.8478
PTPRZ1 AND NOT-SDC1	Astrocytoma	0.7736	0.6833	0.8913
VANGL2 AND NOT-CLDN1	Astrocytoma	0.7723	0.7091	0.8478
VANGL2 AND NOT-TRPM4	Astrocytoma	0.78	0.7222	0.8478
PTPRZ1 AND NOT-EPCAM	Astrocytoma	0.7593	0.6613	0.8913
PTPRZ1 AND NOT-LGR5	Astrocytoma	0.7547	0.6667	0.8696
MMP16 AND NOT-EPCAM	Astrocytoma	0.7532	0.9355	0.6304
PTPRZ1 AND NOT-TPBG	Astrocytoma	0.7525	0.6909	0.8261
VANGL2 AND NOT-ERBB2	Astrocytoma	0.7473	0.7556	0.7391
PTPRZ1 AND NOT-CLDN1	Astrocytoma	0.7455	0.6406	0.8913
PTPRZ1 AND NOT-PMEL	Astrocytoma	0.7455	0.6406	0.8913
PTPRZ1 AND NOT-TRPM4	Astrocytoma	0.7455	0.6406	0.8913
PTPRZ1 AND NOT-STEAP1	Astrocytoma	0.7455	0.6406	0.8913
PTPRZ1 AND NOT-CD52	Astrocytoma	0.7455	0.6406	0.8913
PTPRZ1 AND NOT-CLDN23	Astrocytoma	0.7455	0.6406	0.8913
MMP16 AND NOT-B4GALNT1	Astrocytoma	0.7368	0.9333	0.6087
MMP16 AND NOT-SDC1	Astrocytoma	0.7342	0.8788	0.6304
PTPRZ1 AND NOT-SLC39A6	Astrocytoma	0.7339	0.6349	0.8696
PTPRZ1 AND NOT-ERBB2	Astrocytoma	0.7222	0.629	0.8478
PTPRZ1 AND NOT-CD34	Astrocytoma	0.7222	0.629	0.8478
PTPRZ1 AND NOT-IL11RA	Astrocytoma	0.7222	0.629	0.8478
PTPRZ1 AND NOT-ERBB3	Astrocytoma	0.7222	0.629	0.8478
MMP16 AND NOT-TPBG	Astrocytoma	0.7179	0.875	0.6087
MMP16 AND NOT-MET	Astrocytoma	0.716	0.8286	0.6304
MMP16 AND NOT-IGF1R	Astrocytoma	0.716	0.8286	0.6304
MMP16 AND NOT-FAP	Astrocytoma	0.716	0.8286	0.6304
TSPAN11 AND NOT-CLDN1	Astrocytoma	0.6923	0.8438	0.587
VANGL2 AND NOT-RNF43	Astrocytoma	0.6923	0.8438	0.587
VANGL2 AND NOT-CD34	Astrocytoma	0.7789	0.7551	0.8043

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
STAB1 AND NOT-TSPAN7	Sarcoma	0.7857	1	0.6471
STAB1 AND NOT-NPY1R	Sarcoma	0.8276	1	0.7059
STAB1 AND NOT-LSR	Sarcoma	0.7857	1	0.6471
STAB1 AND NOT-NAALAD2	Sarcoma	0.7097	0.7857	0.6471
OSMR AND NOT-GGTLC1	Sarcoma	0.8387	0.9286	0.7647
STAB1 AND NOT-KCNC4	Sarcoma	0.7742	0.8571	0.7059
STAB1 AND NOT-ICAM4	Sarcoma	0.7333	0.8462	0.6471
STAB1 AND NOT-SLC6A2	Sarcoma	0.7407	1	0.5882
TGFBI AND NOT-SPINT1	Sarcoma	0.9091	0.9375	0.8824
STAB1 AND NOT-PCDHA10	Sarcoma	0.7742	0.8571	0.7059
MMP14 AND NOT-ERVW-1	Sarcoma	0.7407	1	0.5882
MMP14 AND NOT-SLC22A11	Sarcoma	0.7407	1	0.5882
STAB1 AND NOT-SCN7A	Sarcoma	0.7059	0.7059	0.7059
STAB1 AND NOT-ATP7B	Sarcoma	0.7059	0.7059	0.7059
STAB1 AND NOT-MFSD3	Sarcoma	0.7097	0.7857	0.6471
ITGA11 AND NOT-NFASC	Sarcoma	0.8276	1	0.7059
ADAM12 AND NOT-MRAP	Sarcoma	0.7907	0.6538	1
MMP14 AND NOT-SLC2A1	Sarcoma	0.7407	1	0.5882
FLRT2 AND NOT-NFASC	Sarcoma	0.7407	1	0.5882
OSMR AND NOT-RGR	Sarcoma	0.8387	0.9286	0.7647
STAB1 AND NOT-BCAM	Sarcoma	0.7586	0.9167	0.6471
STAB1 AND NOT-TSPAN32	Sarcoma	0.8276	1	0.7059
STAB1 AND NOT-CACNA1D	Sarcoma	0.7742	0.8571	0.7059
STAB1 AND NOT-KCNH6	Sarcoma	0.7407	1	0.5882
STAB1 AND NOT-LTA	Sarcoma	0.75	0.8	0.7059
STAB1 AND NOT-SYP	Sarcoma	0.7407	1	0.5882
FGFR1 AND NOT-KCNC2	Sarcoma	0.6923	1	0.5294
FGFR1 AND NOT-NPBWR1	Sarcoma	0.6923	1	0.5294
FGFR1 AND NOT-TMEM235	Sarcoma	0.6923	1	0.5294
FGFR1 AND NOT-KIAA0319	Sarcoma	0.6923	1	0.5294
FGFR1 AND NOT-SLC4A8	Sarcoma	0.6923	1	0.5294
FGFR1 AND NOT-CHRNB4	Sarcoma	0.6923	1	0.5294
FGFR1 AND NOT-GPR22	Sarcoma	0.6923	1	0.5294
FGFR1 AND NOT-EPHA10	Sarcoma	0.6923	1	0.5294
FGFR1 AND NOT-GPR26	Sarcoma	0.6923	1	0.5294
FGFR1 AND NOT-TMPRSS11E	Sarcoma	0.6923	1	0.5294
FGFR1 AND NOT-CLRN1	Sarcoma	0.6923	1	0.5294
FGFR1 AND NOT-GRM3	Sarcoma	0.6923	1	0.5294
FGFR1 AND NOT-CDH20	Sarcoma	0.6923	1	0.5294
FGFR1 AND NOT-PCDHB1	Sarcoma	0.6923	1	0.5294
FGFR1 AND NOT-ERVW-1	Sarcoma	0.6923	1	0.5294
FGFR1 AND NOT-VN1R2	Sarcoma	0.6923	1	0.5294
FGFR1 AND NOT-HTR1D	Sarcoma	0.6923	1	0.5294
FGFR1 AND NOT-HTR5A	Sarcoma	0.6923	1	0.5294
FGFR1 AND NOT-ATP13A5	Sarcoma	0.6923	1	0.5294
FGFR1 AND NOT-CATSPER3	Sarcoma	0.6923	1	0.5294
FGFR1 AND NOT-HCN1	Sarcoma	0.6923	1	0.5294
FGFR1 AND NOT-RGSL1	Sarcoma	0.6923	1	0.5294
FGFR1 AND NOT-IL13	Sarcoma	0.6923	1	0.5294
FGFR1 AND NOT-SHISA9	Sarcoma	0.6923	1	0.5294
FGFR1 AND NOT-OR8D1	Sarcoma	0.6923	1	0.5294
FGFR1 AND NOT-XCR1	Sarcoma	0.6923	1	0.5294
FGFR1 AND NOT-KCNK16	Sarcoma	0.6923	1	0.5294
FGFR1 AND NOT-GLRA3	Sarcoma	0.6923	1	0.5294
FGFR1 AND NOT-DSCAM	Sarcoma	0.6923	1	0.5294
STAB1 AND NOT-CNGA1	Sarcoma	0.6923	1	0.5294
FGFR1 AND NOT-ACSL6	Sarcoma	0.6923	1	0.5294
FGFR1 AND NOT-CEACAM7	Sarcoma	0.6923	1	0.5294
FGFR1 AND NOT-CHRNA2	Sarcoma	0.6923	1	0.5294
FGFR1 AND NOT-LPPR3	Sarcoma	0.6923	1	0.5294
FGFR1 AND NOT-NKAIN1	Sarcoma	0.6923	1	0.5294
FGFR1 AND NOT-NTSR2	Sarcoma	0.6923	1	0.5294
FGFR1 AND NOT-CACNG2	Sarcoma	0.6923	1	0.5294
FGFR1 AND NOT-GABRA4	Sarcoma	0.6923	1	0.5294
FGFR1 AND NOT-SLC17A2	Sarcoma	0.6923	1	0.5294
FGFR1 AND NOT-GABRD	Sarcoma	0.6923	1	0.5294
CDH17 AND NOT-AQP8	Stomach	0.8533	1	0.7442
CDH17 AND NOT-CLCA4	Stomach	0.7473	0.7083	0.7907
CDH17 AND NOT-TSPAN7	Stomach	0.7467	0.875	0.6512
TM4SF5 AND NOT-SLC30A10	Stomach	0.8247	0.7407	0.9302
CDH17 AND NOT-CDHR1	Stomach	0.6972	0.5758	0.8837
CDH17 AND NOT-PTGDR	Stomach	0.6909	0.5672	0.8837
CDH17 AND NOT-PARM1	Stomach	0.6891	0.5395	0.9535
CDH17 AND NOT-PTPRD	Stomach	0.6835	0.75	0.6279
CDH17 AND NOT-SLC26A3	Stomach	0.6833	0.5325	0.9535
CDH17 AND NOT-SCNN1B	Stomach	0.6833	0.5325	0.9535
MUC17 AND NOT-MEP1A	Stomach	0.7353	1	0.5814

FIG. 12 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
FAP AND NOT-CAV1	Breast	0.8687	0.9556	0.7963
FAP AND NOT-SLC22A3	Breast	0.8333	0.9524	0.7407
ADAM12 AND NOT-EGFR	Breast	0.8302	0.8462	0.8148
FAP AND NOT-LYVE1	Breast	0.8687	0.9556	0.7963
FAP AND NOT-ITM2A	Breast	0.8421	0.9756	0.7407
ADAM12 AND NOT-GPC3	Breast	0.8073	0.8	0.8148
FAP AND NOT-CAV2	Breast	0.8283	0.9111	0.7593
FAP AND NOT-PVRL3	Breast	0.8041	0.907	0.7222
FAP AND NOT-SLC9A9	Breast	0.8081	0.8889	0.7407
FAP AND NOT-SLC16A7	Breast	0.8298	0.975	0.7222
FAP AND NOT-PPAP2A	Breast	0.8544	0.898	0.8148
FAP AND NOT-F10	Breast	0.86	0.9348	0.7963
FAP AND NOT-BMP2	Breast	0.7959	0.8864	0.7222
FAP AND NOT-GLDN	Breast	0.8317	0.8936	0.7778
FAP AND NOT-EMCN	Breast	0.8713	0.9362	0.8148
FAP AND NOT-FXYD1	Breast	0.88	0.9565	0.8148
FAP AND NOT-CXCR2	Breast	0.7959	0.8864	0.7222
FAP AND NOT-ACPP	Breast	0.8431	0.8958	0.7963
FAP AND NOT-PTGFR	Breast	0.8367	0.9318	0.7593
FAP AND NOT-NT5E	Breast	0.8041	0.907	0.7222
FAP AND NOT-PLP1	Breast	0.7826	0.9474	0.6667
ADAM12 AND NOT-EDNRB	Breast	0.781	0.8039	0.7593
FAP AND NOT-AVPR1A	Breast	0.835	0.8776	0.7963
FAP AND NOT-IL1R2	Breast	0.7835	0.8837	0.7037
FAP AND NOT-STX8	Breast	0.8544	0.898	0.8148
FAP AND NOT-PLA2R1	Breast	0.82	0.8913	0.7593
FAP AND NOT-KL	Breast	0.82	0.8913	0.7593
FAP AND NOT-GPR87	Breast	0.8235	0.875	0.7778
FAP AND NOT-KCNT2	Breast	0.8081	0.8889	0.7407
FAP AND NOT-ABCA8	Breast	0.8081	0.8889	0.7407
FAP AND NOT-SCN3A	Breast	0.8317	0.8936	0.7778
FAP AND NOT-IL1RAP	Breast	0.8	0.8696	0.7407
FAP AND NOT-ABHD6	Breast	0.82	0.8913	0.7593
FAP AND NOT-SLC31A2	Breast	0.8119	0.8723	0.7593
FAP AND NOT-NAALAD2	Breast	0.8544	0.898	0.8148
FAP AND NOT-IL22RA1	Breast	0.8544	0.898	0.8148
FAP AND NOT-CHL1	Breast	0.78	0.8478	0.7222
FAP AND NOT-MPZ	Breast	0.88	0.9565	0.8148
FAP AND NOT-IL20RB	Breast	0.8269	0.86	0.7963
FAP AND NOT-SLMAP	Breast	0.7692	0.9459	0.6481
FAP AND NOT-CLCA4	Breast	0.835	0.8776	0.7963
FAP AND NOT-FZD10	Breast	0.835	0.8776	0.7963
FAP AND NOT-GJB6	Breast	0.8462	0.88	0.8148
FAP AND NOT-DPP4	Breast	0.835	0.8776	0.7963
FAP AND NOT-COL17A1	Breast	0.8381	0.8627	0.8148
FAP AND NOT-SLC46A2	Breast	0.8462	0.88	0.8148
FAP AND NOT-CDHR1	Breast	0.8544	0.898	0.8148
FAP AND NOT-LRFN5	Breast	0.8431	0.8958	0.7963
FAP AND NOT-TSPAN7	Breast	0.766	0.9	0.6667
FAP AND NOT-LIFR	Breast	0.766	0.9	0.6667
FAP AND NOT-SGCG	Breast	0.8317	0.8936	0.7778
FAP AND NOT-MRAP2	Breast	0.7742	0.9231	0.6667
FAP AND NOT-ATP10B	Breast	0.835	0.8776	0.7963
FAP AND NOT-GJB5	Breast	0.8269	0.86	0.7963
FAP AND NOT-ATP1A2	Breast	0.8085	0.95	0.7037
FAP AND NOT-APCDD1	Breast	0.8381	0.8627	0.8148
FAP AND NOT-JAM3	Breast	0.8317	0.8936	0.7778
FAP AND NOT-PERP	Breast	0.8269	0.86	0.7963
FAP AND NOT-DSC3	Breast	0.835	0.8776	0.7963
ADAM12 AND NOT-FOLR2	Breast	0.7619	0.7843	0.7407
FAP AND NOT-SGCA	Breast	0.8889	0.9778	0.8148
FAP AND NOT-SLC8A1	Breast	0.8387	1	0.7222
FAP AND NOT-LRP4	Breast	0.7579	0.878	0.6667
FAP AND NOT-P2RY1	Breast	0.8317	0.8936	0.7778
FAP AND NOT-KCNK7	Breast	0.8381	0.8627	0.8148
FAP AND NOT-PRIMA1	Breast	0.8163	0.9091	0.7407
FAP AND NOT-EREG	Breast	0.7879	0.8667	0.7222
FAP AND NOT-P2RX1	Breast	0.7921	0.8511	0.7407
FAP AND NOT-DPP6	Breast	0.88	0.9565	0.8148
FAP AND NOT-ABCC9	Breast	0.8627	0.9167	0.8148
FAP AND NOT-PCDH20	Breast	0.7917	0.9048	0.7037
FAP AND NOT-SCN9A	Breast	0.8462	0.88	0.8148
FAP AND NOT-DUOXA1	Breast	0.8462	0.88	0.8148
FAP AND NOT-FAM57A	Breast	0.7551	0.8409	0.6852
FAP AND NOT-RECK	Breast	0.7551	0.8409	0.6852
FAP AND NOT-GJB3	Breast	0.8381	0.8627	0.8148
FAP AND NOT-ACVR2A	Breast	0.8269	0.86	0.7963

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
CDH17 AND NOT-CNGA1	Stomach	0.8718	0.9714	0.7907
CDH17 AND NOT-TRPM6	Stomach	0.6783	0.5417	0.907
MUC17 AND NOT-TRPM6	Stomach	0.8378	1	0.7209
MUC17 AND NOT-TPSG1	Stomach	0.8378	1	0.7209
MUC17 AND NOT-PTPRD	Stomach	0.8108	0.9677	0.6977
MUC17 AND NOT-ABCG2	Stomach	0.6769	1	0.5116
MUC17 AND NOT-SLC51B	Stomach	0.7887	1	0.6512
MUC17 AND NOT-GDPD2	Stomach	0.7945	0.9667	0.6744
MUC17 AND NOT-SLC2A13	Stomach	0.6765	0.92	0.5349
MUC17 AND NOT-SLC26A2	Stomach	0.8378	1	0.7209
MUC17 AND NOT-CLCA4	Stomach	0.8378	1	0.7209
FUT3 AND NOT-CLCA4	Stomach	0.6866	0.9583	0.5349
CDH17 AND NOT-SLC26A2	Stomach	0.6723	0.5263	0.9302
MUC17 AND NOT-CDHR5	Stomach	0.7246	0.9615	0.5814
MUC17 AND NOT-SLC26A3	Stomach	0.8378	1	0.7209
MUC17 AND NOT-ACE	Stomach	0.7324	0.9286	0.6047
MUC17 AND NOT-ANO10	Stomach	0.8267	0.9688	0.7209
VSIG1 AND NOT-PCDHA9	Stomach	0.6849	0.8333	0.5814
MUC17 AND NOT-SEMA6A	Stomach	0.7143	0.9259	0.5814
MUC17 AND NOT-SLC39A5	Stomach	0.7606	0.9643	0.6279
MUC17 AND NOT-VIPR1	Stomach	0.7536	1	0.6047
MUC17 AND NOT-SLC6A19	Stomach	0.8	0.9375	0.6977
MUC17 AND NOT-SLC23A1	Stomach	0.75	0.931	0.6279
MUC17 AND NOT-MEP1B	Stomach	0.7838	0.9355	0.6744
MUC17 AND NOT-SLC36A1	Stomach	0.8267	0.9688	0.7209
MUC17 AND NOT-NAALADL1	Stomach	0.7778	0.9655	0.6512
VSIG1 AND NOT-KCNV1	Stomach	0.6757	0.8065	0.5814
MUC17 AND NOT-SLC1A1	Stomach	0.7838	0.9355	0.6744
MUC17 AND NOT-SECTM1	Stomach	0.7838	0.9355	0.6744
MUC17 AND NOT-CDHR2	Stomach	0.8421	0.9697	0.7442
MUC17 AND NOT-ABHD6	Stomach	0.8219	1	0.6977
MUC17 AND NOT-CYP4F12	Stomach	0.7246	0.9615	0.5814
FUT3 AND NOT-SCNN1B	Stomach	0.8081	0.7143	0.9302
MUC17 AND NOT-FLVCR2	Stomach	0.8158	0.9394	0.7209
MUC17 AND NOT-BST1	Stomach	0.7429	0.963	0.6047
MUC17 AND NOT-P2RY2	Stomach	0.7164	1	0.5581
MUC17 AND NOT-SLC51A	Stomach	0.8312	0.9412	0.7442
MUC17 AND NOT-PTPRH	Stomach	0.8158	0.9394	0.7209
MUC17 AND NOT-ABCB1	Stomach	0.8056	1	0.6744
EPHA2 AND NOT-SCNN1B	Stomach	0.68	0.5965	0.7907
VSIG1 AND NOT-CALHM1	Stomach	0.6667	0.7813	0.5814
VSIG1 AND NOT-CLDN19	Stomach	0.6579	0.7576	0.5814
CXCL16 AND NOT-ADRB2	Stomach	0.6571	0.8519	0.5349
NOT-LIFR AND CDH11	Colon	0.9091	0.8333	1
NOT-ADRB2 AND TGFBI	Colon	0.8889	1	0.8
NOT-PCDH9 AND PPAPDC1A	Colon	0.8889	1	0.8
NOT-NFASC AND SYT1	Esophagus	0.6087	0.7778	0.5
NOT-VAPA AND VANGL1	Esophagus	0.6	1	0.4286
NOT-SCNN1B AND FUT3	Esophagus	0.6	1	0.4286
NOT-CSMD3 AND SEZ6L2	Esophagus	0.6	1	0.4286
NOT-ATP8A1 AND TRPC1	Leiomyosarcoma	0.7407	0.7143	0.7692
NOT-F11R AND VANGL1	Leiomyosarcoma	0.7879	0.65	1
NOT-CLDN10 AND CNTNAP1	Leiomyosarcoma	0.6667	0.6429	0.6923
NOT-ATP8A1 AND BVES	Leiomyosarcoma	0.7407	0.7143	0.7692
NOT-ATP8A1 AND CNTNAP1	Leiomyosarcoma	0.6667	0.6429	0.6923
NOT-ATP8A1 AND CHIC1	Leiomyosarcoma	0.7273	0.8889	0.6154
NOT-ATP8A1 AND STX2	Leiomyosarcoma	0.7619	1	0.6154
NOT-CLDN10 AND GRIN2D	Leiomyosarcoma	0.7273	0.8889	0.6154
NOT-ATP8A1 AND TPM1	Leiomyosarcoma	0.6667	0.6429	0.6923
NOT-WLS AND SLC22A16	AML	0.9537	0.9634	0.9442
NOT-PTPRK AND SLC22A16	AML	0.9435	0.9551	0.9323
NOT-PHLDB2 AND HCST	AML	0.9448	0.9051	0.988
NOT-PHLDB2 AND TAAR5	AML	0.9409	0.93	0.9522
NOT-PHLDB2 AND CLEC12A	AML	0.9318	0.9122	0.9522
NOT-PHLDB2 AND PIEZO1	AML	0.9261	0.9049	0.9482
NOT-PHLDB2 AND CXCR4	AML	0.9254	0.8702	0.988
NOT-CXADR AND VANGL1	Liposarcoma	0.72	0.6923	0.75
NOT-CD1A AND SMO	Liposarcoma	0.6774	0.8077	0.5833
NOT-XK AND SLC6A7	Liposarcoma	0.6667	0.619	0.7222
NOT-SCNN1A AND VANGL1	Liposarcoma	0.6316	0.6	0.6667
NOT-ADCY4 AND SLC2A1	Lung Carcinoma	0.6471	0.6111	0.6875
NOT-GPM6A AND GPR19	Anaplastic Lymphoma	0.625	0.5882	0.6667
NOT-CD9 AND IL15RA	Anaplastic Lymphoma	0.6667	0.6111	0.7333
NOT-LRIG3 AND ST14	Mantle-Cell Lymphoma	0.9315	0.9714	0.8947
NOT-FAT1 AND ST14	Mantle-Cell Lymphoma	0.9296	1	0.8684
NOT-LRIG3 AND CELSR1	Mantle-Cell Lymphoma	0.8736	0.7755	1
NOT-FAT1 AND CELSR1	Mantle-Cell Lymphoma	0.8889	0.8372	0.9474

FIG. 12 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
FAP AND NOT-TMEM47	Breast	0.8571	0.9545	0.7778
FAP AND NOT-LY6D	Breast	0.8381	0.8627	0.8148
SLC4A2 AND NOT-TPBG	Liver	0.8	1	0.6667
SMO AND NOT-PTK7	Liver	0.8	1	0.6667
SMO AND NOT-IGF1R	Liver	0.8	1	0.6667
FGFR4 AND NOT-EDNRB	Liver	0.8	1	0.6667
FGFR4 AND NOT-ITGB6	Liver	0.8	1	0.6667
FGFR4 AND NOT-AXL	Liver	0.8	1	0.6667
FGFR4 AND NOT-CLDN7	Liver	0.8	1	0.6667
FGFR4 AND NOT-TPBG	Liver	0.8	1	0.6667
SLC4A2 AND NOT-AXL	Liver	0.8	1	0.6667
SLC4A2 AND NOT-FOLR1	Liver	0.8	1	0.6667
FGFR4 AND NOT-ENG	Liver	0.8	1	0.6667
FGFR4 AND NOT-SLC34A2	Liver	0.8	1	0.6667
FGFR4 AND NOT-GPA33	Liver	0.8	1	0.6667
FGFR4 AND NOT-CLDN5	Liver	0.8	1	0.6667
FGFR4 AND NOT-CLDN18	Liver	0.8	1	0.6667
FGFR4 AND NOT-ROR1	Liver	0.8	1	0.6667

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
NOT-VANGL1 AND ST14	Mantle-Cell Lymphoma	0.9444	1	0.8947
NOT-VANGL1 AND CELSR1	Mantle-Cell Lymphoma	0.987	0.9744	1
NOT-EPHA2 AND CELSR1	Mantle-Cell Lymphoma	0.8636	0.76	1
NOT-CHRNA2 AND SGCD	Melanoma	0.9524	1	0.9091
NOT-NMUR2 AND SGCD	Melanoma	0.9524	1	0.9091
NOT-CLDN19 AND SGCD	Melanoma	0.9524	1	0.9091
NOT-OR51B2 AND GPR19	Melanoma	0.9524	1	0.9091
NOT-CHRNA2 AND NRP2	Melanoma	0.9	1	0.8182
NOT-NMUR2 AND NRP2	Melanoma	0.9	1	0.8182
NOT-CHRNA2 AND SDC3	Melanoma	0.9	1	0.8182
NOT-SCN10A AND ADAM12	Melanoma	0.9	1	0.8182
NOT-TMEM150B AND ICAM1	Melanoma	0.9	1	0.8182
NOT-TM4SF5 AND GPR19	Melanoma	0.9524	1	0.9091
NOT-F11R AND SLC31A1	Sarcoma	0.8571	0.8333	0.8824
NOT-F11R AND VANGL1	Sarcoma	0.7317	0.625	0.8824
NOT-SCNN1A AND VANGL1	Sarcoma	0.8276	1	0.7059
NOT-F11R AND ATP7A	Sarcoma	0.6923	1	0.5294

FIG. 13

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
EPCAM AND ROR1	Breast	0.441379	0.351648	0.592593
EPCAM AND ERBB2	Breast	0.358491	0.365385	0.351852
CD33 AND IL3RA	AML	0.552567	0.71519	0.450199
EPCAM AND NOT-MET	Breast	0.573913	0.540984	0.611111
EPCAM AND NOT-MSLN	Breast	0.587413	0.47191	0.777778
EPCAM AND NOT-TYR	Breast	0.577181	0.452632	0.796296
ERBB2 AND NOT-ROR1	Breast	0.392157	0.416667	0.37037
ERBB2 AND NOT-TYR	Breast	0.38835	0.408163	0.37037
ERBB2 AND NOT-MET	Breast	0.305556	0.611111	0.203704
MSLN AND NOT-ROR1	Breast	0.093137	0.053672	0.351852
MSLN AND NOT-TYR	Breast	0.091127	0.052342	0.351852
MSLN AND NOT-MET	Breast	0.063158	0.03681	0.222222
MSLN AND NOT-ERBB2	Pancreas	0.146341	0.1	0.272727
MSLN AND NOT-ERBB2	Lung Adenocarcinoma	0.354167	0.653846	0.242857
MUC1 AND NOT-ERBB2	Lung Adenocarcinoma	0.385787	0.299213	0.542857
CD19 AND NOT-CD22	B-Cell Diffuse	0.192308	0.333333	0.135135
MS4A1 AND NOT-CD22	B-Cell Diffuse	0.151515	0.172414	0.135135
MLANA AND NOT-KDR	Melanoma	0.9	1	0.818182
NOT-TYR AND ROR1	Breast	0.183486	0.10989	0.555556
NOT-MET AND ROR1	Breast	0.123656	0.072327	0.425926
NOT-MET AND TYR	Breast	0.11588	0.065534	0.5
NOT-GPC3 AND MSLN	Colon	0.352941	0.25	0.6
NOT-GPC3 AND MUC1	Colon	0.47619	0.3125	1
NOT-GPC3 AND ERBB2	Colon	0.454545	0.294118	1
NOT-EPHA2 AND ERBB2	Glioma	0.295918	0.179567	0.84058
NOT-EPHA2 AND MUC1	Glioma	0.281481	0.169643	0.826087
NOT-ROR1 AND MUC1	Lung Carcinoma	0.248996	0.167568	0.484375
NOT-ROR1 AND TNFRSF10A	Lung Carcinoma	0.042553	0.027523	0.09375

FIG. 14

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
CD33	AML	0.762987	0.643836	0.936255
CD19	Mantle-Cell Lymphoma	0.914286	1	0.842105
PMEL	Melanoma	0.952381	1	0.909091
MLANA	Melanoma	1	1	1
VTCN1	Breast	0.690476	0.966667	0.537037
FAP	Breast	0.838095	0.862745	0.814815
TPBG	Breast	0.62	0.673913	0.574074
CLDN12	Liver	0.631579	0.461538	1
FAP	Pancreas	0.842105	1	0.727273
VCAM1	Renal	0.666667	0.6	0.75
RNF43	Colon	0.75	1	0.6
BIRC5	Colon	0.625	0.454545	1
EPHB2	Colon	0.615385	0.5	0.8
SSTR1	Ependymoma	1	1	1
BIRC5	Esophagus	0.692308	0.75	0.642857
ITGAV	Glioma	0.897959	0.846154	0.956522
EDNRB	Glioma	0.707483	0.666667	0.753623
SLC39A6	Glioma	0.777778	0.746667	0.811594
CLDN12	Glioma	0.651852	0.666667	0.637681
FAP	Leiomyosarcoma	0.818182	1	0.692308
CD37	AML	0.71407	0.57561	0.940239
CD38	AML	0.62069	0.497596	0.824701
CD276	Liposarcoma	0.648148	0.486111	0.972222
SLC34A2	Lung Adenocarcinoma	0.607143	0.520408	0.728571
MAGEA4	Lung Carcinoma	0.605505	0.733333	0.515625
IL2RA	Anaplastic Lymphoma	0.62069	0.642857	0.6
TNFRSF8	Anaplastic Lymphoma	0.875	0.823529	0.933333
FCRL2	Mantle-Cell Lymphoma	0.730159	0.92	0.605263
MS4A1	Mantle-Cell Lymphoma	0.71028	0.550725	1
CD70	Mantle-Cell Lymphoma	0.885714	0.96875	0.815789
CXCR5	Mantle-Cell Lymphoma	0.675676	0.694444	0.657895
CD79B	Mantle-Cell Lymphoma	0.666667	0.608696	0.736842
TNFRSF13C	Mantle-Cell Lymphoma	0.77551	0.633333	1
FCRL1	Mantle-Cell Lymphoma	0.606742	0.529412	0.710526
EDNRB	Melanoma	0.9	1	0.818182
CLDN12	Melanoma	0.777778	1	0.636364
SLC7A5	Melanoma	0.705882	1	0.545455
ABCB5	Melanoma	0.705882	1	0.545455
RNF43	Melanoma	0.777778	1	0.636364
BMPR1B	Neuroblastoma	0.666667	0.692308	0.642857
STEAP2	Prostate	0.727273	0.571429	1
STEAP1	Prostate	0.666667	0.6	0.75
CLDN12	Prostate	0.666667	1	0.5
BMPR1B	Prostate	0.666667	1	0.5
FAP	Sarcoma	0.866667	1	0.764706
ROR1	Sarcoma	0.83871	0.928571	0.764706
CD276	Sarcoma	0.740741	1	0.588235
CLDN18	Stomach	0.765957	0.705882	0.837209
MUC13	Stomach	0.705882	0.552632	0.976744
MUC1	Stomach	0.666667	0.956522	0.511628
CLDN2	Stomach	0.621359	0.533333	0.744186
EPCAM	Stomach	0.609929	0.438776	1
MST1R	Stomach	0.688525	0.531646	0.976744
NOT-ITGAV	AML	0.690625	0.568123	0.880478
NOT-EDNRB	AML	0.664865	0.503067	0.98008
NOT-STEAP2	AML	0.620519	0.449821	1
NOT-ERBB3	AML	0.607748	0.436522	1
NOT-FCRL5	Melanoma	0.625	1	0.454545
NKAIN4	Astrocytoma	0.764045	0.790698	0.73913
BEST3	Astrocytoma	0.608696	0.608696	0.608696
ABCG8	Liver	0.8	1	0.666667
SLCO1B1	Liver	0.75	0.6	1
SLC17A2	Liver	0.666667	0.555556	0.833333
SLC2A2	Liver	0.857143	0.75	1
SLC10A1	Liver	0.666667	0.555556	0.833333
SLC30A8	Pancreas	0.777778	1	0.636364
MUC17	Pancreas	0.666667	0.857143	0.545455
IFI6	Colon	0.666667	0.5	1
MUC17	Esophagus	0.785714	0.785714	0.785714
SLC1A5	Esophagus	0.64	0.727273	0.571429
LYPD1	Glioblastoma	0.666667	0.56	0.823529
SYT11	Glioma	0.964539	0.944444	0.985507
CRB1	Glioma	0.813559	0.979592	0.695652
LYPD1	Glioma	0.876923	0.934426	0.826087
DTNA	Glioma	0.711864	0.857143	0.608696
GPR19	Glioma	0.816327	0.769231	0.869565
MEGF11	Glioma	0.682927	0.777778	0.608696

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
PRLR	Breast	0.654206	0.660377	0.648148
SLC4A2	Breast	0.652174	0.789474	0.555556
F2RL2	Breast	0.674419	0.90625	0.537037
SLC5A6	Breast	0.636364	0.625	0.648148
PMEPA1	Breast	0.66055	0.654545	0.666667
MFSD10	Breast	0.640777	0.673469	0.611111
NOX4	Breast	0.643357	0.516854	0.851852
SLC52A2	Breast	0.673267	0.723404	0.62963
KCNK15	Breast	0.629213	0.8	0.518519
FGFR4	Liver	0.8	1	0.666667
ABCC2	Liver	1	1	1
ABCB11	Liver	0.666667	0.666667	0.666667
TMPRSS6	Liver	0.666667	1	0.5
APOB	Liver	0.833333	0.833333	0.833333
HPN	Liver	0.857143	0.75	1
FGG	Liver	0.8	0.666667	1
ABCC6	Liver	0.75	0.6	1
TFR2	Liver	0.666667	0.555556	0.833333
ABCC8	Pancreas	0.777778	1	0.636364
CDH6	Renal	0.857143	1	0.75
LRP2	Renal	0.636364	0.7	0.583333
SLC3A1	Renal	0.72	0.692308	0.75
SLC12A2	Colon	0.75	1	0.6
LY6G6D	Colon	0.75	1	0.6
PMEPA1	Colon	0.666667	0.571429	0.8
KCNA6	Ependymoma	1	1	1
DLL1	Ependymoma	0.666667	0.5	1
SLC7A11	Ependymoma	1	1	1
PRNP	Ependymoma	1	1	1
SLC52A2	Esophagus	0.769231	0.833333	0.714286
SLC6A20	Esophagus	0.62069	0.6	0.642857
TM4SF20	Esophagus	0.666667	0.692308	0.642857
PTPRZ1	Glioblastoma	0.909091	0.9375	0.882353
AQP4	Glioblastoma	0.604651	0.5	0.764706
HRH1	Glioblastoma	0.666667	0.9	0.529412
PTPRZ1	Glioma	0.985507	0.985507	0.985507
NRCAM	Glioma	0.917293	0.953125	0.884058
NLGN1	Glioma	0.938462	1	0.884058
PCDHB10	Glioma	0.904762	1	0.826087
SGCB	Glioma	0.855072	0.855072	0.855072
GRIK3	Glioma	0.754967	0.695122	0.826087
XPR1	Glioma	0.75	0.659341	0.869565
NLGN4X	Glioma	0.730539	0.622449	0.884058
FZD3	Glioma	0.722581	0.651163	0.811594
SLC1A3	Glioma	0.72	0.666667	0.782609
FGFR1	Leiomyosarcoma	0.782609	0.9	0.692308
ROR2	Leiomyosarcoma	0.692308	0.692308	0.692308
ITGA11	Leiomyosarcoma	0.64	0.666667	0.615385
OSMR	Leiomyosarcoma	0.695652	0.8	0.615385
FLT3	AML	0.942857	0.966527	0.920319
P2RX1	AML	0.823729	0.716814	0.968127
ATP8B4	AML	0.796875	0.781609	0.812749
CSF3R	AML	0.778878	0.664789	0.940239
HCST	AML	0.767802	0.627848	0.988048
ADAM12	Liposarcoma	0.62	0.484375	0.861111
TSPAN31	Liposarcoma	0.618182	0.894737	0.472222
IGDCC4	Liposarcoma	0.631579	0.857143	0.5
MMP14	Liposarcoma	0.686567	0.741935	0.638889
CXCL9	B-Cell Diffuse	0.712329	0.722222	0.702703
CD40	B-Cell Diffuse	0.675325	0.65	0.702703
CD80	B-Cell Diffuse	0.611765	0.541667	0.702703
CXCL10	B-Cell Diffuse	0.60274	0.611111	0.594595
CLECL1	Mantle-Cell Lymphoma	0.767677	0.622951	1
QSOX2	Mantle-Cell Lymphoma	0.891566	0.822222	0.973684
TNFRSF13B	Mantle-Cell Lymphoma	0.78481	0.756098	0.815789
CD83	Mantle-Cell Lymphoma	0.648649	0.493151	0.947368
CXCL9	T-Cell, Peripheral	0.733333	0.6875	0.785714
GPR143	Melanoma	0.842105	1	0.727273
STEAP1B	Melanoma	0.842105	1	0.727273
HTRA2	Melanoma	0.777778	1	0.636364
MC1R	Melanoma	0.9	1	0.818182
MARVELD1	Melanoma	0.952381	1	0.909091
C11orf24	Melanoma	0.9	1	0.818182
PTPRM	Melanoma	0.842105	1	0.727273
TNFSF9	Melanoma	0.909091	0.909091	0.909091
SLC4A2	Melanoma	0.777778	1	0.636364
IL12RB2	Melanoma	0.9	1	0.818182

FIG. 14 (Cont.)

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
CSMD2	Glioma	0.686131	0.691176	0.681159
DSCAM	Glioma	0.640777	0.970588	0.478261
BEST3	Glioma	0.671642	0.692308	0.652174
SLCO1C1	Glioma	0.624204	0.556818	0.710145
SLC22A16	AML	0.896552	0.863469	0.932271
FMNL1	AML	0.676783	0.546569	0.888446
SLC39A8	AML	0.648871	0.669492	0.629482
LAPTM5	AML	0.692414	0.529536	1
EMP3	AML	0.667598	0.513978	0.952191
SLC43A1	AML	0.695833	0.729258	0.665339
STAB1	Liposarcoma	0.617284	0.555556	0.694444
TRPM1	Melanoma	0.705882	1	0.545455
SLC6A15	Melanoma	0.705882	1	0.545455
OR7C1	Melanoma	0.636364	0.636364	0.636364
FAT1	Melanoma	0.8	0.888889	0.727273
LYPD1	Melanoma	0.705882	1	0.545455
IFI6	Melanoma	0.75	0.692308	0.818182
NKAIN1	Neuroblastoma	0.797872	0.721154	0.892857
GRM8	Neuroblastoma	0.637037	0.843137	0.511905
PIRT	Neuroblastoma	0.632258	0.690141	0.583333
CHRNB4	Neuroblastoma	0.628205	0.680556	0.583333
GPR19	Neuroblastoma	0.610687	0.449438	0.952381
GPR85	Neuroblastoma	0.609865	0.489209	0.809524
NKAIN4	Oligodendroglioma	0.888889	1	0.8
CRB1	Oligodendroglioma	0.75	1	0.6
KCNQ2	Oligodendroglioma	0.736842	0.608696	0.933333
MDGA2	Oligodendroglioma	0.685714	0.6	0.8
DSCAM	Oligodendroglioma	0.75	1	0.6
LYPD1	Oligodendroglioma	0.733333	0.733333	0.733333
GPR37L1	Oligodendroglioma	0.608696	0.875	0.466667
DTNA	Oligodendroglioma	0.702703	0.590909	0.866667
TRPM8	Prostate	0.727273	0.571429	1
UPK3A	Prostate	0.666667	0.6	0.75
TMEFF2	Prostate	0.857143	1	0.75
STAB1	Sarcoma	0.6	0.521739	0.705882
PTGIS	Sarcoma	0.714286	0.909091	0.588235
MUC17	Stomach	0.831169	0.941176	0.744186
ATP8B1	Stomach	0.831461	0.804348	0.860465
NOT-PHLDB2	AML	0.698592	0.540305	0.988048
NOT-FAT1	AML	0.693642	0.544218	0.956175
NOT-EFNB2	AML	0.673051	0.5125	0.98008
NOT-GPR137B	AML	0.672052	0.565217	0.828685
PTPRZ1	Astrocytoma	0.745455	0.640625	0.891304
PTPRZ1	Astrocytoma	0.745455	0.640625	0.891304
MMP16	Astrocytoma	0.674419	0.725	0.630435
ADCYAP1R1	Astrocytoma	0.62963	0.548387	0.73913
NRCAM	Astrocytoma	0.617886	0.493506	0.826087
KCNN3	Astrocytoma	0.682353	0.74359	0.630435
ADAM12	Breast	0.818182	0.803571	0.833333

Antigen Pair Logic	Cancer (short name)	F1	Precision	Recall
SLC10A4	Neuroblastoma	0.922222	0.864583	0.988095
CHRNA3	Neuroblastoma	0.922156	0.927711	0.916667
DIABLO	Neuroblastoma	0.881988	0.922078	0.845238
ACVR2B	Neuroblastoma	0.831169	0.914286	0.761905
PCDHB10	Neuroblastoma	0.820809	0.797753	0.845238
SLC6A2	Neuroblastoma	0.814815	0.846154	0.785714
PCDHB6	Neuroblastoma	0.797297	0.921875	0.702381
SCN9A	Neuroblastoma	0.776471	0.767442	0.785714
GPR173	Neuroblastoma	0.745763	0.709677	0.785714
PODXL2	Neuroblastoma	0.708995	0.638095	0.797619
NLGN3	Oligodendroglioma	0.933333	0.933333	0.933333
GPR173	Oligodendroglioma	0.769231	0.909091	0.666667
GRIK3	Oligodendroglioma	0.681818	0.517241	1
ADCYAP1R1	Oligodendroglioma	0.846154	1	0.733333
MMP16	Oligodendroglioma	0.888889	1	0.8
PTPRZ1	Oligodendroglioma	0.857143	0.923077	0.8
NRCAM	Oligodendroglioma	0.702703	0.590909	0.866667
GRIK2	Oligodendroglioma	0.666667	0.571429	0.8
KCNN3	Oligodendroglioma	0.64	0.8	0.533333
EPHB1	Oligodendroglioma	0.666667	0.666667	0.666667
KCNK15	Ovarian	0.774194	0.631579	1
OR51E2	Prostate	0.888889	0.8	1
ANO7	Prostate	0.8	0.666667	1
ACPP	Prostate	0.857143	1	0.75
OR51E1	Prostate	0.857143	1	0.75
TSPAN1	Prostate	0.666667	1	0.5
ABCC4	Prostate	0.75	0.75	0.75
SLC2A12	Prostate	0.857143	1	0.75
PMEPA1	Prostate	0.6	0.5	0.75
NDRG1	Prostate	0.666667	1	0.5
FGFR1	Sarcoma	0.666667	0.9	0.529412
MMP14	Sarcoma	0.740741	1	0.588235
OSMR	Sarcoma	0.848485	0.875	0.823529
ADAM12	Sarcoma	0.755556	0.607143	1
VASN	Sarcoma	0.692308	1	0.529412
FLRT2	Sarcoma	0.733333	0.846154	0.647059
BFAR	Sarcoma	0.733333	0.846154	0.647059
ITGA11	Sarcoma	0.787879	0.8125	0.764706
HRH1	Sarcoma	0.615385	0.888889	0.470588
VSIG1	Stomach	0.617284	0.657895	0.581395
LSR	Stomach	0.690909	0.567164	0.883721
TM4SF20	Stomach	0.611111	0.507692	0.767442
B3GNT3	Stomach	0.661538	0.494253	1
SI	Stomach	0.6	0.467532	0.837209
NOT-PTPRK	AML	0.805778	0.674731	1
NOT-HEG1	AML	0.797853	0.724026	0.888446
NOT-WLS	AML	0.782609	0.656757	0.968127
NOT-CADM1	AML	0.766506	0.643243	0.948207
NOT-PTPRM	AML	0.782753	0.670455	0.940239

FIG. 15

Antigen Pair Logic	F1	Precision	Recall
Astrocytoma			
PTPRZ1 AND NOT-SDC1 AND F2R	0.91954	0.97561	0.869565
PTPRZ1 AND NOT-EPCAM AND F2R	0.911111	0.931818	0.891304
PTPRZ1 AND CLDN17 AND TNC	0.891304	0.891304	0.891304
PTPRZ1 AND NOT-SDC1 AND NOT-GABRA2	0.891304	0.891304	0.891304
PCDH10 AND F2R AND NOT-SDC1	0.886364	0.928571	0.847826
PTPRZ1 AND NOT-SDC1 AND ANTXR2	0.886364	0.928571	0.847826
CELSR2 AND NOT-SDC1 AND F2R	0.883721	0.95	0.826087
PTPRZ1 AND NOT-STEAP1 AND NOT-GABRA2	0.88172	0.87234	0.891304
PTPRZ1 AND GABRB2 AND TNC	0.88172	0.87234	0.891304
PTPRZ1 AND NOT-CLDN23 AND NOT-GABRA2	0.88172	0.87234	0.891304
PTPRZ1 AND NOT-CD52 AND NOT-GABRA2	0.88172	0.87234	0.891304
PTPRZ1 AND NCAM1 AND NOT-GRM3	0.880952	0.973684	0.804348
PTPRZ1 AND NOT-SDC1 AND NOT-GRM3	0.880952	0.973684	0.804348
PTPRZ1 AND NOT-PMEL AND NOT-GRM3	0.880952	0.973684	0.804348
PTPRZ1 AND NOT-EPCAM AND NOT-GRM3	0.880952	0.973684	0.804348
PTPRZ1 AND NOT-TRPM4 AND NOT-GRM3	0.880952	0.973684	0.804348
PTPRZ1 AND NOT-STEAP1 AND NOT-GRM3	0.880952	0.973684	0.804348
PTPRZ1 AND NOT-CLDN23 AND NOT-GRM3	0.880952	0.973684	0.804348
PTPRZ1 AND NOT-EPCAM AND NOT-GRM3	0.880952	0.973684	0.804348
PTPRZ1 AND NOT-CLDN1 AND NOT-GRM3	0.880952	0.973684	0.804348
PTPRZ1 AND NOT-TRPM4 AND F2R	0.879121	0.888889	0.869565
PTPRZ1 AND NOT-CLDN1 AND F2R	0.879121	0.888889	0.869565
PTPRZ1 AND NCAM1 AND F2R	0.879121	0.888889	0.869565
PTPRZ1 AND FXYD6 AND NOT-STEAP2	0.869565	0.869565	0.869565
PTPRZ1 AND F2R AND NOT-CD52	0.866667	0.886364	0.847826
PTPRZ1 AND F2R AND NOT-STEAP1	0.866667	0.886364	0.847826
PTPRZ1 AND BEST3 AND NOT-SDC1	0.866667	0.886364	0.847826
PTPRZ1 AND F2R AND NOT-PMEL	0.866667	0.886364	0.847826
PTPRZ1 AND F2R AND NOT-CLDN23	0.866667	0.886364	0.847826
PTPRZ1 AND SLCO1C1 AND CD33	0.863636	0.904762	0.826087
PTPRZ1 AND GPR85 AND CD33	0.860465	0.925	0.804348
PTPRZ1 AND CRB1 AND CD33	0.860465	0.925	0.804348
COMPLEX-TNC/PTPRZ1/PCDH15	0.863158	0.836735	0.891304
PCDH10 AND F2R AND NOT-EPCAM	0.857143	0.866667	0.847826
PTPRZ1 AND NOT-TPBG AND NOT-GABRA2	0.853933	0.883721	0.826087
PTPRZ1 AND NOT-AOC3 AND FOLR2	0.851064	0.833333	0.869565
PTPRZ1 AND NOT-MRGPRF AND FOLR2	0.851064	0.833333	0.869565
PTPRZ1 AND SLC4A8 AND FOLR2	0.851064	0.833333	0.869565
PTPRZ1 AND NOT-SLC43A1 AND FOLR2	0.851064	0.833333	0.869565
PTPRZ1 AND NOT-STEAP1 AND ANTXR2	0.847826	0.847826	0.847826
PTPRZ1 AND NCAM1 AND ANTXR2	0.847826	0.847826	0.847826
PTPRZ1 AND NOT-CD52 AND ANTXR2	0.847826	0.847826	0.847826
PTPRZ1 AND NOT-CLDN23 AND ANTXR2	0.847826	0.847826	0.847826
PTPRZ1 AND NOT-CLDN1 AND ANTXR2	0.847826	0.847826	0.847826
PTPRZ1 AND NOT-TRPM4 AND ANTXR2	0.847826	0.847826	0.847826
PTPRZ1 AND NOT-SDC1 AND NOT-SCN2A	0.845361	0.803922	0.891304
PTPRZ1 AND NCAM1 AND NOT-ATP8A2	0.844444	0.863636	0.826087
PTPRZ1 AND NOT-CLDN23 AND NOT-GPR83	0.844444	0.863636	0.826087
PTPRZ1 AND NOT-STEAP1 AND NOT-GPR83	0.844444	0.863636	0.826087
PTPRZ1 AND NOT-CLDN23 AND NOT-ATP8A2	0.844444	0.863636	0.826087
PTPRZ1 AND NOT-STEAP1 AND NOT-ATP8A2	0.844444	0.863636	0.826087
PTPRZ1 AND NOT-TRPM4 AND NOT-ATP8A2	0.844444	0.863636	0.826087
PTPRZ1 AND NOT-CD52 AND NOT-GPR83	0.844444	0.863636	0.826087
PTPRZ1 AND NOT-CLDN1 AND NOT-GPR83	0.844444	0.863636	0.826087
PTPRZ1 AND NOT-CLDN1 AND NOT-ATP8A2	0.844444	0.863636	0.826087
PTPRZ1 AND NOT-TRPM4 AND NOT-GPR83	0.844444	0.863636	0.826087
PTPRZ1 AND NOT-SDC1 AND NOT-GPR83	0.844444	0.863636	0.826087
PTPRZ1 AND NOT-SDC1 AND NOT-ATP8A2	0.844444	0.863636	0.826087
PTPRZ1 AND NCAM1 AND NOT-GPR83	0.844444	0.863636	0.826087
PTPRZ1 AND NOT-SDC1 AND NOT-CDH18	0.840909	0.880952	0.804348
PTPRZ1 AND NOT-MSMO1 AND FOLR2	0.851064	0.833333	0.869565
PTPRZ1 AND NOT-RNF43 AND NOT-DYSF	0.833333	0.8	0.869565
PTPRZ1 AND CD82 AND VCAM1	0.831461	0.860465	0.804348
COMPLEX-PTGER4/PTPRZ1/CD33	0.829787	0.8125	0.847826
PTPRZ1 AND CD33 AND NOT-KCNK6	0.829787	0.8125	0.847826
PTPRZ1 AND CD33 AND NOT-AOC3	0.829787	0.8125	0.847826
PTPRZ1 AND CD33 AND OR2L13	0.829787	0.8125	0.847826
PTPRZ1 AND CD33 AND NOT-CD36	0.829787	0.8125	0.847826
PTPRZ1 AND CD33 AND SLC4A8	0.829787	0.8125	0.847826
PTPRZ1 AND CD33 AND NOT-DYSF	0.829787	0.8125	0.847826
PTPRZ1 AND CD33 AND NOT-CLDN10	0.829787	0.8125	0.847826
PTPRZ1 AND CD33 AND NOT-MRGPRF	0.829787	0.8125	0.847826
PTPRZ1 AND CD33 AND NOT-SLC43A1	0.829787	0.8125	0.847826
SLC1A2 AND SLC3A2 AND NOT-SDC1	0.833333	0.8	0.869565
PTPRZ1 AND CLDN17 AND NOT-STEAP2	0.828283	0.773585	0.891304
PTPRZ1 AND CLDN17 AND NOT-STEAP2	0.828283	0.773585	0.891304

Antigen Pair Logic	F1	Precision	Recall
(Cont.)			
SELL AND NOT-HLA-DOB AND NOT-KDR	0.917603	0.865724	0.976096
NOT-PTPRK AND CD70 AND NOT-ABCA5	0.899804	0.887597	0.912351
NOT-PODXL AND CD37 AND NOT-FCRL1	0.899628	0.843206	0.964143
NOT-EDNRB AND ITGB2 AND NOT-MS4A1	0.911488	0.864286	0.964143
NOT-EDNRB AND SLC7A5 AND NOT-MAL	0.899431	0.858696	0.944223
P2RY8 AND NOT-PCYT1A AND NOT-FCRL2	0.919325	0.868794	0.976096
NOT-PTPRK AND CD70 AND NOT-ALDH1A1	0.899225	0.875472	0.924303
SELL AND NOT-HLA-DOB AND NOT-AXL	0.911488	0.864286	0.964143
P2RY8 AND NOT-IL11RA AND NOT-CR2	0.898833	0.878327	0.920319
NOT-EDNRB AND ITGB2 AND NOT-FCRL5	0.898711	0.835616	0.972112
P2RY8 AND NOT-CD22 AND NOT-CLDN8	0.897436	0.830508	0.976096
SELL AND NOT-HLA-DOB AND NOT-STEAP1	0.906367	0.855124	0.964143
NOT-PODXL AND CD37 AND NOT-CD22	0.896679	0.835052	0.968127
P2RY8 AND NOT-CD22 AND NOT-AXL	0.896679	0.835052	0.968127
P2RY8 AND NOT-ALDH1A1 AND NOT-CR2	0.901575	0.891051	0.912351
CXCR4 AND NOT-MS4A1 AND NOT-CLDN11	0.897579	0.842657	0.960159
NOT-PTPRK AND CD37 AND NOT-CD22	0.895028	0.832192	0.968127
NOT-SGCB AND NOT-HLA-DOB AND CD70	0.907063	0.850174	0.972112
P2RY8 AND NOT-FCRL1 AND NOT-AXL	0.89464	0.834483	0.964143
P2RY8 AND NOT-FCRL1 AND NOT-AXL	0.89464	0.834483	0.964143
NOT-WLS AND CD70 AND NOT-ABCA5	0.894212	0.896	0.89243
P2RY8 AND NOT-ITGAV AND NOT-CR2	0.893939	0.851986	0.940239
SELL AND NOT-AXL AND NOT-CD160	0.893855	0.839161	0.956175
NOT-EDNRB AND FXYD5 AND NOT-HLA-DOB	0.902752	0.836735	0.98008
P2RY8 AND NOT-FCRL1 AND NOT-EDNRB	0.893458	0.841549	0.952191
P2RY8 AND NOT-FCRL1 AND NOT-EDNRB	0.893458	0.841549	0.952191
P2RY8 AND NOT-CD22 AND NOT-CLDN5	0.897436	0.830508	0.976096
NOT-WLS AND CD70 AND NOT-HLA-DOB	0.892989	0.831615	0.964143
SELL AND NOT-AXL AND NOT-MS4A1	0.901887	0.856631	0.952191
P2RY8 AND NOT-CD22 AND NOT-EDNRB	0.892193	0.836237	0.956175
NOT-EDNRB AND SLC7A5 AND NOT-EPHA4	0.892139	0.824324	0.972112
NOT-EDNRB AND SLC7A5 AND NOT-EPHA4	0.892139	0.824324	0.972112
CXCR4 AND NOT-SLAMF7 AND NOT-CLDN11	0.901887	0.856631	0.952191
NOT-PODXL AND CD33 AND NOT-CD160	0.94902	0.934363	0.964143
NOT-PODXL AND CD33 AND NOT-MS4A1	0.941406	0.923372	0.960159
NOT-PTPRK AND CD33 AND NOT-SLAMF7	0.945098	0.930502	0.960159
NOT-PODXL AND CD33 AND NOT-FCRL5	0.935922	0.912879	0.960159
NOT-PODXL AND CD33 AND NOT-HLA-DOB	0.933078	0.897059	0.972112
NOT-PODXL AND CD33 AND NOT-SLAMF7	0.934109	0.909434	0.960159
NOT-PTPRK AND CD33 AND NOT-MS4A1	0.940945	0.929961	0.952191
NOT-PTPRK AND NOT-HLA-DOB AND CD33	0.930769	0.899628	0.964143
NOT-PTPRK AND CD33 AND NOT-FCRL1	0.930693	0.925197	0.936255
NOT-PODXL AND CD33 AND NOT-FCRL1	0.930502	0.902622	0.960159
NOT-PODXL AND CD33 AND NOT-CD22	0.92543	0.889706	0.964143
NOT-PTPRK AND CD33 AND NOT-FCRL5	0.934109	0.909434	0.960159
NOT-PTPRK AND CD33 AND NOT-CD22	0.921305	0.888889	0.956175
NOT-PODXL AND CD33 AND NOT-CR2	0.920455	0.877256	0.968127
NOT-PTPRK AND CD33 AND NOT-SPON2	0.939806	0.916667	0.964143
NOT-PTPRK AND CD33 AND NOT-ALDH1A1	0.912779	0.929752	0.896414
NOT-PODXL AND CD33 AND NOT-SPON2	0.91215	0.859155	0.972112
NOT-PTPRK AND CD33 AND NOT-OAS1	0.936	0.939759	0.932271
NOT-PTPRK AND CD33 AND NOT-IL11RA	0.910891	0.905512	0.916335
NOT-PTPRK AND CD33 AND NOT-CD79A	0.908745	0.869091	0.952191
NOT-PTPRK AND CD33 AND NOT-PCYT1A	0.907721	0.860714	0.960159
NOT-PODXL AND CD33 AND NOT-OAS1	0.917647	0.903475	0.932271
NOT-PODXL AND CD33 AND NOT-IL11RA	0.907336	0.88015	0.936255
NOT-PODXL AND CD33 AND NOT-ALDH1A1	0.905812	0.91129	0.900398
NOT-PODXL AND CD33 AND NOT-CD79A	0.902622	0.85159	0.960159
NOT-PTPRK AND CD33 AND NOT-CD72	0.926357	0.901887	0.952191
NOT-PTPRK AND CD33 AND NOT-CXCR5	0.899628	0.843206	0.964143
NOT-PODXL AND CD33 AND NOT-ABCA5	0.895582	0.902834	0.888446
NOT-PTPRK AND NOT-CD79B AND CD33	0.893878	0.916318	0.87251
COMPLEX-ALDH1A1/SLC4A1/CD33	0.893617	0.868421	0.920319
NOT-PODXL AND CD33 AND NOT-CD79B	0.892057	0.9125	0.87251
CD33 AND NOT-EPHA4 AND NOT-EDNRB	0.911243	0.902344	0.920319
NOT-PTPRK AND CD33 AND NOT-CD19	0.918812	0.913386	0.924303
CD33 AND NOT-EPHA4 AND NOT-ITGB6	0.909091	0.883459	0.936255
CD33 AND NOT-HLA-DOB AND SLC4A1	0.925	0.969432	0.884462
CD33 AND NOT-TLR1 AND NOT-ERBB2	0.882474	0.91453	0.85259
CD33 AND NOT-CD79B AND SLC4A1	0.874459	0.957346	0.804781
CD33 AND NOT-SLAMF7 AND SLC4A1	0.92562	0.961373	0.89243
NOT-PTPRK AND CD33 AND NOT-IL13RA1	0.867188	0.850575	0.884462
COMPLEX-SPON2/SLC4A1/CD33	0.867031	0.798658	0.948207
CD33 AND NOT-ACVR1C AND NOT-CLDN23	0.899614	0.872659	0.928287
CD33 AND NOT-ABHD6 AND NOT-CLDN5	0.876866	0.824561	0.936255
CD33 AND NOT-CD22 AND SLC4A1	0.914761	0.956522	0.876494

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
PTPRZ1 AND NOT-CLDN1 AND NOT-KCNJ6	0.827586	0.878049	0.782609
PTPRZ1 AND NOT-EPCAM AND NOT-KCNJ6	0.827586	0.878049	0.782609
PTPRZ1 AND NOT-PMEL AND NOT-KCNJ6	0.827586	0.878049	0.782609
PTPRZ1 AND NCAM1 AND NOT-KCNJ6	0.827586	0.878049	0.782609
PTPRZ1 AND NOT-CLDN23 AND NOT-KCNJ6	0.827586	0.878049	0.782609
PTPRZ1 AND NOT-STEAP1 AND NOT-KCNJ6	0.827586	0.878049	0.782609
PTPRZ1 AND NOT-KCNJ6 AND NOT-CD52	0.827586	0.878049	0.782609
PTPRZ1 AND NOT-TRPM4 AND NOT-KCNJ6	0.827586	0.878049	0.782609
PTPRZ1 AND NOT-SDC1 AND NOT-KCNJ6	0.827586	0.878049	0.782609
PTPRZ1 AND NOT-SDC1 AND STAB1	0.826087	0.826087	0.826087
PTPRZ1 AND SYT6 AND NOT-SDC1	0.826087	0.826087	0.826087
VANGL2 AND NOT-EPCAM AND GABRG2	0.826087	0.826087	0.826087
PTPRZ1 AND NOT-SDC1 AND NOT-GPR22	0.823529	0.897436	0.76087
PTPRZ1 AND NOT-SDC1 AND NOT-SYT4	0.823529	0.897436	0.76087
VANGL2 AND NOT-EPCAM AND P2RX4	0.864198	1	0.76087
SLC1A2 AND F2RL1 AND NOT-CLDN1	0.827586	0.878049	0.782609
PTPRZ1 AND SEMA5B AND NOT-CNTNAP2	0.819277	0.918919	0.73913
PTPRZ1 AND NOT-SDC1 AND VANGL1	0.819277	0.918919	0.73913
VANGL2 AND NOT-TPBG AND NOT-CNTNAP2	0.818182	0.857143	0.782609
PTPRZ1 AND NOT-CLDN23 AND SEMA4B	0.817204	0.808511	0.826087
PTPRZ1 AND NOT-SDC1 AND SEMA4B	0.817204	0.808511	0.826087
PTPRZ1 AND NOT-TRPM4 AND SEMA4B	0.817204	0.808511	0.826087
PTPRZ1 AND NCAM1 AND SEMA4B	0.817204	0.808511	0.826087
PTPRZ1 AND NOT-CLDN1 AND SEMA4B	0.817204	0.808511	0.826087
SLC1A2 AND F2R AND NOT-SDC1	0.835165	0.844444	0.826087
SLC1A2 AND F2R AND NOT-CLDN1	0.835165	0.844444	0.826087
PTPRZ1 AND NOT-SDK1 AND HAS2	0.917647	1	0.847826
PTPRZ1 AND NOT-FZD10 AND HAS2	0.917647	1	0.847826
PTPRZ1 AND NOT-ZDHHC5 AND HAS2	0.906977	0.975	0.847826
PTPRZ1 AND NOT-SMAGP AND HAS2	0.906977	0.975	0.847826
PTPRZ1 AND NOT-GPR87 AND HAS2	0.906977	0.975	0.847826
PTPRZ1 AND SLC8A2 AND HAS2	0.906977	0.975	0.847826
PTPRZ1 AND GPM6B AND HAS2	0.906977	0.975	0.847826
PTPRZ1 AND NOT-DUOX1 AND HAS2	0.906977	0.975	0.847826
PTPRZ1 AND SEMA6D AND HAS2	0.906977	0.975	0.847826
PTPRZ1 AND SCAMP5 AND HAS2	0.906977	0.975	0.847826
PTPRZ1 AND NOT-F11R AND HAS2	0.906977	0.975	0.847826
PTPRZ1 AND NOT-ST14 AND HAS2	0.906977	0.975	0.847826
PTPRZ1 AND NOT-SLC46A2 AND HAS2	0.906977	0.975	0.847826
PTPRZ1 AND NOT-BTC AND HAS2	0.906977	0.975	0.847826
PTPRZ1 AND NOT-ACPP AND HAS2	0.906977	0.975	0.847826
PTPRZ1 AND NOT-PERP AND HAS2	0.906977	0.975	0.847826
PTPRZ1 AND NOT-COL17A1 AND HAS2	0.906977	0.975	0.847826
PTPRZ1 AND NOT-RECK AND HAS2	0.906977	0.975	0.847826
PTPRZ1 AND NOT-AQP3 AND HAS2	0.906977	0.975	0.847826
PTPRZ1 AND NOT-SPINT1 AND HAS2	0.906977	0.975	0.847826
PTPRZ1 AND NOT-SLC6A14 AND HAS2	0.906977	0.975	0.847826
PTPRZ1 AND SYNDIG1 AND HAS2	0.906977	0.975	0.847826
PTPRZ1 AND NOT-LRP11 AND HAS2	0.906977	0.975	0.847826
PTPRZ1 AND NOT-IL22RA1 AND HAS2	0.906977	0.975	0.847826
PTPRZ1 AND GPR173 AND HAS2	0.906977	0.975	0.847826
PTPRZ1 AND NOT-ATP2C2 AND HAS2	0.906977	0.975	0.847826
PTPRZ1 AND ADAM22 AND HAS2	0.906977	0.975	0.847826
PTPRZ1 AND NOT-SLC15A1 AND HAS2	0.906977	0.975	0.847826
PTPRZ1 AND NOT-ACVRL1 AND HAS2	0.906977	0.975	0.847826
PTPRZ1 AND LRRTM2 AND HAS2	0.906977	0.975	0.847826
PTPRZ1 AND NOT-MYOF AND HAS2	0.906977	0.975	0.847826
PTPRZ1 AND SLC4A4 AND HAS2	0.906977	0.975	0.847826
PTPRZ1 AND NOT-F11R AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND NOT-SLC35G1 AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND NOT-SLC6A14 AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND NOT-ATP2C2 AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND NOT-FZD10 AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND NOT-SPPL3 AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND ADAM22 AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND CLSTN2 AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND NOT-SDK1 AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND NOT-RECK AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND NOT-SLCO4C1 AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND NOT-ST14 AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND NOT-ACPP AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND LRRTM2 AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND NOT-TNFSF10 AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND NOT-ANO1 AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND NOT-IL1R2 AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND NOT-ATP10D AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND NOT-ZDHHC5 AND MPZL1	0.904762	1	0.826087

Antigen Pair Logic	F1	Precision	Recall
CD33 AND NOT-ABHD6 AND NOT-EDNRB	0.871698	0.827957	0.920319
CD33 AND NOT-LPAR1 AND CD70	0.880455	0.84058	0.924303
CD33 AND NOT-CD79A AND SLC4A1	0.908714	0.948052	0.87251
CD33 AND NOT-TNFRSF17 AND SLC4A1	0.915612	0.973094	0.864542
COMPLEX-IL11RA/SLC4A1/CD33	0.853229	0.838462	0.868526
CD33 AND NOT-CD72 AND SLC4A1	0.91858	0.964912	0.876494
CD33 AND NOT-ABHD6 AND CD70	0.889313	0.85348	0.928287
COMPLEX-PCYT1A/SLC4A1/CD33	0.851211	0.752294	0.98008
CD33 AND NOT-ACVR1C AND NOT-CLDN11	0.847826	0.777409	0.932271
CD33 AND NOT-DAG1 AND CD70	0.87218	0.825623	0.924303
COMPLEX-ST8SIA1/SLC4A1/CD33	0.84375	0.747692	0.968127
CD33 AND NOT-CD19 AND SLC4A1	0.906445	0.947826	0.868526
CD33 AND NOT-EDNRB AND NOT-KIAA1324	0.836364	0.769231	0.916335
COMPLEX-OAS1/SLC4A1/CD33	0.88632	0.858209	0.916335
CD33 AND NOT-ACVR1C AND NOT-ERBB2	0.83274	0.752412	0.932271
COMPLEX-ITPR3/CLDN5/CD33	0.831858	0.748408	0.936255
CD33 AND NOT-SLC25A4 AND SLC7A5	0.857678	0.809187	0.912351
NOT-ABCA5 AND CD33 AND SLC4A1	0.825054	0.900943	0.760956
COMPLEX-SLC4A1/SLC7A5/CD33	0.85461	0.769968	0.960159
NOT-PTPRK AND CD33 AND NOT-P2RX5	0.816415	0.891509	0.752988
NOT-PODXL AND BIRC5 AND CD33	0.814655	0.887324	0.752988
NOT-PODXL AND CD33 AND PROM1	0.8125	0.923858	0.7251
CD33 AND NOT-P2RX5 AND SLC4A1	0.808314	0.961538	0.697211
CD33 AND NOT-ENPP5 AND NOT-EDNRB	0.801394	0.712074	0.916335
NOT-PHLDB2 AND SLC7A5 AND ANXA1	0.953668	0.925094	0.984064
NOT-PHLDB2 AND NOT-ERBB3 AND DDX3X	0.939163	0.898182	0.984064
NOT-PHLDB2 AND SLC7A5 AND DDX3X	0.936416	0.906716	0.968127
NOT-PHLDB2 AND SLC7A5 AND CD52	0.930502	0.902622	0.960159
SLC22A16 AND SLC7A5 AND NOT-EDNRB	0.923695	0.931174	0.916335
NOT-PHLDB2 AND DDX3X AND NOT-BMPR1B	0.923664	0.886447	0.964143
NOT-PHLDB2 AND DDX3X AND NOT-FOLH1	0.923372	0.889299	0.960159
NOT-PHLDB2 AND DDX3X AND NOT-SDC1	0.916667	0.873646	0.964143
NOT-PHLDB2 AND DDX3X AND NOT-CLDN7	0.916667	0.873646	0.964143
NOT-PHLDB2 AND DDX3X AND NOT-ERBB2	0.916667	0.873646	0.964143
NOT-AXL AND EMP3 AND NOT-CD160	0.915254	0.867857	0.968127
NOT-PHLDB2 AND DDX3X AND NOT-CLDN8	0.914934	0.870504	0.964143
NOT-PHLDB2 AND DDX3X AND NOT-RNF43	0.914934	0.870504	0.964143
NOT-PHLDB2 AND DDX3X AND NOT-EPCAM	0.914611	0.873188	0.960159
NOT-PHLDB2 AND DDX3X AND NOT-VTCN1	0.913208	0.867384	0.964143
NOT-AXL AND EMP3 AND NOT-SLAMF7	0.911488	0.864286	0.964143
NOT-PHLDB2 AND DDX3X AND NOT-STEAP1	0.910476	0.872263	0.952191
NOT-PHLDB2 AND DDX3X AND NOT-MET	0.908397	0.871795	0.948207
NOT-AXL AND EMP3 AND NOT-FCRL5	0.908068	0.858156	0.964143
NOT-AXL AND EMP3 AND NOT-HLA-DOB	0.907407	0.847751	0.976096
NOT-PHLDB2 AND DDX3X AND NOT-MUC1	0.902111	0.87037	0.936255
NOT-EDNRB AND EMP3 AND NOT-CD160	0.895238	0.857664	0.936255
NOT-EDNRB AND EMP3 AND NOT-SLAMF7	0.893536	0.854545	0.936255
NOT-ERBB2 AND CD44 AND NOT-SLAMF7	0.892989	0.831615	0.964143
NOT-AXL AND EMP3 AND NOT-TNFRSF17	0.892193	0.836237	0.956175
NOT-AXL AND EMP3 AND NOT-CD22	0.892139	0.824324	0.972112
NOT-KDR AND EMP3 AND NOT-HLA-DOB	0.891304	0.817276	0.98008
NOT-EDNRB AND EMP3 AND NOT-MS4A1	0.886792	0.842294	0.936255
NOT-ALDH1A1 AND SLC30A1 AND NOT-EGFR	0.885496	0.849817	0.924303
NOT-PHLDB2 AND SLC7A5 AND NOT-ERBB4	0.883721	0.801948	0.984064
NOT-PHLDB2 AND SLC7A5 AND NOT-ERBB4	0.883721	0.801948	0.984064
NOT-AXL AND EMP3 AND NOT-CD72	0.883636	0.812709	0.968127
NOT-ERBB2 AND CD44 AND NOT-HLA-DOB	0.882143	0.799353	0.984064
NOT-ALDH1A1 AND SLC30A1 AND NOT-ERBB2	0.878788	0.837545	0.924303
NOT-EDNRB AND SLC7A5 AND NOT-STEAP4	0.878505	0.827465	0.936255
NOT-EDNRB AND EMP3 AND NOT-HLA-DOB	0.878229	0.817869	0.948207
NOT-EDNRB AND EMP3 AND NOT-FCRL5	0.877778	0.820069	0.944223
NOT-EDNRB AND SLC7A5 AND NOT-S1PR1	0.875676	0.799342	0.968127
NOT-EDNRB AND SLC7A5 AND NOT-SYT11	0.875	0.812287	0.948207
NOT-PHLDB2 AND ENG AND NOT-FOLH1	0.873706	0.909483	0.840637
NOT-EDNRB AND EMP3 AND NOT-TNFRSF17	0.870544	0.822695	0.924303
NOT-PHLDB2 AND ENG AND NOT-ERBB3	0.870103	0.901709	0.840637
NOT-ALDH1A1 AND SLC30A1 AND NOT-SLC39A6	0.870544	0.822695	0.924303
NOT-ALDH1A1 AND EMP3 AND NOT-AXL	0.868526	0.868526	0.868526
NOT-ALDH1A1 AND CD44 AND NOT-ERBB2	0.867925	0.824373	0.916335
NOT-ALDH1A1 AND EMP3 AND NOT-KDR	0.865613	0.858824	0.87251
NOT-EDNRB AND EMP3 AND NOT-CD22	0.864964	0.79798	0.944223
LAPTM5 AND NOT-ERBB2 AND NOT-HLA-DOB	0.890071	0.801917	1
NOT-AXL AND EMP3 AND NOT-OAS1	0.863551	0.81338	0.920319
NOT-EDNRB AND SLC7A5 AND SLCO1B3	0.860254	0.79	0.944223
NOT-ERBB2 AND CD44 AND NOT-CD19	0.857143	0.784768	0.944223
NOT-PHLDB2 AND NOT-CEACAM5 AND NOT-ERBB4	0.853701	0.751515	0.988048
NOT-ALDH1A1 AND CD44 AND NOT-CLDN8	0.856611	0.804196	0.916335

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
PTPRZ1 AND SEMA6D AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND SYNDIG1 AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND NOT-SLC16A10 AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND NOT-IL22RA1 AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND NOT-LY6K AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND NOT-ACP1 AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND NOT-IL2RB AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND NOT-SLC46A2 AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND NOT-GPR87 AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND NOT-MYOF AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND NOT-SPINT1 AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND GPM6B AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND NOT-ITPR3 AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND NOT-LRP11 AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND NOT-SMAGP AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND NOT-PERP AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND NOT-APCDD1 AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND SLC4A4 AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND NOT-SLMAP AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND NOT-ACVRL1 AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND NOT-SLC15A1 AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND NOT-AQP9 AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND GPR173 AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND NOT-F2RL1 AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND SLC8A2 AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND NOT-SACM1L AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND NOT-SLC12A7 AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND MPZL1 AND NOT-GDE1	0.904762	1	0.826087
PTPRZ1 AND NOT-KCNK5 AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND NOT-BTC AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND NOT-DUOX1 AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND OR51B6 AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND NOT-CDH1 AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND NRCAM AND F2RL1	0.904762	1	0.826087
PTPRZ1 AND NOT-TMEM30B AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND NOT-AQP3 AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND NOT-P2RY14 AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND NOT-TMEM30B AND HAS2	0.898876	0.930233	0.869565
PTPRZ1 AND NRCAM AND CLCN5	0.898876	0.930233	0.869565
PTPRZ1 AND ITM2B AND NOT-SACM1L	0.896552	0.95122	0.847826
PTPRZ1 AND NOT-LRP11 AND ITM2B	0.896552	0.95122	0.847826
PTPRZ1 AND NRCAM AND TPCN2	0.888889	0.909091	0.869565
PTPRZ1 AND NOT-FZD10 AND F2RL1	0.888889	0.909091	0.869565
PTPRZ1 AND NRCAM AND NOT-LRP11	0.888889	0.909091	0.869565
PTPRZ1 AND NOT-FZD10 AND F2RL1	0.888889	0.909091	0.869565
PTPRZ1 AND ITM2B AND NOT-RECK	0.886364	0.928571	0.847826
PTPRZ1 AND ITM2B AND NOT-COL17A1	0.886364	0.928571	0.847826
PTPRZ1 AND GPR85 AND HAS2	0.906977	0.975	0.847826
PTPRZ1 AND F2R AND NOT-FZD10	0.906977	0.975	0.847826
PTPRZ1 AND NOT-CD36 AND HAS2	0.906977	0.975	0.847826
PTPRZ1 AND CRB1 AND HAS2	0.906977	0.975	0.847826
PTPRZ1 AND SLCO1C1 AND HAS2	0.906977	0.975	0.847826
PTPRZ1 AND NOT-KCNK6 AND HAS2	0.906977	0.975	0.847826
PTPRZ1 AND NOT-LRIG3 AND HAS2	0.906977	0.975	0.847826
PTPRZ1 AND MOG AND HAS2	0.906977	0.975	0.847826
PTPRZ1 AND SLCO1A2 AND HAS2	0.906977	0.975	0.847826
PTPRZ1 AND CLDN17 AND HAS2	0.906977	0.975	0.847826
PTPRZ1 AND ATP2B2 AND HAS2	0.906977	0.975	0.847826
PTPRZ1 AND KCNJ10 AND HAS2	0.906977	0.975	0.847826
PTPRZ1 AND SLC4A8 AND HAS2	0.906977	0.975	0.847826
PTPRZ1 AND NOT-ATP8B1 AND HAS2	0.906977	0.975	0.847826
PTPRZ1 AND SLC1A2 AND HAS2	0.906977	0.975	0.847826
PTPRZ1 AND GABRB2 AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND NOT-SLC43A1 AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND KCNA2 AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND GABBR2 AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND MPZL1 AND NOT-MRGPRF	0.904762	1	0.826087
PTPRZ1 AND NOT-CD36 AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND CRB1 AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND SLCO1C1 AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND GPR85 AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND NOT-EPHA2 AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND NOT-KCNK6 AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND NOT-LRIG3 AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND NOT-AOC3 AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND KCNJ10 AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND CLDN17 AND MPZL1	0.904762	1	0.826087

Antigen Pair Logic	F1	Precision	Recall
NOT-ITGAV AND SLC39A6 AND NOT-BTN3A1	0.851563	0.835249	0.868526
NOT-STEAP1 AND EMP3 AND NOT-HLA-DOB	0.851064	0.766773	0.956175
NOT-STEAP1 AND EMP3 AND NOT-SLAMF7	0.849462	0.771987	0.944223
NOT-PHLDB2 AND SLC7A5 AND NOT-STEAP2	0.847341	0.743976	0.984064
NOT-PHLDB2 AND SLC7A5 AND NOT-STEAP2	0.847341	0.743976	0.984064
LAPTM5 AND NOT-ERBB2 AND NOT-SLAMF7	0.875445	0.790997	0.98008
NOT-ERBB2 AND EMP3 AND NOT-HLA-DOB	0.845754	0.748466	0.972112
NOT-ALDH1A1 AND VAMP8 AND NOT-EGFR	0.845173	0.778523	0.924303
NOT-PHLDB2 AND NOT-ERBB3 AND MUC1	0.844538	0.893333	0.800797
NOT-PHLDB2 AND NOT-ERBB3 AND MUC1	0.844538	0.893333	0.800797
NOT-ERBB2 AND CD44 AND NOT-OAS1	0.844203	0.774086	0.928287
CD37 AND NOT-PHLDB2 AND MUC1	0.843956	0.941176	0.76494
NOT-PHLDB2 AND CD70 AND NOT-FOLH1	0.843478	0.92823	0.772908
NOT-ERBB2 AND CD44 AND NOT-IL11RA	0.843066	0.777778	0.920319
NOT-ABCA5 AND SLC30A1 AND NOT-ERBB2	0.842324	0.878788	0.808765
NOT-ERBB2 AND EMP3 AND SLC7A5	0.840426	0.757188	0.944223
NOT-ERBB2 AND EMP3 AND NOT-SLAMF7	0.839721	0.74613	0.960159
NOT-EDNRB AND EMP3 AND NOT-CD79A	0.839286	0.760518	0.936255
NOT-ABCA5 AND EMP3 AND NOT-AXL	0.837607	0.903226	0.780876
NOT-ATP8B1 AND SLC7A5 AND NOT-EDNRB	0.836555	0.748428	0.948207
NOT-ATP8B1 AND SLC7A5 AND NOT-EDNRB	0.836555	0.748428	0.948207
NOT-PHLDB2 AND NOT-FOLH1 AND MUC1	0.836134	0.884444	0.792829
NOT-PHLDB2 AND NOT-FOLH1 AND MUC1	0.836134	0.884444	0.792829
NOT-ERBB2 AND CD44 AND SLC7A5	0.834783	0.740741	0.956175
LAPTM5 AND NOT-CLDN7 AND NOT-HLA-DOB	0.896429	0.812298	1
NOT-EDNRB AND SLC7A5 AND NOT-TGFBI	0.834559	0.774744	0.904382
NOT-ERBB2 AND EMP3 AND NOT-SPON2	0.831889	0.736196	0.956175
CD44 AND NOT-SPON2 AND NOT-CLDN8	0.831793	0.775862	0.896414
NOT-EDNRB AND SLC43A1 AND SLC7A5	0.829694	0.917874	0.756972
LAPTM5 AND SLC7A5 AND NOT-CLDN5	0.835616	0.732733	0.972112
LAPTM5 AND SLC7A5 AND NOT-CLDN11	0.845754	0.748466	0.972112
NOT-EDNRB AND SLC7A5 AND P2RX4	0.828571	0.750809	0.924303
NOT-ERBB2 AND SLC7A5 AND VAMP8	0.828523	0.721893	0.972112
NOT-PHLDB2 AND SLC7A5 AND CD70	0.828179	0.728097	0.960159
NOT-ABCA5 AND CD44 AND NOT-FAP	0.828157	0.862069	0.796813
NOT-EDNRB AND CD44 AND SLC7A5	0.827709	0.746795	0.928287
LAPTM5 AND SLC7A5 AND NOT-ROR1	0.835616	0.732733	0.972112
NOT-GPR137B AND SLC7A5 AND NOT-EDNRB	0.825911	0.839506	0.812749
LAPTM5 AND NOT-ERBB2 AND SLC7A5	0.844291	0.746177	0.972112
NOT-ERBB2 AND CD44 AND NOT-PCYT1A	0.825503	0.713043	0.98008
NOT-ABCA5 AND CD44 AND NOT-ERBB2	0.824742	0.854701	0.796813
CD44 AND NOT-CLDN8 AND NOT-SLAMF7	0.869565	0.797342	0.956175
NOT-EDNRB AND SLC7A5 AND SLC30A1	0.824324	0.715543	0.972112
NOT-STEAP1 AND EMP3 AND SLC7A5	0.821869	0.737342	0.928287
NOT-PHLDB2 AND ENG AND NOT-ERBB4	0.821012	0.802281	0.840637
NOT-ABCA5 AND EMP3 AND NOT-KDR	0.820833	0.860262	0.784861
CD44 AND NOT-CLDN8 AND NOT-HLA-DOB	0.862676	0.772871	0.976096
NOT-PHLDB2 AND CD33 AND SLC7A5	0.961771	0.971545	0.952191
NOT-PHLDB2 AND CD33 AND NOT-STEAP2	0.942801	0.933594	0.952191
NOT-PHLDB2 AND CD33 AND NOT-CLDN8	0.940945	0.929961	0.952191
NOT-PHLDB2 AND CD33 AND NOT-RNF43	0.939096	0.926357	0.952191
COMPLEX-VTCN1/PHLDB2/CD33	0.939096	0.926357	0.952191
NOT-PHLDB2 AND CD33 AND MUC1	0.871111	0.984925	0.780876
CD33 AND NOT-GHR AND SLC7A5	0.85283	0.810036	0.900398
NOT-EDNRB AND SLC7A5 AND NOT-HLA-DOB	0.872987	0.792208	0.972112
NOT-ABCA5 AND SLC7A5 AND NOT-EDNRB	0.824268	0.867841	0.784861
NOT-EDNRB AND SLC7A5 AND DDX3X	0.813675	0.712575	0.948207
NOT-EDNRB AND SLC7A5 AND NOT-CEACAM5	0.806612	0.689266	0.972112
CD33 AND NOT-EDNRB AND SLC7A5	0.850662	0.809353	0.896414
Liposarcoma			
ADAM12 AND NOT-ITGA6 AND SLC6A18	0.865672	0.935484	0.805556
CNTNAP1 AND CD68 AND NOT-GPR22	0.833333	0.833333	0.833333
NOT-CXADR AND LIFR AND TNFRSF10B	0.80597	0.870968	0.75
NOT-CXADR AND MCAM AND TNFRSF10B	0.84375	0.964286	0.75
SLC2A10 AND PROCR AND NOT-ITGA6	0.860759	0.790698	0.944444
ADAM12 AND KCNV2 AND S100A10	0.837838	0.815789	0.861111
ADAM12 AND FCAMR AND NOT-CLCA2	0.805195	0.756098	0.861111
PLXND1 AND NOT-MUC15 AND GABRA1	0.842105	0.8	0.888889
CNTNAP1 AND NOT-GPR22 AND CHIC2	0.84507	0.857143	0.833333
ADAM12 AND NOT-ITGA6 AND KL	0.828571	0.852941	0.805556
ADAM12 AND NOT-LRP2 AND CHIC2	0.810811	0.789474	0.833333
ADAM12 AND NOT-ITGA6 AND SIRPA	0.861111	0.861111	0.861111
ADAM12 AND NOT-ITGA6 AND DUOX1	0.826667	0.794872	0.861111
ADAM12 AND NOT-ITGA6 AND CD4	0.826667	0.794872	0.861111
ADAM12 AND NOT-ITGA6 AND CLEC2B	0.805195	0.756098	0.861111
ADAM12 AND NOT-ITGA6 AND CD68	0.837838	0.815789	0.861111
ADAM12 AND NOT-ITGA6 AND S100A10	0.869565	0.909091	0.833333

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
PTPRZ1 AND SLC4A8 AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND SLC1A2 AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND MOG AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND CSF1R AND NOT-DYSF	0.896552	0.95122	0.847826
PTPRZ1 AND F2R AND NOT-SDK1	0.896552	0.95122	0.847826
PTPRZ1 AND CSF1R AND NOT-DYSF	0.896552	0.95122	0.847826
PTPRZ1 AND NOT-FZD10 AND ANTXR2	0.896552	0.95122	0.847826
PTPRZ1 AND CLDN17 AND F2RL1	0.894118	0.974359	0.826087
PTPRZ1 AND CLDN17 AND ABCA1	0.894118	0.974359	0.826087
PTPRZ1 AND NOT-TMEM30B AND NOT-GRM3	0.894118	0.974359	0.826087
PTPRZ1 AND NOT-LRP11 AND F2R	0.891304	0.891304	0.891304
PTPRZ1 AND NOT-FZD10 AND NOT-GABRA2	0.891304	0.891304	0.891304
PTPRZ1 AND NOT-IL22RA1 AND F2R	0.888889	0.909091	0.869565
PTPRZ1 AND NOT-ABCC5 AND NOT-GABRA2	0.901099	0.911111	0.891304
PTPRZ1 AND NOT-CDH3 AND NOT-GABRA2	0.88172	0.87234	0.891304
PTPRZ1 AND GPR173 AND NOT-GABRA2	0.88172	0.87234	0.891304
PTPRZ1 AND NOT-SLCO4C1 AND NOT-GABRA2	0.88172	0.87234	0.891304
PTPRZ1 AND NOT-AQP9 AND NOT-GABRA2	0.88172	0.87234	0.891304
PTPRZ1 AND NOT-SLC35G1 AND NOT-GABRA2	0.88172	0.87234	0.891304
PTPRZ1 AND CLSTN2 AND NOT-GABRA2	0.88172	0.87234	0.891304
PTPRZ1 AND NOT-P2RY14 AND NOT-GABRA2	0.88172	0.87234	0.891304
PTPRZ1 AND NOT-APCDD1 AND NOT-GABRA2	0.88172	0.87234	0.891304
PTPRZ1 AND NOT-F11R AND NOT-GABRA2	0.88172	0.87234	0.891304
PTPRZ1 AND NOT-CDH1 AND NOT-GABRA2	0.88172	0.87234	0.891304
PTPRZ1 AND NOT-IL1R2 AND NOT-GABRA2	0.88172	0.87234	0.891304
PTPRZ1 AND SLC4A4 AND NOT-GABRA2	0.88172	0.87234	0.891304
PTPRZ1 AND NOT-LY6K AND NOT-GABRA2	0.88172	0.87234	0.891304
PTPRZ1 AND NOT-SLCO4C1 AND NOT-GABRA2	0.88172	0.87234	0.891304
PTPRZ1 AND NOT-GRM3 AND NOT-TNFSF10	0.880952	0.973684	0.804348
PTPRZ1 AND SLC8A2 AND NOT-GRM3	0.880952	0.973684	0.804348
PTPRZ1 AND NOT-SLC6A14 AND NOT-GRM3	0.880952	0.973684	0.804348
PTPRZ1 AND NOT-GRM3 AND NOT-FCER1A	0.880952	0.973684	0.804348
PTPRZ1 AND NOT-F11R AND NOT-GRM3	0.880952	0.973684	0.804348
PTPRZ1 AND SEMA6D AND NOT-GRM3	0.880952	0.973684	0.804348
PTPRZ1 AND LRRTM2 AND NOT-GRM3	0.880952	0.973684	0.804348
PTPRZ1 AND GPM6B AND NOT-GRM3	0.880952	0.973684	0.804348
PTPRZ1 AND NOT-SDK1 AND NOT-GRM3	0.880952	0.973684	0.804348
PTPRZ1 AND NOT-BTC AND NOT-GRM3	0.880952	0.973684	0.804348
PTPRZ1 AND NOT-IL22RA1 AND NOT-GRM3	0.880952	0.973684	0.804348
PTPRZ1 AND NOT-PERP AND NOT-GRM3	0.880952	0.973684	0.804348
PTPRZ1 AND NOT-LRP11 AND NOT-GRM3	0.880952	0.973684	0.804348
PTPRZ1 AND NOT-GRM3 AND NOT-LSR	0.880952	0.973684	0.804348
PTPRZ1 AND ADAM22 AND NOT-GRM3	0.880952	0.973684	0.804348
PTPRZ1 AND NOT-ZDHHC5 AND NOT-GRM3	0.880952	0.973684	0.804348
PTPRZ1 AND NOT-GRM3 AND NOT-COL17A1	0.880952	0.973684	0.804348
PTPRZ1 AND NOT-GRM3 AND SLC4A4	0.880952	0.973684	0.804348
PTPRZ1 AND NOT-GPR87 AND NOT-GRM3	0.880952	0.973684	0.804348
PTPRZ1 AND NOT-DUOX1 AND NOT-GRM3	0.880952	0.973684	0.804348
PTPRZ1 AND NOT-SMAGP AND NOT-GRM3	0.880952	0.973684	0.804348
PTPRZ1 AND SCAMP5 AND NOT-GRM3	0.880952	0.973684	0.804348
PTPRZ1 AND NOT-ATP2C2 AND NOT-GRM3	0.880952	0.973684	0.804348
PTPRZ1 AND GPRC5B AND NOT-GRM3	0.880952	0.973684	0.804348
PTPRZ1 AND NOT-FZD10 AND NOT-GRM3	0.880952	0.973684	0.804348
PTPRZ1 AND NOT-SLC15A1 AND NOT-GRM3	0.880952	0.973684	0.804348
PTPRZ1 AND NOT-GRM3 AND NOT-RECK	0.880952	0.973684	0.804348
PTPRZ1 AND SYNDIG1 AND NOT-GRM3	0.880952	0.973684	0.804348
PTPRZ1 AND NOT-ST14 AND NOT-GRM3	0.880952	0.973684	0.804348
PTPRZ1 AND NOT-AQP3 AND NOT-GRM3	0.880952	0.973684	0.804348
PTPRZ1 AND NOT-GRM3 AND NOT-ACPP	0.880952	0.973684	0.804348
PTPRZ1 AND NOT-SMAGP AND NOT-GRM3	0.880952	0.973684	0.804348
PTPRZ1 AND NOT-ACVRL1 AND NOT-GRM3	0.880952	0.973684	0.804348
PTPRZ1 AND NOT-GRM3 AND NOT-SLC46A2	0.880952	0.973684	0.804348
PTPRZ1 AND NOT-SPINT1 AND NOT-GRM3	0.880952	0.973684	0.804348
PTPRZ1 AND GPR173 AND NOT-GRM3	0.880952	0.973684	0.804348
PTPRZ1 AND NOT-MYOF AND NOT-GRM3	0.880952	0.973684	0.804348
PTPRZ1 AND NOT-PTPRM AND NOT-GABRA2	0.891304	0.891304	0.891304
PTPRZ1 AND PLXNA4 AND NOT-GABRA2	0.88172	0.87234	0.891304
PTPRZ1 AND NOT-SDC1 AND HAS2	0.917647	1	0.847826
PTPRZ1 AND NOT-STEAP1 AND HAS2	0.906977	0.975	0.847826
PTPRZ1 AND NCAM1 AND HAS2	0.906977	0.975	0.847826
PTPRZ1 AND NOT-CLDN1 AND HAS2	0.906977	0.975	0.847826
PTPRZ1 AND NOT-TRPM4 AND HAS2	0.906977	0.975	0.847826
PTPRZ1 AND NOT-CLDN23 AND HAS2	0.906977	0.975	0.847826
PTPRZ1 AND NOT-PMEL AND HAS2	0.906977	0.975	0.847826
PTPRZ1 AND NOT-CLDN23 AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND NOT-STEAP1 AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND NOT-CD52 AND MPZL1	0.904762	1	0.826087

Antigen Pair Logic	F1	Precision	Recall
ADAM12 AND NOT-ITGA6 AND EMP1	0.873239	0.885714	0.861111
ADAM12 AND NOT-ITGA6 AND CXCR4	0.828571	0.852941	0.805556
ADAM12 AND NOT-LRP2 AND NCAM2	0.8	0.769231	0.833333
SLC2A10 AND EMP1 AND NOT-ITGA6	0.871795	0.809524	0.944444
ADAM12 AND NOT-ITGA6 AND CLPTM1	0.869565	0.909091	0.833333
SLC2A10 AND NOT-IYD AND CD68	0.825	0.75	0.916667
SLC2A10 AND GJA1 AND NOT-ITGA6	0.849315	0.837838	0.861111
ADAM12 AND NOT-ITGA6 AND EDNRB	0.826667	0.794872	0.861111
CD276 AND NOT-ITGA6 AND NOT-ATP12A	0.873239	0.885714	0.861111
EPHB2 AND NOT-IHH AND CD68	0.810811	0.789474	0.833333
CD276 AND NOT-ITGA6 AND MMD	0.933333	0.897436	0.972222
CD276 AND NOT-ITGA6 AND KL	0.891892	0.868421	0.916667
CD276 AND NOT-MUC15 AND CLEC2B	0.873239	0.885714	0.861111
EPHB2 AND NOT-SLC26A3 AND CD68	0.810811	0.789474	0.833333
CD276 AND NOT-ESYT3 AND STOML3	0.868421	0.825	0.916667
EPHB2 AND NOT-SLC26A3 AND S100A10	0.821918	0.810811	0.833333
EPHB2 AND NOT-SLC26A3 AND STOML3	0.821918	0.810811	0.833333
EPHB2 AND NOT-SLC26A3 AND WNT1	0.805556	0.805556	0.805556
CD276 AND NOT-FZD10 AND NOT-PON2	0.814815	0.733333	0.916667
CD276 AND NOT-GJB3 AND NOT-LRP2	0.805195	0.756098	0.861111
CD276 AND NOT-ESYT3 AND CLEC2B	0.868421	0.825	0.916667
EPHB2 AND NOT-IYD AND CD68	0.815789	0.775	0.861111
EPHB2 AND NOT-SLC26A3 AND OR5L2	0.805556	0.805556	0.805556
EPHB2 AND NOT-SLC26A3 AND TAS1R2	0.833333	0.833333	0.833333
EPHB2 AND NOT-IHH AND NOT-FZD10	0.852941	0.90625	0.805556
EPHB2 AND NOT-IHH AND OR10H2	0.810811	0.789474	0.833333
EPHB2 AND NOT-SLC26A3 AND CHIC2	0.821918	0.810811	0.833333
EPHB2 AND RECK AND NOT-GJB3	0.828571	0.852941	0.805556
CD276 AND NOT-DUOX1 AND CLEC2B	0.868421	0.825	0.916667
EPHB2 AND SLC41A1 AND NOT-MUC15	0.811594	0.848485	0.777778
EPHB2 AND NOT-SLC26A3 AND RECK	0.816901	0.828571	0.805556
CNTNAP1 AND NOT-ABCA12 AND TNFRSF10B	0.828571	0.852941	0.805556
CD276 AND NOT-ITGA6 AND NOT-MSMO1	0.868421	0.825	0.916667
EPHB2 AND NOT-IYD AND NOT-GPR22	0.84507	0.857143	0.833333
EPHB2 AND NOT-SLC26A3 AND PCDHA6	0.821918	0.810811	0.833333
EPHB2 AND NOT-IHH AND SLC22A12	0.8	0.769231	0.833333
EPHB2 AND NOT-IHH AND NPHS2	0.8	0.769231	0.833333
EPHB2 AND NOT-SLC26A3 AND CHRNB4	0.810811	0.789474	0.833333
EPHB2 AND NOT-SLC26A3 AND PLVAP	0.821918	0.810811	0.833333
EPHB2 AND NOT-SLC26A3 AND OR2C3	0.833333	0.833333	0.833333
EPHB2 AND NOT-SLC30A10 AND STOML3	0.84507	0.857143	0.833333
EPHB2 AND NOT-SLC26A3 AND ROS1	0.821918	0.810811	0.833333
EPHB2 AND NOT-SLC26A3 AND OR52D1	0.8	0.769231	0.833333
EPHB2 AND SLC41A1 AND NOT-GPR22	0.882353	0.9375	0.833333
EPHB2 AND RECK AND NOT-ABCA12	0.828571	0.852941	0.805556
CD276 AND NOT-GJB3 AND NOT-SLC34A2	0.829268	0.73913	0.944444
ATP8B2 AND NOT-GPR22 AND GPR85	0.84507	0.857143	0.833333
Lung Adenocarcinoma			
NOT-CLCA4 AND SLC6A14 AND NOT-FXYD1	0.81761	0.730337	0.928571
Lung Carcinoma			
NOT-ADCY4 AND GJB2 AND ADAM12	0.815385	0.80303	0.828125
ADAM12 AND NOT-NRG2 AND IGSF9	0.823529	0.777778	0.875
Diffuse B-Cell Lymphoma			
FCRL3 AND NOT-FCER1A AND FPR1	0.871795	0.829268	0.918919
FCRL3 AND NOT-FCER1A AND FPR1	0.871795	0.829268	0.918919
FCRL3 AND ENPP2 AND NOT-ITM2B	0.826667	0.815789	0.837838
FCRL3 AND ENPP2 AND NOT-ITM2B	0.826667	0.815789	0.837838
FCRL3 AND NOT-FCER1A AND IGSF6	0.826667	0.815789	0.837838
FCRL3 AND SLC1A3 AND NOT-FCER1A	0.823529	0.903226	0.756757
FCRL3 AND NOT-FCER1A AND C3AR1	0.84058	0.90625	0.783784
FCRL3 AND NOT-TSPAN32 AND FPR1	0.821918	0.833333	0.810811
FCRL3 AND GJA1 AND NOT-ITM2B	0.820513	0.780488	0.864865
FCRL3 AND GJA1 AND NOT-ITM2B	0.820513	0.780488	0.864865
MCOLN2 AND CD53 AND GJA1	0.818182	0.931034	0.72973
FCRL3 AND NOT-FCER1A AND VAMP5	0.816901	0.852941	0.783784
FCRL3 AND NOT-FCER1A AND CLEC4E	0.816901	0.852941	0.783784
FCRL3 AND ENPP2 AND NOT-SFRP1	0.815789	0.794872	0.837838
FCRL3 AND ENPP2 AND NOT-SLC7A9	0.815789	0.794872	0.837838
FAM26F AND NOT-FCER1A AND NOT-ADRA1A	0.815789	0.794872	0.837838
FCRL3 AND GJA1 AND NOT-NPY5R	0.815789	0.794872	0.837838
TNFRSF9 AND GJA1 AND NOT-ENTPD3	0.830769	0.964286	0.72973
FCRL3 AND NOT-FCER1A AND TMEM97	0.84058	0.90625	0.783784
CD53 AND NOT-FCER1A AND FPR1	0.814815	0.75	0.891892
CD53 AND NOT-FCER1A AND FPR1	0.814815	0.75	0.891892
CD53 AND NOT-FCER1A AND CLEC4E	0.828571	0.878788	0.783784
FCRL3 AND NOT-FCER1A AND TLR2	0.820513	0.780488	0.864865
FCRL3 AND NOT-FCER1A AND TLR2	0.820513	0.780488	0.864865

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
PTPRZ1 AND NCAM1 AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND NOT-CLDN1 AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND NOT-SDC1 AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND NOT-TRPM4 AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND NOT-EPCAM AND MPZL1	0.904762	1	0.826087
PTPRZ1 AND NOT-RNF43 AND F2RL1	0.896552	0.95122	0.847826
PTPRZ1 AND NOT-SDC1 AND ABCA1	0.894118	0.974359	0.826087
PTPRZ1 AND CLSTN2 AND TNC	0.88172	0.87234	0.891304
PTPRZ1 AND OR51B6 AND TNC	0.88172	0.87234	0.891304
PTPRZ1 AND NOT-ABCC5 AND TNC	0.88172	0.87234	0.891304
PTPRZ1 AND NOT-SLCO4C1 AND TNC	0.88172	0.87234	0.891304
PTPRZ1 AND NOT-SDC1 AND F2RL1	0.879121	0.888889	0.869565
PTPRZ1 AND NOT-SDC1 AND F2RL1	0.879121	0.888889	0.869565
PTPRZ1 AND ITM2B AND NOT-STEAP1	0.876404	0.906977	0.847826
COMPLEX-L1CAM/PTPRZ1/OR6A2	0.876404	0.906977	0.847826
PTPRZ1 AND ITM2B AND NOT-PMEL	0.876404	0.906977	0.847826
PTPRZ1 AND ITM2B AND NOT-CD52	0.876404	0.906977	0.847826
PTPRZ1 AND NOT-RNF43 AND CLCN5	0.873563	0.926829	0.826087
PTPRZ1 AND NOT-FZD10 AND FOLR2	0.869565	0.869565	0.869565
PTPRZ1 AND DLL3 AND F2RL1	0.86747	0.972973	0.782609
PTPRZ1 AND ITM2B AND NOT-CLDN23	0.866667	0.886364	0.847826
PTPRZ1 AND NCAM1 AND ITM2B	0.866667	0.886364	0.847826
PTPRZ1 AND CD33 AND PANX1	0.866667	0.886364	0.847826
PTPRZ1 AND NOT-CLDN1 AND ITM2B	0.866667	0.886364	0.847826
PTPRZ1 AND NOT-TRPM4 AND ITM2B	0.866667	0.886364	0.847826
PTPRZ1 AND NOT-EPCAM AND ITM2B	0.866667	0.886364	0.847826
PTPRZ1 AND NOT-SDC1 AND VANGL2	0.863636	0.904762	0.826087
PTPRZ1 AND NOT-SDK1 AND CD33	0.863636	0.904762	0.826087
PTPRZ1 AND NOT-LGR5 AND CSF1R	0.863636	0.904762	0.826087
PTPRZ1 AND NOT-ACVRL1 AND CD33	0.863636	0.904762	0.826087
PTPRZ1 AND NOT-F11R AND CD33	0.863636	0.904762	0.826087
PTPRZ1 AND NOT-EPCAM AND F2RL1	0.863158	0.836735	0.891304
PTPRZ1 AND NOT-EPCAM AND F2RL1	0.863158	0.836735	0.891304
PTPRZ1 AND NOT-ABCA5 AND CLCN5	0.863158	0.836735	0.891304
PTPRZ1 AND SYNDIG1 AND CD33	0.860465	0.925	0.804348
PTPRZ1 AND ADAM22 AND CD33	0.860465	0.925	0.804348
PTPRZ1 AND GPM6B AND CD33	0.860465	0.925	0.804348
PTPRZ1 AND NOT-ST14 AND CD33	0.860465	0.925	0.804348
PTPRZ1 AND NOT-DUOX1 AND CD33	0.860465	0.925	0.804348
PTPRZ1 AND LRRTM2 AND CD33	0.860465	0.925	0.804348
PTPRZ1 AND NOT-AQP3 AND CD33	0.860465	0.925	0.804348
PTPRZ1 AND NOT-ATP2C2 AND CD33	0.860465	0.925	0.804348
PTPRZ1 AND NOT-GPR87 AND CD33	0.860465	0.925	0.804348
PTPRZ1 AND NOT-PERP AND CD33	0.860465	0.925	0.804348
PTPRZ1 AND NOT-LRP11 AND SEMA5B	0.860215	0.851064	0.869565
PTPRZ1 AND NRCAM AND NOT-STEAP2	0.860215	0.851064	0.869565
PTPRZ1 AND NOT-ACP1 AND FOLR2	0.860215	0.851064	0.869565
CNTNAP1 AND NOT-EPCAM AND VANGL2	0.866667	0.886364	0.847826
SLC1A3 AND VANGL2 AND NOT-CLDN1	0.866667	0.886364	0.847826
VANGL2 AND NOT-FZD10 AND CD33	0.860465	0.925	0.804348
PTPRZ1 AND NOT-TRPM4 AND SLC44A2	0.857143	0.866667	0.847826
PTPRZ1 AND NOT-SDC1 AND TSPAN11	0.857143	0.866667	0.847826
CNTNAP1 AND NOT-SDC1 AND VANGL2	0.857143	0.866667	0.847826
PTPRZ1 AND NCAM1 AND SLC44A2	0.857143	0.866667	0.847826
PCDH10 AND NOT-SDC1 AND F2RL1	0.878049	1	0.782609
SLC1A3 AND VANGL2 AND NOT-CD52	0.857143	0.866667	0.847826
PTPRZ1 AND PANX1 AND NOT-STEAP2	0.854167	0.82	0.891304
PTPRZ1 AND NOT-SDC1 AND NOT-TRHDE	0.854167	0.82	0.891304
PTPRZ1 AND NOT-EPCAM AND HAS2	0.854167	0.82	0.891304
PTPRZ1 AND NOT-FZD10 AND VCAM1	0.854167	0.82	0.891304
PTPRZ1 AND PANX1 AND NOT-STEAP2	0.854167	0.82	0.891304
PTPRZ1 AND NOT-FZD10 AND HLA-DOB	0.854167	0.82	0.891304
VANGL2 AND NOT-CLDN1 AND CSF1R	0.853933	0.883721	0.826087
PTPRZ1 AND NOT-EPCAM AND CALCRL	0.853659	0.972222	0.76087
PTPRZ1 AND NOT-SDC1 AND CALCRL	0.853659	0.972222	0.76087
PTPRZ1 AND SLC4A4 AND FOLR2	0.851064	0.833333	0.869565
PTPRZ1 AND NOT-P2RY14 AND FOLR2	0.851064	0.833333	0.869565
SLC1A3 AND VANGL2 AND NOT-STEAP1	0.866667	0.886364	0.847826
PTPRZ1 AND NOT-KIT AND VCAM1	0.851064	0.833333	0.869565
PTPRZ1 AND NOT-SLC35G1 AND FOLR2	0.851064	0.833333	0.869565
PTPRZ1 AND NOT-IL1R2 AND FOLR2	0.851064	0.833333	0.869565
PTPRZ1 AND NOT-LY6K AND FOLR2	0.851064	0.833333	0.869565
PTPRZ1 AND NOT-CDH1 AND FOLR2	0.851064	0.833333	0.869565
PCDH10 AND VANGL2 AND NOT-EPCAM	0.850575	0.902439	0.804348
PTPRZ1 AND SEMA5B AND F2RL1	0.85	1	0.73913
PTPRZ1 AND NOT-ENTPD6 AND FOLR2	0.860215	0.851064	0.869565
PTPRZ1 AND NOT-SDC1 AND CLCN5	0.847826	0.847826	0.847826

Antigen Pair Logic	F1	Precision	Recall
FCRL3 AND NOT-IL1RAP AND PLGRKT	0.8125	0.962963	0.702703
FCRL3 AND NOT-FCER1A AND SIRPA	0.831169	0.8	0.864865
NOT-FCER1A AND IGSF6 AND GPR18	0.811594	0.875	0.756757
FCRL3 AND SLC1A3 AND NOT-LGR6	0.811594	0.875	0.756757
CD53 AND NOT-FCER1A AND SLC1A3	0.810811	0.810811	0.810811
FAM26F AND NOT-FCER1A AND NOT-SEMA4G	0.826667	0.815789	0.837838
FCRL3 AND ENPP2 AND NOT-NPY5R	0.810811	0.810811	0.810811
FAM26F AND NOT-FCER1A AND NOT-TM4SF5	0.826667	0.815789	0.837838
FCRL3 AND GJA1 AND NOT-SFRP1	0.810127	0.761905	0.864865
CD53 AND NOT-FCER1A AND TMEM97	0.84058	0.90625	0.783784
CD53 AND NOT-FCER1A AND FAM26F	0.815789	0.794872	0.837838
GPR18 AND NOT-SLC18A2 AND RAMP3	0.815789	0.794872	0.837838
FCRL3 AND SLC1A3 AND NOT-S1PR5	0.80597	0.9	0.72973
FCRL3 AND NOT-CXCR2 AND FPR1	0.80597	0.9	0.72973
MCOLN2 AND NOT-GDPD2 AND CYBRD1	0.80597	0.9	0.72973
FCRL3 AND KCNMA1 AND NOT-SLC16A2	0.80597	0.9	0.72973
FCRL3 AND SLC1A3 AND NOT-CRIM1	0.80597	0.9	0.72973
CD53 AND NOT-FCER1A AND SIRPA	0.810127	0.761905	0.864865
FCRL3 AND ENPP2 AND NOT-TACSTD2	0.805556	0.828571	0.783784
FAM26F AND NOT-FCER1A AND NOT-CLEC4M	0.849315	0.861111	0.837838
FAM26F AND NOT-FCER1A AND NOT-PTH1R	0.826667	0.815789	0.837838
FAM26F AND NOT-FCER1A AND NOT-SLC17A4	0.805195	0.775	0.837838
FAM26F AND NOT-FCER1A AND NOT-SLC51A	0.805195	0.775	0.837838
FCRL3 AND GJA1 AND NOT-SLC18A2	0.805195	0.775	0.837838
CD53 AND NOT-CXCR2 AND FPR1	0.804878	0.733333	0.891892
CD53 AND NOT-CXCR2 AND FPR1	0.804878	0.733333	0.891892
FCRL3 AND NOT-FCER1A AND SULF2	0.804878	0.733333	0.891892
NOT-FCER1A AND IGSF6 AND NOT-PTH1R	0.8	0.848485	0.756757
FCRL3 AND GJA1 AND NOT-MFAP3L	0.8	0.744186	0.864865
FCRL3 AND NOT-ATP10A AND SLC38A6	0.80597	0.9	0.72973
FCRL3 AND GJA1 AND NOT-MFAP3L	0.8	0.744186	0.864865
NOT-FCER1A AND GPR18 AND SECTM1	0.823529	0.903226	0.756757
FCRL3 AND ENPP2 AND NOT-SLC18A2	0.8	0.789474	0.810811
FAM26F AND NOT-FCER1A AND NOT-CDH16	0.8	0.789474	0.810811
FCRL3 AND GJA1 AND ITGAE	0.8	0.744186	0.864865
FAM26F AND NOT-FCER1A AND NOT-GPR88	0.826667	0.815789	0.837838
FAM26F AND NOT-FCER1A AND NOT-SLC22A7	0.805195	0.775	0.837838
FCRL3 AND NOT-FCER1A AND PKD2L1	0.8	0.744186	0.864865
FCRL3 AND NOT-FCER1A AND PLGRKT	0.816901	0.852941	0.783784
FCRL3 AND SLC1A3 AND NOT-ITM2B	0.852941	0.935484	0.783784
FCRL3 AND TMEM119 AND NOT-KITLG	0.821918	0.833333	0.810811
FCRL3 AND NOT-F2RL1 AND TMEM119	0.814815	0.75	0.891892
FAM26F AND NOT-FCER1A AND NOT-FREM2	0.815789	0.794872	0.837838
FAM26F AND NOT-FCER1A AND NOT-SLC22A1	0.805195	0.775	0.837838
FAM26F AND NOT-FCER1A AND NOT-SLC23A1	0.805195	0.775	0.837838
FAM26F AND NOT-FCER1A AND NOT-SLC47A1	0.805195	0.775	0.837838
NOT-FCER1A AND GPR18 AND TLR2	0.828571	0.878788	0.783784
CD40 AND GJA1 AND NOT-ENTPD3	0.825397	1	0.702703
FCRL3 AND NOT-FCER1A AND CCR1	0.849315	0.861111	0.837838
CD53 AND TMEM119 AND NOT-PROS1	0.8	0.744186	0.864865
KCNJ10 AND CD53 AND NOT-PROS1	0.852941	0.935484	0.783784
KCNJ10 AND CD53 AND NOT-FZD4	0.84507	0.882353	0.810811
FCRL3 AND NOT-FCER1A AND FXYD6	0.843373	0.76087	0.945946
KCNJ10 AND CD53 AND NOT-PCDHB16	0.852941	0.935484	0.783784
FCRL3 AND NOT-FCER1A AND SLC31A1	0.849315	0.861111	0.837838
KCNJ10 AND NOT-PTPRZ1 AND NOT-FZD4	0.84058	0.90625	0.783784
KCNJ10 AND CD53 AND NOT-FRAS1	0.833333	0.857143	0.810811
KCNJ10 AND CD53 AND NOT-TEK	0.833333	0.857143	0.810811
KCNJ10 AND CD53 AND NOT-PLB1	0.852941	0.935484	0.783784
KCNJ10 AND NOT-SLC16A2 AND ICAM3	0.830769	0.964286	0.72973
KCNJ10 AND NOT-SLC16A2 AND IL10RA	0.830769	0.964286	0.72973
KCNJ10 AND NOT-SLC16A2 AND FCRL3	0.830769	0.964286	0.72973
KCNJ10 AND CD53 AND NOT-CXCR2	0.84058	0.90625	0.783784
KCNJ10 AND NOT-PLP1 AND NOT-PROS1	0.828571	0.878788	0.783784
KCNJ10 AND GPR18 AND NOT-TNFRSF19	0.828571	0.878788	0.783784
KCNJ10 AND FCRL3 AND NOT-TNFRSF19	0.828571	0.878788	0.783784
KCNJ10 AND FCRL3 AND NOT-ENTPD3	0.833333	0.857143	0.810811
FCRL3 AND NOT-MFAP3L AND FXYD6	0.825	0.767442	0.891892
KCNJ10 AND CD53 AND NOT-F2RL1	0.852941	0.935484	0.783784
FCRL3 AND NOT-ITGA6 AND FXYD6	0.825	0.767442	0.891892
KCNJ10 AND CD53 AND NOT-ENTPD3	0.833333	0.857143	0.810811
KCNJ10 AND CD53 AND NOT-TSPAN7	0.821918	0.833333	0.810811
KCNJ10 AND CD53 AND NOT-EMCN	0.852941	0.935484	0.783784
KCNJ10 AND CD53 AND NOT-DUOX1	0.818182	0.931034	0.72973
KCNJ10 AND CD53 AND NOT-SLC46A2	0.818182	0.931034	0.72973
KCNJ10 AND FCRL3 AND NOT-FZD4	0.816901	0.852941	0.783784
KCNJ10 AND CD53 AND NOT-CRIM1	0.816901	0.852941	0.783784

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
PTPRZ1 AND NCAM1 AND CSF1R	0.847826	0.847826	0.847826
PTPRZ1 AND NOT-SDC1 AND CSF1R	0.847826	0.847826	0.847826
PTPRZ1 AND NOT-STEAP1 AND CSF1R	0.847826	0.847826	0.847826
PTPRZ1 AND CD33 AND NOT-FZD10	0.847826	0.847826	0.847826
PTPRZ1 AND NOT-SDC1 AND CLCN5	0.847826	0.847826	0.847826
PTPRZ1 AND NOT-SDC1 AND CSF1R	0.847826	0.847826	0.847826
PTPRZ1 AND NOT-CLDN1 AND CSF1R	0.847826	0.847826	0.847826
PTPRZ1 AND NOT-CLDN1 AND CSF1R	0.847826	0.847826	0.847826
PTPRZ1 AND CSF1R AND NOT-CD52	0.847826	0.847826	0.847826
PTPRZ1 AND NOT-CLDN23 AND CSF1R	0.847826	0.847826	0.847826
PTPRZ1 AND NOT-SDC1 AND ITM2B	0.847826	0.847826	0.847826
PTPRZ1 AND CD33 AND CLCN5	0.847826	0.847826	0.847826
PTPRZ1 AND CLDN17 AND NOT-GABRA2	0.891304	0.891304	0.891304
PTPRZ1 AND SLC4A8 AND NOT-GABRA2	0.88172	0.87234	0.891304
PTPRZ1 AND CRB1 AND NOT-GABRA2	0.88172	0.87234	0.891304
PTPRZ1 AND NOT-AOC3 AND NOT-GABRA2	0.88172	0.87234	0.891304
PTPRZ1 AND GABBR2 AND NOT-GABRA2	0.88172	0.87234	0.891304
PTPRZ1 AND NOT-AOC3 AND NOT-GABRA2	0.88172	0.87234	0.891304
PTPRZ1 AND NOT-CD36 AND NOT-GABRA2	0.88172	0.87234	0.891304
PTPRZ1 AND SLCO1C1 AND NOT-GABRA2	0.88172	0.87234	0.891304
PTPRZ1 AND SLC4A8 AND NOT-GRM3	0.880952	0.973684	0.804348
PTPRZ1 AND NOT-ATP8B1 AND NOT-GRM3	0.880952	0.973684	0.804348
PTPRZ1 AND SLC1A2 AND NOT-GRM3	0.880952	0.973684	0.804348
PTPRZ1 AND NOT-GRM3 AND NOT-EPHA2	0.880952	0.973684	0.804348
PTPRZ1 AND NOT-GRM3 AND NOT-CD36	0.880952	0.973684	0.804348
PTPRZ1 AND NOT-KCNK6 AND NOT-GRM3	0.880952	0.973684	0.804348
PTPRZ1 AND NOT-GRM3 AND GABBR2	0.880952	0.973684	0.804348
PTPRZ1 AND NOT-GRM3 AND NOT-AOC3	0.880952	0.973684	0.804348
PTPRZ1 AND KCNJ10 AND NOT-GRM3	0.880952	0.973684	0.804348
PTPRZ1 AND CRB1 AND NOT-GRM3	0.880952	0.973684	0.804348
PTPRZ1 AND NOT-LRIG3 AND NOT-GRM3	0.880952	0.973684	0.804348
PTPRZ1 AND ATP2B2 AND NOT-GRM3	0.880952	0.973684	0.804348
PTPRZ1 AND CLDN17 AND NOT-SCN8A	0.880952	0.973684	0.804348
PTPRZ1 AND CNTNAP2 AND NOT-GRM3	0.880952	0.973684	0.804348
PTPRZ1 AND SLCO1A2 AND NOT-GRM3	0.880952	0.973684	0.804348
PTPRZ1 AND CLDN17 AND NOT-GRM3	0.880952	0.973684	0.804348
PTPRZ1 AND MOG AND NOT-GRM3	0.880952	0.973684	0.804348
PTPRZ1 AND GPR85 AND NOT-GRM3	0.880952	0.973684	0.804348
PTPRZ1 AND SLCO1C1 AND NOT-GRM3	0.880952	0.973684	0.804348
PTPRZ1 AND OR2L13 AND NOT-GABRA2	0.88172	0.87234	0.891304
PTPRZ1 AND NOT-ATP6V0A4 AND NOT-GABRA2	0.88172	0.87234	0.891304
PTPRZ1 AND ATP2B2 AND F2R	0.879121	0.888889	0.869565
PTPRZ1 AND SLC1A2 AND F2R	0.879121	0.888889	0.869565
PTPRZ1 AND KCNJ10 AND F2R	0.879121	0.888889	0.869565
PTPRZ1 AND CLDN17 AND F2R	0.879121	0.888889	0.869565
PTPRZ1 AND MOG AND F2R	0.879121	0.888889	0.869565
PTPRZ1 AND GPR85 AND F2R	0.879121	0.888889	0.869565
PTPRZ1 AND NOT-LRIG3 AND F2R	0.879121	0.888889	0.869565
PTPRZ1 AND CRB1 AND F2R	0.879121	0.888889	0.869565
PTPRZ1 AND CLDN17 AND NOT-GPR83	0.873563	0.926829	0.826087
PTPRZ1 AND NOT-MSMO1 AND NOT-GABRA2	0.88172	0.87234	0.891304
SLC1A3 AND F2R AND NOT-CD36	0.870588	0.948718	0.804348
PTPRZ1 AND F2R AND CNTNAP4	0.866667	0.886364	0.847826
PTPRZ1 AND F2R AND SLCO1A2	0.866667	0.886364	0.847826
PTPRZ1 AND F2R AND SLCO1C1	0.866667	0.886364	0.847826
PTPRZ1 AND F2R AND CNTNAP2	0.866667	0.886364	0.847826
PTPRZ1 AND F2R AND NOT-KCNK6	0.866667	0.886364	0.847826
PTPRZ1 AND F2R AND GABBR2	0.866667	0.886364	0.847826
PTPRZ1 AND F2R AND SLC4A8	0.866667	0.886364	0.847826
PTPRZ1 AND F2R AND NOT-CD36	0.866667	0.886364	0.847826
PTPRZ1 AND F2R AND KCNA2	0.866667	0.886364	0.847826
PTPRZ1 AND F2R AND NOT-AOC3	0.866667	0.886364	0.847826
PTPRZ1 AND F2R AND NOT-EPHA2	0.866667	0.886364	0.847826
PTPRZ1 AND F2R AND NOT-SLC43A1	0.866667	0.886364	0.847826
PTPRZ1 AND F2R AND NOT-MRGPRF	0.866667	0.886364	0.847826
PTPRZ1 AND F2R AND NOT-ATP8B1	0.866667	0.886364	0.847826
PTPRZ1 AND CLDN17 AND NOT-SYT4	0.860465	0.925	0.804348
PTPRZ1 AND BEST3 AND CRB1	0.857143	0.866667	0.847826
PTPRZ1 AND BEST3 AND NOT-MOG	0.857143	0.866667	0.847826
PTPRZ1 AND CLDN17 AND NOT-KCNJ6	0.857143	0.947368	0.782609
PTPRZ1 AND BEST3 AND CLDN17	0.857143	0.866667	0.847826
SLC1A2 AND F2R AND NOT-AQP3	0.853933	0.883721	0.826087
PTPRZ1 AND CLDN17 AND NOT-GRM1	0.853933	0.883721	0.826087
PTPRZ1 AND NOT-GRM3 AND KCNA2	0.853659	0.972222	0.76087
PTPRZ1 AND GPR85 AND ANTXR2	0.847826	0.847826	0.847826
PTPRZ1 AND MOG AND ANTXR2	0.847826	0.847826	0.847826
PTPRZ1 AND SLC1A2 AND ANTXR2	0.847826	0.847826	0.847826

Antigen Pair Logic	F1	Precision	Recall
KCNJ10 AND NOT-PTPRZ1 AND NOT-SLC3A1	0.816901	0.852941	0.783784
KCNJ10 AND FCRL3 AND NOT-PLB1	0.816901	0.852941	0.783784
FAM26F AND NOT-FCER1A AND NOT-DIO1	0.849315	0.861111	0.837838
CD53 AND NOT-FCER1A AND FXYD6	0.814815	0.75	0.891892
CD53 AND NOT-FCER1A AND FXYD6	0.814815	0.75	0.891892
FAM26F AND NOT-FCER1A AND NOT-SLC13A5	0.821918	0.833333	0.810811
KCNJ10 AND CD53 AND NOT-TSPAN6	0.816901	0.852941	0.783784
KCNJ10 AND CD53 AND NOT-PKD2L1	0.828571	0.878788	0.783784
KCNJ10 AND NOT-ROM1 AND FCRL3	0.821918	0.833333	0.810811
KCNJ10 AND FCRL3 AND NOT-FCER1A	0.84058	0.90625	0.783784
KCNJ10 AND CD53 AND NOT-KITLG	0.8125	0.962963	0.702703
KCNJ10 AND NOT-PTPRZ1 AND NOT-SEMA4G	0.821918	0.833333	0.810811
KCNJ10 AND CD53 AND NOT-CD300LB	0.811594	0.875	0.756757
KCNJ10 AND FCRL3 AND NOT-HCRTR1	0.810811	0.810811	0.810811
KCNJ10 AND GPR18 AND NOT-HCRTR1	0.810811	0.810811	0.810811
FAM26F AND NOT-FCER1A AND NOT-SLC10A1	0.849315	0.861111	0.837838
FAM26F AND NOT-FCER1A AND NOT-SLC22A25	0.810811	0.810811	0.810811
KCNJ10 AND NOT-PCDH19 AND FCRL3	0.810811	0.810811	0.810811
KCNJ10 AND CD53 AND NOT-FCER1A	0.878788	1	0.783784
KCNJ10 AND FCRL3 AND NOT-ITGA2B	0.833333	0.857143	0.810811
KCNJ10 AND NOT-ABCA8 AND GPR18	0.84507	0.882353	0.810811
KCNJ10 AND NOT-NTRK2 AND NOT-FCER1A	0.84058	0.90625	0.783784
KCNJ10 AND NOT-SLC18A2 AND GPR18	0.80597	0.9	0.72973
KCNJ10 AND CD53 AND NOT-TSPAN18	0.80597	0.9	0.72973
KCNJ10 AND CD53 AND NOT-KCNQ1	0.80597	0.9	0.72973
KCNJ10 AND CD53 AND NOT-NRP1	0.80597	0.9	0.72973
KCNJ10 AND ICAM3 AND NOT-PROS1	0.805556	0.828571	0.783784
KCNJ10 AND NOT-GPR173 AND FCRL3	0.805556	0.828571	0.783784
KCNJ10 AND GPR18 AND NOT-LY6D	0.805556	0.828571	0.783784
KCNJ10 AND NOT-TENM1 AND FCRL3	0.805556	0.828571	0.783784
KCNJ10 AND GPR18 AND NOT-FZD4	0.805556	0.828571	0.783784
KCNJ10 AND GPR18 AND NOT-PROS1	0.805556	0.828571	0.783784
KCNJ10 AND FCRL3 AND NOT-F2RL1	0.805556	0.828571	0.783784
KCNJ10 AND CD53 AND NOT-LEPR	0.805556	0.828571	0.783784
KCNJ10 AND CD53 AND NOT-SCN7A	0.805556	0.828571	0.783784
FAM26F AND NOT-FCER1A AND NOT-SLC6A13	0.837838	0.837838	0.837838
KCNJ10 AND ITGAL AND NOT-PROS1	0.8	0.848485	0.756757
KCNJ10 AND CD53 AND NOT-ITGA6	0.8	0.848485	0.756757
KCNJ10 AND FCRL3 AND NOT-S1PR5	0.8	0.848485	0.756757
KCNJ10 AND NOT-PLP1 AND NOT-FCER1A	0.878788	1	0.783784
KCNJ10 AND NOT-PTPRZ1 AND NOT-TM4SF5	0.8	0.789474	0.810811
KCNJ10 AND FCRL3 AND NOT-AVPR2	0.8	0.789474	0.810811
KCNJ10 AND CD53 AND NOT-GDPD2	0.869565	0.9375	0.810811
FAM26F AND NOT-FCER1A AND NOT-SLC17A2	0.805195	0.775	0.837838
KCNJ10 AND HLA-A AND NOT-PROS1	0.828571	0.878788	0.783784
KCNJ10 AND CD53 AND NOT-ACVR2A	0.823529	0.903226	0.756757
FAM26F AND NOT-FCER1A AND NOT-ABCG5	0.805195	0.775	0.837838
FCRL3 AND NOT-CLEC1B AND FXYD6	0.853659	0.777778	0.945946
KCNJ10 AND NOT-PLP1 AND GPR18	0.821918	0.833333	0.810811
KCNJ10 AND NOT-ENTPD3 AND GPR18	0.821918	0.833333	0.810811
KCNJ10 AND GPR18 AND NOT-CNGA1	0.8	0.848485	0.756757
KCNJ10 AND NOT-S1PR5 AND GPR18	0.8	0.848485	0.756757
KCNJ10 AND NOT-ENTPD3 AND ADTRP	0.8	0.789474	0.810811
KCNJ10 AND NOT-PCDH19 AND GPR18	0.810811	0.810811	0.810811
CD53 AND NOT-FCER1A AND GPNMB	0.849315	0.861111	0.837838
CD53 AND NOT-FCER1A AND GPNMB	0.849315	0.861111	0.837838
BIRC5 AND FCRL3 AND GJA1	0.830769	0.964286	0.72973
FCRL3 AND GPNMB AND NOT-FCER1A	0.830769	0.964286	0.72973
FCRL3 AND GPNMB AND NOT-S1PR5	0.830769	0.964286	0.72973
CXCR5 AND GJA1 AND NOT-ENTPD3	0.84058	0.90625	0.783784
FCRL3 AND GPNMB AND NOT-ITM2B	0.823529	0.903226	0.756757
FCRL3 AND GPNMB AND NOT-CRIM1	0.818182	0.931034	0.72973
BIRC5 AND CD53 AND CD8A	0.818182	0.931034	0.72973
FCRL3 AND NOT-CD160 AND FPR1	0.816901	0.852941	0.783784
FCRL3 AND GJA1 AND NOT-EGFR	0.815789	0.794872	0.837838
FCRL3 AND GJA1 AND NOT-EGFR	0.815789	0.794872	0.837838
FCRL3 AND VCAM1 AND NOT-TACSTD2	0.815789	0.794872	0.837838
FCRL3 AND GPNMB AND NOT-ATP10A	0.8125	0.962963	0.702703
FCRL3 AND GPNMB AND NOT-GPR35	0.8125	0.962963	0.702703
FCRL3 AND GPNMB AND NOT-LGR6	0.8125	0.962963	0.702703
FCRL3 AND VCAM1 AND NOT-ITM2B	0.810127	0.761905	0.864865
FCRL3 AND VCAM1 AND NOT-SFRP1	0.810127	0.761905	0.864865
FCRL3 AND VCAM1 AND NOT-ITM2B	0.810127	0.761905	0.864865
BIRC5 AND FCRL3 AND FCER1G	0.80597	0.9	0.72973
BIRC5 AND FCRL3 AND TYROBP	0.80597	0.9	0.72973
BIRC5 AND FCRL3 AND IGSF6	0.80597	0.9	0.72973
FCRL3 AND ENPP2 AND NOT-CLDN1	0.805195	0.775	0.837838

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
PTPRZ1 AND NOT-CD36 AND ANTXR2	0.847826	0.847826	0.847826
PTPRZ1 AND GABBR2 AND ANTXR2	0.847826	0.847826	0.847826
PTPRZ1 AND KCNA2 AND ANTXR2	0.847826	0.847826	0.847826
PTPRZ1 AND NOT-KCNK6 AND ANTXR2	0.847826	0.847826	0.847826
PTPRZ1 AND KCNJ10 AND ANTXR2	0.847826	0.847826	0.847826
PTPRZ1 AND NOT-AOC3 AND ANTXR2	0.847826	0.847826	0.847826
PTPRZ1 AND SLC4A8 AND ANTXR2	0.847826	0.847826	0.847826
PTPRZ1 AND NOT-LRIG3 AND ANTXR2	0.847826	0.847826	0.847826
PTPRZ1 AND GABRB2 AND ANTXR2	0.847826	0.847826	0.847826
PTPRZ1 AND CRB1 AND ANTXR2	0.847826	0.847826	0.847826
PTPRZ1 AND SLCO1C1 AND ANTXR2	0.847826	0.847826	0.847826
PTPRZ1 AND NOT-SLC43A1 AND ANTXR2	0.847826	0.847826	0.847826
PTPRZ1 AND NOT-MRGPRF AND ANTXR2	0.847826	0.847826	0.847826
PTPRZ1 AND CLDN17 AND ANTXR2	0.847826	0.847826	0.847826
PTPRZ1 AND CLDN17 AND NOT-SCN2A	0.845361	0.803922	0.891304
PTPRZ1 AND NOT-CD36 AND NOT-GPR83	0.844444	0.863636	0.826087
PTPRZ1 AND NOT-CD36 AND NOT-ATP8A2	0.844444	0.863636	0.826087
PTPRZ1 AND SLC1A2 AND NOT-GPR83	0.844444	0.863636	0.826087
PTPRZ1 AND GABRB2 AND NOT-GPR83	0.844444	0.863636	0.826087
PTPRZ1 AND GPR85 AND NOT-GPR83	0.844444	0.863636	0.826087
PTPRZ1 AND GABBR2 AND NOT-ATP8A2	0.844444	0.863636	0.826087
PTPRZ1 AND CRB1 AND NOT-GPR83	0.844444	0.863636	0.826087
PTPRZ1 AND SLC1A2 AND NOT-ATP8A2	0.844444	0.863636	0.826087
PTPRZ1 AND CRB1 AND NOT-ATP8A2	0.844444	0.863636	0.826087
PTPRZ1 AND NOT-AOC3 AND NOT-ATP8A2	0.844444	0.863636	0.826087
PTPRZ1 AND SLC4A8 AND NOT-GPR83	0.844444	0.863636	0.826087
PTPRZ1 AND GABBR2 AND NOT-GPR83	0.844444	0.863636	0.826087
PTPRZ1 AND NOT-KCNK6 AND NOT-ATP8A2	0.844444	0.863636	0.826087
PTPRZ1 AND SLC4A8 AND NOT-ATP8A2	0.844444	0.863636	0.826087
PTPRZ1 AND SLCO1C1 AND NOT-GPR83	0.844444	0.863636	0.826087
PTPRZ1 AND NOT-LRIG3 AND NOT-ATP8A2	0.844444	0.863636	0.826087
PTPRZ1 AND SLCO1C1 AND NOT-ATP8A2	0.844444	0.863636	0.826087
PTPRZ1 AND NOT-SDC1 AND TNC	0.921348	0.953488	0.891304
PTPRZ1 AND NOT-CLDN23 AND CD33	0.863636	0.904762	0.826087
PTPRZ1 AND NOT-CLDN1 AND CD33	0.860465	0.925	0.804348
PTPRZ1 AND NCAM1 AND CD33	0.860465	0.925	0.804348
PTPRZ1 AND NOT-TRPM4 AND CD33	0.860465	0.925	0.804348
PTPRZ1 AND SEMA5B AND NOT-STEAP2	0.844444	0.863636	0.826087
PTPRZ1 AND CD33 AND NOT-STEAP2	0.83871	0.829787	0.847826
PTPRZ1 AND NOT-SDC1 AND NOT-STEAP2	0.836735	0.788462	0.891304
PTPRZ1 AND NOT-SDC1 AND NOT-STEAP2	0.836735	0.788462	0.891304
PTPRZ1 AND CD33 AND NOT-STEAP1	0.829787	0.8125	0.847826
COMPLEX-L1CAM/PTPRZ1/CD33	0.829787	0.8125	0.847826
PTPRZ1 AND CD33 AND NOT-CD52	0.829787	0.8125	0.847826
COMPLEX-PTPRZ1/VCAM1/CD33	0.829787	0.8125	0.847826
PTPRZ1 AND CD33 AND NOT-PMEL	0.829787	0.8125	0.847826
PTPRZ1 AND NOT-SDC1 AND SEMA5B	0.824742	0.784314	0.869565
PTPRZ1 AND DLL3 AND NOT-STEAP2	0.821053	0.795918	0.847826
COMPLEX-MET/PTPRZ1/CD33	0.821053	0.795918	0.847826
VANGL2 AND NOT-TPBG AND L1CAM	0.818182	0.857143	0.782609
PCDH10 AND TNC AND NOT-SDC1	0.816327	0.769231	0.869565
PTPRZ1 AND NOT-EPCAM AND SEMA5B	0.808081	0.754717	0.869565
COMPLEX-FOLR2/MET/PTPRZ1	0.808081	0.754717	0.869565
PTPRZ1 AND NOT-TRPM4 AND NOT-STEAP2	0.803922	0.732143	0.891304
PTPRZ1 AND NOT-SPON2 AND NOT-STEAP2	0.803922	0.732143	0.891304
PTPRZ1 AND NOT-CLDN1 AND NOT-STEAP2	0.803922	0.732143	0.891304
PTPRZ1 AND NOT-CD52 AND SLC39A6	0.803922	0.732143	0.891304
PTPRZ1 AND NOT-CD52 AND NOT-STEAP2	0.803922	0.732143	0.891304
PTPRZ1 AND NCAM1 AND NOT-STEAP2	0.803922	0.732143	0.891304
PTPRZ1 AND BCAN AND NOT-STEAP2	0.803922	0.732143	0.891304
PTPRZ1 AND NOT-CD52 AND NOT-STEAP2	0.803922	0.732143	0.891304
PTPRZ1 AND NCAM1 AND NOT-STEAP2	0.803922	0.732143	0.891304
PTPRZ1 AND NOT-TRPM4 AND NOT-STEAP2	0.803922	0.732143	0.891304
PTPRZ1 AND NOT-CLDN1 AND NOT-STEAP2	0.803922	0.732143	0.891304
COMPLEX-FOLH1/EPCAM/VANGL2	0.821053	0.795918	0.847826
VANGL2 AND NOT-EPCAM AND L1CAM	0.808989	0.837209	0.782609
Breast Cancer			
ADAM12 AND NOT-NAALAD2 AND NOT-GPBAR1	0.882353	0.9375	0.833333
ADAM12 AND NOT-NAALAD2 AND TMEM47	0.857143	0.882353	0.833333
ADAM12 AND NOT-PAG1 AND NOT-IL1RL1	0.854369	0.897959	0.814815
ADAM12 AND NOT-NAALAD2 AND TMPRSS2	0.841121	0.849057	0.833333
ADAM12 AND NOT-GPBAR1 AND LY75	0.84	0.913043	0.777778
GPRC5A AND NOT-ATP10B AND NOT-FPR1	0.857143	0.882353	0.833333
ADAM12 AND NOT-SCN9A AND TMPRSS2	0.833333	0.833333	0.833333
ADAM12 AND NOT-PAG1 AND APCDD1	0.830189	0.846154	0.814815
ADAM12 AND NOT-C15orf27 AND KCNS1	0.830189	0.846154	0.814815
GPRC5A AND NOT-IL22RA1 AND NOT-HHIP	0.873786	0.918367	0.833333

Antigen Pair Logic	F1	Precision	Recall
BIRC5 AND CD53 AND GJA1	0.830769	0.964286	0.72973
BIRC5 AND CD53 AND IGSF6	0.80597	0.9	0.72973
GPR18 AND NOT-SLC18A2 AND GPNMB	0.8	0.789474	0.810811
CD53 AND GPNMB AND NOT-PROS1	0.8	0.744186	0.864865
CD53 AND GPNMB AND NOT-PROS1	0.8	0.744186	0.864865
FCRL3 AND TMEM119 AND NOT-EGFR	0.860759	0.809524	0.918919
GPR18 AND NOT-S1PR5 AND GPNMB	0.805195	0.775	0.837838
BIRC5 AND CD53 AND ADTRP	0.811594	0.875	0.756757
CD53 AND GPNMB AND NOT-HAS1	0.831169	0.8	0.864865
FCRL3 AND NOT-NCAM1 AND BRCA1	0.8	0.789474	0.810811
BIRC5 AND CD53 AND FPR1	0.80597	0.9	0.72973
NOT-SLC6A8 AND NOT-SDC1 AND TMEM119	0.8	0.789474	0.810811
FCRL3 AND TMEM119 AND NOT-IL20RA	0.813953	0.714286	0.945946
CXCR5 AND PPAP2B AND NOT-ENTPD3	0.8	0.848485	0.756757
KCNJ10 AND CD53 AND NOT-LRIG3	0.828571	0.878788	0.783784
KCNJ10 AND CD53 AND NOT-GJA3	0.857143	0.909091	0.810811
KCNJ10 AND NOT-FXYD7 AND FCRL3	0.818182	0.931034	0.72973
KCNJ10 AND NOT-ASTN1 AND NOT-FCER1A	0.823529	0.903226	0.756757
KCNJ10 AND CD53 AND NOT-CD93	0.811594	0.875	0.756757
KCNJ10 AND FCRL3 AND NOT-KCNF1	0.810811	0.810811	0.810811
KCNJ10 AND FCRL3 AND NOT-CABP7	0.810811	0.810811	0.810811
KCNJ10 AND CD53 AND NOT-CCKAR	0.811594	0.875	0.756757
KCNJ10 AND CD53 AND NOT-STEAP4	0.8	0.848485	0.756757
KCNJ10 AND CD53 AND NOT-AXL	0.828571	0.878788	0.783784
FCRL3 AND CDH11 AND NOT-EGFR	0.818182	0.931034	0.72973
KCNJ10 AND NOT-EGFR AND FCRL3	0.818182	0.931034	0.72973
KCNJ10 AND NOT-EGFR AND IL10RA	0.818182	0.931034	0.72973
KCNJ10 AND CD53 AND NOT-MSLN	0.810811	0.810811	0.810811
KCNJ10 AND NOT-NCAM1 AND FCRL3	0.8	0.848485	0.756757
FCRL3 AND NOT-SLC6A13 AND THY1	0.805556	0.828571	0.783784
KCNJ10 AND NOT-IL20RA AND GPR18	0.8	0.789474	0.810811
BIRC5 AND FCRL3 AND VCAM1	0.830769	0.964286	0.72973
FCRL3 AND THY1 AND NOT-EGFR	0.811594	0.875	0.756757
FCRL3 AND VCAM1 AND NOT-EGFR	0.805195	0.775	0.837838
FCRL3 AND TNC AND NOT-EGFR	0.8	0.848485	0.756757
FCRL3 AND VCAM1 AND NOT-CLDN1	0.8	0.744186	0.864865
Anaplastic Lymphoma			
NOT-MRAP2 AND NOT-FXYD1 AND RAMP3	0.8	1	0.666667
NOT-MRAP2 AND NOT-FXYD1 AND MCAM	0.846154	1	0.733333
GPR18 AND NOT-CA4 AND APLNR	0.8125	0.764706	0.866667
NOT-MRAP2 AND NOT-FXYD1 AND OSMR	0.846154	1	0.733333
NOT-MRAP2 AND NOT-FXYD1 AND CDH5	0.827586	0.857143	0.8
NOT-MRAP2 AND NOT-FXYD1 AND GJA1	0.857143	0.923077	0.8
NOT-MRAP2 AND NOT-GHR AND APLNR	0.903226	0.875	0.933333
NOT-FXYD1 AND CDH5 AND NOT-SLC39A2	0.814815	0.916667	0.733333
NOT-ERBB3 AND NOT-SCN4B AND APLNR	0.848485	0.777778	0.933333
NOT-ERBB3 AND NOT-SCN4B AND RAMP3	0.848485	0.777778	0.933333
NOT-MRAP2 AND NOT-FXYD1 AND TNC	0.8	1	0.666667
NOT-EGFR AND CDH5 AND NOT-FXYD1	0.846154	1	0.733333
NOT-ERBB3 AND NOT-FXYD1 AND APLNR	0.965517	1	0.933333
NOT-ERBB3 AND NOT-FXYD1 AND AQP1	0.8125	0.764706	0.866667
NOT-MRAP2 AND NOT-EGFR AND APLNR	0.896552	0.928571	0.866667
NOT-FCER1A AND SECTM1 AND NOT-EGFR	0.8	0.8	0.8
NOT-FXYD1 AND THY1 AND NOT-CNTN1	0.8125	0.764706	0.866667
NOT-FXYD1 AND THY1 AND NOT-PRRG1	0.83871	0.8125	0.866667
NOT-FXYD1 AND NOT-ITGB6 AND CDH5	0.846154	1	0.733333
NOT-EGFR AND NOT-FXYD1 AND GJA4	0.903226	0.875	0.933333
COMPLEX-ANO4/ATP8B1/TNFRSF8	0.823529	0.736842	0.933333
NOT-ERBB3 AND NOT-GHR AND APLNR	0.875	0.823529	0.933333
NOT-GHR AND NOT-EFNB3 AND THY1	0.814815	0.916667	0.733333
NOT-FXYD1 AND THY1 AND NOT-GHR	0.8125	0.764706	0.866667
NOT-EGFR AND THY1 AND P2RY8	0.814815	0.916667	0.733333
NOT-ERBB3 AND NOT-SCN4B AND THY1	0.83871	0.8125	0.866667
NOT-EGFR AND THY1 AND FAM26F	0.814815	0.916667	0.733333
NOT-ERBB2 AND NOT-GLRB AND THY1	0.827586	0.857143	0.8
NOT-EGFR AND THY1 AND SLC7A7	0.814815	0.916667	0.733333
NOT-EGFR AND THY1 AND NOT-ATP1A2	0.83871	0.8125	0.866667
NOT-EGFR AND THY1 AND NOT-AQP4	0.8125	0.764706	0.866667
NOT-ERBB2 AND THY1 AND NOT-GLDN	0.846154	1	0.733333
NOT-ERBB2 AND THY1 AND NOT-TSPAN7	0.8	0.8	0.8
NOT-ERBB2 AND NOT-GPRC5B AND THY1	0.866667	0.866667	0.866667
NOT-ERBB3 AND NOT-GHR AND THY1	0.896552	0.928571	0.866667
NOT-EGFR AND THY1 AND NOT-CNTNAP4	0.8125	0.764706	0.866667
NOT-ERBB3 AND THY1 AND NOT-EGFR	0.866667	0.866667	0.866667
Mantle-Cell Lymphoma			
CLECL1 AND NOT-IL12RB2 AND LRRC32	1	1	1
CLECL1 AND NOT-IL12RB2 AND NOT-CLEC4A	1	1	1

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
ADAM12 AND NOT-SCN9A AND RHCG	0.825688	0.818182	0.833333
GPRC5A AND NOT-IL22RA1 AND NOT-AQP4	0.862745	0.916667	0.814815
ADAM12 AND OR51B2 AND NOT-ASGR1	0.823529	0.875	0.777778
ADAM12 AND NOT-MME AND NOT-IL1RL1	0.823529	0.875	0.777778
ADAM12 AND NOT-TSPAN3 AND NOT-IL1RL1	0.823529	0.875	0.777778
KCNS3 AND NOT-IL22RA1 AND NOT-AQP4	0.851852	0.851852	0.851852
ADAM12 AND NOT-DPP4 AND NOT-DUOXA1	0.818182	0.803571	0.833333
GPRC5A AND NOT-ATP10B AND NOT-DUOXA1	0.873786	0.918367	0.833333
ADAM12 AND NOT-DPP4 AND NOT-COL17A1	0.818182	0.803571	0.833333
GPRC5A AND NOT-IL22RA1 AND NOT-TREM1	0.849057	0.865385	0.833333
ADAM12 AND NOT-MME AND NOT-PRPH2	0.815534	0.857143	0.777778
ADAM12 AND NOT-MME AND NOT-STAB2	0.815534	0.857143	0.777778
ADAM12 AND NOT-PAG1 AND RHCG	0.814815	0.814815	0.814815
ADAM12 AND NOT-DPP4 AND NOT-PAG1	0.814815	0.814815	0.814815
GPRC5A AND NOT-IL22RA1 AND NOT-FPR1	0.865385	0.9	0.833333
ADAM12 AND NOT-PAG1 AND GPR15	0.811321	0.826923	0.796296
GPRC5A AND NOT-IL22RA1 AND NOT-CA4	0.873786	0.918367	0.833333
GPRC5A AND NOT-DUOXA1 AND NOT-MEP1A	0.857143	0.882353	0.833333
PROM2 AND NOT-DUOXA1 AND NOT-CA4	0.815534	0.857143	0.777778
ADAM12 AND NOT-C15orf27 AND TMPRSS2	0.808081	0.888889	0.740741
ADAM12 AND NOT-DPP4 AND NOT-UPK1B	0.807339	0.8	0.814815
GPRC5A AND NOT-DUOXA1 AND NOT-IHH	0.857143	0.882353	0.833333
GPRC5A AND NOT-DUOXA1 AND NOT-TM4SF5	0.857143	0.882353	0.833333
GPRC5A AND NOT-DUOXA1 AND NOT-SLC17A4	0.857143	0.882353	0.833333
GPRC5A AND NOT-DUOXA1 AND NOT-GPR35	0.857143	0.882353	0.833333
GPRC5A AND NOT-IL22RA1 AND NOT-AQP9	0.834951	0.877551	0.796296
GPRC5A AND NOT-CYP4F12 AND NOT-DUOXA1	0.851485	0.914894	0.796296
GPRC5A AND NOT-DUOXA1 AND NOT-SLC39A5	0.857143	0.882353	0.833333
GPRC5A AND NOT-CA4 AND NOT-DUOXA1	0.862745	0.916667	0.814815
GPRC5A AND NOT-IL22RA1 AND NOT-GLDN	0.808081	0.888889	0.740741
KCNS3 AND NOT-CDHR1 AND NOT-AQP4	0.87619	0.901961	0.851852
GPRC5A AND NOT-CA4 AND NOT-IL20RB	0.834951	0.877551	0.796296
GPRC5A AND NOT-IL22RA1 AND NOT-IL1RL1	0.862745	0.916667	0.814815
GPRC5A AND NOT-CA4 AND NOT-DSC3	0.82243	0.830189	0.814815
GPRC5A AND NOT-VSIG2 AND NOT-LY6D	0.873786	0.918367	0.833333
KCNS3 AND NOT-CA4 AND NOT-IL20RB	0.803571	0.775862	0.833333
CDH3 AND NOT-DUOXA1 AND NOT-DPP6	0.803279	0.720588	0.907407
ADAM12 AND NOT-MME AND NOT-SLC8A3	0.84	0.913043	0.777778
JAG2 AND NOT-DUOXA1 AND VAMP8	0.824742	0.930233	0.740741
LRRC8E AND NOT-IL22RA1 AND NOT-FNDC5	0.82	0.891304	0.759259
ADAM12 AND NOT-C15orf27 AND NOT-GYPC	0.814815	0.814815	0.814815
GPRC5A AND NOT-DUOXA1 AND NOT-CDH17	0.857143	0.882353	0.833333
LRRC8E AND NOT-DUOXA1 AND NOT-IL1RL1	0.808081	0.888889	0.740741
JAG2 AND NOT-DUOXA1 AND NOT-GABRG2	0.803922	0.854167	0.759259
GPRC5A AND NOT-DUOXA1 AND NOT-PTPRR	0.857143	0.882353	0.833333
JAG2 AND NOT-IL22RA1 AND VAMP8	0.808081	0.888889	0.740741
GPRC5A AND NOT-ATP10B AND NOT-GYPC	0.841121	0.849057	0.833333
GPRC5A AND NOT-IL22RA1 AND NOT-GYPC	0.849057	0.865385	0.833333
GPRC5A AND NOT-PIGR AND NOT-SLC39A2	0.831683	0.893617	0.777778
SDC1 AND NOT-DUOXA1 AND NOT-TM4SF5	0.865385	0.9	0.833333
GPRC5A AND NOT-ATP10B AND NOT-CD52	0.857143	0.882353	0.833333
SDC1 AND NOT-DUOXA1 AND NOT-SLC39A5	0.873786	0.918367	0.833333
SDC1 AND NOT-DUOXA1 AND NOT-SLC17A4	0.833333	0.833333	0.833333
ADAM12 AND NOT-DPP4 AND NOT-GPA33	0.833333	0.833333	0.833333
ADAM12 AND NOT-SCN9A AND SLAMF7	0.825688	0.818182	0.833333
FAP AND NOT-NAALAD2 AND NOT-IL22RA1	0.871287	0.93617	0.814815
GPRC5A AND NOT-GPA33 AND NOT-DUOXA1	0.865385	0.9	0.833333
GPRC5A AND NOT-SSTR1 AND NOT-DUOXA1	0.857143	0.882353	0.833333
FAP AND NOT-NAALAD2 AND NOT-COL17A1	0.854369	0.897959	0.814815
SDC1 AND NOT-CYP4F12 AND NOT-DUOXA1	0.819048	0.843137	0.796296
GPRC5A AND NOT-GUCY2C AND NOT-DUOXA1	0.873786	0.918367	0.833333
GPRC5A AND NOT-IL22RA1 AND NOT-CD52	0.865385	0.9	0.833333
FAP AND NOT-COL17A1 AND NOT-DPP6	0.88	0.956522	0.814815
ADAM12 AND NOT-MME AND NOT-GPA33	0.815534	0.857143	0.777778
ADAM12 AND NOT-MME AND NOT-CD52	0.815534	0.857143	0.777778
FAP AND NOT-IL22RA1 AND NOT-DPP6	0.88	0.956522	0.814815
COMPLEX-C15orf27/PMEL/ADAM12	0.814815	0.814815	0.814815
FAP AND NOT-NAALAD2 AND NOT-DUOXA1	0.862745	0.916667	0.814815
FAP AND NOT-APCDD1 AND NOT-SCN9A	0.838095	0.862745	0.814815
FAP AND NOT-NAALAD2 AND NOT-LY6D	0.854369	0.897959	0.814815
FAP AND NOT-NAALAD2 AND NOT-GJB3	0.854369	0.897959	0.814815
GPRC5A AND NOT-DUOX1 AND NOT-SST	0.849057	0.865385	0.833333
FAP AND NOT-COL17A1 AND NOT-SCN9A	0.838095	0.862745	0.814815
CLDN7 AND NOT-AQP3 AND NOT-CA4	0.807692	0.84	0.777778
SDC1 AND NOT-ATP10B AND NOT-DUOXA1	0.807339	0.8	0.814815
FXYD3 AND NOT-DUOX1 AND NOT-SST	0.836066	0.75	0.944444
FAP AND NOT-NAALAD2 AND NOT-CHIC2	0.854369	0.897959	0.814815

Antigen Pair Logic	F1	Precision	Recall
CLECL1 AND NOT-IL12RB2 AND NOT-RECK	1	1	1
CLECL1 AND NOT-IL12RB2 AND NOT-DNAJC5	1	1	1
CLECL1 AND NOT-IL12RB2 AND NOT-IL10RB	0.987013	0.974359	1
CLECL1 AND NOT-IL12RB2 AND NOT-SLC30A5	1	1	1
CLECL1 AND NOT-IL12RB2 AND NOT-CYBB	1	1	1
CLECL1 AND NOT-IL12RB2 AND NOT-PTGER2	1	1	1
CLECL1 AND NOT-IL12RB2 AND NOT-PAG1	1	1	1
CLECL1 AND NOT-IL12RB2 AND NOT-CYBB	1	1	1
CLECL1 AND NOT-KCNH6 AND NOT-PTGER2	0.987013	0.974359	1
CLECL1 AND NOT-P2RY2 AND STRA6	0.974359	0.95	1
CLECL1 AND NOT-IL12RB2 AND NOT-AQP9	0.974359	0.95	1
COMPLEX-CLECL1/FPR2/LNPEP	0.974359	0.95	1
CLECL1 AND NOT-P2RY2 AND MRGPRX4	0.974359	0.95	1
CLECL1 AND NOT-KCNH6 AND NOT-CYBB	0.974359	0.95	1
CLECL1 AND NOT-KCNH6 AND NOT-CLEC4A	0.974359	0.95	1
CLECL1 AND NOT-ABCG2 AND NOT-FCER1A	0.973684	0.973684	0.973684
CLECL1 AND NOT-TREM2 AND NOT-FCER1A	0.973684	0.973684	0.973684
CLECL1 AND NOT-ERVFRD-1 AND NOT-FCER1A	0.973684	0.973684	0.973684
CLECL1 AND NOT-TAS2R16 AND NOT-NKG7	0.973684	0.973684	0.973684
CLECL1 AND NOT-SLC46A2 AND SLC25A3	0.973684	0.973684	0.973684
CLECL1 AND NOT-SLC46A2 AND MRGPRX4	0.973684	0.973684	0.973684
CLECL1 AND NOT-TAS2R16 AND NOT-CD28	0.973684	0.973684	0.973684
CLECL1 AND NOT-SLC46A2 AND STRA6	0.973684	0.973684	0.973684
COMPLEX-CLECL1/LNPEP/SLC46A2	0.973684	0.973684	0.973684
CLECL1 AND NOT-IL12RB2 AND NOT-NPC1	0.973684	0.973684	0.973684
CLECL1 AND NOT-ATP1A4 AND NOT-FCER1A	0.973684	0.973684	0.973684
CLECL1 AND NOT-TAS2R5 AND NOT-CLEC4A	0.973684	0.973684	0.973684
CLECL1 AND NOT-TAS2R16 AND NOT-TRAT1	0.973684	0.973684	0.973684
CLECL1 AND NOT-KCNG2 AND NOT-FCER1A	0.973684	0.973684	0.973684
CLECL1 AND NOT-SLC46A2 AND NOT-GPR1	0.973684	0.973684	0.973684
CLECL1 AND NOT-IL12RB2 AND NOT-SLAMF6	0.972973	1	0.947368
CLECL1 AND NOT-IL12RB2 AND ST14	0.972973	1	0.947368
CLECL1 AND NOT-SLC24A4 AND NOT-DEGS1	0.972973	1	0.947368
CLECL1 AND NOT-IL12RB2 AND NOT-IFNAR1	0.972973	1	0.947368
CLECL1 AND NOT-IL12RB2 AND NOT-CLEC12A	0.972973	1	0.947368
CLECL1 AND NOT-IL12RB2 AND NOT-CD86	0.972973	1	0.947368
CLECL1 AND NOT-ABCG2 AND NOT-SLC16A6	0.974359	0.95	1
CLECL1 AND NOT-KCNG2 AND NOT-IL10RB	0.974359	0.95	1
CLECL1 AND NOT-TREM2 AND NOT-NKG7	0.987013	0.974359	1
CLECL1 AND NOT-ABCG2 AND NOT-FURIN	0.972973	1	0.947368
CLECL1 AND NOT-ATP1A4 AND NOT-CD28	0.987013	0.974359	1
CLECL1 AND NOT-ATP1A4 AND NOT-CYBB	0.974359	0.95	1
CLECL1 AND NOT-ABCG2 AND NOT-CYBB	0.974359	0.95	1
CLECL1 AND NOT-TREM2 AND NOT-PILRA	0.974359	0.95	1
CLECL1 AND NOT-KCNG2 AND NOT-SLC44A1	0.974359	0.95	1
CLECL1 AND NOT-IL12RB2 AND NOT-LDLRAD3	1	1	1
CLECL1 AND NOT-ATP1A4 AND NOT-CLEC4D	0.974359	0.95	1
CLECL1 AND NOT-ABCG2 AND NOT-AMICA1	0.974359	0.95	1
CLECL1 AND NOT-KCNG2 AND NOT-IGSF6	0.974359	0.95	1
CLECL1 AND NOT-TREM2 AND NOT-SLC44A1	0.974359	0.95	1
CLECL1 AND NOT-TREM2 AND NOT-FLT3LG	0.987013	0.974359	1
COMPLEX-CD226/CLECL1/AQP9	0.987013	0.974359	1
CLECL1 AND NOT-TREM2 AND NOT-STOM	0.987013	0.974359	1
CLECL1 AND NOT-TREM2 AND NOT-SLC31A2	0.974359	0.95	1
CLECL1 AND NOT-ABCG2 AND NOT-RECK	0.974359	0.95	1
CLECL1 AND NOT-TREM2 AND NOT-RECK	0.974359	0.95	1
CLECL1 AND NOT-KCNG2 AND NOT-PILRA	0.974359	0.95	1
CLECL1 AND NOT-KCNG2 AND NOT-LILRB2	0.974359	0.95	1
CLECL1 AND NOT-ATP1A4 AND NOT-KLRD1	0.987013	0.974359	1
CLECL1 AND NOT-ATP1A4 AND NOT-LILRB2	0.974359	0.95	1
CLECL1 AND NOT-ATP1A4 AND NOT-GPR171	0.987013	0.974359	1
CLECL1 AND NOT-SHH AND NOT-CD28	0.973684	0.973684	0.973684
CLECL1 AND NOT-TREM2 AND NOT-DNAJC5	0.974359	0.95	1
CLECL1 AND NOT-ATP1A4 AND NOT-IGSF6	0.974359	0.95	1
CLECL1 AND NOT-TREM2 AND NOT-TRAT1	0.987013	0.974359	1
CLECL1 AND NOT-ABCG2 AND NOT-FCER1G	0.973684	0.973684	0.973684
CLECL1 AND NOT-KCNG2 AND NOT-NKG7	0.987013	0.974359	1
CLECL1 AND NOT-TREM2 AND NOT-CD3G	0.987013	0.974359	1
CLECL1 AND NOT-ATP1A4 AND NOT-NUCB2	0.974359	0.95	1
CLECL1 AND NOT-KCNG2 AND NOT-CD244	0.974359	0.95	1
CLECL1 AND NOT-TREM2 AND NOT-CD28	0.987013	0.974359	1
CLECL1 AND NOT-KCNG2 AND NOT-PRCP	0.974359	0.95	1
CLECL1 AND NOT-IL12RB2 AND NOT-SLC9A6	0.986667	1	0.973684
CLECL1 AND NOT-LRP12 AND NOT-DNAJC5	0.974359	0.95	1
CLECL1 AND NOT-IL12RB2 AND NOT-SIRPB1	1	1	1
CLECL1 AND NOT-LRP12 AND NOT-CYBB	0.974359	0.95	1
CLECL1 AND NOT-IL12RB2 AND NOT-CLEC4E	1	1	1

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
FAP AND NOT-DPP6 AND NOT-DUOXA1	0.88	0.956522	0.814815
FAP AND NOT-DPP6 AND NOT-LY6D	0.88	0.956522	0.814815
FAP AND NOT-NAALAD2 AND NOT-P2RX1	0.854369	0.897959	0.814815
FAP AND NOT-DPP6 AND NOT-GJB3	0.88	0.956522	0.814815
FAP AND NOT-IL22RA1 AND NOT-KCNMB2	0.871287	0.93617	0.814815
FAP AND NOT-SLC46A2 AND NOT-KCNMB2	0.84	0.913043	0.777778
GPRC5A AND NOT-IL22RA1 AND NOT-SSTR1	0.857143	0.882353	0.833333
FAP AND NOT-IL22RA1 AND KCNS1	0.854369	0.897959	0.814815
FAP AND NOT-NAALAD2 AND NOT-TMEM47	0.857143	0.954545	0.777778
GPRC5A AND NOT-DUOXA1 AND NOT-SST	0.833333	0.833333	0.833333
FAP AND NOT-CDHR1 AND NOT-KCNMB2	0.848485	0.933333	0.777778
GPRC5A AND NOT-GUCY2C AND NOT-SLC46A2	0.818182	0.803571	0.833333
FAP AND NOT-COL17A1 AND NOT-DPP4	0.838095	0.862745	0.814815
FAP AND NOT-FXYD1 AND NOT-DUOXA1	0.88	0.956522	0.814815
FAP AND NOT-IL22RA1 AND CYP4F12	0.831683	0.893617	0.777778
FAP AND NOT-IL22RA1 AND NOT-TREM1	0.843137	0.895833	0.796296
FAP AND NOT-IL22RA1 AND NOT-PRIMA1	0.851485	0.914894	0.796296
FAP AND NOT-COL17A1 AND KCNS1	0.838095	0.862745	0.814815
FAP AND NOT-IL22RA1 AND NOT-FPR1	0.854369	0.897959	0.814815
FAP AND NOT-APCDD1 AND NOT-GPBAR1	0.843137	0.895833	0.796296
FAP AND NOT-APCDD1 AND NOT-PAG1	0.815534	0.857143	0.777778
FAP AND NOT-DUOXA1 AND NOT-KCNMB2	0.84	0.913043	0.777778
FAP AND NOT-COL17A1 AND NOT-GPBAR1	0.843137	0.895833	0.796296
FAP AND NOT-IL22RA1 AND NOT-PAG1	0.843137	0.895833	0.796296
FAP AND NOT-COL17A1 AND NOT-TREM1	0.834951	0.877551	0.796296
FAP AND NOT-ATP10B AND LY75	0.804348	0.973684	0.685185
FAP AND NOT-IL22RA1 AND NOT-GPBAR1	0.843137	0.895833	0.796296
FAP AND NOT-IL20RB AND NOT-PAG1	0.803922	0.854167	0.759259
FAP AND NOT-SGCA AND NOT-GJB3	0.888889	0.977778	0.814815
FAP AND NOT-IL22RA1 AND ICAM3	0.868687	0.955556	0.796296
FAP AND NOT-CDHR1 AND NOT-ADRB1	0.851485	0.914894	0.796296
GPRC5A AND NOT-SSTR1 AND NOT-PTPRZ1	0.841121	0.849057	0.833333
FAP AND NOT-COL17A1 AND NOT-PAG1	0.815534	0.857143	0.777778
GPRC5A AND NOT-SSTR1 AND NOT-IL20RB	0.830189	0.846154	0.814815
FAP AND NOT-DPP4 AND NOT-DUOXA1	0.834951	0.877551	0.796296
FAP AND NOT-NAALAD2 AND NOT-TREM1	0.851485	0.914894	0.796296
FAP AND NOT-IL20RB AND NOT-PDGFRA	0.82	0.891304	0.759259
FAP AND NOT-COL17A1 AND NOT-IL1RL1	0.854369	0.897959	0.814815
FAP AND NOT-COL17A1 AND NOT-CHRNB3	0.815534	0.857143	0.777778
FAP AND NOT-COL17A1 AND NOT-C15orf27	0.826923	0.86	0.796296
FAP AND NOT-SLC46A2 AND ATP12A	0.82	0.891304	0.759259
FAP AND NOT-SGCA AND NOT-GPBAR1	0.877551	0.977273	0.796296
FAP AND NOT-APCDD1 AND NOT-PDGFRA	0.831683	0.893617	0.777778
FAP AND NOT-COL17A1 AND ICAM3	0.851485	0.914894	0.796296
FAP AND NOT-NAALAD2 AND NOT-CR1	0.8125	0.928571	0.722222
FAP AND NOT-ATP10B AND NOT-FPR1	0.843137	0.895833	0.796296
FAP AND NOT-IL22RA1 AND GRIK5	0.854369	0.897959	0.814815
FAP AND NOT-SCN9A AND NOT-GJB3	0.846154	0.88	0.814815
FAP AND NOT-SCN9A AND NOT-LY6D	0.846154	0.88	0.814815
GPRC5A AND NOT-SSTR1 AND NOT-DSC3	0.818182	0.803571	0.833333
FAP AND NOT-DUOXA1 AND NOT-PRLHR	0.831683	0.893617	0.777778
FAP AND NOT-SCN9A AND NOT-DUOXA1	0.854369	0.897959	0.814815
FAP AND NOT-COL17A1 AND NOT-SNAP23	0.838095	0.862745	0.814815
FAP AND NOT-NAALAD2 AND TMPRSS2	0.851485	0.914894	0.796296
FAP AND NOT-ATP10B AND ICAM3	0.848485	0.933333	0.777778
FAP AND NOT-DPP6 AND NOT-AQP3	0.88	0.956522	0.814815
FAP AND NOT-COL17A1 AND NOT-HHIP	0.838095	0.862745	0.814815
FAP AND NOT-COL17A1 AND RHCG	0.846154	0.88	0.814815
FAP AND NOT-PPAP2A AND KCND2	0.88	0.956522	0.814815
FAP AND NOT-IL22RA1 AND NOT-GYPC	0.871287	0.93617	0.814815
FAP AND NOT-IL22RA1 AND SEMA4D	0.851064	1	0.740741
FAP AND NOT-COL17A1 AND NOT-GYPC	0.854369	0.897959	0.814815
FAP AND NOT-CDHR1 AND NOT-GYPC	0.871287	0.93617	0.814815
FAP AND NOT-IL22RA1 AND VAMP8	0.888889	0.977778	0.814815
FAP AND NOT-ATP10B AND NOT-GYPC	0.851485	0.914894	0.796296
FAP AND NOT-IL22RA1 AND NOT-PTPRR	0.871287	0.93617	0.814815
FAP AND NOT-AVPR1A AND KCND2	0.86	0.934783	0.796296
FAP AND NOT-CDHR1 AND VAMP8	0.888889	0.977778	0.814815
FAP AND NOT-SLC46A2 AND VAMP8	0.88	0.956522	0.814815
FAP AND NOT-SLC46A2 AND NOT-PTPRR	0.862745	0.916667	0.814815
FAP AND NOT-APCDD1 AND NOT-GYPC	0.854369	0.897959	0.814815
SDC1 AND NOT-DUOXA1 AND NOT-CDH17	0.810811	0.789474	0.833333
FAP AND NOT-GJB3 AND NOT-GYPC	0.854369	0.897959	0.814815
FAP AND NOT-LY6D AND NOT-GYPC	0.854369	0.897959	0.814815
FAP AND NOT-DUOXA1 AND NOT-GYPC	0.862745	0.916667	0.814815
FAP AND NOT-ESYT3 AND NOT-GYPC	0.871287	0.93617	0.814815
FAP AND NOT-DUOXA1 AND NOT-PTPRR	0.862745	0.916667	0.814815

Antigen Pair Logic	F1	Precision	Recall
CLECL1 AND NOT-ERVFRD-1 AND NOT-SIRPB1	0.974359	0.95	1
CLECL1 AND NOT-LRP12 AND NOT-CCR6	0.974359	0.95	1
CLECL1 AND NOT-ABCG2 AND NOT-SLC39A11	0.974359	0.95	1
CLECL1 AND NOT-TREM2 AND NOT-ACVR2A	0.974359	0.95	1
CLECL1 AND NOT-GALR3 AND NOT-CD28	0.973684	0.973684	0.973684
CLECL1 AND NOT-IL12RB2 AND NOT-LILRB2	1	1	1
CLECL1 AND NOT-TREM2 AND NOT-CD244	0.974359	0.95	1
CLECL1 AND NOT-IL12RB2 AND NOT-CPD	1	1	1
CLECL1 AND NOT-ATP1A4 AND NOT-IL2RB	0.974359	0.95	1
COMPLEX-PTGDR2/CLECL1/LNPEP	0.974359	0.95	1
CLECL1 AND NOT-IL12RB2 AND NOT-IGSF6	1	1	1
CLECL1 AND NOT-TREM2 AND NOT-C5AR1	0.974359	0.95	1
CLECL1 AND NOT-LRP12 AND NOT-RECK	0.974359	0.95	1
CLECL1 AND NOT-ABCG2 AND NOT-CD28	0.987013	0.974359	1
CLECL1 AND NOT-TREM2 AND NOT-PTGDR	0.974359	0.95	1
CLECL1 AND NOT-KCNG2 AND NOT-PAG1	0.974359	0.95	1
CLECL1 AND NOT-ATP1A4 AND NOT-SLC24A4	0.974359	0.95	1
CLECL1 AND NOT-ABCG2 AND LRRC32	0.987013	0.974359	1
CLECL1 AND NOT-KCNG2 AND NOT-PTGDR	0.974359	0.95	1
CLECL1 AND NOT-TREM2 AND NOT-PAG1	0.974359	0.95	1
CLECL1 AND NOT-ABCG2 AND NOT-CLEC4E	0.974359	0.95	1
CLECL1 AND NOT-CHRNA4 AND NOT-CLEC4A	0.986667	1	0.973684
CLECL1 AND NOT-CHRNA4 AND NOT-TRPV2	0.974359	0.95	1
CLECL1 AND NOT-CHRNA4 AND NOT-TRPV2	0.974359	0.95	1
CLECL1 AND NOT-GUCY2D AND NOT-TRPV2	0.973684	0.973684	0.973684
CLECL1 AND NOT-CRB2 AND NOT-CD28	0.973684	0.973684	0.973684
CLECL1 AND NOT-CRB2 AND NOT-TRAT1	0.973684	0.973684	0.973684
CLECL1 AND NOT-SLC46A2 AND SEMA4D	0.973684	0.973684	0.973684
COMPLEX-CLECL1/MLC1/LNPEP	0.973684	0.973684	0.973684
CLECL1 AND NOT-CRB2 AND NOT-NKG7	0.973684	0.973684	0.973684
CLECL1 AND NOT-IL12RB2 AND NOT-EBP	0.972973	1	0.947368
CLECL1 AND NOT-FCER1A AND NOT-CHRNA4	0.972973	1	0.947368
CLECL1 AND NOT-CHRNA4 AND NOT-IL18RAP	1	1	1
CLECL1 AND NOT-CHRNA4 AND NOT-C5AR1	1	1	1
CLECL1 AND NOT-CHRNA4 AND NOT-SLC22A4	0.974359	0.95	1
CLECL1 AND NOT-CHRNA4 AND NOT-CD3D	1	1	1
CLECL1 AND NOT-ERVFRD-1 AND NOT-CD93	0.974359	0.95	1
CLECL1 AND NOT-CHRNA4 AND NOT-PTGDR	1	1	1
CLECL1 AND NOT-CHRNA4 AND NOT-CD28	1	1	1
CLECL1 AND NOT-CHRNA4 AND NOT-NFAM1	1	1	1
CLECL1 AND NOT-CHRNA4 AND NOT-PAG1	1	1	1
CLECL1 AND NOT-CHRNA4 AND NOT-CX3CR1	0.972973	1	0.947368
CLECL1 AND NOT-CHRNA4 AND NOT-LAMP2	0.974359	0.95	1
CLECL1 AND NOT-CHRNA4 AND NOT-ATP8B4	1	1	1
CLECL1 AND NOT-CHRNA4 AND NOT-KLRD1	1	1	1
CLECL1 AND NOT-CHRNA4 AND NOT-PRCP	0.987013	0.974359	1
CLECL1 AND NOT-SLC9B1 AND NOT-TRPV2	0.974359	0.95	1
CLECL1 AND NOT-KCNC2 AND NOT-CD28	0.973684	0.973684	0.973684
CLECL1 AND NOT-CHRNA4 AND NOT-CD244	1	1	1
CLECL1 AND NOT-MAG AND NOT-CYBB	0.974359	0.95	1
CLECL1 AND NOT-CHRNA4 AND NOT-TRAT1	1	1	1
CLECL1 AND NOT-CHRNA4 AND NOT-NKG7	1	1	1
CLECL1 AND NOT-CHRNA4 AND NOT-TLR2	0.974359	0.95	1
CLECL1 AND NOT-GUCY2D AND NOT-CYBB	0.973684	0.973684	0.973684
CLECL1 AND NOT-ATP1B4 AND NOT-CYBB	0.974359	0.95	1
CLECL1 AND NOT-KCNC2 AND NOT-BST1	0.973684	0.973684	0.973684
CLECL1 AND NOT-CHRNA4 AND NOT-TRAT1	1	1	1
CLECL1 AND NOT-TREM2 AND NOT-STEAP4	0.974359	0.95	1
CLECL1 AND NOT-MAG AND NOT-RECK	0.974359	0.95	1
CLECL1 AND NOT-MAG AND NOT-DNAJC5	0.974359	0.95	1
CLECL1 AND NOT-TREM2 AND NOT-CXCL16	0.973684	0.973684	0.973684
CLECL1 AND NOT-KCNC2 AND MRGPRX4	0.973684	0.973684	0.973684
CLECL1 AND NOT-ABCG2 AND NOT-ZDHHC2	0.974359	0.95	1
CLECL1 AND NOT-CHRNA4 AND NOT-SPN	1	1	1
CLECL1 AND NOT-CHRNA4 AND NOT-IL7R	0.987013	0.974359	1
CLECL1 AND NOT-CHRNA4 AND NOT-KLRB1	1	1	1
CLECL1 AND NOT-CHRNA4 AND NOT-NPC1	0.973684	0.973684	0.973684
CLECL1 AND NOT-TNFSF10 AND NOT-GPR61	0.974359	0.95	1
CLECL1 AND NOT-SHH AND NOT-CD93	0.973684	0.973684	0.973684
CLECL1 AND NOT-CHRNA4 AND NOT-P2RY13	1	1	1
CLECL1 AND NOT-CHRNA4 AND NOT-CD86	0.972973	1	0.947368
CLECL1 AND NOT-CHRNA4 AND NOT-ACVR2A	1	1	1
CLECL1 AND NOT-CHRNA4 AND NOT-CD3G	1	1	1
CLECL1 AND NOT-KCNH6 AND NOT-CD93	0.974359	0.95	1
CLECL1 AND NOT-IL12RB2 AND NOT-BTN3A3	1	1	1
CLECL1 AND NOT-CHRNA4 AND NOT-C10orf54	0.973684	0.973684	0.973684
CLECL1 AND NOT-IL12RB2 AND NOT-CD36	0.986667	1	0.973684

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
FAP AND NOT-DUOXA1 AND VAMP8	0.88	0.956522	0.814815
FAP AND NOT-IL22RA1 AND KCND2	0.871287	0.93617	0.814815
FAP AND NOT-COL17A1 AND NOT-GRM7	0.838095	0.862745	0.814815
FAP AND NOT-IL20RB AND KCND2	0.86	0.934783	0.796296
FAP AND OR51B2 AND NOT-GRM7	0.808511	0.95	0.703704
GPRC5A AND NOT-SSTR1 AND NOT-SLC39A2	0.849057	0.865385	0.833333
FAP AND NOT-APCDD1 AND NOT-GRM7	0.838095	0.862745	0.814815
FAP AND NOT-SLC46A2 AND KCND2	0.871287	0.93617	0.814815
FAP AND NOT-DSC3 AND NOT-GRM7	0.86	0.934783	0.796296
FAP AND NOT-COL17A1 AND PIRT	0.823529	0.875	0.777778
FAP AND NOT-COL17A1 AND KCND2	0.871287	0.93617	0.814815
FAP AND NOT-MLANA AND NOT-PAG1	0.834951	0.877551	0.796296
FAP AND NOT-SCN9A AND NOT-GYPC	0.862745	0.916667	0.814815
FAP AND NOT-P2RX1 AND CACNG1	0.808081	0.888889	0.740741
FAP AND NOT-APCDD1 AND KCND2	0.871287	0.93617	0.814815
FAP AND NOT-SGCA AND KCND2	0.888889	0.977778	0.814815
FAP AND NOT-ABCC9 AND KCND2	0.888889	0.977778	0.814815
FAP AND NOT-GJB3 AND NOT-GRM7	0.838095	0.862745	0.814815
FAP AND NOT-DUOXA1 AND CACNG1	0.862745	0.916667	0.814815
FAP AND NOT-P2RX1 AND KCND2	0.824742	0.930233	0.740741
FAP AND NOT-APCDD1 AND CACNG6	0.803922	0.854167	0.759259
FAP AND NOT-KCNK7 AND KCND2	0.871287	0.93617	0.814815
FAP AND NOT-LY6D AND NOT-GRM7	0.838095	0.862745	0.814815
FAP AND NOT-DUOXA1 AND PIRT	0.831683	0.893617	0.777778
FAP AND NOT-DUOXA1 AND KCND2	0.871287	0.93617	0.814815
FAP AND NOT-S1PR5 AND NOT-PTPRR	0.848485	0.933333	0.777778
FAP AND NOT-COL17A1 AND SLC30A10	0.843137	0.895833	0.796296
FAP AND NOT-AQP3 AND NOT-GYPC	0.854369	0.897959	0.814815
FAP AND NOT-GJB3 AND KCND2	0.871287	0.93617	0.814815
FAP AND NOT-ACVR2A AND CACNG1	0.843137	0.895833	0.796296
FAP AND NOT-DUOXA1 AND NOT-CLDN19	0.846154	0.88	0.814815
LRRC8E AND NOT-AQP3 AND NOT-SST	0.811881	0.87234	0.759259
FAP AND NOT-GJB3 AND NOT-GABRA4	0.846154	0.88	0.814815
FAP AND NOT-LY6D AND KCND2	0.871287	0.93617	0.814815
FAP AND NOT-APCDD1 AND SLC30A10	0.843137	0.895833	0.796296
FAP AND NOT-DUOXA1 AND CACNG6	0.811881	0.87234	0.759259
FAP AND NOT-COL17A1 AND SPAM1	0.831683	0.893617	0.777778
FAP AND NOT-DUOXA1 AND SLC13A2	0.823529	0.875	0.777778
FAP AND NOT-DUOXA1 AND SLC12A5	0.84	0.913043	0.777778
FAP AND NOT-SLC46A2 AND SLC30A10	0.851485	0.914894	0.796296
GPRC5A AND NOT-GPA33 AND NOT-SLC39A2	0.849057	0.865385	0.833333
COMPLEX-COL17A1/FAP/FXYD7	0.838095	0.862745	0.814815
FAP AND NOT-S1PR5 AND KCND2	0.86	0.934783	0.796296
FAP AND NOT-TENM1 AND KCND2	0.871287	0.93617	0.814815
FAP AND NOT-SCN9A AND KCND2	0.871287	0.93617	0.814815
FAP AND NOT-TNFRSF19 AND KCND2	0.871287	0.93617	0.814815
FAP AND NOT-DUOXA1 AND OR1C1	0.816327	0.909091	0.740741
FAP AND NOT-CADM3 AND KCND2	0.871287	0.93617	0.814815
FAP AND NOT-DUOXA1 AND SPAM1	0.84	0.913043	0.777778
COMPLEX-FAP/SLC5A2/DUOXA1	0.846154	0.88	0.814815
FAP AND NOT-CAV2 AND NOT-SST	0.828283	0.911111	0.759259
FAP AND NOT-IL22RA1 AND NOT-NCAM1	0.886598	1	0.796296
SDC1 AND NOT-AQP3 AND NOT-SSTR1	0.849057	0.865385	0.833333
FAP AND NOT-COL17A1 AND NOT-NCAM1	0.868687	0.955556	0.796296
FAP AND NOT-CAV2 AND NOT-GPA33	0.828283	0.911111	0.759259
FAP AND NOT-APCDD1 AND NOT-NCAM1	0.868687	0.955556	0.796296
FAP AND NOT-DUOXA1 AND NOT-NCAM1	0.877551	0.977273	0.796296
FAP AND NOT-GJB3 AND NOT-NCAM1	0.868687	0.955556	0.796296
FAP AND NOT-LY6D AND NOT-NCAM1	0.868687	0.955556	0.796296
FAP AND NOT-PMEL AND NOT-GPBAR1	0.843137	0.895833	0.796296
FAP AND NOT-NAALAD2 AND NOT-ABCA5	0.831683	0.893617	0.777778
FAP AND NOT-APCDD1 AND NOT-SSTR1	0.838095	0.862745	0.814815
FAP AND NOT-PMEL AND NOT-C15orf27	0.826923	0.86	0.796296
FAP AND NOT-SLC46A2 AND NOT-ABCA5	0.823529	0.875	0.777778
FAP AND NOT-TNFRSF19 AND NOT-NCAM1	0.868687	0.955556	0.796296
MUC1 AND NOT-SSTR1 AND NOT-AQP3	0.803419	0.746032	0.87037
FAP AND NOT-APCDD1 AND NOT-GPA33	0.838095	0.862745	0.814815
FAP AND NOT-COL17A1 AND CD52	0.838095	0.862745	0.814815
FAP AND NOT-LY6D AND SLAMF7	0.838095	0.862745	0.814815
FAP AND NOT-GJB3 AND SLAMF7	0.838095	0.862745	0.814815
FAP AND NOT-P2RX1 AND NOT-NCAM1	0.833333	0.952381	0.740741
FAP AND NOT-ATP10B AND CD52	0.834951	0.877551	0.796296
FAP AND NOT-DSC3 AND NOT-SSTR1	0.843137	0.895833	0.796296
FAP AND NOT-DUOXA1 AND NOT-SSTR1	0.846154	0.88	0.814815
FAP AND NOT-PTPRZ1 AND NOT-SSTR1	0.838095	0.862745	0.814815
FAP AND NOT-DUOXA1 AND NOT-ABCA5	0.823529	0.875	0.777778
MUC1 AND NOT-AQP3 AND NOT-SST	0.817391	0.770492	0.87037

Antigen Pair Logic	F1	Precision	Recall
CLECL1 AND NOT-CHRNA4 AND NOT-C3AR1	1	1	1
CLECL1 AND NOT-CHRNA4 AND NOT-GOLM1	0.974359	0.95	1
CLECL1 AND NOT-CHRNA4 AND NOT-PTGER2	1	1	1
CLECL1 AND NOT-OR3A2 AND NOT-NKG7	0.972973	1	0.947368
CLECL1 AND NOT-MAG AND NOT-CYBB	0.974359	0.95	1
CLECL1 AND NOT-IL12RB2 AND NOT-STEAP4	1	1	1
CLECL1 AND NOT-ERVFRD-1 AND NOT-STEAP4	0.974359	0.95	1
CLECL1 AND NOT-CHRNA4 AND NOT-SELPLG	1	1	1
CLECL1 AND NOT-ERVFRD-1 AND NOT-CXCL16	0.973684	0.973684	0.973684
CLECL1 AND NOT-OR3A2 AND NOT-TRAT1	0.972973	1	0.947368
CLECL1 AND NOT-KCNC2 AND NOT-SLC24A4	0.973684	0.973684	0.973684
CLECL1 AND NOT-KCNH6 AND NOT-CD93	0.974359	0.95	1
CLECL1 AND NOT-GRID2 AND NOT-CYBB	0.974359	0.95	1
CLECL1 AND NOT-KLRD1 AND NOT-GPR61	0.973684	0.973684	0.973684
CLECL1 AND NOT-SLC9B1 AND NOT-TRPV2	0.974359	0.95	1
CLECL1 AND NOT-KCNC2 AND NOT-TRAT1	0.986667	1	0.973684
CLECL1 AND NOT-ATP1B4 AND NOT-PTGER2	0.987013	0.974359	1
CLECL1 AND NOT-KCNH6 AND NOT-CXCL16	0.973684	0.973684	0.973684
COMPLEX-CLECL1/MLC1/KCNG2	0.973684	0.973684	0.973684
CLECL1 AND NOT-KCNC2 AND NOT-CYBB	0.973684	0.973684	0.973684
CLECL1 AND NOT-GRID2 AND NOT-PAG1	0.974359	0.95	1
CLECL1 AND NOT-KCNG2 AND NOT-STEAP4	0.974359	0.95	1
CLECL1 AND NOT-KCNG2 AND NOT-CXCL16	0.973684	0.973684	0.973684
CLECL1 AND NOT-TREM2 AND NOT-CD93	0.974359	0.95	1
CLECL1 AND NOT-CHRNA4 AND NOT-BST1	1	1	1
CLECL1 AND NOT-CHRNA4 AND NOT-SMPD1	0.987013	0.974359	1
CLECL1 AND NOT-IL12RB2 AND TMEFF2	0.973684	0.973684	0.973684
CLECL1 AND NOT-CHRNA4 AND NOT-TNFSF10	1	1	1
CLECL1 AND NOT-KCNG2 AND NOT-CD93	0.974359	0.95	1
CLECL1 AND NOT-KCNC2 AND NOT-NKG7	0.986667	1	0.973684
CLECL1 AND NOT-CHRNA4 AND NOT-CYBB	1	1	1
CLECL1 AND NOT-CHRNA4 AND NOT-KLRF1	0.972973	1	0.947368
CLECL1 AND NOT-CHRNA4 AND NOT-IL10RB	0.987013	0.974359	1
CLECL1 AND NOT-GALR2 AND NOT-PAG1	0.974359	0.95	1
CLECL1 AND NOT-CHRNA4 AND NOT-PILRA	1	1	1
CLECL1 AND NOT-MAG AND NOT-PAG1	0.974359	0.95	1
CLECL1 AND NOT-CHRNA4 AND NOT-CLEC7A	1	1	1
CLECL1 AND NOT-KCNC2 AND STRA6	0.973684	0.973684	0.973684
CLECL1 AND NOT-CHRNA4 AND NOT-SMPD1	0.987013	0.974359	1
CLECL1 AND NOT-CHRNA4 AND NOT-BST1	1	1	1
CLECL1 AND NOT-CHRNA4 AND NOT-TLR5	1	1	1
CLECL1 AND NOT-CHRNA4 AND NOT-TYROBP	0.986667	1	0.973684
CLECL1 AND NOT-CHRNA4 AND NOT-SLAMF6	0.972973	1	0.947368
CLECL1 AND NOT-GALR2 AND NOT-RECK	0.974359	0.95	1
MS4A1 AND PKD2L1 AND NOT-P2RY2	0.986667	1	0.973684
MS4A1 AND PKD2L1 AND NOT-PTGDR2	0.986667	1	0.973684
MS4A1 AND PKD2L1 AND NOT-TGOLN2	0.986667	1	0.973684
MS4A1 AND GDE1 AND NOT-PTGDR2	0.986667	1	0.973684
MS4A1 AND STRA6 AND NOT-TGOLN2	0.974359	0.95	1
MS4A1 AND STRA6 AND NOT-P2RY2	0.974359	0.95	1
MS4A1 AND STRA6 AND NOT-PTGDR2	0.974359	0.95	1
MS4A1 AND STRA6 AND NOT-PTGDR2	0.974359	0.95	1
MS4A1 AND STRA6 AND NOT-TGOLN2	0.974359	0.95	1
MS4A1 AND STRA6 AND NOT-P2RY2	0.974359	0.95	1
MS4A1 AND PIGR AND NOT-PSEN1	0.973684	0.973684	0.973684
MS4A1 AND STRA6 AND NOT-SLC46A2	0.973684	0.973684	0.973684
MS4A1 AND MARVELD2 AND NOT-SLC46A2	0.973684	0.973684	0.973684
MS4A1 AND STRA6 AND NOT-IL12RB2	0.973684	0.973684	0.973684
MS4A1 AND STRA6 AND NOT-IL12RB2	0.973684	0.973684	0.973684
MS4A1 AND PIGR AND NOT-PTGDR2	0.973684	0.973684	0.973684
MS4A1 AND STRA6 AND NOT-FPR2	0.973684	0.973684	0.973684
MS4A1 AND PKD2L1 AND NOT-IL12RB2	0.972973	1	0.947368
CLECL1 AND NOT-IL12RB2 AND NOT-CD33	0.987013	0.974359	1
CLECL1 AND NOT-CLDN2 AND NOT-PTGER2	0.987013	0.974359	1
CLECL1 AND NOT-SLC24A4 AND NOT-DDX3X	0.974359	0.95	1
CLECL1 AND NOT-DNAJB8 AND NOT-BST1	0.974359	0.95	1
CLECL1 AND NOT-DNAJB8 AND NOT-RECK	0.974359	0.95	1
CLECL1 AND NOT-CLDN2 AND NOT-DNAJC5	0.974359	0.95	1
CLECL1 AND NOT-IL12RB2 AND NOT-HSPA5	0.972973	1	0.947368
CLECL1 AND NOT-CLDN2 AND NOT-RECK	0.974359	0.95	1
MS4A1 AND PIGR AND NOT-SERINC3	0.973684	0.973684	0.973684
MS4A1 AND MARVELD2 AND NOT-PTGDR2	0.974359	0.95	1
CLECL1 AND NOT-CLDN2 AND NOT-PAG1	0.974359	0.95	1
CLECL1 AND NOT-DNAJB8 AND NOT-LILRB2	0.974359	0.95	1
COMPLEX-CLECL1/ITGB3/LNPEP	0.987013	0.974359	1
CLECL1 AND NOT-DNAJB8 AND NOT-CD28	0.987013	0.974359	1
CLECL1 AND NOT-ATP1A4 AND NOT-CD160	0.974359	0.95	1

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
FAP AND NOT-DUOXA1 AND CD52	0.846154	0.88	0.814815
FAP AND NOT-GJB3 AND CD52	0.838095	0.862745	0.814815
FAP AND NOT-LY6D AND CD52	0.838095	0.862745	0.814815
FAP AND NOT-LY6D AND NOT-SSTR1	0.846154	0.88	0.814815
FAP AND NOT-CADM3 AND NOT-ABCA5	0.831683	0.893617	0.777778
FAP AND NOT-EPHB6 AND CD52	0.826923	0.86	0.796296
FAP AND NOT-ABCA5 AND NOT-IL1RL1	0.831683	0.893617	0.777778
FAP AND NOT-COL17A1 AND SLAMF7	0.8	0.926829	0.703704
FAP AND NOT-TNFRSF19 AND NOT-SSTR1	0.854369	0.897959	0.814815
FAP AND NOT-ABCA5 AND KCNS1	0.815534	0.857143	0.777778
FAP AND NOT-ABCA5 AND NOT-CYSLTR2	0.815534	0.857143	0.777778
SDC1 AND NOT-AQP3 AND NOT-GPA33	0.849057	0.865385	0.833333
FAP AND NOT-AQP3 AND NOT-SSTR1	0.846154	0.88	0.814815
FAP AND NOT-AQP3 AND CD52	0.838095	0.862745	0.814815
FAP AND NOT-FNDC5 AND NOT-DDX3X	0.871287	0.93617	0.814815
MUC1 AND NOT-MUC4 AND NOT-DPP6	0.814159	0.779661	0.851852
FAP AND NOT-CHIC2 AND NOT-GPA33	0.84	0.913043	0.777778
SDC1 AND NOT-AQP3 AND NOT-SST	0.833333	0.833333	0.833333
FAP AND MEP1A AND NOT-GPA33	0.851485	0.914894	0.796296
PPAPDC1A AND NOT-OMG AND NOT-IL22RA1	0.808511	0.95	0.703704
FAP AND NOT-PMEL AND KCND2	0.871287	0.93617	0.814815
FAP AND NOT-ABCA5 AND NOT-GYPC	0.831683	0.893617	0.777778
COMPLEX-FAP/SLC30A10/PMEL	0.838095	0.862745	0.814815
FAP AND NOT-CDH22 AND CD52	0.815534	0.857143	0.777778
COMPLEX-FAP/ABCA5/GABRG2	0.815534	0.857143	0.777778
FAP AND NOT-GUCY2C AND KCND2	0.88	0.956522	0.814815
FAP AND NOT-SSTR1 AND KCND2	0.871287	0.93617	0.814815
FAP AND NOT-DDX3X AND GLRA2	0.838095	0.862745	0.814815
FAP AND KCND2 AND NOT-CLDN18	0.871287	0.93617	0.814815
FAP AND KCND2 AND NOT-SST	0.871287	0.93617	0.814815
FAP AND SLC10A1 AND NOT-CLDN18	0.838095	0.862745	0.814815
FAP AND KCND2 AND NOT-GPA33	0.868687	0.955556	0.796296
FAP AND GLRA2 AND NOT-TNFRSF13C	0.838095	0.862745	0.814815
COMPLEX-FAP/ZACN/CEACAM6	0.838095	0.862745	0.814815
FAP AND NOT-SLC39A2 AND NOT-GYPC	0.862745	0.916667	0.814815
FAP AND NOT-MLANA AND NOT-PTPRR	0.862745	0.916667	0.814815
FAP AND NOT-SLC39A2 AND KCND2	0.871287	0.93617	0.814815
FAP AND NOT-CDH22 AND KCND2	0.848485	0.933333	0.777778
FAP AND NOT-CDH22 AND GLRA2	0.815534	0.857143	0.777778
FAP AND ADAM29 AND NOT-PIK3IP1	0.838095	0.862745	0.814815
FAP AND NOT-GABRA4 AND KCND2	0.871287	0.93617	0.814815
FAP AND NOT-GYPC AND NOT-KCNA1	0.851485	0.914894	0.796296
FAP AND NOT-PIK3IP1 AND GLRA2	0.838095	0.862745	0.814815
FAP AND SLC10A1 AND NOT-SLC13A5	0.838095	0.862745	0.814815
FAP AND NOT-GPR22 AND NOT-PIK3IP1	0.862745	0.916667	0.814815
FAP AND NOT-HCN1 AND NOT-PIK3IP1	0.838095	0.862745	0.814815
FAP AND KCND2 AND NOT-TMEFF2	0.871287	0.93617	0.814815
FAP AND KCND2 AND NOT-CALN1	0.871287	0.93617	0.814815
FAP AND KCND2 AND NOT-SLC13A5	0.871287	0.93617	0.814815
FAP AND KCND2 AND NOT-HCN1	0.871287	0.93617	0.814815
FAP AND KCND2 AND NOT-OPCML	0.871287	0.93617	0.814815
FAP AND KCND2 AND NOT-MPL	0.871287	0.93617	0.814815
FAP AND NOT-CLDN16 AND GLRA2	0.815534	0.857143	0.777778
FAP AND KCND2 AND NOT-BEST2	0.871287	0.93617	0.814815
FAP AND KCND2 AND CACNG3	0.8	0.926829	0.703704
FAP AND KCND2 AND JPH3	0.8	0.926829	0.703704
FAP AND KCND2 AND NOT-CACNA1E	0.871287	0.93617	0.814815
FAP AND NOT-PIK3IP1 AND NOT-FXYD7	0.846154	0.88	0.814815
FAP AND KCND2 AND NOT-HTR1E	0.868687	0.955556	0.796296
FAP AND KCND2 AND NOT-FSHR	0.888889	0.977778	0.814815
FAP AND KCND2 AND NOT-GLRA2	0.871287	0.93617	0.814815
FAP AND KCND2 AND NOT-AGER	0.871287	0.93617	0.814815
FAP AND GLRA2 AND SLC13A2	0.815534	0.857143	0.777778
FAP AND GLRA2 AND NOT-AMHR2	0.826923	0.86	0.796296
FAP AND GLRA2 AND NOT-BEST2	0.838095	0.862745	0.814815
FAP AND GLRA2 AND NOT-HTR1E	0.834951	0.877551	0.796296
FAP AND GLRA2 AND NOT-FSHR	0.854369	0.897959	0.814815
COMPLEX-FAP/SLC24A2/KCND2	0.871287	0.93617	0.814815
COMPLEX-FAP/SLC24A2/GLRA2	0.838095	0.862745	0.814815
COMPLEX-FAP/GLRA2/KIRREL3	0.838095	0.862745	0.814815
COMPLEX-AGER/FAP/GLRA2	0.838095	0.862745	0.814815
PPAPDC1A AND CD52 AND NOT-OR1Q1	0.808511	0.95	0.703704
EPCAM AND NOT-VSIG2 AND NOT-DPP6	0.815534	0.857143	0.777778
EPCAM AND NOT-CA4 AND NOT-AQP3	0.807692	0.84	0.777778
EPCAM AND NOT-ATP10B AND ICAM3	0.826087	1	0.703704
FAP AND NOT-TYR AND NOT-PAG1	0.826923	0.86	0.796296
FAP AND NOT-TYR AND NOT-PDGFRA	0.831683	0.893617	0.777778

Antigen Pair Logic	F1	Precision	Recall
CLECL1 AND NOT-DNAJB8 AND NOT-NKG7	0.987013	0.974359	1
CLECL1 AND NOT-CLDN2 AND NOT-CYBB	0.974359	0.95	1
CLECL1 AND NOT-DNAJB8 AND NOT-TNFSF10	0.974359	0.95	1
CLECL1 AND NOT-DNAJB8 AND NOT-CD226	0.974359	0.95	1
CLECL1 AND NOT-KCNG2 AND NOT-CD33	0.974359	0.95	1
CLECL1 AND NOT-CLDN2 AND NOT-CYBB	0.974359	0.95	1
CLECL1 AND NOT-ERVFRD-1 AND NOT-CD160	0.974359	0.95	1
COMPLEX-CLECL1/KCNG2/ITGB3	0.987013	0.974359	1
CLECL1 AND NOT-SSTR1 AND NOT-DNAJC5	0.974359	0.95	1
CLECL1 AND NOT-KCNH6 AND NOT-CD160	0.974359	0.95	1
MS4A1 AND STRA6 AND NOT-PTPRA	0.974359	0.95	1
CLECL1 AND NOT-IL12RB2 AND NOT-ALDH1A1	0.974359	0.95	1
CLECL1 AND NOT-DNAJB8 AND NOT-TRAT1	0.987013	0.974359	1
CLECL1 AND NOT-DNAJB8 AND NOT-SELPLG	0.974359	0.95	1
CLECL1 AND NOT-IL12RB2 AND NOT-CD160	1	1	1
COMPLEX-CLECL1/IFNGR1/ITGB3	0.974359	0.95	1
CLECL1 AND NOT-CLDN2 AND NOT-RECK	0.974359	0.95	1
CLECL1 AND NOT-DNAJB8 AND NOT-SLC24A4	0.974359	0.95	1
COMPLEX-ADAM8/CLECL1/ITGB3	0.974359	0.95	1
CLECL1 AND NOT-DNAJB8 AND NOT-CYBB	0.974359	0.95	1
CLECL1 AND NOT-ALDH1A1 AND NOT-TGFBR1	0.974359	0.95	1
MS4A1 AND MARVELD2 AND NOT-PTGDR2	0.974359	0.95	1
CLECL1 AND NOT-SSTR1 AND NOT-PAG1	0.974359	0.95	1
CLECL1 AND NOT-ABCG2 AND NOT-HSPA5	0.973684	0.973684	0.973684
CLECL1 AND NOT-TREM2 AND NOT-CD160	0.974359	0.95	1
CLECL1 AND NOT-KCNG2 AND NOT-CD160	0.974359	0.95	1
MS4A1 AND MARVELD2 AND NOT-TGOLN2	0.974359	0.95	1
MS4A1 AND PIGR AND NOT-M6PR	0.973684	0.973684	0.973684
CLECL1 AND NOT-ITGB3 AND NOT-TRPV2	1	1	1
CLECL1 AND NOT-ITGB3 AND TSPAN15	0.987013	0.974359	1
CLECL1 AND NOT-ITGB3 AND SLC25A3	0.987013	0.974359	1
CLECL1 AND NOT-ITGB3 AND MRGPRX4	1	1	1
MS4A1 AND MARVELD2 AND NOT-FPR2	0.973684	0.973684	0.973684
MS4A1 AND STRA6 AND NOT-ACPP	0.974359	0.95	1
CLECL1 AND NOT-ITGB3 AND PPAP2A	0.973684	0.973684	0.973684
CLECL1 AND NOT-PSCA AND NOT-TRPV2	0.974359	0.95	1
CLECL1 AND NOT-ITGB3 AND STRA6	1	1	1
CLECL1 AND NOT-ITGB3 AND NOT-IL12RB2	0.987013	0.974359	1
CLECL1 AND NOT-ITGB3 AND NOT-GPR1	0.987013	0.974359	1
MS4A1 AND STRA6 AND NOT-FAM26E	0.973684	0.973684	0.973684
CLECL1 AND NOT-ITGB3 AND NOT-KCNH6	0.987013	0.974359	1
CLECL1 AND NOT-ITGB3 AND PPAP2A	0.973684	0.973684	0.973684
CLECL1 AND NOT-ITGB3 AND NOT-ERVFRD-1	0.973684	0.973684	0.973684
CLECL1 AND NOT-ITGB3 AND HTR7	0.973684	0.973684	0.973684
MS4A1 AND MARVELD2 AND NOT-P2RY2	0.962025	0.926829	1
MS4A1 AND NOT-MS4A2 AND MARVELD2	0.962025	0.926829	1
CLECL1 AND NOT-ATP1A4 AND NOT-CD33	0.962025	0.926829	1
CLECL1 AND NOT-IL12RB2 AND TNFRSF13C	0.962025	0.926829	1
CLECL1 AND NOT-DNAJB8 AND NOT-CD300LF	0.962025	0.926829	1
MS4A1 AND MRGPRX4 AND NOT-P2RY2	0.962025	0.926829	1
MS4A1 AND STRA6 AND NOT-KCNG2	0.962025	0.926829	1
CLECL1 AND NOT-DNAJB8 AND NOT-MGST2	0.962025	0.926829	1
CLECL1 AND NOT-MST1R AND NOT-DNAJC5	0.962025	0.926829	1
CLECL1 AND NOT-CLDN2 AND NOT-IL10RB	0.962025	0.926829	1
MS4A1 AND MRGPRX4 AND NOT-P2RY2	0.962025	0.926829	1
CLECL1 AND NOT-KCNG2 AND NOT-ALDH1A1	0.962025	0.926829	1
CLECL1 AND NOT-ERVFRD-1 AND NOT-CD33	0.962025	0.926829	1
MS4A1 AND MARVELD2 AND NOT-P2RY2	0.962025	0.926829	1
CLECL1 AND NOT-MST1R AND NOT-TRPV2	0.962025	0.926829	1
MS4A1 AND STRA6 AND NOT-MS4A2	0.962025	0.926829	1
CLECL1 AND NOT-ABCG2 AND NOT-CD33	0.962025	0.926829	1
CLECL1 AND NOT-KCNH6 AND NOT-CD33	0.962025	0.926829	1
CLECL1 AND NOT-MST1R AND NOT-TRPV2	0.962025	0.926829	1
CLECL1 AND NOT-DNAJB8 AND NOT-SMPD1	0.962025	0.926829	1
CLECL1 AND NOT-DNAJB8 AND NOT-LRRC8D	0.962025	0.926829	1
CLECL1 AND NOT-DLL3 AND NOT-TRPV2	0.962025	0.926829	1
CLECL1 AND NOT-TREM2 AND NOT-CD33	0.962025	0.926829	1
CLECL1 AND NOT-CHRNA4 AND NOT-STEAP4	1	1	1
CLECL1 AND NOT-CHRNA4 AND NOT-CD93	1	1	1
CLECL1 AND NOT-CHRNA4 AND NOT-CXCL16	0.986667	1	0.973684
CLECL1 AND NOT-CHRNA4 AND NOT-CD36	0.986667	1	0.973684
CLECL1 AND NOT-CHRNA4 AND NOT-CD93	1	1	1
CLECL1 AND NOT-CHRNA4 AND NOT-CD163	0.987013	0.974359	1
CLECL1 AND NOT-KCNC2 AND NOT-CD93	0.973684	0.973684	0.973684
CLECL1 AND NOT-GPR78 AND NOT-CD93	0.986667	1	0.973684
CLECL1 AND NOT-MEGF11 AND NOT-CD93	0.974359	0.95	1
CLECL1 AND NOT-KCNK4 AND NOT-CD93	0.987013	0.974359	1

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
EPCAM AND NOT-MUC4 AND NOT-DPP6	0.807692	0.84	0.777778
GPRC5A AND NOT-GPA33 AND NOT-TYR	0.810811	0.789474	0.833333
FAP AND NOT-CLDN1 AND NOT-ABCA5	0.828283	0.911111	0.759259
FAP AND NOT-ABCA5 AND NOT-SSTR1	0.815534	0.857143	0.777778
EPCAM AND NOT-BEST2 AND KCND2	0.803922	0.854167	0.759259
EPCAM AND NOT-NOX1 AND KCND2	0.803922	0.854167	0.759259
EPCAM AND NOT-GRM7 AND KCND2	0.824742	0.930233	0.740741
LRRC8E AND NOT-TYR AND KCND2	0.813187	1	0.685185
TGFBI AND NOT-TYR AND KCND2	0.815534	0.857143	0.777778
BST2 AND KCND2 AND NOT-TYR	0.804124	0.906977	0.722222
ATP8B1 AND NOT-TYR AND KCND2	0.824742	0.930233	0.740741
FAP AND NOT-TYR AND KCND2	0.871287	0.93617	0.814815
FAP AND NOT-TYR AND SLC30A10	0.843137	0.895833	0.796296
EPCAM AND NOT-SST AND KCND2	0.803738	0.811321	0.796296
SDC1 AND NOT-TYR AND KCND2	0.857143	0.954545	0.777778
MUC1 AND NOT-TYR AND KCND2	0.851064	1	0.740741
CLDN4 AND NOT-TYR AND KCND2	0.826087	1	0.703704
Liver Cancer			
ABCG5 AND NOT-MUC17 AND HM13	1	1	1
ABCG5 AND NOT-MUC17 AND HM13	1	1	1
SLCO1B1 AND NOT-MARCO AND AGER	1	1	1
SLCO1B1 AND NOT-MARCO AND SLC5A2	1	1	1
SLCO1B1 AND NOT-MARCO AND DSCAM	1	1	1
SLCO1B1 AND NOT-MARCO AND CSPG5	1	1	1
SLCO1B1 AND NOT-MARCO AND NPFFR1	1	1	1
SLCO1B1 AND NOT-MARCO AND ZACN	1	1	1
SLCO1B1 AND NOT-MARCO AND FLVCR1	1	1	1
SLCO1B1 AND NOT-MARCO AND FAT1	1	1	1
SLCO1B1 AND NOT-MARCO AND ALCAM	1	1	1
SLCO1B1 AND NOT-MARCO AND MIP	1	1	1
SLCO1B1 AND NOT-MARCO AND PTGIR	1	1	1
ABCG5 AND FGFRL1 AND NOT-SCARA5	1	1	1
ABCG5 AND HM13 AND NOT-CDH17	1	1	1
ABCG5 AND HM13 AND NOT-SLC13A2	1	1	1
ABCG5 AND HM13 AND NOT-SLC16A5	1	1	1
ABCG5 AND FLVCR1 AND NOT-MEP1B	1	1	1
ABCG5 AND HM13 AND NOT-SLC16A5	1	1	1
ABCG5 AND HM13 AND NOT-CDH17	1	1	1
ABCG5 AND FLVCR1 AND NOT-MEP1B	1	1	1
ABCG5 AND HM13 AND NOT-PTPRR	1	1	1
ABCG5 AND HM13 AND NOT-SLC13A2	1	1	1
ABCG5 AND HM13 AND NOT-FUT3	1	1	1
ABCG5 AND HM13 AND NOT-PTPRR	1	1	1
ABCG5 AND FGFRL1 AND NOT-SLC31A1	1	1	1
SLC43A1 AND RGR AND NOT-CLEC4G	1	1	1
SLC43A1 AND FLVCR1 AND NOT-LRRC8B	1	1	1
SLCO1B1 AND NOT-MARCO AND CD58	1	1	1
SLCO1B1 AND NOT-MARCO AND DYSF	1	1	1
ABCG5 AND FGFRL1 AND NOT-GPR83	1	1	1
ABCG5 AND FGFRL1 AND NOT-MUC17	1	1	1
ABCG5 AND FGFRL1 AND NOT-MUC17	1	1	1
SLC43A1 AND NOT-GPR22 AND SLC26A6	1	1	1
SLCO1B1 AND NOT-MARCO AND P2RX4	1	1	1
SLC43A1 AND RGR AND NOT-MARCO	1	1	1
SLCO1B1 AND SLC7A11 AND NOT-MUC17	1	1	1
ABCG5 AND FGFRL1 AND NOT-IL15RA	1	1	1
SLCO1B1 AND HM13 AND NOT-OR2C3	1	1	1
ABCG5 AND NOT-MARCO AND DYSF	1	1	1
SLCO1B1 AND HM13 AND NOT-TRHR	1	1	1
SLCO1B1 AND HM13 AND NOT-GABRA5	1	1	1
SLCO1B1 AND NOT-MARCO AND CHRNB2	1	1	1
SLCO1B1 AND HM13 AND NOT-SLC31A1	1	1	1
SLCO1B1 AND HM13 AND NOT-MUC17	1	1	1
SLCO1B1 AND SLC7A11 AND NOT-MUC17	1	1	1
SLCO1B1 AND HM13 AND NOT-MUC17	1	1	1
SLCO1B1 AND HM13 AND NOT-SLC17A2	1	1	1
SLCO1B1 AND SLC7A11 AND NOT-SLC17A2	1	1	1
SLC43A1 AND RGR AND NOT-MARCO	1	1	1
SLCO1B1 AND NOT-MARCO AND SLC30A1	1	1	1
COMPLEX-P2RY4/SLC2A2/SLC4A2	1	1	1
SLC2A2 AND SLC26A6 AND C8B	1	1	1
SLC2A2 AND SLC26A6 AND SLC6A12	1	1	1
SLC2A2 AND SLC26A6 AND NOT-PAQR8	1	1	1
SLCO1B1 AND NOT-MARCO AND CALY	1	1	1
SLC2A2 AND SLC26A6 AND DYSF	1	1	1
SLCO1B1 AND NOT-MARCO AND TNFRSF12A	1	1	1
SLC2A2 AND SLC26A6 AND S1PR1	1	1	1

Antigen Pair Logic	F1	Precision	Recall
CLECL1 AND NOT-GPR78 AND NOT-PAQR8	0.986667	1	0.973684
CLECL1 AND NOT-MEGF11 AND NOT-PAQR8	0.962025	0.926829	1
CLECL1 AND NOT-GRID2 AND NOT-STEAP4	0.974359	0.95	1
CLECL1 AND NOT-MAG AND NOT-CD93	0.974359	0.95	1
CLECL1 AND NOT-MAG AND NOT-PAQR8	0.962025	0.926829	1
CLECL1 AND NOT-GALR2 AND NOT-STEAP4	0.974359	0.95	1
CLECL1 AND NOT-ATP1B4 AND NOT-CXCL16	0.973684	0.973684	0.973684
CLECL1 AND NOT-GUCY2D AND NOT-CD93	0.973684	0.973684	0.973684
CLECL1 AND NOT-CHRNA4 AND NOT-EBP	0.972973	1	0.947368
CLECL1 AND NOT-GALR2 AND NOT-DYSF	0.962025	0.926829	1
CLECL1 AND NOT-GALR2 AND NOT-CD93	0.974359	0.95	1
CLECL1 AND NOT-ATP1B4 AND NOT-CD93	0.974359	0.95	1
CLECL1 AND NOT-GALR2 AND NOT-CD163	0.962025	0.926829	1
CLECL1 AND NOT-CHRNA4 AND NOT-PAQR7	0.986667	1	0.973684
CLECL1 AND NOT-ATP1B4 AND NOT-CD93	0.974359	0.95	1
CLECL1 AND NOT-GRID2 AND NOT-CXCL16	0.973684	0.973684	0.973684
CLECL1 AND NOT-GRID2 AND NOT-CD163	0.962025	0.926829	1
CLECL1 AND NOT-GALR2 AND NOT-DYSF	0.962025	0.926829	1
CLECL1 AND NOT-GALR2 AND NOT-CD93	0.974359	0.95	1
CLECL1 AND NOT-MAG AND NOT-PAQR8	0.962025	0.926829	1
CLECL1 AND NOT-MAG AND NOT-STEAP4	0.974359	0.95	1
CLECL1 AND NOT-MAG AND NOT-CXCL16	0.973684	0.973684	0.973684
CLECL1 AND NOT-GPR37L1 AND NOT-ZDHHC2	0.987013	0.974359	1
CLECL1 AND NOT-GALR2 AND NOT-CXCL16	0.973684	0.973684	0.973684
CLECL1 AND NOT-GUCY2D AND NOT-PAQR8	0.973684	0.973684	0.973684
CLECL1 AND NOT-GALR2 AND NOT-PAQR8	0.974359	0.95	1
CLECL1 AND NOT-GRID2 AND NOT-CD93	0.974359	0.95	1
CLECL1 AND NOT-GALR2 AND NOT-PAQR8	0.974359	0.95	1
CLECL1 AND NOT-MAG AND NOT-CD93	0.974359	0.95	1
CLECL1 AND NOT-ATP1B4 AND NOT-CD36	0.961039	0.948718	0.973684
CLECL1 AND NOT-MAG AND NOT-CD36	0.961039	0.948718	0.973684
CLECL1 AND NOT-LPPR3 AND NOT-CD93	0.961039	0.948718	0.973684
CLECL1 AND NOT-CD36 AND NOT-GPR61	0.961039	0.948718	0.973684
CLECL1 AND NOT-GRID2 AND NOT-CD36	0.961039	0.948718	0.973684
CLECL1 AND NOT-CRB2 AND NOT-CD93	0.961039	0.948718	0.973684
CLECL1 AND NOT-CALHM3 AND NOT-FMNL1	0.961039	0.948718	0.973684
CLECL1 AND NOT-GALR2 AND NOT-CD36	0.961039	0.948718	0.973684
CLECL1 AND NOT-CHRNA4 AND NOT-ACSL6	1	1	1
CLECL1 AND NOT-CD207 AND NOT-CD93	0.974359	0.95	1
CLECL1 AND NOT-GGTLC1 AND NOT-CD93	0.961039	0.948718	0.973684
CLECL1 AND NOT-CHRNA4 AND NOT-PAQR8	0.987013	0.974359	1
CLECL1 AND NOT-TRPM3 AND NOT-CD93	0.961039	0.948718	0.973684
CLECL1 AND NOT-CHRNA4 AND NOT-PAQR8	0.987013	0.974359	1
CLECL1 AND NOT-SLC22A6 AND NOT-CD93	0.961039	0.948718	0.973684
CLECL1 AND NOT-SLC22A13 AND NOT-CD93	0.961039	0.948718	0.973684
CLECL1 AND NOT-GABRD AND NOT-CD93	0.986667	1	0.973684
CLECL1 AND NOT-GABRA4 AND NOT-CD93	0.974359	0.95	1
CLECL1 AND NOT-CD207 AND NOT-PAQR8	0.962025	0.926829	1
CLECL1 AND NOT-GABRD AND NOT-PAQR8	0.973684	0.973684	0.973684
CLECL1 AND NOT-GRM4 AND NOT-CD93	0.974359	0.95	1
CLECL1 AND NOT-CLDN16 AND NOT-CD93	1	1	1
CLECL1 AND NOT-AJAP1 AND NOT-CD93	0.973684	0.973684	0.973684
CLECL1 AND NOT-CLDN16 AND NOT-PAQR8	0.987013	0.974359	1
CLECL1 AND NOT-AJAP1 AND NOT-PAQR8	0.961039	0.948718	0.973684
CLECL1 AND NOT-CALY AND NOT-PAQR8	0.96	0.972973	0.947368
CLECL1 AND NOT-OR3A2 AND NOT-PAQR8	0.96	0.972973	0.947368
COMPLEX-SLC9B1/CLECL1/MLC1	0.96	0.972973	0.947368
CLECL1 AND NOT-CALY AND NOT-CD93	0.96	0.972973	0.947368
CLECL1 AND NOT-SLC5A8 AND NOT-CD93	0.96	0.972973	0.947368
CLECL1 AND NOT-OR3A2 AND NOT-CD93	0.96	0.972973	0.947368
CLECL1 AND NOT-CHRNA4 AND NOT-IFNAR2	0.958904	1	0.921053
CLECL1 AND NOT-GALR2 AND NOT-PAQR7	0.961039	0.948718	0.973684
CLECL1 AND NOT-GPR61 AND NOT-CD93	0.974359	0.95	1
CLECL1 AND NOT-CD36 AND NOT-HTR2C	0.961039	0.948718	0.973684
CLECL1 AND NOT-CD36 AND NOT-GPR37L1	0.961039	0.948718	0.973684
CLECL1 AND NOT-CD36 AND NOT-GABRB1	0.961039	0.948718	0.973684
CLECL1 AND NOT-CD36 AND NOT-SLC9B1	0.961039	0.948718	0.973684
CLECL1 AND NOT-GALR2 AND OR1Q1	0.962025	0.926829	1
CLECL1 AND NOT-CHRNA4 AND NOT-BTN3A3	1	1	1
CLECL1 AND NOT-MAG AND NOT-PAQR7	0.961039	0.948718	0.973684
CLECL1 AND NOT-CD36 AND NOT-SLC13A1	0.961039	0.948718	0.973684
CLECL1 AND NOT-SLC9B1 AND NOT-SLC31A1	0.974359	0.95	1
CLECL1 AND NOT-CALHM3 AND NOT-CD93	0.973684	0.973684	0.973684
CLECL1 AND NOT-CD36 AND NOT-KCNK16	0.961039	0.948718	0.973684
CLECL1 AND NOT-CD36 AND NOT-NPHS2	0.961039	0.948718	0.973684
CLECL1 AND NOT-CD36 AND NOT-GABRG2	0.961039	0.948718	0.973684
CLECL1 AND NOT-CD36 AND NOT-GABRG1	0.961039	0.948718	0.973684

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
SLCO1B1 AND TLCD1 AND NOT-SLC17A2	1	1	1
SLCO1B1 AND NOT-KCNN2 AND CLDN15	1	1	1
SLCO1B1 AND NOT-KCNN2 AND SEZ6	1	1	1
SLC2A2 AND SLC26A6 AND NOT-SLC13A2	1	1	1
SLC2A2 AND SLC26A6 AND ENPP1	1	1	1
SLC2A2 AND SLC26A6 AND PHLDB2	1	1	1
SLC2A2 AND SLC26A6 AND SLC13A5	1	1	1
SLC2A2 AND SLC26A6 AND NOT-FUT3	1	1	1
SLC2A2 AND SLC26A6 AND DIO1	1	1	1
SLC2A2 AND SLC26A6 AND SLC23A2	1	1	1
SLCO1B1 AND SMPD2 AND NOT-IL2RG	1	1	1
SLCO1B1 AND SMPD2 AND NOT-FCGR3B	1	1	1
ABCG5 AND FGFRL1 AND NOT-EPHA2	1	1	1
SLC2A2 AND SLC26A6 AND NOT-CD58	1	1	1
SLC2A2 AND SLC26A6 AND SLC43A1	1	1	1
SLC2A2 AND SLC26A6 AND NOT-ATP8A1	1	1	1
SLC2A2 AND SLC26A6 AND NOT-MUC12	1	1	1
SLC2A2 AND SLC26A6 AND NOT-MUC17	1	1	1
SLCO1B1 AND SMPD2 AND NOT-NGFR	1	1	1
SLC2A2 AND SLC26A6 AND NOT-SLC16A5	1	1	1
SLCO1B1 AND NOT-KCNN2 AND PCDHB1	1	1	1
SLCO1B1 AND FGFRL1 AND NOT-SCARA5	1	1	1
SLC2A2 AND SLC26A6 AND NOT-CD82	1	1	1
SLC2A2 AND SLC26A6 AND SLC22A9	1	1	1
SLC2A2 AND SLC26A6 AND SLC17A3	1	1	1
SLCO1B1 AND NOT-KCNN2 AND SLCO1A2	1	1	1
SLCO1B1 AND NOT-MARCO AND ABCG4	1	1	1
SLCO1B1 AND HM13 AND NOT-CD163	1	1	1
SLC2A2 AND SLC26A6 AND NOT-SLC5A11	1	1	1
SLC2A2 AND SLC26A6 AND CLDN14	1	1	1
SLCO1B1 AND SMPD2 AND NOT-CXCR2	1	1	1
SLCO1B1 AND SMPD2 AND NOT-GPR171	1	1	1
SLCO1B1 AND SMPD2 AND NOT-LYVE1	1	1	1
SLCO1B1 AND TLCD1 AND NOT-SLC17A2	1	1	1
SLCO1B1 AND NOT-MARCO AND HTR5A	1	1	1
SLC2A2 AND SLC26A6 AND NOT-CDH17	1	1	1
SLC2A2 AND SLC26A6 AND NOT-MUC17	1	1	1
SLC2A2 AND SLC26A6 AND NOT-LCT	1	1	1
SLC2A2 AND SLC26A6 AND NOT-SLC9A1	1	1	1
SLC2A2 AND SLC26A6 AND NOT-SLC10A2	1	1	1
SLC2A2 AND SLC26A6 AND NOT-PTPRR	1	1	1
SLCO1B1 AND HM13 AND NOT-KCNN2	1	1	1
SLCO1B1 AND NOT-MARCO AND DST	1	1	1
SLCO1B1 AND NOT-MARCO AND PVRL2	1	1	1
SLCO1B1 AND NOT-MARCO AND DAGLB	1	1	1
SLCO1B1 AND NOT-MARCO AND NCSTN	1	1	1
SLC43A1 AND NOT-ENTPD3 AND SLC26A6	1	1	1
ABCG5 AND FGFRL1 AND NOT-MARCO	1	1	1
ABCG5 AND FGFRL1 AND NOT-CLEC4M	1	1	1
SLCO1B1 AND NOT-MARCO AND FAIM2	1	1	1
SLCO1B1 AND NOT-MARCO AND RHOT2	1	1	1
SLC43A1 AND NOT-PARM1 AND SLC26A6	1	1	1
ABCG5 AND FGFRL1 AND NOT-FNDC4	1	1	1
SLCO1B1 AND NOT-MARCO AND HM13	1	1	1
SLCO1B1 AND NOT-MARCO AND ATP9B	1	1	1
SLCO1B1 AND NOT-MARCO AND SLC26A11	1	1	1
SLC43A1 AND NOT-SLC4A3 AND SLC26A6	1	1	1
SLCO1B1 AND NOT-MARCO AND ANO10	1	1	1
SLCO1B1 AND NOT-MARCO AND CLPTM1	1	1	1
SLCO1B1 AND NOT-MARCO AND MERTK	1	1	1
SLCO1B1 AND NOT-MARCO AND BSG	1	1	1
SLCO1B1 AND NOT-MARCO AND HM13	1	1	1
SLCO1B1 AND NOT-MARCO AND SLC10A3	1	1	1
SLCO1B1 AND NOT-MARCO AND STX6	1	1	1
SLCO1B1 AND NOT-MARCO AND ORAI1	1	1	1
SLCO1B1 AND NOT-MARCO AND ADAM15	1	1	1
SLCO1B1 AND NOT-MARCO AND CHRNA6	1	1	1
SLCO1B1 AND NOT-MARCO AND RYK	1	1	1
ABCG5 AND FGFRL1 AND NOT-PRIMA1	1	1	1
ABCG5 AND FGFRL1 AND NOT-CLEC4G	1	1	1
SLCO1B1 AND NOT-MARCO AND ADIPOR1	1	1	1
SLCO1B1 AND NOT-MARCO AND ASPH	1	1	1
SLCO1B1 AND NOT-MARCO AND CD46	1	1	1
SLCO1B1 AND NOT-MARCO AND SLC38A6	1	1	1
SLCO1B1 AND NOT-MARCO AND CSF1	1	1	1
SLCO1B1 AND HM13 AND NOT-ADRA1B	1	1	1
SLCO1B1 AND NOT-MARCO AND OR51M1	1	1	1

Antigen Pair Logic	F1	Precision	Recall
CLECL1 AND NOT-CD36 AND NOT-OR8B2	0.961039	0.948718	0.973684
CLECL1 AND NOT-CHRNA4 AND NOT-SLC12A9	0.986667	1	0.973684
CLECL1 AND NOT-CHRNA4 AND TMEFF2	0.973684	0.973684	0.973684
CLECL1 AND NOT-CD36 AND NOT-LRRC52	0.973684	0.973684	0.973684
CLECL1 AND NOT-CD36 AND NOT-GABRA5	0.961039	0.948718	0.973684
CLECL1 AND NOT-CD36 AND NOT-ATP13A5	0.961039	0.948718	0.973684
CLECL1 AND NOT-CD36 AND NOT-GABRA3	0.961039	0.948718	0.973684
CLECL1 AND NOT-CD36 AND NOT-GRM6	0.961039	0.948718	0.973684
CLECL1 AND NOT-CD36 AND NOT-SLC39A12	0.961039	0.948718	0.973684
CLECL1 AND NOT-CD36 AND LAPTM5	0.961039	0.948718	0.973684
CLECL1 AND NOT-GRID2 AND NOT-PAQR7	0.961039	0.948718	0.973684
CLECL1 AND NOT-CD36 AND NOT-GABRA2	0.961039	0.948718	0.973684
MS4A1 AND SLC22A18 AND NOT-SLC46A2	0.973684	0.973684	0.973684
CLECL1 AND NOT-CHRNA4 AND NOT-CD33	0.987013	0.974359	1
MS4A1 AND SLC22A18 AND NOT-PTGDR2	0.974359	0.95	1
CLECL1 AND NOT-CHRNA4 AND NOT-ALDH1A1	0.974359	0.95	1
CLECL1 AND NOT-CHRNA4 AND NOT-CD160	1	1	1
CLECL1 AND NOT-DNAJB8 AND NOT-CD93	0.974359	0.95	1
MS4A1 AND SLC22A18 AND NOT-PTGDR2	0.974359	0.95	1
MS4A1 AND SLC22A18 AND NOT-FPR2	0.973684	0.973684	0.973684
CLECL1 AND NOT-ITGB3 AND SEMA4D	0.987013	0.974359	1
MS4A1 AND HCN2 AND NOT-P2RY2	0.962025	0.926829	1
MS4A1 AND MARVELD2 AND NOT-ANPEP	0.962025	0.926829	1
CLECL1 AND NOT-DNAJB8 AND NOT-PAQR8	0.962025	0.926829	1
MS4A1 AND STRA6 AND NOT-ANPEP	0.962025	0.926829	1
MS4A1 AND ADCY10 AND NOT-SLC46A2	0.986667	1	0.973684
CLECL1 AND NOT-MAG AND NOT-CD160	0.974359	0.95	1
CLECL1 AND NOT-GRID2 AND NOT-CD33	0.962025	0.926829	1
CLECL1 AND NOT-CLDN2 AND NOT-CD93	0.974359	0.95	1
CLECL1 AND NOT-CLDN2 AND NOT-STEAP4	0.974359	0.95	1
CLECL1 AND NOT-ATP1B4 AND NOT-CD160	0.974359	0.95	1
CLECL1 AND NOT-CLDN2 AND NOT-CXCL16	0.973684	0.973684	0.973684
CLECL1 AND NOT-GALR2 AND NOT-CD33	0.974359	0.95	1
CLECL1 AND NOT-CHRNA4 AND TNFRSF13C	0.962025	0.926829	1
CLECL1 AND NOT-CLDN2 AND NOT-CD93	0.974359	0.95	1
CLECL1 AND NOT-CHRNA4 AND NOT-HSPA5	0.972973	1	0.947368
CLECL1 AND NOT-GRID2 AND NOT-CD160	0.974359	0.95	1
MS4A1 AND ADCY10 AND NOT-P2RY2	0.987013	0.974359	1
CLECL1 AND NOT-ATP1B4 AND NOT-CD33	0.962025	0.926829	1
CLECL1 AND NOT-MAG AND NOT-CD33	0.962025	0.926829	1
CLECL1 AND NOT-GALR2 AND NOT-CD160	0.974359	0.95	1
CLECL1 AND NOT-CLDN2 AND NOT-PAQR8	0.962025	0.926829	1
CLECL1 AND NOT-GALR2 AND NOT-ALDH1A1	0.962025	0.926829	1
CLECL1 AND NOT-CLDN2 AND NOT-CD36	0.961039	0.948718	0.973684
MS4A1 AND SLC22A18 AND NOT-ASGR1	0.961039	0.948718	0.973684
MS4A1 AND STRA6 AND NOT-CD36	0.961039	0.948718	0.973684
MS4A1 AND SLC22A18 AND NOT-SLC16A10	0.961039	0.948718	0.973684
MS4A1 AND HCN2 AND NOT-IL12RB2	0.961039	0.948718	0.973684
MS4A1 AND JPH3 AND NOT-P2RY2	0.961039	0.948718	0.973684
MS4A1 AND SLC22A18 AND NOT-PTPRA	0.961039	0.948718	0.973684
CLECL1 AND NOT-SST AND NOT-CD93	0.974359	0.95	1
CLECL1 AND NOT-SSTR5 AND NOT-CD93	0.974359	0.95	1
MS4A1 AND JPH3 AND NOT-PTGDR2	0.961039	0.948718	0.973684
MS4A1 AND HCN2 AND NOT-PTGDR2	0.974359	0.95	1
MS4A1 AND JPH3 AND NOT-TGOLN2	0.961039	0.948718	0.973684
MS4A1 AND NOT-MS4A2 AND HRH3	0.962025	0.926829	1
MS4A1 AND NOT-MS4A2 AND ADCY10	0.974359	0.95	1
COMPLEX-CLECL1/ITGB3/LRRC52	0.987013	0.974359	1
CLECL1 AND NOT-SSTR5 AND NOT-PAQR8	0.962025	0.926829	1
CLECL1 AND NOT-SST AND NOT-PAQR8	0.962025	0.926829	1
COMPLEX-CLECL1/GABRA2/ITGB3	0.987013	0.974359	1
MS4A1 AND STRA6 AND NOT-SLC28A1	0.962025	0.926829	1
CLECL1 AND NOT-ALK AND NOT-CD93	0.974359	0.95	1
COMPLEX-CLECL1/OR8B2/ITGB3	0.987013	0.974359	1
COMPLEX-CLECL1/ITGB3/LAPTM5	0.987013	0.974359	1
MS4A1 AND HCN2 AND NOT-TGOLN2	0.962025	0.926829	1
COMPLEX-CLECL1/ABHD3/ITGB3	0.987013	0.974359	1
CLECL1 AND NOT-ULBP3 AND NOT-CD93	0.961039	0.948718	0.973684
COMPLEX-CLECL1/MPL/MST1R	0.962025	0.926829	1
MS4A1 AND SLC22A18 AND NOT-S1PR5	0.961039	0.948718	0.973684
MS4A1 AND STRA6 AND NOT-MPL	0.974359	0.95	1
CLECL1 AND NOT-MAG AND NOT-HSPA5	0.96	0.972973	0.947368
CLECL1 AND NOT-ITGB3 AND GALR2	0.96	0.972973	0.947368
MS4A1 AND JPH3 AND NOT-IL12RB2	0.96	0.972973	0.947368
CLECL1 AND NOT-GALR2 AND NOT-HSPA5	0.96	0.972973	0.947368
CLECL1 AND NOT-ATP1B4 AND NOT-HSPA5	0.96	0.972973	0.947368
MS4A1 AND STRA6 AND NOT-PTGER4	0.974359	0.95	1

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
ABCG5 AND FGFRL1 AND NOT-PLP1	1	1	1
SLCO1B1 AND NOT-MARCO AND RHOT2	1	1	1
SLCO1B1 AND NOT-MARCO AND CSF1	1	1	1
SLCO1B1 AND NOT-MARCO AND PTPRA	1	1	1
SLCO1B1 AND NOT-MARCO AND ATP13A1	1	1	1
ABCG5 AND NOTCH4 AND NOT-MARCO	1	1	1
SLCO1B1 AND NOT-MARCO AND SYNDIG1	1	1	1
SLCO1B1 AND NOT-MARCO AND CLPTM1	1	1	1
ABCG5 AND FGFRL1 AND NOT-PLP1	1	1	1
SLCO1B1 AND NOT-MARCO AND ABHD12	1	1	1
SLCO1B1 AND HM13 AND NOT-KCNN2	1	1	1
SLCO1B1 AND NOT-MARCO AND ATP13A1	1	1	1
ABCG5 AND NOTCH4 AND NOT-MARCO	1	1	1
SLC43A1 AND NOT-KIAA1324 AND SLC26A6	1	1	1
SLCO1B1 AND NOT-MARCO AND ITGAX	1	1	1
SLCO1B1 AND NOT-MARCO AND SLC41A3	1	1	1
SLCO1B1 AND NOT-MARCO AND CD151	1	1	1
SLCO1B1 AND NOT-MARCO AND ORAI1	1	1	1
SLCO1B1 AND NOT-MARCO AND FAM73A	1	1	1
SLCO1B1 AND NOT-MARCO AND OR51M1	1	1	1
ABCG5 AND FGFRL1 AND NOT-CLEC1B	1	1	1
SLCO1B1 AND NOT-MARCO AND ABCC1	1	1	1
ABCG5 AND FGFRL1 AND NOT-MARCO	1	1	1
ABCG5 AND FGFRL1 AND NOT-PRIMA1	1	1	1
SLCO1B1 AND NOT-MARCO AND ABCC1	1	1	1
ABCG5 AND HM13 AND NOT-SLC5A1	1	1	1
ABCG5 AND HM13 AND NOT-SPINT1	1	1	1
ABCG5 AND HM13 AND NOT-TNFRSF11A	1	1	1
ABCG5 AND HM13 AND NOT-ADTRP	1	1	1
ABCG5 AND HM13 AND NOT-SLC28A2	1	1	1
ABCG5 AND HM13 AND NOT-SLC6A19	1	1	1
ABCG5 AND HM13 AND NOT-UGT8	1	1	1
ABCG5 AND HM13 AND NOT-ADTRP	1	1	1
ABCG5 AND HM13 AND NOT-ATP2C2	1	1	1
ABCG5 AND HM13 AND NOT-SLC5A1	1	1	1
ABCG5 AND HM13 AND NOT-SLC6A20	1	1	1
ABCG5 AND HM13 AND NOT-TNFRSF11A	1	1	1
ABCG5 AND HM13 AND NOT-KCNJ3	1	1	1
ABCG5 AND HM13 AND NOT-HEPH	1	1	1
ABCG5 AND HM13 AND NOT-GPRC5A	1	1	1
ABCG5 AND HM13 AND NOT-TSPAN1	1	1	1
ABCG5 AND HM13 AND NOT-SI	1	1	1
ABCG5 AND HM13 AND NOT-SLC26A3	1	1	1
ABCG5 AND HM13 AND NOT-B3GNT3	1	1	1
ABCG5 AND HM13 AND NOT-KCNJ3	1	1	1
ABCG5 AND HM13 AND NOT-GPR160	1	1	1
ABCG5 AND HM13 AND NOT-KIAA1324	1	1	1
ABCG5 AND HM13 AND NOT-GDPD2	1	1	1
ABCG5 AND HM13 AND NOT-HEPH	1	1	1
ABCG5 AND HM13 AND NOT-ATP10B	1	1	1
ABCG5 AND HM13 AND NOT-SLC28A2	1	1	1
ABCG5 AND HM13 AND NOT-KIAA1324	1	1	1
ABCG5 AND HM13 AND NOT-MGAM	1	1	1
ABCG5 AND HM13 AND NOT-ACE	1	1	1
ABCG5 AND HM13 AND NOT-SPINT1	1	1	1
ABCG5 AND HM13 AND NOT-TSPAN1	1	1	1
ABCG5 AND HM13 AND NOT-MGAM	1	1	1
ABCG5 AND HM13 AND NOT-SLC6A20	1	1	1
ABCG5 AND HM13 AND NOT-SLC6A19	1	1	1
ABCG5 AND HM13 AND NOT-GPR35	1	1	1
ABCG5 AND HM13 AND NOT-KCNQ1	1	1	1
ABCG5 AND HM13 AND NOT-MEP1B	1	1	1
ABCG5 AND HM13 AND NOT-ATP10B	1	1	1
SLCO1B1 AND NOT-MARCO AND CLDN8	1	1	1
ABCG5 AND HM13 AND NOT-SST	1	1	1
ABCG5 AND HM13 AND NOT-DPEP1	1	1	1
ABCG5 AND HM13 AND NOT-GPA33	1	1	1
ABCG5 AND HM13 AND NOT-GUCY2C	1	1	1
ABCG5 AND HM13 AND NOT-GPA33	1	1	1
ABCG5 AND HM13 AND NOT-HHLA2	1	1	1
ABCG5 AND HM13 AND NOT-CEACAM5	1	1	1
ABCG5 AND HM13 AND NOT-GUCY2C	1	1	1
ABCG5 AND HM13 AND NOT-MST1R	1	1	1
ABCG5 AND HM13 AND NOT-MST1R	1	1	1
ABCG5 AND HM13 AND NOT-CEACAM6	1	1	1
SLCO1B1 AND HM13 AND NOT-CLDN5	1	1	1
SLC43A1 AND NOT-IL20RA AND SLC26A6	1	1	1

Antigen Pair Logic	F1	Precision	Recall
MS4A1 AND MARVELD2 AND NOT-PTGER4	0.974359	0.95	1
MS4A1 AND SLC22A18 AND NOT-ACPP	0.961039	0.948718	0.973684
MS4A1 AND SLC22A18 AND NOT-PSEN1	0.974359	0.95	1
CLECL1 AND NOT-SSTR1 AND NOT-CXCL16	0.973684	0.973684	0.973684
CLECL1 AND NOT-GPR61 AND NOT-CD160	0.974359	0.95	1
MS4A1 AND NOT-TRPV2 AND ADCY10	0.962025	0.926829	1
MS4A1 AND ADCY10 AND NOT-SLC16A10	0.973684	0.973684	0.973684
MS4A1 AND NOT-TGOLN2 AND ADCY10	0.987013	0.974359	1
MS4A1 AND MARVELD2 AND NOT-CD36	0.961039	0.948718	0.973684
MS4A1 AND SLC22A18 AND NOT-DHRS3	0.961039	0.948718	0.973684
MS4A1 AND ADCY10 AND NOT-PTGDR2	0.987013	0.974359	1
CLECL1 AND NOT-MST1R AND NOT-CD163	0.974359	0.95	1
MS4A1 AND ADCY10 AND NOT-CD300C	0.973684	0.973684	0.973684
CLECL1 AND NOT-GPR61 AND NOT-CD33	0.962025	0.926829	1
CLECL1 AND NOT-GALR2 AND TNFRSF13C	0.974359	0.95	1
MS4A1 AND SLC22A18 AND NOT-PAG1	0.974359	0.95	1
MS4A1 AND ADCY10 AND NOT-IL12RB2	0.987013	0.974359	1
CLECL1 AND NOT-SSTR1 AND NOT-STEAP4	0.974359	0.95	1
MS4A1 AND SLC22A18 AND NOT-PTGER2	0.961039	0.948718	0.973684
MS4A1 AND SLC22A18 AND NOT-PSEN1	0.974359	0.95	1
CLECL1 AND NOT-MAG AND TNFRSF13C	0.962025	0.926829	1
MS4A1 AND NOT-TGOLN2 AND ADCY10	0.987013	0.974359	1
CLECL1 AND NOT-GRID2 AND TNFRSF13C	0.962025	0.926829	1
MS4A1 AND ADCY10 AND NOT-HM13	0.962025	0.926829	1
MS4A1 AND ADCY10 AND NOT-ASGR1	0.973684	0.973684	0.973684
CLECL1 AND NOT-SSTR1 AND NOT-CD93	0.974359	0.95	1
MS4A1 AND SLC30A8 AND NOT-IL12RB2	0.962025	0.926829	1
CLECL1 AND NOT-CLDN2 AND NOT-BTN3A3	0.974359	0.95	1
CLECL1 AND NOT-CD36 AND NOT-TPBG	0.961039	0.948718	0.973684
MS4A1 AND SLC30A8 AND NOT-SLC46A2	0.961039	0.948718	0.973684
CLECL1 AND NOT-SSTR1 AND NOT-PAQR8	0.962025	0.926829	1
CLECL1 AND NOT-CLDN2 AND NOT-PAQR7	0.961039	0.948718	0.973684
CLECL1 AND NOT-SSTR1 AND NOT-PAQR8	0.962025	0.926829	1
CD19 AND CLECL1 AND NOT-SLC2A5	0.914286	1	0.842105
COMPLEX-CLECL1/SLC22A3/CD19	0.914286	1	0.842105
CD19 AND CLECL1 AND NOT-TAS2R5	0.914286	1	0.842105
CD19 AND CLECL1 AND NOT-CTNS	0.914286	1	0.842105
CD19 AND CLECL1 AND NOT-LNPEP	0.914286	1	0.842105
COMPLEX-CLECL1/EQTN/CD19	0.914286	1	0.842105
CD19 AND CLECL1 AND NOT-SLC16A3	0.914286	1	0.842105
COMPLEX-CLECL1/SLCO2A1/CD19	0.914286	1	0.842105
CD19 AND CLECL1 AND NOT-TGFBR2	0.914286	1	0.842105
COMPLEX-SLC26A7/CLECL1/CD19	0.914286	1	0.842105
COMPLEX-CLECL1/GPRC5B/CD19	0.914286	1	0.842105
CD19 AND LRRC32 AND PPAP2A	0.914286	1	0.842105
CD19 AND CLECL1 AND NOT-TREML2	0.914286	1	0.842105
CD19 AND CLECL1 AND NOT-SLC4A7	0.914286	1	0.842105
CD19 AND PRPH2 AND NOT-SLC2A12	0.914286	1	0.842105
CD19 AND KIR2DL4 AND NOT-SLC2A12	0.914286	1	0.842105
CD19 AND SLC26A6 AND NOT-SLC2A12	0.914286	1	0.842105
CD19 AND OR51M1 AND NOT-SLC2A12	0.914286	1	0.842105
CD19 AND HRH1 AND NOT-SLC2A12	0.914286	1	0.842105
CD19 AND CLECL1 AND NOT-MMP15	0.914286	1	0.842105
CD19 AND CLECL1 AND NOT-ABCG2	0.914286	1	0.842105
COMPLEX-CLECL1/OR7A5/CD19	0.914286	1	0.842105
CD19 AND OR10A4 AND NOT-SLC2A12	0.914286	1	0.842105
CD19 AND CLECL1 AND NOT-TPCN1	0.914286	1	0.842105
CD19 AND STRA6 AND NOT-ADCY2	0.914286	1	0.842105
CD19 AND OR51B4 AND SLC25A3	0.914286	1	0.842105
CD19 AND LRP4 AND LRRC8B	0.914286	1	0.842105
CD19 AND EDNRA AND NOT-SLC2A12	0.914286	1	0.842105
CD19 AND BTN1A1 AND SLC25A3	0.914286	1	0.842105
CD19 AND CLECL1 AND NOT-LRP12	0.914286	1	0.842105
CD19 AND SLC5A9 AND NOT-SLC2A12	0.914286	1	0.842105
COMPLEX-CLECL1/SEMA5A/CD19	0.914286	1	0.842105
CD19 AND NOT-TRPV2 AND CLECL1	0.914286	1	0.842105
CD19 AND CLECL1 AND NOT-TGOLN2	0.914286	1	0.842105
CD19 AND CLECL1 AND NOT-ADAM8	0.914286	1	0.842105
COMPLEX-CLECL1/LY6G6D/CD19	0.914286	1	0.842105
CD19 AND CLECL1 AND NOT-IL1RAPL1	0.914286	1	0.842105
CD19 AND CLECL1 AND NOT-KCNH6	0.914286	1	0.842105
COMPLEX-RXFP2/CLECL1/CD19	0.914286	1	0.842105
CD19 AND CLECL1 AND NOT-SMPD1	0.914286	1	0.842105
CD19 AND CLECL1 AND NOT-CYBB	0.914286	1	0.842105
CD19 AND NOT-CYP4F12 AND NOT-SLC2A12	0.914286	1	0.842105
CD19 AND SLC22A7 AND NOT-SLC2A12	0.914286	1	0.842105
CD19 AND CLECL1 AND NOT-TCTN3	0.914286	1	0.842105

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
SLCO1B1 AND HM13 AND NOT-STEAP1	1	1	1
SLCO1B1 AND HM13 AND NOT-MAGEA4	1	1	1
ABCG5 AND FGFRL1 AND NOT-FOLH1	1	1	1
SLCO1B1 AND EPHB2 AND NOT-HMOX1	1	1	1
SLC2A2 AND SLC26A6 AND NOT-CLDN7	1	1	1
SLC2A2 AND SLC26A6 AND NOT-GPA33	1	1	1
SLCO1B1 AND NOT-MARCO AND CD22	1	1	1
SLC2A2 AND SLC26A6 AND CD70	1	1	1
SLC2A2 AND SLC26A6 AND NOT-DPEP1	1	1	1
SLC2A2 AND SLC26A6 AND NOT-CLDN3	1	1	1
SLC2A2 AND SLC26A6 AND NOT-GUCY2C	1	1	1
SLC2A2 AND SLC26A6 AND NOT-MUC13	1	1	1
SLC2A2 AND SLC26A6 AND NOT-CLDN23	1	1	1
SLC2A2 AND SLC26A6 AND NOT-PROM1	1	1	1
ABCG5 AND FGFRL1 AND NOT-IL13RA2	1	1	1
SLCO1B1 AND EPHB2 AND NOT-STEAP3	1	1	1
SLCO1B1 AND TLCD1 AND NOT-CLDN5	1	1	1
SLC2A2 AND SLC26A6 AND CLDN1	1	1	1
SLC2A2 AND SLC26A6 AND NOT-CLDN4	1	1	1
SLCO1B1 AND EPHB2 AND NOT-P2RY13	1	1	1
SLC2A2 AND SLC26A6 AND NOT-ENPP3	1	1	1
SLC2A2 AND SLC26A6 AND NOT-MST1R	1	1	1
SLC2A2 AND SLC26A6 AND NOT-SST	1	1	1
SLCO1B1 AND NOT-MARCO AND SEMA5B	1	1	1
SLC2A2 AND SLC26A6 AND KDR	1	1	1
SLC2A2 AND SLC26A6 AND NOT-CEACAM5	1	1	1
SLC2A2 AND SLC26A6 AND NOT-CEACAM6	1	1	1
SLC2A2 AND SLC26A6 AND CD70	1	1	1
SLCO1B1 AND NOT-MARCO AND MST1R	1	1	1
SLC2A2 AND SLC26A6 AND NOT-CD160	1	1	1
SLC2A2 AND SLC26A6 AND NOT-HHLA2	1	1	1
SLCO1B1 AND TLCD1 AND NOT-STEAP1	1	1	1
ABCG5 AND SLC7A11 AND NOT-CEACAM5	1	1	1
ABCG5 AND HM13 AND NOT-MUC13	1	1	1
ABCG5 AND SLC7A11 AND NOT-CEACAM6	1	1	1
ABCG5 AND FGFRL1 AND NOT-BMPR1B	1	1	1
SLC43A1 AND NOT-IL20RA AND NOT-MARCO	1	1	1
SLCO1B1 AND HM13 AND NOT-KDR	1	1	1
ABCG5 AND FGFRL1 AND NOT-CD19	1	1	1
ABCG5 AND FGFRL1 AND NOT-SSTR3	1	1	1
ABCG5 AND FGFRL1 AND NOT-SSTR4	1	1	1
SLCO1B1 AND HM13 AND NOT-TNFRSF13C	1	1	1
SLCO1B1 AND HM13 AND NOT-ANXA1	1	1	1
ABCG5 AND FGFRL1 AND NOT-L1CAM	1	1	1
SLCO1B1 AND HM13 AND NOT-WT1	1	1	1
SLCO1B1 AND HM13 AND NOT-ITGB6	1	1	1
SLCO1B1 AND HM13 AND NOT-CTAG2	1	1	1
ABCG5 AND FGFRL1 AND NOT-THY1	1	1	1
ABCG5 AND HM13 AND NOT-ITGB6	1	1	1
ABCG5 AND FGFRL1 AND NOT-MS4A1	1	1	1
SLCO1B1 AND HM13 AND NOT-SLC39A6	1	1	1
SLCO1B1 AND HM13 AND NOT-FLOT2	1	1	1
SLCO1B1 AND HM13 AND NOT-PMEL	1	1	1
SLCO1B1 AND HM13 AND NOT-CD160	1	1	1
SLCO1B1 AND HM13 AND NOT-CLDN3	1	1	1
ABCG5 AND FGFRL1 AND NOT-ITGB3	1	1	1
ABCG5 AND FGFRL1 AND NOT-L1CAM	1	1	1
SLCO1B1 AND HM13 AND NOT-CXCR5	1	1	1
ABCG5 AND HM13 AND NOT-PROM1	1	1	1
SLCO1B1 AND HM13 AND NOT-THY1	1	1	1
SLCO1B1 AND HM13 AND NOT-VTCN1	1	1	1
ABCG5 AND FGFRL1 AND NOT-TNFRSF8	1	1	1
SLC43A1 AND NOT-MUC1 AND SLC26A6	1	1	1
SLC43A1 AND NOT-MUC1 AND NOT-CLEC4G	1	1	1
SLC43A1 AND NOT-MUC1 AND NOT-MARCO	1	1	1
SLC43A1 AND NOT-MUC1 AND NOT-MARCO	1	1	1
SLC43A1 AND NOT-MUC1 AND HM13	1	1	1
SLCO1B1 AND SLC26A6 AND NOT-MSLN	0.923077	0.857143	1
SLCO1B1 AND SLC26A6 AND NOT-MUC1	0.923077	0.857143	1
SLC43A1 AND CELSR3 AND NOT-MUC1	1	1	1
SLC43A1 AND NOT-MUC1 AND CD200	1	1	1
SLC43A1 AND CELSR3 AND NOT-MUC1	1	1	1
SLCO1B1 AND C6orf89 AND NOT-MSLN	1	1	1
ABCG5 AND SLC38A6 AND NOT-MSLN	0.909091	1	0.833333
SLC43A1 AND NOT-MUC1 AND ANO10	0.909091	1	0.833333
SLC43A1 AND SLC38A6 AND NOT-MUC1	0.909091	1	0.833333
SLCO1B1 AND NOT-MARCO AND GPC3	0.909091	1	0.833333

Antigen Pair Logic	F1	Precision	Recall
CD19 AND AQP4 AND PPAP2A	0.914286	1	0.842105
CD19 AND CLECL1 AND NOT-ATP1A4	0.914286	1	0.842105
CD19 AND CORIN AND SLC38A2	0.914286	1	0.842105
CD19 AND NOT-MS4A2 AND PTH1R	0.914286	1	0.842105
CD19 AND NOT-MS4A2 AND CXADR	0.914286	1	0.842105
CD19 AND NOT-MS4A2 AND TMEM30B	0.914286	1	0.842105
CD19 AND NOT-TRPV2 AND SLC2A6	0.914286	1	0.842105
CD19 AND EFNB3 AND PPAP2A	0.914286	1	0.842105
CD19 AND STRA6 AND NOT-FPR3	0.914286	1	0.842105
CD19 AND CLECL1 AND NOT-GPR15	0.914286	1	0.842105
CD19 AND CLECL1 AND NOT-RECK	0.914286	1	0.842105
CD19 AND CLECL1 AND NOT-TFR2	0.914286	1	0.842105
CD19 AND NOT-MS4A2 AND USP48	0.914286	1	0.842105
COMPLEX-PPAP2A/CD19/CELSR1	0.914286	1	0.842105
CD19 AND PTH1R AND NOT-HM13	0.914286	1	0.842105
CD19 AND NDRG4 AND NOT-PAG1	0.914286	1	0.842105
CD19 AND CORIN AND PTH1R	0.914286	1	0.842105
CD19 AND CLEC4M AND NOT-SLC2A12	0.914286	1	0.842105
CD19 AND NOT-MS4A2 AND SLC7A2	0.914286	1	0.842105
CD19 AND CLECL1 AND NOT-KCNG2	0.914286	1	0.842105
CD19 AND CXADR AND NOT-SLC2A12	0.914286	1	0.842105
COMPLEX-CLECL1/TGFBR3/CD19	0.914286	1	0.842105
CD19 AND CLECL1 AND NOT-TAS2R50	0.914286	1	0.842105
COMPLEX-SLC5A12/CLECL1/CD19	0.914286	1	0.842105
CD19 AND PCDHB12 AND NOT-SLC2A12	0.914286	1	0.842105
CD19 AND EMCN AND NOT-SLC2A12	0.914286	1	0.842105
COMPLEX-CLECL1/NLGN1/CD19	0.914286	1	0.842105
CD19 AND GPBAR1 AND NOT-SLC2A12	0.914286	1	0.842105
CD19 AND CLECL1 AND NOT-GLRB	0.914286	1	0.842105
CD19 AND CLECL1 AND NOT-TMC1	0.914286	1	0.842105
COMPLEX-CLECL1/CD19/CD151	0.914286	1	0.842105
CD19 AND CORIN AND NOT-HTR4	0.914286	1	0.842105
CD19 AND NDRG4 AND NOT-CATSPER1	0.914286	1	0.842105
CD19 AND CLECL1 AND NOT-COL25A1	0.914286	1	0.842105
CD19 AND ITPR2 AND NOT-NTRK1	0.914286	1	0.842105
CD19 AND NOT-MS4A2 AND NOT-CYP4F12	0.914286	1	0.842105
CD19 AND CLECL1 AND NOT-TGFBR1	0.914286	1	0.842105
CD19 AND NDRG4 AND NOT-FPR3	0.914286	1	0.842105
COMPLEX-SLC24A4/CLECL1/CD19	0.914286	1	0.842105
COMPLEX-CLECL1/HTR7/CD19	0.914286	1	0.842105
COMPLEX-CLECL1/GPR88/CD19	0.914286	1	0.842105
CD19 AND CORIN AND NOT-ADCY2	0.914286	1	0.842105
CD19 AND SLC6A9 AND NOT-SLC2A12	0.914286	1	0.842105
CD19 AND NOT-TREML2 AND SLC2A6	0.914286	1	0.842105
COMPLEX-CFTR/ITGAE/CD19	0.914286	1	0.842105
CD19 AND HRH1 AND NOT-GLRB	0.914286	1	0.842105
CD19 AND PCDH7 AND PPAP2A	0.898551	1	0.815789
CD19 AND ADCY2 AND NOT-HM13	0.898551	1	0.815789
CD19 AND PCDH7 AND NOT-HTR4	0.898551	1	0.815789
CD19 AND ADCY2 AND NOT-HTR4	0.898551	1	0.815789
CD19 AND PCDH7 AND NOT-TNFSF14	0.898551	1	0.815789
CD19 AND ABCA8 AND NOT-FLT3LG	0.898551	1	0.815789
CD19 AND PCDH7 AND NOT-CD46	0.898551	1	0.815789
COMPLEX-PCDH7/UGT8/CD19	0.898551	1	0.815789
CD19 AND ABCA8 AND ST6GALNAC6	0.898551	1	0.815789
MS4A1 AND NOT-KCNC2 AND ADCY10	0.961039	0.948718	0.973684
MS4A1 AND SLC22A18 AND NOT-CD36	0.961039	0.948718	0.973684
MS4A1 AND SLC22A18 AND NOT-MPL	0.974359	0.95	1
MS4A1 AND ADCY10 AND NOT-CHRNA4	0.948718	0.925	0.973684
MS4A1 AND SLC22A18 AND NOT-MLC1	0.944444	1	0.894737
MS4A1 AND ADCY10 AND NOT-MLC1	0.944444	1	0.894737
MS4A1 AND ADCY10 AND NOT-SLC22A6	0.961039	0.948718	0.973684
MS4A1 AND ADCY10 AND NOT-CD36	0.973684	0.973684	0.973684
MS4A1 AND ADCY10 AND NOT-AJAP1	0.948718	0.925	0.973684
MS4A1 AND ADCY10 AND NOT-SLC28A1	0.974359	0.95	1
MS4A1 AND ADCY10 AND NOT-SYT6	0.948718	0.925	0.973684
MS4A1 AND ADCY10 AND NOT-CALY	0.947368	0.947368	0.947368
MS4A1 AND SLC22A18 AND NOT-PTGER4	0.938272	0.883721	1
MS4A1 AND ADCY10 AND NOT-MPL	0.987013	0.974359	1
MS4A1 AND NOT-KCNC2 AND HRH3	0.936709	0.902439	0.973684
MS4A1 AND HCN2 AND NOT-KCNK4	0.936709	0.902439	0.973684
MS4A1 AND NOT-KCNC2 AND SLC30A8	0.936709	0.902439	0.973684
MS4A1 AND ADCY10 AND NOT-PTGER4	0.987013	0.974359	1
MS4A1 AND PCDHAC2 AND NOT-PTGER4	0.95	0.904762	1
MS4A1 AND SLC30A8 AND NOT-PTGER4	0.935065	0.923077	0.947368
MS4A1 AND JPH3 AND NOT-KCNK4	0.935065	0.923077	0.947368
MS4A1 AND ADCY10 AND NOT-OR3A2	0.935065	0.923077	0.947368

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
SLC43A1 AND NOT-MUC1 AND NOT-CLEC1B	0.909091	1	0.833333
SLC43A1 AND NOT-MUC1 AND MRAP2	1	1	1
SLC43A1 AND NOT-MUC1 AND TMEM67	1	1	1
SLC43A1 AND NOT-MUC1 AND TMPRSS5	1	1	1
SLCO1B1 AND CD200 AND NOT-MSLN	0.923077	0.857143	1
SLC43A1 AND NOT-MUC1 AND TMPRSS5	1	1	1
SLC43A1 AND NOT-MUC1 AND TMEM67	1	1	1
SLC43A1 AND NOT-MUC1 AND SLC7A11	1	1	1
SLC2A2 AND NOT-CLEC1B AND MSLN	0.909091	1	0.833333
SLC2A2 AND NOT-MARCO AND MSLN	0.923077	0.857143	1
SLC2A2 AND SLC26A6 AND MSLN	1	1	1
SLC43A1 AND NOT-MUC1 AND SLC4A11	0.909091	1	0.833333
SLCO1B1 AND TLCD1 AND NOT-MSLN	1	1	1
SLC2A2 AND NOT-MARCO AND MSLN	0.923077	0.857143	1
SLCO1B1 AND TLCD1 AND NOT-MSLN	1	1	1
SLC2A2 AND SLC26A6 AND MSLN	1	1	1
SLC43A1 AND NOT-MUC1 AND MMP14	1	1	1
SLC43A1 AND DAGLB AND NOT-MUC1	0.909091	1	0.833333
SLC43A1 AND PANX2 AND NOT-MUC1	1	1	1
SLC43A1 AND NOT-MUC1 AND TREML2	0.909091	1	0.833333
SLC43A1 AND TMUB1 AND NOT-MUC1	0.909091	1	0.833333
SLCO1B1 AND SLC39A1 AND NOT-MSLN	0.909091	1	0.833333
SLC43A1 AND FGFRL1 AND NOT-MUC1	1	1	1
SLC43A1 AND NOT-MUC1 AND CACNA1I	0.909091	1	0.833333
SLC43A1 AND CNNM3 AND NOT-MUC1	1	1	1
SLC43A1 AND NOT-MUC1 AND SLC2A5	0.909091	1	0.833333
SLCO1B1 AND LRRC8D AND NOT-MSLN	0.909091	1	0.833333
SLC43A1 AND NOT-MUC1 AND NIPA1	0.909091	1	0.833333
SLC43A1 AND SLC39A1 AND NOT-MUC1	0.909091	1	0.833333
SLC43A1 AND NOT-MUC1 AND ADORA2B	0.909091	1	0.833333
SLCO1B1 AND NCSTN AND NOT-MSLN	1	1	1
SLC43A1 AND NOT-MUC1 AND SLC52A2	1	1	1
SLCO1B1 AND SLC52A2 AND NOT-MSLN	1	1	1
SLC43A1 AND NOT-MUC1 AND ADAM23	0.909091	1	0.833333
SLC43A1 AND ATP2A2 AND NOT-MUC1	0.909091	1	0.833333
SLCO1B1 AND LRP11 AND NOT-MSLN	1	1	1
ABCG5 AND CELSR3 AND NOT-MUC1	0.923077	0.857143	1
SLC43A1 AND NOT-MUC1 AND LRP11	1	1	1
SLC43A1 AND C6orf89 AND NOT-MUC1	1	1	1
SLC43A1 AND NOT-MUC1 AND ASIC5	1	1	1
SLCO1B1 AND SLC7A11 AND NOT-MUC1	1	1	1
SLC43A1 AND NOT-MUC1 AND ASIC5	1	1	1
SLC2A2 AND MSLN AND SLC5A6	1	1	1
SLC43A1 AND BSND AND NOT-MUC1	0.909091	1	0.833333
SLC43A1 AND GABRE AND NOT-MUC1	0.909091	1	0.833333
SLC2A2 AND NOT-CLEC4G AND MSLN	0.909091	1	0.833333
SLC43A1 AND NOT-MUC1 AND KCNG3	1	1	1
SLC43A1 AND TMEM150A AND NOT-MUC1	0.909091	1	0.833333
ABCG5 AND SLC5A5 AND NOT-MUC1	0.909091	1	0.833333
SLC43A1 AND NOT-MUC1 AND KCNG3	1	1	1
SLCO1B1 AND ADCY6 AND NOT-TNFRSF10A	1	1	1
SLCO1B1 AND NOT-KCNN2 AND MSLN	1	1	1
ABCG5 AND SLC39A1 AND NOT-MUC1	0.909091	1	0.833333
SLC43A1 AND ADCY6 AND NOT-MUC1	1	1	1
SLCO1B1 AND MMP14 AND NOT-MSLN	0.923077	0.857143	1
SLCO1B1 AND NOT-KCNN2 AND MSLN	1	1	1
SLC2A2 AND CD200 AND MSLN	1	1	1
SLC43A1 AND ADCY6 AND NOT-MUC1	1	1	1
SLC2A2 AND SLC12A7 AND MSLN	0.909091	1	0.833333
SLCO1B1 AND SLC2A5 AND NOT-MSLN	0.909091	1	0.833333
SLC43A1 AND PDCD1 AND NOT-MUC1	0.909091	1	0.833333
SLC43A1 AND NOT-MUC1 AND TAS2R4	0.909091	1	0.833333
ALCAM AND NOT-MUC1 AND SLC40A1	0.909091	1	0.833333
SLC2A2 AND HM13 AND MSLN	1	1	1
SLC43A1 AND IGF2R AND NOT-MUC1	0.909091	1	0.833333
SLC43A1 AND SLC5A5 AND NOT-MUC1	0.909091	1	0.833333
ASGR1 AND HM13 AND NOT-CD160	1	1	1
ASGR1 AND HM13 AND NOT-CD33	1	1	1
ASGR1 AND FGFRL1 AND NOT-IL13RA2	1	1	1
HPN AND SLC26A6 AND NOT-IL3RA	1	1	1
HPN AND SLC26A6 AND NOT-PMEL	1	1	1
HPN AND SLC26A6 AND NOT-PCYT1A	1	1	1
HPN AND SLC26A6 AND NOT-ENPP3	1	1	1
FGG AND NOT-MARCO AND MET	1	1	1
HPN AND SLC26A6 AND NOT-TYR	1	1	1
HPN AND SLC26A6 AND NOT-PTK7	1	1	1
FGG AND HM13 AND NOT-MAGEA4	1	1	1

Antigen Pair Logic	F1	Precision	Recall
MS4A1 AND ADCY10 AND NOT-SLC5A8	0.935065	0.923077	0.947368
MS4A1 AND NOT-SLC5A11 AND ADCY10	0.933333	0.945946	0.921053
MS4A1 AND ADCY10 AND NOT-MAG	0.95	0.904762	1
MS4A1 AND SLC30A8 AND NOT-SLC22A6	0.936709	0.902439	0.973684
MS4A1 AND CLDN20 AND NOT-CHRNA4	0.936709	0.902439	0.973684
MS4A1 AND PCDHAC2 AND NOT-CD36	0.948718	0.925	0.973684
MS4A1 AND SLC30A8 AND NOT-CD36	0.948718	0.925	0.973684
MS4A1 AND SLC22A18 AND NOT-CD93	0.974359	0.95	1
MS4A1 AND PRND AND NOT-MLC1	0.931507	0.971429	0.894737
MS4A1 AND ADAM20 AND NOT-MLC1	0.931507	0.971429	0.894737
MS4A1 AND SLC30A8 AND NOT-MLC1	0.931507	0.971429	0.894737
MS4A1 AND PCDHAC2 AND NOT-MLC1	0.931507	0.971429	0.894737
MS4A1 AND ADCY10 AND NOT-KCNF1	0.962025	0.926829	1
MS4A1 AND ADCY10 AND NOT-GRM1	0.962025	0.926829	1
MS4A1 AND SLC30A8 AND NOT-SLC28A1	0.95	0.904762	1
MS4A1 AND SLC22A18 AND NOT-TGFBI	0.933333	0.945946	0.921053
MS4A1 AND ADCY10 AND NOT-ATP6V0A4	0.948718	0.925	0.973684
MS4A1 AND ADCY10 AND NOT-LPPR3	0.948718	0.925	0.973684
MS4A1 AND CCKBR AND NOT-MLC1	0.931507	0.971429	0.894737
MS4A1 AND CLDN20 AND NOT-MLC1	0.931507	0.971429	0.894737
MS4A1 AND FGF6 AND NOT-MLC1	0.931507	0.971429	0.894737
MS4A1 AND GPR156 AND NOT-MLC1	0.931507	0.971429	0.894737
MS4A1 AND SLC22A18 AND NOT-CD163	0.974359	0.95	1
MS4A1 AND ADCY10 AND NOT-UNC5A	0.929577	1	0.868421
MS4A1 AND SLC22A18 AND NOT-KCNK6	0.95	0.904762	1
MS4A1 AND PRND AND NOT-PTGER4	0.935065	0.923077	0.947368
MS4A1 AND ADCY10 AND NOT-LRFN2	0.926829	0.863636	1
MS4A1 AND SLC30A8 AND NOT-MAG	0.926829	0.863636	1
MS4A1 AND PRND AND NOT-CD36	0.961039	0.948718	0.973684
MS4A1 AND ADCY10 AND NOT-CALHM3	0.961039	0.948718	0.973684
MS4A1 AND ADAM20 AND NOT-CD36	0.948718	0.925	0.973684
MS4A1 AND NOT-CHRNA4 AND HRH3	0.925	0.880952	0.973684
MS4A1 AND ADCY10 AND NOT-SLC22A13	0.925	0.880952	0.973684
MS4A1 AND SLC30A8 AND NOT-AJAP1	0.925	0.880952	0.973684
MS4A1 AND ADCY10 AND NOT-GABRD	0.925	0.880952	0.973684
MS4A1 AND SLC30A8 AND NOT-CHRNA4	0.925	0.880952	0.973684
MS4A1 AND ADCY10 AND NOT-GRIA4	0.925	0.880952	0.973684
MS4A1 AND SLC30A8 AND NOT-SYT6	0.925	0.880952	0.973684
MS4A1 AND NOT-CHRNA4 AND DCC	0.925	0.880952	0.973684
MS4A1 AND PRND AND NOT-SLC28A1	0.926829	0.863636	1
MS4A1 AND CLDN20 AND NOT-SYT6	0.936709	0.902439	0.973684
MS4A1 AND SLC30A8 AND NOT-ATP6V0A4	0.925	0.880952	0.973684
MS4A1 AND SLC30A8 AND NOT-LPPR3	0.925	0.880952	0.973684
MS4A1 AND ADCY10 AND NOT-GRM3	0.926829	0.863636	1
MS4A1 AND SLC30A8 AND NOT-CALY	0.923077	0.9	0.947368
MS4A1 AND PRND AND NOT-CALY	0.923077	0.9	0.947368
MS4A1 AND SLC30A8 AND NOT-OR3A2	0.923077	0.9	0.947368
MS4A1 AND KCNA1 AND NOT-MLC1	0.931507	0.971429	0.894737
MS4A1 AND DTNA AND NOT-MLC1	0.931507	0.971429	0.894737
MS4A1 AND DCC AND NOT-MLC1	0.931507	0.971429	0.894737
MS4A1 AND ADCY10 AND NOT-CACNG7	0.935065	0.923077	0.947368
MS4A1 AND CATSPERD AND NOT-MLC1	0.931507	0.971429	0.894737
MS4A1 AND ADCY10 AND NOT-OXTR	0.931507	0.971429	0.894737
MS4A1 AND OR52D1 AND NOT-MLC1	0.931507	0.971429	0.894737
MS4A1 AND SLC22A18 AND NOT-VANGL1	0.938272	0.883721	1
MS4A1 AND ADCY10 AND NOT-TRPM1	0.926829	0.863636	1
MS4A1 AND FGF6 AND NOT-PTGER4	0.95	0.904762	1
MS4A1 AND TMEFF2 AND NOT-KCNK4	0.921053	0.921053	0.921053
MS4A1 AND ADCY10 AND NOT-SCN1A	0.921053	0.921053	0.921053
MS4A1 AND CDH22 AND NOT-KCNK4	0.921053	0.921053	0.921053
MS4A1 AND CCKBR AND NOT-PTGER4	0.935065	0.923077	0.947368
MS4A1 AND GPR156 AND NOT-PTGER4	0.947368	0.947368	0.947368
MS4A1 AND CLDN20 AND NOT-PTGER4	0.947368	0.947368	0.947368
MS4A1 AND ADCY10 AND NOT-CASR	0.935065	0.923077	0.947368
MS4A1 AND ADCY10 AND NOT-CCR9	0.945946	0.972222	0.921053
MS4A1 AND PCDHAC2 AND NOT-KCNA4	0.925	0.880952	0.973684
MS4A1 AND ADCY10 AND NOT-KCNA4	0.961039	0.948718	0.973684
MS4A1 AND NOT-KCNC2 AND CCKBR	0.947368	0.947368	0.947368
MS4A1 AND PVRL1 AND NOT-KCNK4	0.918919	0.944444	0.894737
MS4A1 AND MC2R AND NOT-MLC1	0.918919	0.944444	0.894737
MS4A1 AND PCDHAC2 AND NOT-CALHM3	0.925	0.880952	0.973684
MS4A1 AND SLC30A8 AND NOT-CALHM3	0.936709	0.902439	0.973684
MS4A1 AND GPR156 AND NOT-SLC22A6	0.925	0.880952	0.973684
MS4A1 AND DCC AND NOT-SYT6	0.925	0.880952	0.973684
MS4A1 AND GPR156 AND NOT-CALY	0.923077	0.9	0.947368
MS4A1 AND NOT-MLC1 AND OR7C1	0.931507	0.971429	0.894737
MS4A1 AND CCKBR AND NOT-CALY	0.923077	0.9	0.947368

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
HPN AND SLC26A6 AND NOT-GUCY2C	1	1	1
HPN AND SLC26A6 AND NOT-FOLR1	1	1	1
HPN AND SLC26A6 AND NOT-ERBB4	1	1	1
HPN AND SLC26A6 AND NOT-CLDN3	1	1	1
HPN AND SLC26A6 AND NOT-RAET1E	1	1	1
HPN AND SLC26A6 AND NOT-CD52	1	1	1
HPN AND SLC26A6 AND NOT-SLC34A2	1	1	1
HPN AND SLC26A6 AND NOT-CLEC14A	1	1	1
HPN AND SLC26A6 AND NOT-ERBB3	1	1	1
FGG AND NOT-MARCO AND THY1	1	1	1
HPN AND SLC26A6 AND NOT-FOLR2	1	1	1
HPN AND SLC26A6 AND NOT-B4GALNT1	1	1	1
HPN AND SLC26A6 AND NOT-IL11RA	1	1	1
HPN AND SLC26A6 AND NOT-ROR1	1	1	1
HPN AND SLC26A6 AND NOT-LGR5	1	1	1
HPN AND SLC26A6 AND NOT-CLDN11	1	1	1
HPN AND SLC26A6 AND NOT-MOK	1	1	1
HPN AND SLC26A6 AND NOT-ST8SIA1	1	1	1
HPN AND SLC26A6 AND NOT-FCRL2	1	1	1
HPN AND SLC26A6 AND NOT-CD19	1	1	1
HPN AND SLC26A6 AND NOT-KDR	1	1	1
HPN AND SLC26A6 AND NOT-CD37	1	1	1
HPN AND SLC26A6 AND NOT-CD180	1	1	1
FGG AND NCSTN AND NOT-FOLH1	1	1	1
HPN AND SLC26A6 AND NOT-FCRL5	1	1	1
HPN AND SLC26A6 AND NOT-ULBP2	1	1	1
FGG AND NOT-MARCO AND TNC	1	1	1
HPN AND SLC26A6 AND NOT-CA9	1	1	1
HPN AND SLC26A6 AND NOT-MUC4	1	1	1
FGG AND NOT-MARCO AND CD34	1	1	1
HPN AND SLC26A6 AND NOT-CLDN9	1	1	1
HPN AND SLC26A6 AND NOT-SEMA5B	1	1	1
HPN AND SLC26A6 AND NOT-CD160	1	1	1
HPN AND SLC26A6 AND NOT-CD38	1	1	1
HPN AND SLC26A6 AND NOT-GAGE1	1	1	1
ASGR1 AND HM13 AND NOT-FCRL1	1	1	1
ASGR1 AND HM13 AND NOT-CD22	1	1	1
ASGR1 AND HM13 AND NOT-CD79B	1	1	1
ASGR1 AND HM13 AND NOT-CD180	1	1	1
ASGR1 AND HM13 AND NOT-P2RX5	1	1	1
ASGR1 AND HM13 AND ALDH1A1	1	1	1
ASGR1 AND HM13 AND NOT-MS4A1	1	1	1
ASGR1 AND HM13 AND NOT-TNFRSF8	1	1	1
ASGR1 AND HM13 AND NOT-CD52	1	1	1
ASGR1 AND HM13 AND NOT-CD37	1	1	1
ASGR1 AND HM13 AND NOT-CD72	1	1	1
ASGR1 AND HM13 AND NOT-CD79A	1	1	1
ASGR1 AND HM13 AND NOT-CD19	1	1	1
FGG AND SLC26A6 AND NOT-GUCY2C	0.923077	0.857143	1
FGG AND SPON2 AND NOT-CD79B	0.909091	1	0.833333
ASGR1 AND CD34 AND NOT-ITGB6	0.909091	1	0.833333
ASGR1 AND CD34 AND NOT-ABCB5	0.909091	1	0.833333
ASGR1 AND EPHB2 AND NOT-ITGB6	0.909091	1	0.833333
SLC7A9 AND SEMA5B AND NOT-TPBG	0.857143	0.75	1
SLC7A9 AND ERBB3 AND NOT-TPBG	0.833333	0.833333	0.833333
SDC1 AND NOT-FXYD3 AND NOT-IL13RA2	0.833333	0.833333	0.833333
FGG AND ERBB3 AND NOT-CD79B	0.833333	0.833333	0.833333
ASGR1 AND MAGEA1 AND NOT-ITGB6	0.909091	1	0.833333
SEMA5B AND SDC1 AND NOT-FXYD3	1	1	1
AFP AND HM13 AND NOT-IL20RA	0.909091	1	0.833333
ERBB3 AND NOT-IL20RA AND SDC4	0.833333	0.833333	0.833333
ERBB3 AND NOT-SCNN1B AND SDC1	0.909091	1	0.833333
SLC7A9 AND NOT-TPBG AND EPHB2	0.8	0.666667	1
CLDN1 AND NOT-IL20RA AND HM13	0.8	1	0.666667
SEMA5B AND SDC1 AND NOT-ACPP	1	1	1
SEMA5B AND SDC1 AND NOT-SCNN1B	1	1	1
SEMA5B AND NOT-MAL AND SDC1	0.923077	0.857143	1
ERBB3 AND NOT-PLP1 AND EDNRB	0.833333	0.833333	0.833333
SDC1 AND NOT-SCNN1B AND NOT-IL13RA2	0.909091	1	0.833333
SLC7A9 AND CD72 AND NOT-TPBG	0.833333	0.833333	0.833333
SLC7A9 AND SEMA5B AND CD70	0.923077	0.857143	1
SLC2A8 AND CLDN1 AND NOT-IL20RA	0.8	1	0.666667
SDC1 AND NOT-SCNN1B AND SSTR5	0.8	1	0.666667
FGG AND CD79B AND NOT-CEACAM5	0.909091	1	0.833333
SLC2A8 AND NOT-ITGB6 AND ERBB3	0.909091	1	0.833333
CD70 AND NOT-KIAA1324 AND CLDN3	0.833333	0.833333	0.833333
FGG AND DDX3X AND CD79B	0.8	1	0.666667

Antigen Pair Logic	F1	Precision	Recall
MS4A1 AND SLC22A18 AND NOT-ITGB3	0.974359	0.95	1
MS4A1 AND SLC22A18 AND NOT-TNFRSF8	0.961039	0.948718	0.973684
MS4A1 AND SLC22A18 AND NOT-NCAM1	0.987013	0.974359	1
MS4A1 AND ADCY10 AND NOT-ITGB3	0.987013	0.974359	1
MS4A1 AND SLC30A8 AND NOT-ITGB3	0.962025	0.926829	1
MS4A1 AND ADCY10 AND NOT-TNFRSF8	0.973684	0.973684	0.973684
MS4A1 AND PCDHAC2 AND NOT-ITGB3	0.95	0.904762	1
MS4A1 AND PRND AND NOT-ITGB3	0.962025	0.926829	1
MS4A1 AND ADAM20 AND NOT-ITGB3	0.95	0.904762	1
MS4A1 AND SLC30A8 AND NOT-TNFRSF8	0.948718	0.925	0.973684
MS4A1 AND SLC22A18 AND NOT-DDX3X	0.938272	0.883721	1
MS4A1 AND PCDHAC2 AND NOT-TNFRSF8	0.936709	0.902439	0.973684
MS4A1 AND PRND AND NOT-CLDN8	0.936709	0.902439	0.973684
MS4A1 AND ADCY10 AND NOT-CLDN8	0.986667	1	0.973684
MS4A1 AND SLC22A18 AND NOT-CD33	0.974359	0.95	1
MS4A1 AND SLC22A18 AND NOT-CD160	0.974359	0.95	1
MS4A1 AND FGF6 AND NOT-ITGB3	0.95	0.904762	1
MS4A1 AND ADAM20 AND NOT-TNFRSF8	0.936709	0.902439	0.973684
MS4A1 AND GPR156 AND NOT-ITGB3	0.962025	0.926829	1
MS4A1 AND CLDN20 AND NOT-ITGB3	0.974359	0.95	1
MS4A1 AND CCKBR AND NOT-ITGB3	0.962025	0.926829	1
MS4A1 AND PRND AND NOT-TNFRSF8	0.948718	0.925	0.973684
MS4A1 AND SLC22A18 AND NOT-ULBP1	0.938272	0.883721	1
MS4A1 AND ADCY10 AND NOT-DNAJB8	0.926829	0.863636	1
MS4A1 AND NOT-CLDN8 AND DCC	0.961039	0.948718	0.973684
MS4A1 AND NOT-CLDN8 AND MC2R	0.948718	0.925	0.973684
MS4A1 AND NOT-CLDN8 AND NMUR2	0.948718	0.925	0.973684
MS4A1 AND NOT-CLDN8 AND CLDN20	0.973684	0.973684	0.973684
MS4A1 AND NOT-CLDN8 AND KCNA1	0.936709	0.902439	0.973684
MS4A1 AND NOT-CLDN8 AND GPR156	0.948718	0.925	0.973684
MS4A1 AND SLC30A8 AND NOT-CLDN8	0.961039	0.948718	0.973684
MS4A1 AND NOT-CLDN8 AND TMEM235	0.936709	0.902439	0.973684
MS4A1 AND NOT-CLDN8 AND CATSPERD	0.936709	0.902439	0.973684
MS4A1 AND NOT-CLDN8 AND DTNA	0.948718	0.925	0.973684
MS4A1 AND GPR156 AND NOT-TNFRSF8	0.948718	0.925	0.973684
MS4A1 AND ADCY10 AND NOT-DDX3X	0.95	0.904762	1
MS4A1 AND CLDN20 AND NOT-TNFRSF8	0.961039	0.948718	0.973684
MS4A1 AND CCKBR AND NOT-TNFRSF8	0.948718	0.925	0.973684
MS4A1 AND FGF6 AND NOT-TNFRSF8	0.936709	0.902439	0.973684
MS4A1 AND ADCY10 AND NOT-MUC4	0.925	0.880952	0.973684
MS4A1 AND SLC22A18 AND NOT-CLDN8	0.925	0.880952	0.973684
MS4A1 AND ERBB3 AND NOT-CD36	0.925	0.880952	0.973684
MS4A1 AND ADCY10 AND NOT-SSTR5	0.925	0.880952	0.973684
MS4A1 AND NOT-CLDN8 AND CCKBR	0.925	0.880952	0.973684
MS4A1 AND ADCY10 AND NOT-ULBP3	0.925	0.880952	0.973684
MS4A1 AND ADCY10 AND NOT-DPEP1	0.923077	0.9	0.947368
MS4A1 AND NOT-SSTR2 AND PCDHAC2	0.926829	0.863636	1
MS4A1 AND NOT-SSTR2 AND HRH3	0.95	0.904762	1
MS4A1 AND ADCY10 AND NOT-CTAG2	0.926829	0.863636	1
MS4A1 AND ADCY10 AND NOT-SSTR2	0.962025	0.926829	1
MS4A1 AND ERBB3 AND NOT-MLC1	0.918919	0.944444	0.894737
MS4A1 AND PCDHAC2 AND NOT-DDX3X	0.926829	0.863636	1
MS4A1 AND CATSPERD AND NOT-TNFRSF8	0.948718	0.925	0.973684
MS4A1 AND MC2R AND NOT-TNFRSF8	0.936709	0.902439	0.973684
MS4A1 AND KCNA1 AND NOT-TNFRSF8	0.948718	0.925	0.973684
MS4A1 AND TMEM235 AND NOT-TNFRSF8	0.925	0.880952	0.973684
MS4A1 AND ADCY10 AND NOT-CD160	0.987013	0.974359	1
MS4A1 AND DCC AND NOT-TNFRSF8	0.948718	0.925	0.973684
MS4A1 AND DTNA AND NOT-TNFRSF8	0.948718	0.925	0.973684
MS4A1 AND SLC30A8 AND NOT-DDX3X	0.926829	0.863636	1
MS4A1 AND NMUR2 AND NOT-TNFRSF8	0.936709	0.902439	0.973684
MS4A1 AND NOT-CLDN8 AND OR52D1	0.936709	0.902439	0.973684
MS4A1 AND PCDHAC2 AND NOT-CLDN8	0.925	0.880952	0.973684
MS4A1 AND NOT-CLDN8 AND FGF6	0.925	0.880952	0.973684
MS4A1 AND NOT-CLDN8 AND SLC22A14	0.948718	0.925	0.973684
MS4A1 AND NOT-SSTR2 AND FGF6	0.926829	0.863636	1
COMPLEX-NMUR2/MS4A1/CD70	0.95	0.904762	1
MS4A1 AND SDC1 AND NOT-PTGER4	0.915663	0.844444	1
MS4A1 AND ADCY10 AND NOT-PSCA	0.915663	0.844444	1
MS4A1 AND ADCY10 AND NOT-SST	0.915663	0.844444	1
MS4A1 AND SLC22A18 AND NOT-SSTR2	0.915663	0.844444	1
MS4A1 AND ERBB3 AND NOT-SLC28A1	0.915663	0.844444	1
MS4A1 AND ADCY10 AND NOT-ALK	0.914286	1	0.842105
MS4A1 AND ADAM20 AND NOT-CLDN8	0.91358	0.860465	0.973684
MS4A1 AND NOT-CLDN8 AND OR7C1	0.91358	0.860465	0.973684
MS4A1 AND SLC30A8 AND NOT-MUC4	0.91358	0.860465	0.973684
MS4A1 AND NOT-SSTR2 AND SLC30A8	0.962025	0.926829	1

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
SDC1 AND NOT-SCNN1B AND CD34	0.8	1	0.666667
SLC2A8 AND ERBB3 AND NOT-IL20RA	0.909091	1	0.833333
AFP AND EPHB2 AND SLC40A1	0.909091	1	0.833333
SLC7A9 AND CD70 AND BIRC5	0.8	0.666667	1
FGG AND CD79B AND ITGAV	0.8	1	0.666667
ERBB3 AND NOT-CLDN8 AND SDC4	0.909091	1	0.833333
SLC7A9 AND NOT-TPBG AND DDX3X	0.8	1	0.666667
SLC2A8 AND ERBB3 AND NOT-CLDN8	0.909091	1	0.833333
SLC7A9 AND CD276 AND NOT-TPBG	0.909091	1	0.833333
AFP AND SLC2A8 AND NOT-IL20RA	0.909091	1	0.833333
AFP AND NOT-PRIMA1 AND CD70	0.833333	0.833333	0.833333
AFP AND SLC2A8 AND NOT-ERBB4	0.909091	1	0.833333
SLC2A8 AND NOT-ITGB6 AND CLDN3	0.833333	0.833333	0.833333
SDC1 AND NOT-SCNN1B AND CLEC14A	0.8	1	0.666667
SDC1 AND NOT-SCNN1B AND THY1	0.8	1	0.666667
SDC1 AND NOT-SCNN1B AND VTCN1	0.8	1	0.666667
SLC7A9 AND NOT-EPCAM AND CD70	0.8	1	0.666667
SDC1 AND NOT-SCNN1B AND TNFRSF8	0.8	1	0.666667
SLC2A8 AND CLDN1 AND NOT-CLDN8	0.8	1	0.666667
ERBB3 AND NOT-PRIMA1 AND EDNRB	0.833333	0.833333	0.833333
ASGR1 AND GPC3 AND ULBP2	0.8	1	0.666667
SLC2A8 AND NOT-MUC1 AND CLDN3	0.833333	0.833333	0.833333
SLC2A8 AND ERBB3 AND NOT-MUC1	0.909091	1	0.833333
SLC2A8 AND CLDN1 AND NOT-MUC1	0.8	1	0.666667
CLDN2 AND SLC2A8 AND NOT-MUC1	0.909091	1	0.833333
SLC2A8 AND NOT-MUC1 AND SDC1	1	1	1
SLC2A8 AND NOT-MUC1 AND CLDN12	0.8	1	0.666667
SLC2A8 AND NOT-MUC1 AND CLDN7	0.833333	0.833333	0.833333
SDC1 AND NOT-ACPP AND MSLN	0.8	0.666667	1
SLC2A8 AND NOT-MUC1 AND ERBB2	0.8	0.666667	1
SDC1 AND SLC20A1 AND NOT-MUC1	1	1	1
SLC40A1 AND NOT-MUC1 AND SSTR1	0.8	1	0.666667
SDC1 AND SLC40A1 AND NOT-MUC1	0.857143	0.75	1
CLDN1 AND SLC40A1 AND NOT-MUC1	0.8	1	0.666667
ASGR1 AND SLC26A6 AND ATP1B1	1	1	1
ASGR1 AND TMPRSS5 AND NOT-CLEC4G	1	1	1
ASGR1 AND HM13 AND NOT-CYBB	1	1	1
ASGR1 AND HM13 AND NOT-EVI2B	1	1	1
ASGR1 AND SLC7A11 AND NOT-KCNN2	1	1	1
ASGR1 AND SLC7A11 AND NOT-SLC41A2	1	1	1
ASGR1 AND HM13 AND NOT-IGSF6	1	1	1
ASGR1 AND SLC52A2 AND NOT-CD300A	1	1	1
ASGR1 AND SLC52A2 AND NOT-EVI2B	1	1	1
SLC7A9 AND SLC26A6 AND NOT-ITGA3	1	1	1
ASGR1 AND HM13 AND NOT-CR1	1	1	1
ASGR1 AND SLC7A11 AND NOT-SLC41A2	1	1	1
FGG AND NOT-MARCO AND SLC39A1	1	1	1
HPN AND NOT-MARCO AND DAGLB	1	1	1
HPN AND SLC26A6 AND NOT-NTSR1	1	1	1
HPN AND SLC26A6 AND NOT-NPY2R	1	1	1
HPN AND SLC26A6 AND NOT-TREM2	1	1	1
HPN AND SLC26A6 AND NOT-P2RY13	1	1	1
HPN AND SLC26A6 AND NOT-PMP22	1	1	1
HPN AND SLC26A6 AND NOT-TMEM8B	1	1	1
HPN AND SLC26A6 AND NOT-SLC22A17	1	1	1
HPN AND SLC26A6 AND NOT-MAL	1	1	1
HPN AND SLC26A6 AND NOT-CLEC1A	1	1	1
HPN AND SLC26A6 AND NOT-SDF4	1	1	1
HPN AND SLC26A6 AND NOT-CLEC4A	1	1	1
HPN AND SLC26A6 AND NOT-LRP5	1	1	1
HPN AND NOT-MARCO AND IFNGR2	1	1	1
HPN AND SLC26A6 AND NOT-SLC7A11	1	1	1
HPN AND SLC26A6 AND NOT-TPSG1	1	1	1
HPN AND SLC26A6 AND NOT-PCDH17	1	1	1
HPN AND SLC26A6 AND NOT-OR5J2	1	1	1
HPN AND SLC26A6 AND NOT-DLL1	1	1	1
HPN AND SLC26A6 AND NOT-GRIA2	1	1	1
HPN AND SLC26A6 AND NOT-GYPA	1	1	1
HPN AND SLC26A6 AND NOT-MR1	1	1	1
HPN AND SLC26A6 AND NOT-IL1RAP	1	1	1
HPN AND SLC26A6 AND NOT-AQP1	1	1	1
HPN AND SLC26A6 AND NOT-AQP3	1	1	1
HPN AND SLC26A6 AND NOT-KCNJ3	1	1	1
HPN AND SLC26A6 AND NOT-KCNS1	1	1	1
HPN AND SLC26A6 AND NOT-CLEC12B	1	1	1
HPN AND SLC26A6 AND NOT-SYT15	1	1	1
HPN AND SLC26A6 AND NOT-TM2D1	1	1	1

Antigen Pair Logic	F1	Precision	Recall
MS4A1 AND ADCY10 AND NOT-ULBP1	1	1	1
MS4A1 AND ADCY10 AND NOT-MST1R	0.933333	0.945946	0.921053
MS4A1 AND ERBB3 AND NOT-CALHM3	0.91358	0.860465	0.973684
MS4A1 AND P2RX3 AND NOT-TNFRSF8	0.961039	0.948718	0.973684
MS4A1 AND NOT-TNFRSF8 AND OR7C1	0.936709	0.902439	0.973684
MS4A1 AND NOT-TNFRSF8 AND ZACN	0.925	0.880952	0.973684
MS4A1 AND PRND AND NOT-DDX3X	0.938272	0.883721	1
MS4A1 AND NOT-TNFRSF8 AND KCNJ9	0.925	0.880952	0.973684
MS4A1 AND ERBB3 AND NOT-PTGER4	0.911392	0.878049	0.947368
MS4A1 AND ADCY10 AND NOT-SSTR1	0.911392	0.878049	0.947368
MS4A1 AND ADCY10 AND NOT-IGF1R	0.911392	0.878049	0.947368
MS4A1 AND NOT-CLDN8 AND GRM6	0.936709	0.902439	0.973684
MS4A1 AND ADCY10 AND NOT-SLC39A6	0.926829	0.863636	1
MS4A1 AND NOT-CLDN8 AND OPALIN	0.925	0.880952	0.973684
MS4A1 AND NOT-SSTR2 AND P2RX3	0.915663	0.844444	1
MS4A1 AND PCDHAC2 AND NOT-MST1R	0.909091	0.897436	0.921053
MS4A1 AND NOT-SSTR2 AND PRND	0.926829	0.863636	1
MS4A1 AND ADCY10 AND NOT-ANXA1	0.916667	0.970588	0.868421
MS4A1 AND SDC1 AND NOT-MLC1	0.906667	0.918919	0.894737
MS4A1 AND NOT-SSTR2 AND ADAM20	0.926829	0.863636	1
MS4A1 AND GPR156 AND NOT-DDX3X	0.938272	0.883721	1
MS4A1 AND CCKBR AND NOT-DDX3X	0.938272	0.883721	1
MS4A1 AND NOT-TNFRSF8 AND OPALIN	0.936709	0.902439	0.973684
COMPLEX-EDNRB/LAPTM5/CD19	0.898551	1	0.815789
CD19 AND PTGIS AND NOT-ITGB3	0.830769	1	0.710526
MS4A1 AND STRA6 AND NOT-ITGB3	0.974359	0.95	1
MS4A1 AND MARVELD2 AND NOT-ITGB3	0.974359	0.95	1
MS4A1 AND STRA6 AND NOT-NCAM1	0.987013	0.974359	1
CLECL1 AND NOT-CLDN2 AND NOT-CD160	0.974359	0.95	1
CLECL1 AND NOT-CLDN2 AND NOT-CD33	0.962025	0.926829	1
MS4A1 AND MRGPRX4 AND NOT-ITGB3	0.962025	0.926829	1
MS4A1 AND STRA6 AND NOT-TNFRSF8	0.961039	0.948718	0.973684
COMPLEX-CLECL1/ITGB3/CD34	0.987013	0.974359	1
MS4A1 AND MARVELD2 AND NOT-NCAM1	0.987013	0.974359	1
MS4A1 AND MARVELD2 AND NOT-TNFRSF8	0.961039	0.948718	0.973684
COMPLEX-CLECL1/ITGB3/TPBG	0.987013	0.974359	1
COMPLEX-CLECL1/ITGB3/CLDN2	0.987013	0.974359	1
COMPLEX-CLECL1/ITGB3/CA9	0.987013	0.974359	1
CLECL1 AND NOT-CLDN2 AND NOT-HSPA5	0.96	0.972973	0.947368
MS4A1 AND DPP4 AND NOT-ITGB3	0.962025	0.926829	1
CLECL1 AND NOT-SSTR1 AND NOT-CD33	0.962025	0.926829	1
MS4A1 AND SLC12A4 AND NOT-ITGB3	0.962025	0.926829	1
CLECL1 AND NOT-SSTR1 AND NOT-CD160	0.974359	0.95	1
CLECL1 AND NOT-CLDN2 AND TNFRSF13C	0.962025	0.926829	1
MS4A1 AND MRGPRX4 AND NOT-NCAM1	0.987013	0.974359	1
MS4A1 AND NDRG4 AND NOT-NCAM1	0.974359	0.95	1
CLECL1 AND NOT-CLDN2 AND NOT-ALDH1A1	0.95	0.904762	1
MS4A1 AND NDRG4 AND NOT-ITGB3	0.95	0.904762	1
MS4A1 AND CLECL1 AND NOT-CLDN2	0.95	0.904762	1
MS4A1 AND MRGPRX4 AND NOT-TNFRSF8	0.948718	0.925	0.973684
CLECL1 AND NOT-MST1R AND NOT-CD160	0.974359	0.95	1
CLECL1 AND NOT-MST1R AND NOT-CD33	0.974359	0.95	1
MS4A1 AND SLC47A2 AND NOT-ITGB3	0.962025	0.926829	1
MS4A1 AND TNFRSF18 AND NOT-ITGB3	0.974359	0.95	1
MS4A1 AND DPP4 AND NOT-TNFRSF8	0.948718	0.925	0.973684
MS4A1 AND SLC12A4 AND NOT-TNFRSF8	0.948718	0.925	0.973684
CLECL1 AND NOT-MST1R AND NOT-HSPA5	0.945946	0.972222	0.921053
CLECL1 AND NOT-SSTR1 AND NOT-HSPA5	0.945946	0.972222	0.921053
CLECL1 AND NOT-ITGB3 AND EDNRB	0.945946	0.972222	0.921053
CLECL1 AND NOT-NCAM1 AND EDNRB	0.945946	0.972222	0.921053
MS4A1 AND STRA6 AND NOT-PSCA	0.962025	0.926829	1
MS4A1 AND ST14 AND ERBB3	0.944444	1	0.894737
MS4A1 AND STRA6 AND NOT-SSTR2	0.962025	0.926829	1
MS4A1 AND RHBDL1 AND NOT-ITGB3	0.962025	0.926829	1
MS4A1 AND SLC5A9 AND NOT-ITGB3	0.95	0.904762	1
MS4A1 AND LRRC32 AND NOT-ITGB3	0.95	0.904762	1
MS4A1 AND TIE1 AND NOT-ITGB3	0.962025	0.926829	1
MS4A1 AND TNFRSF18 AND NOT-TNFRSF8	0.961039	0.948718	0.973684
MS4A1 AND NGFR AND NOT-ITGB3	0.95	0.904762	1
MS4A1 AND STRA6 AND NOT-CD33	0.974359	0.95	1
MS4A1 AND EDNRA AND NOT-ITGB3	0.974359	0.95	1
MS4A1 AND SLC47A2 AND NOT-TNFRSF8	0.948718	0.925	0.973684
MS4A1 AND STRA6 AND NOT-CD160	0.974359	0.95	1
MS4A1 AND ERBB2 AND NOT-PTGDR2	0.961039	0.948718	0.973684
MS4A1 AND MARVELD2 AND NOT-DDX3X	0.938272	0.883721	1
MS4A1 AND ERBB3 AND NOT-PTGDR2	0.938272	0.883721	1
MS4A1 AND STRA6 AND NOT-DDX3X	0.938272	0.883721	1

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
HPN AND SLC26A6 AND NOT-SLC41A2	1	1	1
HPN AND SLC26A6 AND NOT-DISP1	1	1	1
HPN AND SLC26A6 AND NOT-DCHS1	1	1	1
HPN AND SLC26A6 AND NOT-SYT8	1	1	1
HPN AND SLC26A6 AND NOT-SLC16A3	1	1	1
ASGR1 AND ADCY6 AND NOT-SIGLEC1	1	1	1
HPN AND SLC26A6 AND NOT-CD101	1	1	1
HPN AND SLC26A6 AND NOT-NCR2	1	1	1
HPN AND NOT-MARCO AND GPC3	1	1	1
FGG AND SLC26A6 AND NOT-MUC1	0.923077	0.857143	1
FGG AND NCSTN AND NOT-MSLN	1	1	1
FGG AND NCSTN AND NOT-TNFRSF10A	1	1	1
FGG AND NCSTN AND NOT-MUC1	1	1	1
FGG AND FGFRL1 AND NOT-MSLN	1	1	1
ASGR1 AND MMP14 AND NOT-MSLN	0.923077	0.857143	1
HPN AND NOT-MARCO AND MSLN	0.857143	0.75	1
COMPLEX-FGG/MUC1/MARCO	0.857143	0.75	1
HPN AND MRAP2 AND NOT-MSLN	0.857143	0.75	1
ASGR1 AND SLC26A6 AND GPC3	0.909091	1	0.833333
HPN AND TMPRSS5 AND NOT-MUC1	0.857143	0.75	1
FGG AND NOT-MARCO AND MSLN	0.857143	0.75	1
FGG AND SLC7A11 AND NOT-MUC1	1	1	1
HPN AND ASIC5 AND NOT-MSLN	0.857143	0.75	1
HPN AND ASIC5 AND NOT-MUC1	0.857143	0.75	1
HPN AND ASIC5 AND NOT-TNFRSF10A	0.857143	0.75	1
SLC7A9 AND SLC26A6 AND NOT-MUC1	1	1	1
ASGR1 AND NOT-KCNN2 AND MSLN	1	1	1
HPN AND NOT-SLC41A2 AND MSLN	0.857143	0.75	1
ASGR1 AND NOT-KCNN2 AND MSLN	1	1	1
ASGR1 AND SLC26A6 AND MSLN	0.923077	0.857143	1
ASGR1 AND SLC26A6 AND MSLN	0.923077	0.857143	1
ASGR1 AND SLC4A11 AND NOT-MSLN	0.833333	0.833333	0.833333
FGG AND SLC7A11 AND TNFRSF10A	1	1	1
HPN AND SLC5A5 AND NOT-MUC1	1	1	1
FGG AND SLC7A11 AND NOT-MSLN	1	1	1
COMPLEX-FGG/GPC3/KCNT1	0.909091	1	0.833333
ASGR1 AND HM13 AND MSLN	1	1	1
FGG AND ATP13A1 AND NOT-MSLN	1	1	1
COMPLEX-FGG/GPC3/CHRNA10	0.909091	1	0.833333
ASGR1 AND HM13 AND MSLN	1	1	1
FGG AND SLC2A8 AND MSLN	1	1	1
HPN AND NOT-ASGR1 AND MSLN	0.857143	0.75	1
HPN AND NOT-ASGR1 AND MSLN	0.857143	0.75	1
APOB AND SLC2A8 AND NOT-MUC1	0.909091	1	0.833333
SEMA4G AND NOT-MUC1 AND SLC2A8	0.909091	1	0.833333
ASGR1 AND NOT-FCGR3B AND MSLN	0.857143	0.75	1
ASGR1 AND SLC12A7 AND MSLN	0.909091	1	0.833333
ASGR1 AND ATP6V0A1 AND NOT-MUC1	1	1	1
FGG AND VN1R1 AND NOT-MUC1	0.857143	0.75	1
ASGR1 AND SLC52A2 AND NOT-TNFRSF10A	0.8	0.666667	1
ASGR1 AND NOT-GPR182 AND MSLN	0.8	1	0.666667
ASGR1 AND ATP13A1 AND NOT-TNFRSF10A	0.8	0.666667	1
SLC7A9 AND TMEM67 AND NOT-MUC1	0.8	0.666667	1
ASGR1 AND GPC3 AND CD200	0.8	1	0.666667
ASGR1 AND GPC3 AND HM13	0.8	1	0.666667
ASGR1 AND NOT-KCNN2 AND NOT-TNFRSF10A	0.8	0.666667	1
FGG AND MRAP2 AND GPC3	0.8	1	0.666667
ASGR1 AND NOT-STAB2 AND MSLN	0.8	1	0.666667
ASGR1 AND TNFRSF19 AND NOT-MSLN	0.8	1	0.666667
ASGR1 AND GPC3 AND NOT-VSIG2	0.8	1	0.666667
ASGR1 AND NOT-CLEC4M AND MSLN	0.8	1	0.666667
FGG AND TNFSF4 AND MSLN	0.923077	0.857143	1
FGG AND SLC7A11 AND GPC3	0.8	1	0.666667
ASGR1 AND NOT-CLEC4G AND MSLN	0.8	1	0.666667
FGG AND ATP6V0A1 AND TNFRSF10A	0.8	1	0.666667
ASGR1 AND HM13 AND NOT-TNFRSF10A	0.8	0.666667	1
ASGR1 AND NOT-CLEC1B AND MSLN	0.8	1	0.666667
ASGR1 AND GPC3 AND NOT-KCNN2	0.8	1	0.666667
ASGR1 AND VN1R1 AND NOT-TNFRSF10A	0.8	0.666667	1
CNNM3 AND MERTK AND NOT-MUC1	0.8	0.666667	1
ASGR1 AND NOT-PTH1R AND MSLN	0.8	1	0.666667
FGG AND NOT-MARCO AND GPC3	0.8	1	0.666667
ASGR1 AND GPC3 AND WDR19	0.8	1	0.666667
FGG AND TNFRSF10A AND OR51M1	0.8	1	0.666667
ASGR1 AND GPC3 AND GRPR	0.8	1	0.666667
FGG AND ATP13A1 AND TNFRSF10A	0.8	1	0.666667
ITPR2 AND NOT-MUC1 AND SLC2A8	0.909091	1	0.833333

Antigen Pair Logic	F1	Precision	Recall
MS4A1 AND ERBB3 AND NOT-PTGDR2	0.938272	0.883721	1
CLECL1 AND NOT-MST1R AND NOT-DDX3X	0.938272	0.883721	1
MS4A1 AND CLECL1 AND NOT-BMPR1B	0.95	0.904762	1
MS4A1 AND NOT-KCNG2 AND SDC1	0.987013	0.974359	1
MS4A1 AND NDRG4 AND NOT-TNFRSF8	0.936709	0.902439	0.973684
MS4A1 AND ERBB2 AND NOT-HM13	0.936709	0.902439	0.973684
MS4A1 AND NOT-TGOLN2 AND IL11RA	0.936709	0.902439	0.973684
MS4A1 AND MARVELD2 AND NOT-SSTR2	0.95	0.904762	1
MS4A1 AND MRGPRX4 AND NOT-SSTR2	0.962025	0.926829	1
MS4A1 AND NOT-TGOLN2 AND ERBB2	0.948718	0.925	0.973684
MS4A1 AND STRA6 AND NOT-ULBP1	0.987013	0.974359	1
MS4A1 AND STRA6 AND NOT-CLDN8	0.973684	0.973684	0.973684
MS4A1 AND TIE1 AND NOT-CLDN8	0.961039	0.948718	0.973684
MS4A1 AND SLC47A2 AND NOT-CLDN8	0.936709	0.902439	0.973684
MS4A1 AND LRRC32 AND NOT-CLDN8	0.961039	0.948718	0.973684
MS4A1 AND NOT-CLDN8 AND KCNK1	0.961039	0.948718	0.973684
MS4A1 AND MARVELD2 AND NOT-CLDN8	0.961039	0.948718	0.973684
MS4A1 AND ERBB3 AND NOT-GP9	0.938272	0.883721	1
MS4A1 AND KIR2DL4 AND NOT-ITGB3	0.95	0.904762	1
MS4A1 AND NGFR AND NOT-TNFRSF8	0.936709	0.902439	0.973684
MS4A1 AND ERBB3 AND NOT-PTPRA	0.938272	0.883721	1
MS4A1 AND MARVELD2 AND NOT-CD160	0.974359	0.95	1
MS4A1 AND RHBDL1 AND NOT-TNFRSF8	0.948718	0.925	0.973684
MS4A1 AND MARVELD2 AND NOT-CD33	0.974359	0.95	1
MS4A1 AND OR51E1 AND NOT-ITGB3	0.962025	0.926829	1
MS4A1 AND LRRC32 AND NOT-TNFRSF8	0.936709	0.902439	0.973684
MS4A1 AND NOT-CYBB AND PCYT1A	0.944444	1	0.894737
MS4A1 AND SDC1 AND NOT-PTPRA	0.936709	0.902439	0.973684
MS4A1 AND TIE1 AND NOT-TNFRSF8	0.948718	0.925	0.973684
MS4A1 AND ERBB3 AND NOT-GP9	0.938272	0.883721	1
MS4A1 AND AOC2 AND NOT-ITGB3	0.95	0.904762	1
MS4A1 AND ITPR2 AND NOT-ITGB3	0.974359	0.95	1
MS4A1 AND SDC1 AND NOT-PTGDR2	0.95	0.904762	1
MS4A1 AND CLDN2 AND NOT-PTGDR2	0.947368	0.947368	0.947368
MS4A1 AND SDC1 AND NOT-SLC16A10	0.936709	0.902439	0.973684
MS4A1 AND SDC1 AND NOT-PTGDR2	0.95	0.904762	1
MS4A1 AND ERBB3 AND NOT-PAG1	0.938272	0.883721	1
MS4A1 AND EDNRA AND NOT-TNFRSF8	0.961039	0.948718	0.973684
MS4A1 AND KCNK1 AND NOT-ITGB3	0.962025	0.926829	1
CLECL1 AND NOT-CLDN2 AND NOT-ANXA1	0.931507	0.971429	0.894737
MS4A1 AND ERBB2 AND NOT-GP9	0.961039	0.948718	0.973684
MS4A1 AND PPAP2A AND NOT-ITGB3	0.947368	0.947368	0.947368
MS4A1 AND KCNN4 AND ERBB3	0.987013	0.974359	1
MS4A1 AND MUC16 AND NOT-PTGDR2	0.938272	0.883721	1
MS4A1 AND CELSR1 AND ERBB3	0.987013	0.974359	1
MS4A1 AND MARVELD2 AND NOT-ULBP1	0.974359	0.95	1
MS4A1 AND NOT-TGOLN2 AND TNFRSF10A	0.926829	0.863636	1
NOT-LRIG3 AND NOT-DYSF AND SEMA4B	0.883117	0.871795	0.894737
LAPTM5 AND NOT-SORL1 AND NOT-GHR	0.813187	0.698113	0.973684
NOT-LRIG3 AND CLDN15 AND NOT-DYSF	0.818182	0.964286	0.710526
LAPTM5 AND NOT-SORL1 AND NOT-EFNB2	0.813187	0.698113	0.973684
LAPTM5 AND NOT-SORL1 AND NOT-PHLDB2	0.822222	0.711538	0.973684
CLDN15 AND NOT-GHR AND NOT-DYSF	0.80597	0.931034	0.710526
NOT-SYT6 AND ADCY10 AND NOT-DYSF	0.864865	0.888889	0.842105
NOT-SLC39A2 AND NOT-DYSF AND ADCY10	0.918919	0.944444	0.894737
CD19 AND NOT-CD1A AND NOT-PAG1	0.914286	1	0.842105
CD19 AND CLECL1 AND NOT-GALR2	0.914286	1	0.842105
CD19 AND CLECL1 AND NOT-CD82	0.914286	1	0.842105
COMPLEX-CLECL1/LAPTM5/CD19	0.914286	1	0.842105
COMPLEX-MTUS1/HCN2/CD19	0.914286	1	0.842105
CD19 AND CLECL1 AND NOT-IL17RA	0.914286	1	0.842105
CD19 AND PTH1R AND NOT-LYPD1	0.914286	1	0.842105
CD19 AND CLECL1 AND NOT-GPR61	0.914286	1	0.842105
CD19 AND ADCY10 AND NOT-ADCY2	0.914286	1	0.842105
CD19 AND CLECL1 AND NOT-GRID2	0.914286	1	0.842105
CD19 AND ADCY10 AND NOT-SLC39A1	0.914286	1	0.842105
CD19 AND ZP4 AND NOT-SLC2A12	0.914286	1	0.842105
CD19 AND NOT-MS4A2 AND DTNA	0.914286	1	0.842105
CD19 AND DTNA AND NOT-SLC2A12	0.914286	1	0.842105
CD19 AND CLECL1 AND NOT-SLC9B1	0.914286	1	0.842105
CD19 AND CLECL1 AND NOT-ATP1B4	0.914286	1	0.842105
CD19 AND ADAM20 AND NOT-SLC2A12	0.914286	1	0.842105
COMPLEX-CLECL1/HTR2C/CD19	0.914286	1	0.842105
COMPLEX-CLECL1/LRRC55/CD19	0.914286	1	0.842105
COMPLEX-TRPV2/LAPTM5/CD19	0.914286	1	0.842105
CD19 AND FGF6 AND NOT-SLC2A12	0.914286	1	0.842105
CD19 AND CLECL1 AND NOT-FMNL1	0.914286	1	0.842105

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
FGG AND TNFRSF10A AND MFSD10	0.8	1	0.666667
ASGR1 AND GPC3 AND NOT-SCN4B	0.8	1	0.666667
FGG AND TNFRSF10A AND SLC44A2	0.8	1	0.666667
FGG AND TNFRSF10A AND SLC2A8	0.8	1	0.666667
ASGR1 AND GPC3 AND NOT-SLC41A2	0.8	1	0.666667
ASGR1 AND GPC3 AND ASIC5	0.8	1	0.666667
FGG AND TNFRSF10A AND CLPTM1	0.8	1	0.666667
SLC7A9 AND HM13 AND NOT-MUC1	1	1	1
ASGR1 AND KCNJ12 AND NOT-TNFRSF10A	0.8	0.666667	1
ASGR1 AND GPC3 AND DIO3	0.8	1	0.666667
SLC7A9 AND NOTCH4 AND NOT-MUC1	0.923077	0.857143	1
ASGR1 AND GPC3 AND SLC7A14	0.8	1	0.666667
SLC7A9 AND WDR19 AND NOT-MUC1	1	1	1
SLC7A9 AND ASIC5 AND NOT-MUC1	0.8	0.666667	1
ASGR1 AND ACVRL1 AND TNFRSF10A	0.8	1	0.666667
SLCO1B1 AND SMPD2 AND NOT-CD79A	1	1	1
ABCG5 AND FLVCR1 AND NOT-DPEP1	1	1	1
ABCG5 AND FLVCR1 AND NOT-CEACAM5	1	1	1
ABCG5 AND FLVCR1 AND NOT-GPA33	1	1	1
ABCG5 AND FLVCR1 AND NOT-SST	1	1	1
ABCG5 AND FLVCR1 AND NOT-MST1R	1	1	1
ABCG5 AND FLVCR1 AND NOT-GUCY2C	1	1	1
ABCG5 AND FLVCR1 AND NOT-HHLA2	1	1	1
ABCG5 AND FLVCR1 AND NOT-MST1R	1	1	1
ABCG5 AND FLVCR1 AND NOT-CEACAM6	1	1	1
ABCG5 AND FLVCR1 AND NOT-GUCY2C	1	1	1
SLC43A1 AND FLVCR1 AND ALDH1A1	1	1	1
SLCO1B1 AND SMPD2 AND NOT-CEACAM5	1	1	1
ABCG5 AND NOT-CEACAM5 AND SMPD2	1	1	1
SLCO1B1 AND SMPD2 AND NOT-SLAMF7	1	1	1
SLCO1B1 AND SMPD2 AND NOT-TNFRSF17	1	1	1
SLCO1B1 AND SMPD2 AND NOT-CEACAM6	1	1	1
ABCG5 AND NOT-CEACAM6 AND SMPD2	1	1	1
SLCO1B1 AND SMPD2 AND NOT-TNFRSF17	1	1	1
SLCO1B1 AND SMPD2 AND NOT-FCRL5	1	1	1
SLCO1B1 AND SMPD2 AND NOT-CD38	1	1	1
SLCO1B1 AND SMPD2 AND NOT-ENG	1	1	1
SLCO1B1 AND SMPD2 AND NOT-PSCA	1	1	1
SLCO1B1 AND SMPD2 AND NOT-P2RX5	1	1	1
SLCO1B1 AND SMPD2 AND NOT-EDNRB	1	1	1
ABCG5 AND SMPD2 AND NOT-PROM1	1	1	1
SLC43A1 AND BIRC5 AND NOT-ABCA12	1	1	1
SLCO1B1 AND SMPD2 AND NOT-AXL	1	1	1
SLCO1B1 AND SMPD2 AND NOT-SLC34A2	1	1	1
SLCO1B1 AND SMPD2 AND NOT-PROM1	1	1	1
ABCG5 AND SPON2 AND NOT-SLC9A1	0.909091	1	0.833333
ABCG5 AND SPON2 AND NOT-CRB1	0.909091	1	0.833333
ABCG5 AND SPON2 AND NOT-OXTR	0.909091	1	0.833333
ABCG5 AND SPON2 AND NOT-SLC1A5	0.909091	1	0.833333
ABCG5 AND SPON2 AND NOT-GABRD	0.909091	1	0.833333
ABCG5 AND SPON2 AND NOT-OR3A2	0.909091	1	0.833333
ABCG5 AND SPON2 AND NOT-RXFP3	0.909091	1	0.833333
SLC2A2 AND SPON2 AND LRRC55	0.909091	1	0.833333
ABCG5 AND SPON2 AND NOT-EMP3	0.909091	1	0.833333
SLCO1B1 AND SPON2 AND NOT-CCKAR	0.909091	1	0.833333
ABCG5 AND SPON2 AND NOT-SLC39A2	0.909091	1	0.833333
SLCO1B1 AND SPON2 AND NOT-OR1Q1	0.909091	1	0.833333
SLC43A1 AND SPON2 AND NOT-GPR22	0.909091	1	0.833333
ABCG5 AND SPON2 AND NOT-KCNQ2	0.909091	1	0.833333
SLC2A2 AND SPON2 AND DRD5	0.909091	1	0.833333
ABCG5 AND SPON2 AND NOT-UMODL1	0.909091	1	0.833333
ABCG5 AND SPON2 AND NOT-CCKBR	0.909091	1	0.833333
ABCG5 AND SPON2 AND NOT-DCC	0.909091	1	0.833333
SLCO1B1 AND SPON2 AND NOT-RGR	0.909091	1	0.833333
SLC43A1 AND NOT-IL20RA AND FGF6	0.909091	1	0.833333
ABCG5 AND SPON2 AND NOT-AJAP1	0.909091	1	0.833333
SLCO1B1 AND SPON2 AND NOT-SLC6A7	0.909091	1	0.833333
SLCO1B1 AND SPON2 AND NOT-PCDHGC4	0.909091	1	0.833333
ABCG5 AND SPON2 AND NOT-KCNK10	0.909091	1	0.833333
ABCG5 AND SPON2 AND NOT-GSG1L	0.909091	1	0.833333
ABCG5 AND SPON2 AND NOT-KCNH1	0.909091	1	0.833333
ABCG5 AND SPON2 AND NOT-GRID2	0.909091	1	0.833333
ABCG5 AND SPON2 AND NOT-NRG3	0.909091	1	0.833333
SLCO1B1 AND SPON2 AND NOT-GPR37L1	0.909091	1	0.833333
SLCO1B1 AND SPON2 AND NOT-SLC12A9	0.909091	1	0.833333
ABCG5 AND DTNA AND NOT-CD79A	0.909091	1	0.833333
ABCG5 AND SPON2 AND NOT-PCDHB1	0.909091	1	0.833333

Antigen Pair Logic	F1	Precision	Recall
CD19 AND P2RX3 AND NOT-SLC2A12	0.914286	1	0.842105
CD19 AND CLECL1 AND NOT-MAG	0.914286	1	0.842105
CD19 AND CCKBR AND NOT-SLC2A12	0.914286	1	0.842105
COMPLEX-SLC6A16/CD19/SLC22A14	0.914286	1	0.842105
CD19 AND SLC17A1 AND SLC25A3	0.914286	1	0.842105
COMPLEX-TIE1/LAPTM5/CD19	0.914286	1	0.842105
CD19 AND NOT-CD1A AND NOT-CATSPER1	0.914286	1	0.842105
CD19 AND NOT-GPR61 AND SLC2A6	0.914286	1	0.842105
CD19 AND NOT-GPR61 AND OR51B2	0.914286	1	0.842105
COMPLEX-PCDH7/LAPTM5/CD19	0.898551	1	0.815789
CD19 AND ADCY2 AND NOT-SLC13A5	0.898551	1	0.815789
CD19 AND ABCA8 AND NOT-MOG	0.898551	1	0.815789
CD19 AND ADCY2 AND NOT-CD207	0.898551	1	0.815789
CD19 AND ABCA8 AND NOT-GPR61	0.898551	1	0.815789
COMPLEX-CFTR/LAPTM5/CD19	0.898551	1	0.815789
COMPLEX-IPH1/LAPTM5/CD19	0.898551	1	0.815789
CD19 AND PCDH7 AND NOT-SLC39A8	0.898551	1	0.815789
CD19 AND CLECL1 AND NOT-LRRC52	0.898551	1	0.815789
CD19 AND KCNJ16 AND NOT-LYPD1	0.898551	1	0.815789
CD19 AND SLC22A14 AND SLC25A3	0.914286	1	0.842105
COMPLEX-LAPTM5/TREML2/CD19	0.914286	1	0.842105
CD19 AND NOT-PTGER4 AND NOT-PARM1	0.914286	1	0.842105
COMPLEX-LAPTM5/SEMA6D/CD19	0.882353	1	0.789474
CD19 AND CD207 AND NOT-HTR4	0.882353	1	0.789474
COMPLEX-SLC6A16/GPR61/CD19	0.914286	1	0.842105
CD19 AND PVRL1 AND NOT-HM13	0.882353	1	0.789474
CD19 AND CLECL1 AND NOT-CHRNA4	0.882353	1	0.789474
CD19 AND ABCA8 AND CATSPER3	0.882353	1	0.789474
CD19 AND SEMA6D AND NOT-PTPRT	0.882353	1	0.789474
CD19 AND PVRL1 AND NOT-CDH26	0.882353	1	0.789474
CD19 AND GJA3 AND ABCA8	0.882353	1	0.789474
CD19 AND KCNK5 AND NOT-GABRA4	0.882353	1	0.789474
CD19 AND HTR2B AND NOT-LYPD1	0.882353	1	0.789474
COMPLEX-ECSCR/LAPTM5/CD19	0.848485	1	0.736842
CD19 AND LRRC52 AND NOT-SLC2A12	0.848485	1	0.736842
CD19 AND ECSCR AND NOT-CD207	0.848485	1	0.736842
COMPLEX-GGT5/LAPTM5/CD19	0.848485	1	0.736842
CD19 AND GGT5 AND NOT-LYPD1	0.848485	1	0.736842
CD19 AND PCDH7 AND NOT-SLC2A1	0.848485	1	0.736842
CD19 AND MLANA AND NOT-CATSPER1	0.848485	1	0.736842
COMPLEX-LAPTM5/CD19/CD28	0.848485	1	0.736842
CD19 AND PTGIS AND NOT-CNGA3	0.830769	1	0.710526
CD19 AND SGCG AND NOT-CD207	0.830769	1	0.710526
CD19 AND PTGIS AND NOT-CDH26	0.830769	1	0.710526
CD19 AND NAALAD2 AND NOT-CD207	0.830769	1	0.710526
CD19 AND SLC6A1 AND NOT-LYPD1	0.830769	1	0.710526
CD19 AND PTGIS AND NOT-PAG1	0.830769	1	0.710526
CD19 AND PTGIS AND NOT-HM13	0.830769	1	0.710526
CD19 AND PTGIS AND NOT-ADCY2	0.830769	1	0.710526
CD19 AND PTGIS AND NOT-HTR4	0.830769	1	0.710526
CD19 AND PTGIS AND NOT-CORIN	0.830769	1	0.710526
CD19 AND PTGIS AND LRRC8B	0.830769	1	0.710526
COMPLEX-ROBO2/LAPTM5/CD19	0.830769	1	0.710526
COMPLEX-EMB/LAPTM5/CD19	0.830769	1	0.710526
CD19 AND NOT-CD1A AND NOT-SLC2A6	0.8125	1	0.684211
COMPLEX-LAPTM5/PPAP2A/CD19	0.8125	1	0.684211
CD19 AND CDH19 AND NOT-ITGB3	0.914286	1	0.842105
CD19 AND CLECL1 AND NOT-CLDN2	0.914286	1	0.842105
COMPLEX-CLECL1/AXL/CD19	0.914286	1	0.842105
COMPLEX-CLECL1/CD19/CD34	0.914286	1	0.842105
CD19 AND LRP4 AND NOT-ITGB3	0.914286	1	0.842105
CD19 AND CD34 AND NOT-SLC2A12	0.914286	1	0.842105
COMPLEX-CLECL1/TPBG/CD19	0.914286	1	0.842105
CD19 AND PCDH7 AND NOT-ITGB3	0.898551	1	0.815789
CD19 AND EDNRB AND NOT-PVR	0.898551	1	0.815789
CD19 AND ABCA8 AND NOT-ITGB3	0.898551	1	0.815789
CD19 AND ABCA8 AND NOT-SLC39A6	0.898551	1	0.815789
CD19 AND ADCY2 AND NOT-ITGB3	0.898551	1	0.815789
CD19 AND EDNRB AND NOT-ADCY2	0.898551	1	0.815789
CD19 AND CLECL1 AND NOT-SSTR1	0.882353	1	0.789474
CD19 AND SEMA6D AND NOT-IL20RA	0.882353	1	0.789474
CD19 AND KCNK5 AND NOT-ITGB3	0.882353	1	0.789474
CD19 AND CLECL1 AND NOT-MST1R	0.865672	1	0.763158
CD19 AND GGT5 AND NOT-SLC39A6	0.848485	1	0.736842
CD19 AND SEMA6D AND NOT-SLC34A2	0.848485	1	0.736842
CD19 AND SLC35G1 AND NOT-ENG	0.830769	1	0.710526
CD19 AND NOT-CD1A AND LYPD1	0.898551	1	0.815789

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
ABCG5 AND SPON2 AND NOT-SLC12A9	0.909091	1	0.833333
ABCG5 AND SPON2 AND NOT-LRRC55	0.909091	1	0.833333
ABCG5 AND SPON2 AND NOT-OPCML	0.909091	1	0.833333
SLCO1B1 AND SPON2 AND NOT-SCN8A	0.909091	1	0.833333
ABCG5 AND SPON2 AND NOT-CDH22	0.909091	1	0.833333
SLCO1B1 AND SPON2 AND NOT-CACNG7	0.909091	1	0.833333
ABCG5 AND SPON2 AND NOT-HCN4	0.909091	1	0.833333
SLC2A2 AND SPON2 AND SLCO1A2	0.909091	1	0.833333
SLC43A1 AND NOT-IL13RA2 AND SLC17A3	0.909091	1	0.833333
SLCO1B1 AND EPHB2 AND NOT-SEMA4B	0.909091	1	0.833333
SLCO1B1 AND SPON2 AND NOT-SLC1A6	0.909091	1	0.833333
SLCO1B1 AND SPON2 AND NOT-LPPR3	0.909091	1	0.833333
ABCG5 AND FLVCR1 AND NOT-PROM1	0.909091	1	0.833333
SLCO1B1 AND SPON2 AND NOT-UMODL1	0.909091	1	0.833333
SLCO1B1 AND SPON2 AND NOT-GABRG2	0.909091	1	0.833333
SLCO1B1 AND SPON2 AND NOT-SLC22A6	0.909091	1	0.833333
ABCG5 AND SPON2 AND NOT-SMPD2	0.909091	1	0.833333
SLC2A2 AND SPON2 AND SEZ6	0.909091	1	0.833333
ABCG5 AND SPON2 AND NOT-GRIA4	0.909091	1	0.833333
SLCO1B1 AND SPON2 AND NOT-SLC1A5	0.909091	1	0.833333
SLCO1B1 AND SPON2 AND NOT-P2RY4	0.909091	1	0.833333
ABCG5 AND SPON2 AND NOT-OTOF	0.909091	1	0.833333
ABCG5 AND SPON2 AND NOT-CACNA1E	0.909091	1	0.833333
SLCO1B1 AND SPON2 AND NOT-SLC17A2	0.909091	1	0.833333
ABCG5 AND SPON2 AND NOT-LPPR3	0.909091	1	0.833333
SLCO1B1 AND SPON2 AND NOT-HCN4	0.909091	1	0.833333
ABCG5 AND SPON2 AND NOT-GABRG2	0.909091	1	0.833333
SLCO1B1 AND SPON2 AND NOT-AJAP1	0.909091	1	0.833333
ABCG5 AND SPON2 AND NOT-SYT6	0.909091	1	0.833333
SLCO1B1 AND SPON2 AND NOT-CALN1	0.909091	1	0.833333
ABCG5 AND SPON2 AND NOT-ADAM2	0.909091	1	0.833333
SLCO1B1 AND SPON2 AND NOT-KCNK10	0.909091	1	0.833333
SLCO1B1 AND SPON2 AND NOT-TTYH3	0.909091	1	0.833333
ABCG5 AND SPON2 AND NOT-KCNK12	0.909091	1	0.833333
SLCO1B1 AND SPON2 AND NOT-SLCO1A2	0.909091	1	0.833333
ABCG5 AND SPON2 AND NOT-PAQR8	0.909091	1	0.833333
ABCG5 AND SPON2 AND NOT-ROS1	0.909091	1	0.833333
ABCG5 AND SPON2 AND NOT-KCNJ10	0.909091	1	0.833333
SLC43A1 AND BIRC5 AND NOT-IL20RA	1	1	1
SLC43A1 AND BIRC5 AND NOT-FOLR1	1	1	1
ABCG5 AND SPON2 AND NOT-MUC4	0.909091	1	0.833333
ABCG5 AND SPON2 AND NOT-ITGB6	0.909091	1	0.833333
ABCG5 AND SPON2 AND NOT-L1CAM	0.909091	1	0.833333
ABCG5 AND SPON2 AND NOT-FCRL2	0.909091	1	0.833333
SLCO1B1 AND SPON2 AND NOT-ULBP2	0.909091	1	0.833333
ABCG5 AND SPON2 AND NOT-CLDN18	0.909091	1	0.833333
ABCG5 AND SPON2 AND NOT-ULBP2	0.909091	1	0.833333
SLC43A1 AND SPON2 AND NOT-CSPG4	0.909091	1	0.833333
ABCG5 AND SPON2 AND NOT-GUCY2C	0.909091	1	0.833333
SLC43A1 AND NOT-IL13RA2 AND NOT-IL20RA	0.909091	1	0.833333
ABCG5 AND SPON2 AND NOT-EPHB2	0.909091	1	0.833333
SLCO1B1 AND SPON2 AND NOT-GUCY2C	0.909091	1	0.833333
ABCG5 AND SPON2 AND NOT-TPBG	0.909091	1	0.833333
SLCO1B1 AND EPHB2 AND NOT-FOLR2	0.909091	1	0.833333
SLC2A2 AND SPON2 AND ITGB6	0.909091	1	0.833333
ABCG5 AND SPON2 AND NOT-ENG	0.909091	1	0.833333
SLCO1B1 AND SPON2 AND NOT-CD72	0.909091	1	0.833333
SLCO1B1 AND SPON2 AND NOT-ITGB6	0.909091	1	0.833333
SLCO1B1 AND SPON2 AND NOT-TNFRSF17	0.909091	1	0.833333
ABCG5 AND SPON2 AND NOT-MST1R	0.909091	1	0.833333
SLCO1B1 AND SPON2 AND NOT-PMEL	0.909091	1	0.833333
SLC43A1 AND SPON2 AND NOT-IL20RA	0.909091	1	0.833333
SLCO1B1 AND SPON2 AND NOT-MUC4	0.909091	1	0.833333
SLCO1B1 AND SPON2 AND NOT-MUC16	0.909091	1	0.833333
ABCG5 AND SPON2 AND NOT-CLDN7	0.909091	1	0.833333
ABCG5 AND SPON2 AND NOT-CD79B	0.909091	1	0.833333
ABCG5 AND SPON2 AND NOT-TNFRSF13C	0.909091	1	0.833333
ABCG5 AND SPON2 AND NOT-CD70	0.909091	1	0.833333
ABCG5 AND SPON2 AND NOT-TNFRSF17	0.909091	1	0.833333
ABCG5 AND SPON2 AND NOT-EGFR	0.909091	1	0.833333
ABCG5 AND SPON2 AND NOT-IL20RA	0.909091	1	0.833333
ABCG5 AND SPON2 AND NOT-CLDN23	0.909091	1	0.833333
ABCG5 AND SPON2 AND NOT-MUC13	0.909091	1	0.833333
ABCG5 AND SPON2 AND NOT-IL2RA	0.909091	1	0.833333
ABCG5 AND SPON2 AND NOT-CD19	0.909091	1	0.833333
ABCG5 AND SPON2 AND NOT-GPA33	0.909091	1	0.833333
ABCG5 AND SPON2 AND NOT-FCRL5	0.909091	1	0.833333

Antigen Pair Logic	F1	Precision	Recall
CD19 AND MEGF10 AND NOT-LYPD1	0.865672	1	0.763158
CD19 AND MLANA AND NOT-LYPD1	0.848485	1	0.736842
CD19 AND PTGIS AND NOT-LYPD1	0.830769	1	0.710526
CD19 AND PTGIS AND NOT-GABRA4	0.830769	1	0.710526
CD19 AND PTGIS AND NOT-GRM3	0.830769	1	0.710526
CD19 AND PTGIS AND NOT-CD207	0.830769	1	0.710526
MS4A1 AND ERBB3 AND NOT-ITGB3	0.938272	0.883721	1
MS4A1 AND SDC1 AND NOT-ITGB3	0.938272	0.883721	1
MS4A1 AND ERBB3 AND NOT-CLDN8	0.925	0.880952	0.973684
MS4A1 AND SDC1 AND NOT-TNFRSF8	0.925	0.880952	0.973684
MS4A1 AND ERBB3 AND NOT-TNFRSF8	0.925	0.880952	0.973684
MS4A1 AND SDC1 AND NOT-DDX3X	0.915663	0.844444	1
MS4A1 AND NOT-SSTR2 AND SDC1	0.904762	0.826087	1
MS4A1 AND ERBB3 AND NOT-DDX3X	0.904762	0.826087	1
MS4A1 AND NOT-SSTR2 AND ERBB3	0.906667	0.918919	0.894737
MS4A1 AND ERBB3 AND NOT-ULBP1	0.95	0.904762	1
MS4A1 AND SDC1 AND NOT-ULBP1	0.904762	0.826087	1
MS4A1 AND NOT-CLDN8 AND PMEL	0.936709	0.902439	0.973684
MS4A1 AND SDC1 AND NOT-ANXA1	0.891892	0.916667	0.868421
MS4A1 AND ERBB3 AND NOT-MST1R	0.886076	0.853659	0.921053
MS4A1 AND NOT-SSTR2 AND P2RX5	0.880952	0.804348	0.973684
MS4A1 AND ERBB3 AND NOT-ANXA1	0.88	0.891892	0.868421
MS4A1 AND SEMA5B AND NOT-BMPR1B	0.883721	0.791667	1
MS4A1 AND NOT-CLDN8 AND SDC1	0.875	0.833333	0.921053
MS4A1 AND SDC1 AND NOT-MST1R	0.875	0.833333	0.921053
MS4A1 AND ERBB3 AND NOT-DNAJB8	0.873563	0.77551	1
MS4A1 AND ERBB3 AND NOT-CTAG2	0.873563	0.77551	1
MS4A1 AND ERBB3 AND NOT-PSCA	0.873563	0.77551	1
MS4A1 AND ERBB3 AND NOT-SLC39A6	0.873563	0.77551	1
MS4A1 AND ERBB3 AND NOT-BMPR1B	0.904762	0.826087	1
MS4A1 AND ERBB3 AND NOT-BMPR1B	0.904762	0.826087	1
MS4A1 AND ERBB2 AND NOT-BMPR1B	0.902439	0.840909	0.973684
MS4A1 AND ERBB3 AND NOT-ULBP3	0.870588	0.787234	0.973684
MS4A1 AND ERBB3 AND NOT-SSTR5	0.870588	0.787234	0.973684
MS4A1 AND STEAP2 AND NOT-BMPR1B	0.868421	0.868421	0.868421
MS4A1 AND ERBB3 AND NOT-DPEP1	0.86747	0.8	0.947368
MS4A1 AND SDC1 AND NOT-BMPR1B	0.873563	0.77551	1
MS4A1 AND SDC1 AND NOT-BMPR1B	0.873563	0.77551	1
MS4A1 AND ERBB3 AND NOT-SST	0.863636	0.76	1
MS4A1 AND NOT-MST1R AND PMEL	0.921053	0.921053	0.921053
MS4A1 AND ERBB3 AND NOT-SLAMF7	0.88	0.891892	0.868421
MS4A1 AND ERBB3 AND NOT-CLDN6	0.873239	0.939394	0.815789
MS4A1 AND ERBB3 AND NOT-IGF1R	0.857143	0.782609	0.947368
MS4A1 AND SLC39A6 AND NOT-BMPR1B	0.860759	0.829268	0.894737
MS4A1 AND SDC1 AND NOT-SLAMF7	0.891892	0.916667	0.868421
MS4A1 AND ERBB3 AND NOT-CLDN2	0.853933	0.745098	1
MS4A1 AND MAGEA1 AND NOT-BMPR1B	0.853333	0.864865	0.842105
MS4A1 AND CLDN3 AND NOT-BMPR1B	0.880952	0.804348	0.973684
MS4A1 AND SDC1 AND NOT-SSTR5	0.850575	0.755102	0.973684
COMPLEX-PMEL/MS4A1/CD70	0.85	0.809524	0.894737
MS4A1 AND CLDN4 AND NOT-BMPR1B	0.846154	0.825	0.868421
MS4A1 AND SDC1 AND NOT-CTAG2	0.844444	0.730769	1
MS4A1 AND SDC1 AND NOT-PSCA	0.844444	0.730769	1
MS4A1 AND SDC1 AND NOT-DNAJB8	0.844444	0.730769	1
MS4A1 AND SDC1 AND NOT-SLC39A6	0.844444	0.730769	1
MS4A1 AND ERBB3 AND NOT-ITGB6	0.840909	0.74	0.973684
MS4A1 AND TYR AND NOT-BMPR1B	0.839506	0.790698	0.894737
CD52 AND NOT-ITGB3 AND NOT-KDR	0.911392	0.878049	0.947368
MS4A1 AND CBX3 AND NOT-BMPR1B	0.911392	0.878049	0.947368
MS4A1 AND GPC3 AND NOT-BMPR1B	0.837209	0.75	0.947368
MS4A1 AND SDC1 AND NOT-SST	0.835165	0.716981	1
MS4A1 AND SDC1 AND NOT-CLDN3	0.835165	0.716981	1
MS4A1 AND ERBB3 AND NOT-BCAN	0.835165	0.716981	1
MS4A1 AND SDC1 AND NOT-ULBP3	0.833333	0.76087	0.921053
MS4A1 AND SDC1 AND NOT-IGF1R	0.827586	0.734694	0.947368
MS4A1 AND LGR5 AND NOT-BMPR1B	0.827586	0.734694	0.947368
COMPLEX-PMEL/BMPR1B/MS4A1	0.902439	0.840909	0.973684
COMPLEX-SEMA5B/MS4A1/CD37	0.826087	0.703704	1
MS4A1 AND SDC1 AND NOT-CLDN2	0.826087	0.703704	1
COMPLEX-P2RX5/SEMA5B/MS4A1	0.826087	0.703704	1
MS4A1 AND CD52 AND NOT-BMPR1B	0.902439	0.840909	0.973684
MS4A1 AND CLDN12 AND NOT-BMPR1B	0.825	0.785714	0.868421
MS4A1 AND ERBB3 AND CD70	0.823529	0.744681	0.921053
MS4A1 AND ERBB3 AND NOT-SSTR1	0.822222	0.711538	0.973684
MS4A1 AND NOT-DPEP1 AND SDC1	0.821918	0.857143	0.789474
MS4A1 AND ERBB3 AND NOT-GPA33	0.817204	0.690909	1
MS4A1 AND SDC1 AND NOT-CLDN6	0.815789	0.815789	0.815789

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
SLCO1B1 AND SPON2 AND NOT-EPHB2	0.909091	1	0.833333
ABCG5 AND SPON2 AND NOT-ABCB5	0.909091	1	0.833333
SLCO1B1 AND NOT-IL13RA2 AND EPHB2	0.909091	1	0.833333
ABCG5 AND SPON2 AND NOT-CSPG4	0.909091	1	0.833333
SLC43A1 AND NOT-IL13RA2 AND NOT-FOLR1	0.909091	1	0.833333
SLCO1B1 AND SPON2 AND NOT-GPA33	0.909091	1	0.833333
ABCG5 AND SPON2 AND NOT-CA9	0.909091	1	0.833333
ABCG5 AND SPON2 AND NOT-CD79A	0.909091	1	0.833333
SLCO1B1 AND SPON2 AND NOT-FCRL2	0.909091	1	0.833333
SLC43A1 AND ERBB3 AND NOT-FOLR1	0.909091	1	0.833333
ABCG5 AND SEMA5B AND NOT-MST1R	1	1	1
SLC43A1 AND ERBB3 AND NOT-CSPG4	0.909091	1	0.833333
SLC43A1 AND NOT-IL20RA AND CD34	0.909091	1	0.833333
SLCO1B1 AND SEMA5B AND NOT-CD72	1	1	1
ABCG5 AND SEMA5B AND NOT-GPA33	1	1	1
SLCO1B1 AND EPHB2 AND NOT-MAGEA4	1	1	1
SLC43A1 AND NOT-IL13RA2 AND NOT-VTCN1	0.909091	1	0.833333
SLCO1B1 AND EPHB2 AND NOT-OAS1	1	1	1
SLC43A1 AND ERBB3 AND NOT-MUC16	0.909091	1	0.833333
SLC43A1 AND ERBB3 AND NOT-IL20RA	0.909091	1	0.833333
SLCO1B1 AND SEMA5B AND NOT-GPA33	1	1	1
SLCO1B1 AND SEMA5B AND NOT-ULBP2	1	1	1
SLCO1B1 AND SEMA5B AND NOT-TNFRSF17	1	1	1
ABCG5 AND SEMA5B AND NOT-GUCY2C	1	1	1
SLCO1B1 AND EPHB2 AND NOT-EDNRB	1	1	1
SLCO1B1 AND EPHB2 AND NOT-AXL	1	1	1
SLC2A2 AND SEMA5B AND CD70	1	1	1
SLCO1B1 AND EPHB2 AND NOT-HHLA2	1	1	1
SLCO1B1 AND EPHB2 AND NOT-L1CAM	1	1	1
SLCO1B1 AND EPHB2 AND NOT-ERBB3	1	1	1
SLCO1B1 AND EPHB2 AND NOT-CD180	1	1	1
SLCO1B1 AND EPHB2 AND NOT-CD160	1	1	1
SLCO1B1 AND EPHB2 AND NOT-TNFRSF13C	1	1	1
SLC43A1 AND NOT-FOLR1 AND EPHB2	1	1	1
ABCG5 AND SEMA5B AND NOT-MUC13	1	1	1
SLCO1B1 AND EPHB2 AND NOT-GUCY2C	1	1	1
SLC43A1 AND NOT-IL13RA2 AND CD72	0.909091	1	0.833333
SLC43A1 AND NOT-FOLR1 AND EPHB2	1	1	1
SLC43A1 AND NOT-IL20RA AND EPHB2	1	1	1
SLCO1B1 AND EPHB2 AND NOT-CLDN9	1	1	1
SLCO1B1 AND EPHB2 AND NOT-TNC	1	1	1
SLCO1B1 AND EPHB2 AND NOT-SSTR1	1	1	1
SLCO1B1 AND EPHB2 AND NOT-PMEL	1	1	1
SLCO1B1 AND EPHB2 AND NOT-BCAN	1	1	1
SLCO1B1 AND EPHB2 AND NOT-CLDN18	1	1	1
SLCO1B1 AND EPHB2 AND NOT-ANXA1	1	1	1
SLCO1B1 AND EPHB2 AND NOT-GUCY2C	1	1	1
SLCO1B1 AND EPHB2 AND NOT-ERBB4	1	1	1
SLC43A1 AND BIRC5 AND NOT-MUC1	1	1	1
SLC43A1 AND NOT-IL13RA2 AND NOT-MUC1	0.909091	1	0.833333
SLC43A1 AND SPON2 AND NOT-MUC1	0.909091	1	0.833333
SLCO1B1 AND SPON2 AND NOT-MSLN	0.909091	1	0.833333
ABCG5 AND SPON2 AND NOT-MUC1	0.909091	1	0.833333
ABCG5 AND SPON2 AND NOT-MSLN	0.909091	1	0.833333
SLCO1B1 AND SEMA5B AND NOT-MSLN	1	1	1
SLC43A1 AND ERBB3 AND NOT-MUC1	0.909091	1	0.833333
SLC43A1 AND NOT-MUC1 AND CD34	0.909091	1	0.833333
SLCO1B1 AND EPHB2 AND NOT-MSLN	1	1	1
SLCO1B1 AND EPHB2 AND NOT-MSLN	1	1	1
SLC43A1 AND NOT-MUC1 AND EPHB2	1	1	1
SLCO1B1 AND EPHB2 AND NOT-TNFRSF10A	1	1	1
SLC43A1 AND NOT-MUC1 AND EPHB2	1	1	1
SLC43A1 AND SEMA5B AND NOT-MUC1	1	1	1
SLC43A1 AND NOT-MUC1 AND LGR5	0.909091	1	0.833333
SLC43A1 AND NOT-MUC1 AND IL3RA	0.909091	1	0.833333
SLC2A2 AND SEMA5B AND MSLN	1	1	1
SLC2A2 AND BIRC5 AND MSLN	0.857143	0.75	1
SLC2A2 AND EPHB2 AND MSLN	1	1	1
SLC43A1 AND NOT-MUC1 AND ULBP1	0.909091	1	0.833333
SLC2A2 AND NOT-IL13RA2 AND MSLN	0.833333	0.833333	0.833333
SLC2A2 AND MSLN AND NOT-AXL	0.857143	0.75	1
SLC43A1 AND NOT-MUC1 AND GPA33	1	1	1
SLC43A1 AND NOT-MUC1 AND ITGB6	1	1	1
SLC43A1 AND NOT-MUC1 AND GPA33	1	1	1
ABCG5 AND SEMA5B AND NOT-MUC1	1	1	1
SLC2A2 AND VTCN1 AND MSLN	0.8	1	0.666667
ABCG5 AND GPC3 AND NOT-MST1R	0.8	1	0.666667

Antigen Pair Logic	F1	Precision	Recall
MS4A1 AND SDC1 AND NOT-ITGB6	0.813187	0.698113	0.973684
MS4A1 AND CD52 AND P2RX5	0.897436	0.875	0.921053
MS4A1 AND SLC34A2 AND P2RX5	0.8125	1	0.684211
MS4A1 AND HSPA5 AND NOT-BMPR1B	0.815789	0.815789	0.815789
MS4A1 AND MOK AND NOT-BMPR1B	0.810127	0.780488	0.842105
MS4A1 AND TYR AND NOT-ERBB3	0.809524	0.73913	0.894737
MS4A1 AND SDC1 AND NOT-BCAN	0.808511	0.678571	1
MS4A1 AND FOLH1 AND NOT-BMPR1B	0.805556	0.852941	0.763158
COMPLEX-ROR1/PMEL/MS4A1	0.804878	0.75	0.868421
MS4A1 AND ERBB3 AND HSPA5	0.804348	0.685185	0.973684
MS4A1 AND ERBB3 AND IL11RA	0.804348	0.685185	0.973684
MS4A1 AND ERBB3 AND CLDN3	0.8	0.666667	1
MS4A1 AND CD70 AND NOT-BMPR1B	0.898551	1	0.815789
Peripheral T-cell lymphoma			
CXCL9 AND NOT-BMPR1A AND NOT-SDC2	0.830189	0.88	0.785714
GPR174 AND CDH5 AND NOT-CXADR	0.807018	0.793103	0.821429
CXCL9 AND NOT-CRIM1 AND NOT-CYP4F12	0.830189	0.88	0.785714
CD96 AND CDH5 AND NOT-EPHB4	0.80597	0.692308	0.964286
CXCL9 AND NOT-APCDD1 AND NOT-GJB2	0.862745	0.956522	0.785714
CXCL9 AND NOT-PMP22 AND NOT-MGST2	0.814815	0.846154	0.785714
CXCL9 AND NOT-FAM57A AND NOT-SDC2	0.846154	0.916667	0.785714
CXCL9 AND NOT-PMP22 AND CD96	0.814815	0.846154	0.785714
CXCL9 AND NOT-FAM57A AND NOT-AQP9	0.830189	0.88	0.785714
CXCL9 AND NOT-CRIM1 AND NOT-MGST2	0.830189	0.88	0.785714
P2RY10 AND CDH5 AND NOT-CXADR	0.806452	0.735294	0.892857
TRAT1 AND CDH5 AND NOT-CXADR	0.851852	0.884615	0.821429
CXCL9 AND NOT-SLC3A1 AND NOT-SLC11A2	0.814815	0.846154	0.785714
CXCL9 AND NOT-SDC2 AND NOT-MGST2	0.862745	0.956522	0.785714
CXCL9 AND NOT-CRIM1 AND NOT-ITGA2	0.814815	0.846154	0.785714
CXCL9 AND NOT-CRIM1 AND NOT-SPINT1	0.814815	0.846154	0.785714
CXCL9 AND NOT-CRIM1 AND NOT-SLC22A23	0.814815	0.846154	0.785714
CXCL9 AND NOT-GJB2 AND NOT-TMEM219	0.833333	1	0.714286
CXCL9 AND NOT-GJB2 AND NOT-TMEM150A	0.816327	0.952381	0.714286
CXCL9 AND NOT-GJB2 AND NOT-CAV1	0.833333	1	0.714286
CXCL9 AND NOT-GJB2 AND NOT-CSF3R	0.833333	1	0.714286
CXCL9 AND NOT-GJB2 AND NOT-TMEM231	0.816327	0.952381	0.714286
CXCL9 AND NOT-GJB2 AND NOT-RAMP2	0.816327	0.952381	0.714286
CXCL9 AND NOT-GJB2 AND NOT-TUSC3	0.833333	1	0.714286
CXCL9 AND NOT-BMPR1A AND NOT-TNFRSF1A	0.846154	0.916667	0.785714
CXCL9 AND NOT-CRIM1 AND NOT-PTPRK	0.814815	0.846154	0.785714
CXCL9 AND NOT-GJB2 AND NOT-PCDH12	0.833333	1	0.714286
CXCL9 AND NOT-GJB2 AND NOT-COMT	0.833333	1	0.714286
FAM26F AND CDH5 AND NOT-CXADR	0.814815	0.846154	0.785714
CXCL9 AND NOT-CRIM1 AND NOT-FAM57A	0.814815	0.846154	0.785714
CXCL9 AND NOT-APCDD1 AND NOT-AQP9	0.830189	0.88	0.785714
CXCL9 AND NOT-SDC2 AND NOT-MS4A2	0.88	1	0.785714
CXCL9 AND NOT-APLP2 AND CD96	0.846154	0.916667	0.785714
CD96 AND CDH11 AND NOT-KIT	0.806452	0.735294	0.892857
CXCL9 AND NOT-FAM57A AND NOT-ANPEP	0.830189	0.88	0.785714
CXCL9 AND NOT-SDC2 AND NOT-LRIG3	0.846154	0.916667	0.785714
CXCL9 AND NOT-CRIM1 AND NOT-LRIG3	0.814815	0.846154	0.785714
CXCL9 AND NOT-CRIM1 AND NOT-SEMA4B	0.814815	0.846154	0.785714
CXCL9 AND NOT-SEMA4B AND NOT-ERMAP	0.814815	0.846154	0.785714
CXCL9 AND NOT-SEMA4B AND NOT-LMBRD1	0.830189	0.88	0.785714
CD96 AND THY1 AND NOT-EMP2	0.830189	0.88	0.785714
CD96 AND THY1 AND NOT-KIT	0.807692	0.875	0.75
CD96 AND THY1 AND NOT-BMP2	0.84	0.954545	0.75
CXCL9 AND NOT-GJB2 AND NOT-MUC1	0.816327	0.952381	0.714286
CD96 AND THY1 AND NOT-SEMA6A	0.8	0.909091	0.714286
CD96 AND LIFR AND NOT-STEAP2	0.806452	0.735294	0.892857
NOT-EMP2 AND THY1 AND GPR183	0.816327	0.952381	0.714286
CD96 AND THY1 AND NOT-ATP8B1	0.830189	0.88	0.785714
Melanoma			
NOT-CHRNA2 AND LGR4 AND NOT-KCNK5	1	1	1
NOT-CHRNA2 AND LGR4 AND NOT-CFTR	1	1	1
NOT-CLDN19 AND TIE1 AND NOT-CADM2	1	1	1
NOT-CHRNA2 AND PCDHB14 AND NOT-CADM2	1	1	1
GPR137B AND NOT-NPR3 AND TYRO3	1	1	1
NOT-CHRNA2 AND NOT-CADM2 AND TYRO3	1	1	1
NOT-CLDN19 AND LGR4 AND NOT-CADM2	1	1	1
NOT-CHRNA2 AND ZNRF3 AND NOT-CADM2	1	1	1
NOT-CHRNA2 AND ZNRF3 AND NOT-CADM2	1	1	1
NOT-CHRNA2 AND TIE1 AND NOT-KCNK5	1	1	1
NOT-CLDN19 AND PCDHB14 AND NOT-CADM2	1	1	1
NOT-CHRNA2 AND NOT-CADM2 AND TYRO3	1	1	1
NOT-CHRNA2 AND PCDHB14 AND NOT-CADM2	1	1	1
NOT-CHRNA2 AND LGR4 AND NOT-CADM2	1	1	1

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
ABCG5 AND GPC3 AND NOT-CD19	0.8	1	0.666667
ABCG5 AND GPC3 AND NOT-CA9	0.8	1	0.666667
ABCG5 AND THY1 AND NOT-MUC1	0.8	1	0.666667
ABCG5 AND GPC3 AND EPHB2	0.8	1	0.666667
ABCG5 AND GPC3 AND NOT-L1CAM	0.8	1	0.666667
ABCG5 AND GPC3 AND NOT-CLDN23	0.8	1	0.666667
SLC2A2 AND MSLN AND NOT-EDNRB	0.857143	0.75	1
SLC43A1 AND NOT-MUC1 AND VTCN1	0.8	1	0.666667
SLC43A1 AND NOT-MUC1 AND THY1	0.8	1	0.666667
SLC43A1 AND NOT-MUC1 AND SSTR5	0.8	1	0.666667
ABCG5 AND GPC3 AND NOT-CLDN7	0.8	1	0.666667
SLC43A1 AND NOT-MUC1 AND CR2	0.8	1	0.666667
ABCG5 AND GPC3 AND NOT-GUCY2C	0.8	1	0.666667
ABCG5 AND GPC3 AND NOT-ITGB6	0.8	1	0.666667
ABCG5 AND GPC3 AND NOT-FCRL2	0.8	1	0.666667
SLC43A1 AND NOT-MUC1 AND DLL3	0.8	1	0.666667
SLC2A2 AND GPC3 AND CD70	0.8	1	0.666667
SLCO1B1 AND GPC3 AND ULBP2	0.8	1	0.666667
ABCG5 AND GPC3 AND ULBP2	0.8	1	0.666667
SLC2A2 AND THY1 AND MSLN	0.8	1	0.666667
ABCG5 AND GPC3 AND NOT-TPBG	0.8	1	0.666667
ABCG5 AND GPC3 AND ABCB5	0.8	1	0.666667
SLC43A1 AND NOT-MUC1 AND MAGEA11	0.909091	1	0.833333
ABCG5 AND GPC3 AND NOT-TNFRSF13C	0.8	1	0.666667
ABCG5 AND GPC3 AND NOT-MUC13	0.8	1	0.666667
ABCG5 AND GPC3 AND NOT-TNFRSF17	0.8	1	0.666667
ABCG5 AND GPC3 AND NOT-IL20RA	0.8	1	0.666667
SLC2A2 AND MSLN AND FLOT2	0.8	1	0.666667
SLC43A1 AND NOT-MUC1 AND CLDN18	0.8	1	0.666667
ABCG5 AND GPC3 AND NOT-CLDN18	0.8	1	0.666667
ABCG5 AND GPC3 AND NOT-IL2RA	0.8	1	0.666667
ABCG5 AND GPC3 AND NOT-CD79A	0.8	1	0.666667
SLC2A2 AND MSLN AND CBX3	0.8	1	0.666667
SLCO1B1 AND GPC3 AND NOT-TNFRSF17	0.8	1	0.666667
ABCG5 AND GPC3 AND NOT-EGFR	0.8	1	0.666667
ABCG5 AND GPC3 AND NOT-CSPG4	0.8	1	0.666667
SLCO1B1 AND THY1 AND NOT-MSLN	0.8	1	0.666667
ABCG5 AND GPC3 AND NOT-CD70	0.8	1	0.666667
ABCG5 AND MAGEA1 AND NOT-MSLN	0.8	1	0.666667
SLC43A1 AND NOT-MUC1 AND VTCN1	0.8	1	0.666667
SLC43A1 AND NOT-MUC1 AND CSPG4	0.8	1	0.666667
SLC43A1 AND NOT-IL20RA AND TNFRSF10A	0.8	1	0.666667
SLC2A2 AND GPC3 AND ITGB6	0.8	1	0.666667
SLC43A1 AND NOT-MUC1 AND CSPG4	0.8	1	0.666667
ABCG5 AND GPC3 AND NOT-CD79B	0.8	1	0.666667
ABCG5 AND GPC3 AND NOT-GPA33	0.8	1	0.666667
ABCG5 AND GPC3 AND NOT-FCRL5	0.8	1	0.666667
ABCG5 AND GPC3 AND NOT-MUC4	0.8	1	0.666667
SLC43A1 AND NOT-MUC1 AND PSCA	0.8	1	0.666667
SLC2A2 AND MSLN AND RNF43	0.857143	0.75	1
SLC2A2 AND MSLN AND ITGB6	0.857143	0.75	1
SLC2A2 AND MSLN AND NOT-PROM1	0.857143	0.75	1
SLC43A1 AND NOT-MUC1 AND CLEC14A	0.8	1	0.666667
SLC43A1 AND NOT-MUC1 AND GAGE1	0.909091	1	0.833333
SLC2A2 AND CLEC14A AND MSLN	0.8	1	0.666667
SLCO1B1 AND GPC3 AND NOT-GPA33	0.8	1	0.666667
SLCO1B1 AND GPC3 AND NOT-MUC4	0.8	1	0.666667
SLCO1B1 AND GPC3 AND NOT-GUCY2C	0.8	1	0.666667
SLCO1B1 AND GPC3 AND NOT-MUC16	0.8	1	0.666667
SLCO1B1 AND GPC3 AND NOT-PMEL	0.8	1	0.666667
SLCO1B1 AND GPC3 AND NOT-FCRL2	0.8	1	0.666667
SLC2A2 AND MSLN AND NOT-CEACAM6	0.857143	0.75	1
SLC43A1 AND NOT-IL20RA AND MSLN	1	1	1
SLC43A1 AND NOT-FOLR1 AND MSLN	1	1	1
ABCG5 AND NOT-MUC17 AND FLVCR1	1	1	1
ABCG5 AND NOT-MUC17 AND FLVCR1	1	1	1
SLC43A1 AND NOT-MUC17 AND PCDHB1	1	1	1
SLC43A1 AND NOT-MUC17 AND PCDHB1	1	1	1
ABCG5 AND FLVCR1 AND NOT-CDH17	1	1	1
ABCG5 AND FLVCR1 AND NOT-SLC16A5	1	1	1
ABCG5 AND FLVCR1 AND NOT-CDH17	1	1	1
ABCG5 AND FLVCR1 AND NOT-PTPRR	1	1	1
ABCG5 AND FLVCR1 AND NOT-PTPRR	1	1	1
ABCG5 AND FLVCR1 AND NOT-SLC13A2	1	1	1
ABCG5 AND FLVCR1 AND NOT-SLC13A2	1	1	1
ABCG5 AND FLVCR1 AND NOT-FUT3	1	1	1
ABCG5 AND FLVCR1 AND NOT-SLC16A5	1	1	1

Antigen Pair Logic	F1	Precision	Recall
GPR137B AND NOT-NPR3 AND TYRO3	1	1	1
NOT-CHRNA2 AND NOT-CADM2 AND IGDCC4	1	1	1
NOT-CHRNA2 AND PCDHB14 AND NOT-KCNK5	1	1	1
NOT-TM4SF5 AND TTYH2 AND ANO1	1	1	1
NOT-TM4SF5 AND TTYH2 AND KCNK15	1	1	1
NOT-TM4SF5 AND TTYH2 AND SLCO2A1	1	1	1
NOT-TM4SF5 AND TTYH2 AND LGR4	1	1	1
NOT-TM4SF5 AND TTYH2 AND TSPAN11	1	1	1
NOT-TM4SF5 AND TTYH2 AND FZD7	1	1	1
NOT-CHRNA2 AND TIE1 AND NOT-CFTR	1	1	1
NOT-CHRNA2 AND TSPAN11 AND NOT-CFTR	1	1	1
NOT-CHRNA2 AND PCDHB14 AND NOT-CFTR	1	1	1
NOT-CHRNA2 AND TSPAN11 AND NOT-KCNK5	1	1	1
NOT-CHRNA2 AND NOT-CADM2 AND APLNR	1	1	1
NOT-CHRNA2 AND NOT-CADM2 AND CNTNAP1	1	1	1
NOT-CHRNA2 AND TIE1 AND NOT-CADM2	1	1	1
GPR137B AND NOT-ATP1A2 AND TYRO3	1	1	1
GPR137B AND NOT-ATP1A2 AND TYRO3	1	1	1
NOT-CLDN19 AND TSPAN11 AND NOT-CADM2	1	1	1
GPR137B AND NOT-CADM2 AND KCNS3	1	1	1
GPR137B AND NOT-CADM2 AND FXYD3	1	1	1
NOT-CHRNA2 AND NOT-ATP1A2 AND TYRO3	1	1	1
GPR137B AND NOT-CADM2 AND TYRO3	1	1	1
NOT-CHRNA2 AND TIE1 AND NOT-CADM2	1	1	1
GPR137B AND NOT-CADM2 AND VANGL2	1	1	1
GPR137B AND NOT-FGFR2 AND TYRO3	1	1	1
NOT-CLDN19 AND ZNRF3 AND NOT-CADM2	1	1	1
GPR137B AND NOT-CADM2 AND FXYD3	1	1	1
GPR137B AND NOT-CADM2 AND PCDH7	1	1	1
NOT-CHRNA2 AND NOT-CADM2 AND ADRA2C	1	1	1
NOT-CHRNA2 AND NOT-ATP1A2 AND TYRO3	1	1	1
GPR137B AND NOT-CADM2 AND TYRO3	1	1	1
NOT-CHRNA2 AND NOT-CADM2 AND ADRA2C	1	1	1
NOT-CHRNA2 AND FZD7 AND NOT-KCNK5	1	1	1
GPR137B AND NOT-CADM2 AND VANGL2	1	1	1
NOT-CHRNA2 AND NOT-CADM2 AND ANO10	1	1	1
NOT-CHRNA2 AND ZNRF3 AND NOT-KCNK5	1	1	1
GPR137B AND NOT-FGFR2 AND TYRO3	1	1	1
NOT-CHRNA2 AND NOT-CADM2 AND PCDHB16	1	1	1
NOT-CHRNA2 AND TMEM231 AND NOT-CFTR	1	1	1
NOT-CHRNA2 AND WLS AND NOT-KCNK5	1	1	1
NOT-GIPR AND GPR161 AND NOT-PCDH8	1	1	1
GPR137B AND NOT-CXADR AND TYRO3	1	1	1
NOT-TM4SF5 AND NOT-CADM2 AND LRIG3	1	1	1
NOT-CHRNA2 AND NOT-CADM2 AND WLS	1	1	1
NOT-CLDN19 AND TYRO3 AND NOT-CADM2	0.956522	0.916667	1
NOT-CACNG6 AND NOT-CADM2 AND PCDHB14	1	1	1
NOT-CHRNA2 AND NOT-CADM2 AND WLS	1	1	1
NOT-CHRNA2 AND NOT-CADM2 AND LSAMP	0.956522	0.916667	1
COMPLEX-CHRNA2/OR7A5/SGCD	1	1	1
NOT-CACNG6 AND LGR4 AND NOT-CADM2	1	1	1
NOT-SLC10A2 AND GPR161 AND NOT-ATP1A2	1	1	1
NOT-CHRNA2 AND LGR4 AND NOT-SCNN1G	1	1	1
NOT-CHRNA2 AND NOT-CADM2 AND NTM	1	1	1
NOT-CHRNA2 AND TMEM231 AND NOT-KCNK5	1	1	1
NOT-GPR21 AND TTYH2 AND ATP1A1	1	1	1
NOT-CHRNA2 AND TMEM231 AND NOT-SCNN1G	1	1	1
NOT-TM4SF5 AND TTYH2 AND PCDHB14	1	1	1
NOT-CHRNA2 AND SGCD AND PCDHB14	0.952381	1	0.909091
GPR137B AND NOT-CADM2 AND BCAP31	0.952381	1	0.909091
NOT-CLDN19 AND LGR4 AND NOT-ATP1A2	0.952381	1	0.909091
GPR137B AND NOT-CADM2 AND RAMP2	0.952381	1	0.909091
NOT-CHRNA2 AND SLC16A4 AND TNFRSF19	0.952381	1	0.909091
NOT-CLDN19 AND SGCD AND TNFRSF10D	0.952381	1	0.909091
NOT-CHRNA2 AND LGR4 AND NOT-TSPAN8	0.952381	1	0.909091
NOT-CHRNA2 AND SGCD AND PCDHB16	0.952381	1	0.909091
COMPLEX-CD300LB/PSEN2/GPR137B	0.952381	1	0.909091
NOT-CLDN19 AND SGCD AND NOT-SLC7A14	0.952381	1	0.909091
NOT-SLC34A3 AND TTYH2 AND ANO1	0.952381	1	0.909091
GPR137B AND NOT-TSPAN8 AND TYRO3	0.952381	1	0.909091
GPR137B AND SLC5A6 AND NOT-NPR3	0.952381	1	0.909091
NOT-CHRNA2 AND SGCD AND APLNR	0.952381	1	0.909091
NOT-CHRNA2 AND SLC16A4 AND IGDCC4	0.952381	1	0.909091
NOT-CHRNA2 AND SGCD AND DAGLB	0.952381	1	0.909091
NOT-CHRNA2 AND SLC16A4 AND PCDHB16	0.952381	1	0.909091
GPR137B AND NOT-NPR3 AND KCNS3	0.952381	1	0.909091
NOT-CHRNA2 AND SLC16A4 AND LGR4	0.952381	1	0.909091

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
SLC43A1 AND FLVCR1 AND NOT-AOC3	1	1	1
COMPLEX-PCDHB1/SLC2A2/PLVAP	0.923077	0.857143	1
SLCO1B1 AND SMPD2 AND NOT-ABCA12	1	1	1
SLCO1B1 AND SMPD2 AND NOT-STEAP4	1	1	1
SLCO1B1 AND SMPD2 AND NOT-PCDH11Y	1	1	1
SLCO1B1 AND SMPD2 AND NOT-CD163	1	1	1
SLCO1B1 AND PVRL1 AND NOT-SLC17A2	0.923077	0.857143	1
SLCO1B1 AND SMPD2 AND NOT-GYPC	1	1	1
COMPLEX-P2RY4/SLC2A2/PLVAP	0.909091	1	0.833333
ABCG5 AND NOT-MUC17 AND KCNJ6	0.909091	1	0.833333
SLCO1B1 AND SMPD2 AND NOT-PPAPDC1A	0.909091	1	0.833333
SLCO1B1 AND SMPD2 AND NOT-CLDN10	0.909091	1	0.833333
SLC2A2 AND PCDHB1 AND GYPC	0.909091	1	0.833333
SLC43A1 AND FLVCR1 AND NOT-MMP24	1	1	1
SLC43A1 AND FLVCR1 AND NOT-SLC9B1	1	1	1
SLC43A1 AND FLVCR1 AND NOT-CD44	1	1	1
SLC43A1 AND FLVCR1 AND NOT-DGKE	1	1	1
SLC43A1 AND FLVCR1 AND NOT-SLC22A5	1	1	1
SLC43A1 AND FLVCR1 AND NOT-KCNK6	1	1	1
SLC43A1 AND FLVCR1 AND NOT-MPL	1	1	1
SLC43A1 AND FLVCR1 AND NOT-ZACN	1	1	1
SLC43A1 AND FLVCR1 AND NOT-GUCY2D	1	1	1
SLC43A1 AND FLVCR1 AND NOT-LAT	1	1	1
SLC43A1 AND FLVCR1 AND NOT-PAQR7	1	1	1
SLC43A1 AND FLVCR1 AND NOT-SELP	0.909091	1	0.833333
SLC43A1 AND FLVCR1 AND NOT-GPR22	1	1	1
SLC43A1 AND RGR AND NOT-CD163	0.909091	1	0.833333
SLC43A1 AND FLVCR1 AND NOT-SYT11	1	1	1
SLC43A1 AND FLVCR1 AND NOT-ATP8B2	0.909091	1	0.833333
SLC43A1 AND FLVCR1 AND NOT-LRRC8E	1	1	1
SLC43A1 AND FLVCR1 AND NOT-SLC16A5	1	1	1
ABCG5 AND FLVCR1 AND NOT-SLC9A1	1	1	1
SLC43A1 AND FLVCR1 AND NOT-SLC9A1	1	1	1
SLC43A1 AND FLVCR1 AND NOT-NFASC	1	1	1
SLC43A1 AND FLVCR1 AND NOT-SEMA4D	1	1	1
SLC43A1 AND FLVCR1 AND NOT-FXYD6	1	1	1
SLC43A1 AND FLVCR1 AND NOT-GPR22	1	1	1
SLCO1B1 AND SMPD2 AND NOT-KCNH4	1	1	1
SLCO1B1 AND SMPD2 AND NOT-ATP8B2	1	1	1
SLCO1B1 AND SMPD2 AND NOT-PORCN	1	1	1
SLCO1B1 AND SMPD2 AND NOT-UPK3A	0.909091	1	0.833333
SLCO1B1 AND SMPD2 AND NOT-CD1A	1	1	1
SLCO1B1 AND SMPD2 AND NOT-CD93	1	1	1
SLCO1B1 AND SMPD2 AND NOT-ATP1B4	1	1	1
SLCO1B1 AND SMPD2 AND NOT-AOC3	1	1	1
SLCO1B1 AND SMPD2 AND NOT-NOX1	1	1	1
SLCO1B1 AND SMPD2 AND NOT-EPHA10	1	1	1
SLCO1B1 AND SMPD2 AND NOT-ICAM1	1	1	1
SLCO1B1 AND SMPD2 AND NOT-SLC12A9	1	1	1
SLCO1B1 AND SMPD2 AND NOT-OXTR	1	1	1
SLCO1B1 AND SMPD2 AND NOT-DCC	1	1	1
SLCO1B1 AND SMPD2 AND NOT-CABP7	1	1	1
SLCO1B1 AND SMPD2 AND NOT-SV2C	1	1	1
SLCO1B1 AND SMPD2 AND NOT-CLDN15	0.909091	1	0.833333
SLCO1B1 AND SMPD2 AND NOT-SLC24A2	1	1	1
SLCO1B1 AND SMPD2 AND NOT-CDH20	1	1	1
SLCO1B1 AND SMPD2 AND NOT-OR4N4	1	1	1
SLCO1B1 AND SMPD2 AND NOT-GPR22	1	1	1
SLCO1B1 AND SMPD2 AND NOT-GPR26	1	1	1
SLCO1B1 AND SMPD2 AND NOT-GRM7	1	1	1
SLCO1B1 AND SMPD2 AND NOT-SLC39A2	1	1	1
SLCO1B1 AND SMPD2 AND NOT-ATP13A5	1	1	1
SLCO1B1 AND SMPD2 AND NOT-KCNA4	1	1	1
SLCO1B1 AND SMPD2 AND NOT-LRRC52	1	1	1
SLCO1B1 AND SMPD2 AND NOT-OR4N4	1	1	1
SLC43A1 AND NOT-MUC1 AND FGF6	0.909091	1	0.833333
SLC43A1 AND FLVCR1 AND NOT-MUC1	1	1	1
SLCO1B1 AND SMPD2 AND NOT-MSLN	1	1	1
SLCO1B1 AND SMPD2 AND NOT-TNFRSF10A	1	1	1
SLCO1B1 AND SMPD2 AND NOT-MSLN	1	1	1
SLC43A1 AND NOT-MUC1 AND KCNJ6	0.909091	1	0.833333
SLC43A1 AND NOT-MUC1 AND SMPD2	1	1	1
SLCO1B1 AND ZACN AND NOT-MSLN	0.909091	1	0.833333
DIO1 AND FLVCR1 AND NOT-MUC1	1	1	1
SLC43A1 AND NOT-MUC1 AND SLC22A12	0.909091	1	0.833333
SLCO1B1 AND DTNA AND NOT-MSLN	0.909091	1	0.833333
SLC43A1 AND NOT-MUC1 AND SCN2A	0.909091	1	0.833333

Antigen Pair Logic	F1	Precision	Recall
NOT-CLDN19 AND SGCD AND SLC31A2	0.952381	1	0.909091
NOT-CLDN19 AND SGCD AND ZNRF3	0.952381	1	0.909091
COMPLEX-CHRNA2/CNIH3/MCOLN1	0.952381	1	0.909091
NOT-CHRNA2 AND SGCD AND PMEPA1	0.952381	1	0.909091
NOT-CHRNA2 AND SGCD AND BRCA1	0.952381	1	0.909091
NOT-SLC5A5 AND TTYH2 AND FZD7	0.952381	1	0.909091
NOT-CHRNA2 AND SGCD AND FZD4	0.952381	1	0.909091
NOT-CHRNA2 AND SLC16A4 AND PCDHB14	0.952381	1	0.909091
NOT-CHRNA2 AND SGCD AND AKAP1	0.952381	1	0.909091
NOT-CHRNA2 AND TTYH2 AND EMP1	1	1	1
NOT-CHRNA2 AND TTYH2 AND TIE1	1	1	1
GPR137B AND TTYH2 AND NOT-CD300LB	1	1	1
NOT-CHRNA2 AND TTYH2 AND OSMR	1	1	1
GPR137B AND TTYH2 AND NOT-CCR3	1	1	1
GPR137B AND TTYH2 AND NOT-CD300LB	1	1	1
NOT-CHRNA2 AND TTYH2 AND ZNRF3	1	1	1
NOT-CHRNA2 AND TTYH2 AND ANO1	1	1	1
GPR137B AND TTYH2 AND ATP1A1	1	1	1
NOT-CHRNA2 AND TTYH2 AND TSPAN11	1	1	1
NOT-CHRNA2 AND TTYH2 AND WNT5A	1	1	1
NOT-CHRNA2 AND TTYH2 AND CAV2	1	1	1
NOT-CHRNA2 AND TTYH2 AND KCNK15	1	1	1
NOT-CHRNA2 AND TTYH2 AND FZD7	1	1	1
NOT-CHRNA2 AND TTYH2 AND TIE1	1	1	1
NOT-CHRNA2 AND TTYH2 AND TMEM231	1	1	1
GPR137B AND TTYH2 AND NOT-SPN	1	1	1
GPR137B AND TTYH2 AND NOT-AQP9	1	1	1
GPR137B AND TTYH2 AND NOT-CD300C	1	1	1
NOT-CHRNA2 AND TTYH2 AND FZD7	1	1	1
NOT-CHRNA2 AND TTYH2 AND FAM57A	1	1	1
NOT-CHRNA2 AND TTYH2 AND OSMR	1	1	1
NOT-CHRNA2 AND TTYH2 AND SGCD	1	1	1
NOT-CHRNA2 AND TTYH2 AND EMP1	1	1	1
NOT-CHRNA2 AND TTYH2 AND GPR161	1	1	1
NOT-CHRNA2 AND TTYH2 AND GJB2	1	1	1
NOT-CHRNA2 AND TTYH2 AND TMEM231	1	1	1
NOT-CHRNA2 AND TTYH2 AND TNFRSF19	1	1	1
NOT-CHRNA2 AND TTYH2 AND LGR4	1	1	1
NOT-CHRNA2 AND TTYH2 AND TSPAN11	1	1	1
NOT-CHRNA2 AND TTYH2 AND FXYD3	1	1	1
NOT-CHRNA2 AND TTYH2 AND XG	1	1	1
NOT-CHRNA2 AND TTYH2 AND PERP	1	1	1
NOT-CHRNA2 AND TTYH2 AND ANO1	1	1	1
NOT-CHRNA2 AND TTYH2 AND KCNK15	1	1	1
NOT-CHRNA2 AND TTYH2 AND SLCO2A1	1	1	1
NOT-CHRNA2 AND TTYH2 AND GJA4	1	1	1
NOT-CHRNA2 AND TTYH2 AND SCARB1	1	1	1
NOT-CHRNA2 AND TTYH2 AND WNT5A	1	1	1
NOT-CHRNA2 AND TTYH2 AND CAV1	1	1	1
NOT-CHRNA2 AND TTYH2 AND ZNRF3	1	1	1
NOT-CHRNA2 AND TTYH2 AND ITGA3	1	1	1
NOT-CHRNA2 AND TTYH2 AND GPR153	1	1	1
NOT-CHRNA2 AND TTYH2 AND XG	1	1	1
NOT-CHRNA2 AND TTYH2 AND PERP	1	1	1
NOT-CHRNA2 AND TTYH2 AND GJA4	1	1	1
NOT-CHRNA2 AND TTYH2 AND VANGL2	1	1	1
NOT-CHRNA2 AND TTYH2 AND FZD4	1	1	1
NOT-CHRNA2 AND TTYH2 AND PCDH18	1	1	1
NOT-CHRNA2 AND TTYH2 AND EMP2	1	1	1
NOT-CHRNA2 AND TTYH2 AND ADRA2C	1	1	1
NOT-CHRNA2 AND TTYH2 AND PCDHB16	1	1	1
NOT-CHRNA2 AND FAT1 AND NOT-KCNK5	1	1	1
NOT-CHRNA2 AND TTYH2 AND ANO10	1	1	1
NOT-CHRNA2 AND TTYH2 AND ANTXR1	1	1	1
NOT-CHRNA2 AND TTYH2 AND SDC3	1	1	1
NOT-CHRNA2 AND TTYH2 AND EMP2	1	1	1
NOT-CHRNA2 AND TTYH2 AND TSPAN6	1	1	1
NOT-CHRNA2 AND TTYH2 AND ADRA2C	1	1	1
NOT-CHRNA2 AND TTYH2 AND CNTNAP1	1	1	1
NOT-CHRNA2 AND TTYH2 AND TSPAN6	1	1	1
NOT-CHRNA2 AND FAT1 AND NOT-CADM2	1	1	1
NOT-CHRNA2 AND TTYH2 AND ADCY6	1	1	1
GPR137B AND TTYH2 AND NOT-SLC46A2	1	1	1
NOT-CHRNA2 AND TTYH2 AND PCDHB14	1	1	1
NOT-CHRNA2 AND TTYH2 AND FZD6	1	1	1
NOT-CHRNA2 AND OSMR AND NOT-ATP13A5	1	1	1
NOT-CHRNA2 AND TSPAN11 AND NOT-ATP13A5	1	1	1

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
ABCG5 AND ALCAM AND NOT-MSLN	1	1	1
ABCG5 AND ALCAM AND NOT-MUC1	1	1	1
SLC43A1 AND NOT-MUC1 AND DTNA	0.909091	1	0.833333
SLC2A2 AND DTNA AND MSLN	0.909091	1	0.833333
SLC43A1 AND NOT-MUC1 AND SLC22A5	0.909091	1	0.833333
SLC43A1 AND NOT-MUC1 AND PCDHB1	1	1	1
SLC43A1 AND NOT-MUC1 AND PCDHB1	1	1	1
SLC43A1 AND NOT-MUC1 AND GRM3	0.909091	1	0.833333
SLC43A1 AND GPR158 AND NOT-MUC1	0.909091	1	0.833333
ABCG5 AND GPR158 AND NOT-MUC1	0.909091	1	0.833333
SLC43A1 AND NOT-MUC1 AND MRGPRX2	0.909091	1	0.833333
ABCG5 AND DTNA AND NOT-MUC1	0.909091	1	0.833333
ABCG5 AND DTNA AND NOT-MSLN	0.909091	1	0.833333
SLC43A1 AND NOT-MUC1 AND PVRL1	1	1	1
SLC43A1 AND ZACN AND NOT-MUC1	0.909091	1	0.833333
SLC43A1 AND NOT-MUC1 AND DRD5	1	1	1
SLC43A1 AND NOT-MUC1 AND NOT-CD163	0.909091	1	0.833333
ABCG5 AND FGF6 AND NOT-MUC1	0.833333	0.833333	0.833333
SLCO1B1 AND SLC22A11 AND NOT-MSLN	0.833333	0.833333	0.833333
SLCO1B1 AND FGF6 AND NOT-MSLN	0.833333	0.833333	0.833333
SLC43A1 AND NOT-MUC1 AND KCNK16	0.909091	1	0.833333
SLC43A1 AND NOT-MUC1 AND KCNK16	0.909091	1	0.833333
DIO1 AND NOT-MUC1 AND ALCAM	1	1	1
SLC2A2 AND KCNJ6 AND MSLN	0.909091	1	0.833333
SLCO1B1 AND SLC12A9 AND NOT-MUC1	0.857143	0.75	1
SLC43A1 AND NOT-MUC1 AND RGSL1	0.909091	1	0.833333
SLC43A1 AND NOT-MUC1 AND LHFPL5	1	1	1
SLC2A2 AND TMEFF2 AND MSLN	0.833333	0.833333	0.833333
SLC43A1 AND NOT-MUC1 AND LHFPL5	1	1	1
SLC43A1 AND HTR1E AND NOT-MUC1	0.909091	1	0.833333
SLC2A2 AND SMPD2 AND MSLN	1	1	1
SLC2A2 AND ZACN AND MSLN	0.909091	1	0.833333
SLC43A1 AND NOT-MUC1 AND MMP24	0.909091	1	0.833333
SLC43A1 AND NOT-MUC1 AND ROS1	1	1	1
SLC43A1 AND NOT-MUC1 AND GABRA5	1	1	1
SLC43A1 AND NOT-MUC1 AND FUT3	0.909091	1	0.833333
SLC43A1 AND NOT-MUC1 AND OR7C1	0.909091	1	0.833333
SLC2A2 AND SLC22A11 AND MSLN	0.909091	1	0.833333
SLC43A1 AND NOT-MUC1 AND SLC30A8	0.909091	1	0.833333
SLCO1B1 AND SLC50A1 AND NOT-MSLN	1	1	1
SLC2A2 AND GABRA5 AND MSLN	0.857143	0.75	1
SLC43A1 AND NOT-MUC1 AND CALN1	0.909091	1	0.833333
SLC2A2 AND MSLN AND GPR61	0.909091	1	0.833333
SLC2A2 AND MSLN AND NOT-C8B	0.857143	0.75	1
SLC43A1 AND NOT-MUC1 AND CRB2	0.909091	1	0.833333
SLC2A2 AND MSLN AND SLCO1A2	0.857143	0.75	1
SLC2A2 AND MSLN AND NOT-MUC17	0.857143	0.75	1
SLC43A1 AND NOT-MUC1 AND OPCML	1	1	1
SLC43A1 AND NOT-MUC17 AND MSLN	1	1	1
SLC43A1 AND NOT-MUC1 AND SEZ6	1	1	1
SLC2A2 AND MSLN AND SLCO1A2	0.857143	0.75	1
SLC2A2 AND MSLN AND NOT-MUC17	0.857143	0.75	1
ABCG5 AND GPC3 AND NOT-OR1C1	0.8	1	0.666667
ABCG5 AND GPC3 AND NOT-SLC10A2	0.8	1	0.666667
ABCG5 AND GPC3 AND NOT-SLC22A6	0.8	1	0.666667
ABCG5 AND GPC3 AND NOT-LHFPL5	0.8	1	0.666667
ABCG5 AND GPC3 AND FLVCR1	0.8	1	0.666667
SLCO1B1 AND GPC3 AND SCN8A	0.8	1	0.666667
SLC2A2 AND GPC3 AND NOX1	0.8	1	0.666667
ABCG5 AND GPC3 AND NOT-PCDH11Y	0.8	1	0.666667
SLC2A2 AND GPC3 AND DRD5	0.8	1	0.666667
ABCG5 AND GPC3 AND NOT-ATP8A1	0.8	1	0.666667
ABCG5 AND GPC3 AND NOT-GPR6	0.8	1	0.666667
SLCO1B1 AND MSLN AND NOT-SLC17A2	0.8	0.666667	1
ABCG5 AND GPC3 AND NOT-ADAM21	0.8	1	0.666667
ABCG5 AND GPC3 AND NOT-SLC9A1	0.8	1	0.666667
ABCG5 AND GPC3 AND NOT-GSG1L	0.8	1	0.666667
ABCG5 AND GPC3 AND NOT-KCNK10	0.8	1	0.666667
ABCG5 AND GPC3 AND PVRL1	0.8	1	0.666667
ABCG8 AND SLC2A2 AND MSLN	0.8	1	0.666667
ABCG5 AND GPC3 AND NOT-CCR9	0.8	1	0.666667
SLC2A2 AND MSLN AND CSPG5	1	1	1
DIO1 AND GPC3 AND FLVCR1	0.8	1	0.666667
ABCG5 AND GPC3 AND NOT-CATSPER3	0.8	1	0.666667
ABCG5 AND GPC3 AND NOT-CD1A	0.8	1	0.666667
ABCG5 AND GPC3 AND NOT-AJAP1	0.8	1	0.666667
ABCG5 AND GPC3 AND NOT-ROS1	0.8	1	0.666667

Antigen Pair Logic	F1	Precision	Recall
NOT-CHRNA2 AND EMP1 AND NOT-ATP13A5	1	1	1
NOT-CHRNA2 AND TTYH2 AND PCDHB14	1	1	1
NOT-CHRNA2 AND TIE1 AND NOT-ATP13A5	1	1	1
NOT-CHRNA2 AND TTYH2 AND FZD6	1	1	1
NOT-CHRNA2 AND FZD7 AND NOT-ATP13A5	1	1	1
NOT-CHRNA2 AND TTYH2 AND CNIH3	1	1	1
NOT-TM4SF5 AND TTYH2 AND LRIG3	1	1	1
NOT-CHRNA2 AND TTYH2 AND MCOLN3	1	1	1
GPR137B AND NOT-ATP13A5 AND VANGL2	1	1	1
NOT-CHRNA2 AND LGR4 AND NOT-ATP13A5	1	1	1
GPR137B AND NOT-ATP13A5 AND TSPAN11	1	1	1
NOT-CHRNA2 AND CAV1 AND NOT-ATP13A5	1	1	1
NOT-CHRNA2 AND GJA4 AND NOT-ATP13A5	1	1	1
NOT-CHRNA2 AND FAT1 AND NOT-CFTR	1	1	1
NOT-CHRNA2 AND TTYH2 AND IGDCC4	1	1	1
NOT-CHRNA2 AND TTYH2 AND PDGFRA	1	1	1
NOT-CHRNA2 AND TTYH2 AND PDGFRA	1	1	1
NOT-CHRNA2 AND TTYH2 AND PMP22	1	1	1
NOT-CHRNA2 AND TTYH2 AND GJA1	1	1	1
NOT-CHRNA2 AND TTYH2 AND APCDD1	1	1	1
NOT-CHRNA2 AND TTYH2 AND FZD3	0.956522	0.916667	1
NOT-CHRNA2 AND TTYH2 AND ADAM12	0.956522	0.916667	1
GPR137B AND TTYH2 AND SLC12A1	0.956522	0.916667	1
NOT-CHRNA2 AND PCDHB14 AND NOT-ATP13A5	1	1	1
NOT-CHRNA2 AND SORT1 AND NOT-PCDH8	1	1	1
NOT-CHRNA2 AND ADRA2C AND NOT-ATP13A5	1	1	1
NOT-CHRNA2 AND TTYH2 AND ITGA2	1	1	1
NOT-TM4SF5 AND TTYH2 AND NOT-S1PR1	1	1	1
GPR137B AND NOT-ATP13A5 AND NOT-SLC7A7	1	1	1
NOT-CHRNA2 AND TTYH2 AND WLS	0.956522	0.916667	1
NOT-CHRNA2 AND FAT1 AND NOT-SCNN1G	1	1	1
NOT-CHRNA2 AND TTYH2 AND KITLG	1	1	1
NOT-CHRNA2 AND LGR4 AND NOT-CLDN8	1	1	1
NOT-CLDN19 AND GPNMB AND NOT-KCNK5	1	1	1
NOT-TM4SF5 AND TTYH2 AND GPNMB	1	1	1
NOT-CLDN19 AND GPNMB AND NOT-CFTR	1	1	1
NOT-CHRNA2 AND SDC1 AND NOT-CFTR	1	1	1
NOT-CHRNA2 AND TIE1 AND NOT-CLDN8	1	1	1
NOT-CLDN19 AND GPNMB AND NOT-CADM2	1	1	1
NOT-CHRNA2 AND TMEM231 AND NOT-ERBB2	1	1	1
GPR137B AND NOT-NPR3 AND ERBB3	1	1	1
GPR137B AND NOT-CLDN8 AND TYRO3	1	1	1
NOT-CHRNA2 AND GPNMB AND NOT-KCNK5	1	1	1
NOT-CHRNA2 AND SDC1 AND NOT-KCNK5	1	1	1
GPR137B AND NOT-CLDN8 AND TYRO3	1	1	1
GPR137B AND NOT-CLDN8 AND KCNS3	1	1	1
NOT-CHRNA2 AND PCDHB14 AND NOT-CLDN8	1	1	1
NOT-CHRNA2 AND TMEM231 AND NOT-ERBB2	1	1	1
GPR137B AND NOT-CLDN8 AND ATP1A1	1	1	1
NOT-TM4SF5 AND TTYH2 AND SDC1	0.956522	0.916667	1
NOT-TM4SF5 AND GPNMB AND NOT-ATP13A5	1	1	1
NOT-TM4SF5 AND TTYH2 AND TPBG	1	1	1
NOT-SLC5A5 AND TTYH2 AND TPBG	0.952381	1	0.909091
NOT-GIPR AND TTYH2 AND GPNMB	0.952381	1	0.909091
NOT-CHRNA2 AND NOT-CADM2 AND THY1	0.952381	1	0.909091
GPR137B AND NOT-CLDN8 AND APOLD1	0.952381	1	0.909091
NOT-CLDN19 AND ENG AND NOT-GLDN	0.952381	1	0.909091
NOT-CHRNA2 AND LGR4 AND NOT-VTCN1	0.952381	1	0.909091
NOT-SLC5A5 AND TTYH2 AND GPNMB	0.952381	1	0.909091
GPR137B AND NOT-CADM2 AND ERBB2	0.952381	1	0.909091
NOT-CLDN19 AND GPNMB AND NOT-LYPD6B	0.952381	1	0.909091
GPR137B AND SLC5A6 AND NOT-CLDN8	0.952381	1	0.909091
NOT-CLDN19 AND GPNMB AND NOT-TSPAN8	0.952381	1	0.909091
GPR137B AND NOT-NTRK3 AND ERBB3	0.952381	1	0.909091
GPR137B AND NOT-CADM2 AND ERBB2	0.952381	1	0.909091
COMPLEX-CHRNA2/MST1R/SGCD	0.952381	1	0.909091
GPR137B AND NOT-TSPAN8 AND ERBB3	0.952381	1	0.909091
NOT-CHRNA2 AND SLC16A4 AND SDC1	0.952381	1	0.909091
NOT-CHRNA2 AND SGCD AND IL11RA	0.952381	1	0.909091
COMPLEX-CHRNA2/SGCD/TPBG	0.952381	1	0.909091
GPR137B AND NOT-CLDN8 AND IGSF8	0.952381	1	0.909091
GPR137B AND NOT-NPR3 AND L1CAM	0.952381	1	0.909091
NOT-CHRNA2 AND TMEM231 AND NOT-EGFR	0.952381	1	0.909091
NOT-CHRNA2 AND SDC1 AND NOT-TSPAN8	0.952381	1	0.909091
NOT-CHRNA2 AND IGSF8 AND NOT-ERBB2	0.952381	1	0.909091
NOT-CHRNA2 AND SORT1 AND NOT-ERBB2	1	1	1
NOT-CHRNA2 AND PVRL2 AND NOT-ERBB2	0.952381	1	0.909091

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
ABCG5 AND GPC3 AND NOT-SLC39A2	0.8	1	0.666667
SLC43A1 AND CNTNAP4 AND NOT-MUC1	0.8	1	0.666667
SLC43A1 AND MLANA AND NOT-MUC1	0.8	1	0.666667
SLCO1B1 AND CALN1 AND NOT-MUC1	0.8	0.666667	1
SLC43A1 AND NOT-MUC1 AND NPBWR1	0.8	1	0.666667
SLC43A1 AND FAT1 AND NOT-MUC1	0.8	1	0.666667
ABCG5 AND GPC3 AND NOT-MSLN	0.8	1	0.666667
SLC43A1 AND NOT-MUC1 AND TNFRSF10A	0.8	1	0.666667
ABCG5 AND GPC3 AND NOT-MUC1	0.8	1	0.666667
SLC2A2 AND GPC3 AND MSLN	0.8	1	0.666667
SLCO1B1 AND GPC3 AND NOT-MSLN	0.8	1	0.666667
ASGR1 AND GPC3 AND NOT-MSLN	0.8	1	0.666667
Pancreatic Cancer			
KCNE4 AND IFNAR2 AND NOT-TSPAN14	0.842105	1	0.727273
KCNE4 AND IFNAR2 AND NOT-P2RY8	0.842105	1	0.727273
SYT13 AND ITGAM AND NOT-GRM8	0.869565	0.833333	0.909091
KCNE4 AND IFNAR2 AND NOT-ITGAD	0.842105	1	0.727273
KCNE4 AND IFNAR2 AND NOT-TMEM5	0.842105	1	0.727273
NOX4 AND NOT-SLC47A1 AND GJB4	0.818182	0.818182	0.818182
SYT13 AND IFNAR2 AND PCDHB16	1	1	1
NTM AND NOT-RTN3 AND IFNAR2	0.842105	1	0.727273
NOX4 AND NOT-SLC47A1 AND GRM8	0.857143	0.9	0.818182
NOX4 AND NOT-SLC47A1 AND CSMD2	0.9	1	0.818182
NALCN AND NOT-LRP4 AND ATP8B1	0.857143	0.9	0.818182
SYT13 AND IFNAR2 AND NOT-RTN3	0.956522	0.916667	1
SYT13 AND IFNAR2 AND NOT-PTPRD	0.857143	0.9	0.818182
SYT13 AND ITGAM AND NOT-GJB4	0.869565	0.833333	0.909091
TNFRSF12A AND PCDH7 AND VNN2	0.818182	0.818182	0.818182
NALCN AND LRIG3 AND NOT-XG	0.857143	0.9	0.818182
SYT13 AND VNN2 AND NOT-ADAM20	0.9	1	0.818182
NTM AND NOT-CELSR2 AND CSMD2	0.842105	1	0.727273
SLC39A14 AND NOT-LRP4 AND GRM8	0.857143	0.9	0.818182
NTM AND OTOF AND IFNAR2	0.842105	1	0.727273
NOX4 AND OTOF AND IFNAR2	0.9	1	0.818182
GP2 AND IFNAR2 AND NOT-SHISA9	0.842105	1	0.727273
NOX4 AND USH2A AND IFNAR2	0.842105	1	0.727273
TNFRSF12A AND IFNAR2 AND NOT-RTN3	0.9	1	0.818182
TNFRSF12A AND OR7C1 AND FAM26F	0.9	1	0.818182
TNFRSF12A AND CACNG2 AND FAM26F	0.9	1	0.818182
PTGIS AND NOT-XG AND LRIG3	0.814815	0.6875	1
TNFRSF12A AND GJB4 AND FAM26F	0.9	1	0.818182
GP2 AND IFNAR2 AND NOT-PCDH8	0.842105	1	0.727273
NTM AND USH2A AND LRIG3	0.842105	1	0.727273
TNFRSF12A AND GPR1 AND IFNAR2	0.9	1	0.818182
FAP AND NOT-GJC1 AND OTOF	0.842105	1	0.727273
FAP AND NOT-GJC1 AND CSMD2	0.842105	1	0.727273
FAP AND NOT-GJC1 AND GJB4	0.842105	1	0.727273
FAP AND NOT-GJC1 AND SHISA9	0.842105	1	0.727273
FAP AND SLC44A3 AND NOT-SLC6A15	0.842105	1	0.727273
SST AND ITGAM AND NOT-GJB4	0.842105	1	0.727273
FAP AND NOT-SLC47A1 AND GRIN2B	0.842105	1	0.727273
FAP AND NOT-SLC47A1 AND GJB4	0.842105	1	0.727273
THY1 AND IFNAR2 AND NOT-RTN3	0.9	1	0.818182
FAP AND SLC44A3 AND NOT-GRM3	0.842105	1	0.727273
THY1 AND VNN2 AND FCAMR	0.842105	1	0.727273
FAP AND NOT-SLC47A1 AND LCT	0.842105	1	0.727273
FAP AND NOT-SLC47A1 AND GRM8	0.842105	1	0.727273
FAP AND SLC44A3 AND NOT-ADCY10	0.842105	1	0.727273
FAP AND SLC44A3 AND NOT-CSMD2	0.842105	1	0.727273
SST AND DCBLD2 AND NOT-GRM8	0.842105	1	0.727273
FAP AND IL2RG AND NOT-PTPRT	0.842105	1	0.727273
DCBLD2 AND NOT-RAET1E AND OTOF	0.842105	1	0.727273
FAP AND LTB AND NOT-PTPRT	0.842105	1	0.727273
FAP AND LTB AND NOT-SLC39A2	0.842105	1	0.727273
FAP AND IL2RG AND NOT-SLC6A15	0.842105	1	0.727273
THY1 AND VNN2 AND DIO1	0.842105	1	0.727273
THY1 AND VNN2 AND GRM8	0.842105	1	0.727273
THY1 AND VNN2 AND GRIN2B	0.842105	1	0.727273
THY1 AND VNN2 AND CNTNAP4	0.842105	1	0.727273
THY1 AND VNN2 AND SLC30A10	0.842105	1	0.727273
FAP AND SLC44A3 AND NOT-IFNAR2	0.842105	1	0.727273
FAP AND SLC44A3 AND NOT-LCT	0.842105	1	0.727273
FAP AND SLC44A3 AND NOT-GLRA3	0.842105	1	0.727273
FAP AND SLC44A3 AND NOT-GPRC6A	0.842105	1	0.727273
FAP AND SLC44A3 AND NOT-SLC22A6	0.842105	1	0.727273
FAP AND SLC44A3 AND NOT-VN1R2	0.842105	1	0.727273
FAP AND SLC44A3 AND NOT-GRM8	0.842105	1	0.727273

Antigen Pair Logic	F1	Precision	Recall
NOT-CHRNA2 AND ITGB5 AND NOT-ERBB2	0.952381	1	0.909091
NOT-CHRNA2 AND LGR4 AND NOT-ERBB2	1	1	1
GPR137B AND NOT-ERBB2 AND TYRO3	1	1	1
NOT-CHRNA2 AND GPNMB AND NOT-CFTR	1	1	1
NOT-CHRNA2 AND SORT1 AND NOT-ERBB2	1	1	1
NOT-CHRNA2 AND LGR4 AND NOT-ERBB2	1	1	1
NOT-CLDN19 AND GPNMB AND NOT-ACSL1	0.952381	1	0.909091
GPR137B AND NOT-CLDN8 AND VANGL2	1	1	1
NOT-CHRNA2 AND ZNRF3 AND NOT-CLDN8	1	1	1
GPR137B AND NOT-CLDN8 AND PCDH7	1	1	1
COMPLEX-CLDN19/SGCD/TPBG	0.952381	1	0.909091
GPR137B AND NOT-CADM2 AND ERBB3	1	1	1
GPR137B AND NOT-ATP1A2 AND ERBB3	1	1	1
NOT-CHRNA2 AND NOT-CADM2 AND VCAM1	1	1	1
NOT-CHRNA2 AND FZD7 AND NOT-CLDN8	1	1	1
NOT-CLDN19 AND GPNMB AND NOT-NPR3	1	1	1
NOT-CHRNA2 AND GPNMB AND NOT-CADM2	1	1	1
NOT-CHRNA2 AND GPNMB AND NOT-TSPAN8	0.952381	1	0.909091
GPR137B AND NOT-CADM2 AND ERBB3	1	1	1
NOT-CLDN19 AND GPNMB AND NOT-SYT8	0.952381	1	0.909091
NOT-CHRNA2 AND VCAM1 AND NOT-TSPAN8	0.952381	1	0.909091
GPR137B AND NOT-CLDN8 AND FAM57A	0.952381	1	0.909091
GPR137B AND NOT-CLDN8 AND VANGL2	1	1	1
GPR137B AND NOT-CLDN8 AND NOTCH4	0.952381	1	0.909091
NOT-CLDN19 AND LGR4 AND NOT-CLDN8	1	1	1
NOT-CHRNA2 AND NOT-ATP1A2 AND THY1	0.952381	1	0.909091
GPR137B AND SLC5A6 AND NOT-ERBB2	0.952381	1	0.909091
NOT-CHRNA2 AND TSPAN11 AND NOT-CLDN8	1	1	1
NOT-CHRNA2 AND VCAM1 AND NOT-KCNK5	1	1	1
NOT-CHRNA2 AND NOT-CADM2 AND CD34	0.952381	1	0.909091
NOT-CHRNA2 AND NOT-CADM2 AND VCAM1	1	1	1
NOT-CHRNA2 AND SDC1 AND NOT-SCNN1G	1	1	1
NOT-CACNG6 AND SDC1 AND NOT-TSPAN8	0.952381	1	0.909091
GPR137B AND NOT-CLDN8 AND JAG1	0.952381	1	0.909091
NOT-CLDN19 AND VCAM1 AND NOT-CADM2	1	1	1
GPR137B AND NOT-NPR3 AND ERBB2	0.952381	1	0.909091
GPR137B AND NOT-KCNK5 AND ERBB2	0.952381	1	0.909091
GPR137B AND NOT-ATP1A2 AND ERBB2	0.952381	1	0.909091
GPR137B AND NOT-KCNK5 AND ERBB2	0.952381	1	0.909091
GPR137B AND NOT-ATP1A2 AND ERBB2	0.952381	1	0.909091
NOT-CACNG6 AND LGR4 AND NOT-CLDN8	1	1	1
GPR137B AND NOT-IL20RA AND TYRO3	0.952381	1	0.909091
NOT-CHRNA2 AND PTPRJ AND GPNMB	0.952381	1	0.909091
GPR137B AND NOT-CLDN8 AND SLC19A2	0.952381	1	0.909091
NOT-CLDN19 AND GPNMB AND NOT-NTRK3	0.952381	1	0.909091
GPR137B AND NOT-NPR3 AND ERBB2	0.952381	1	0.909091
NOT-CLDN19 AND GPNMB AND NOT-ATP1A2	0.952381	1	0.909091
NOT-CLDN19 AND VCAM1 AND NOT-TSPAN8	0.952381	1	0.909091
GPR137B AND NOT-IL20RA AND VANGL2	0.952381	1	0.909091
NOT-CHRNA2 AND BEST1 AND GPNMB	0.952381	1	0.909091
NOT-CHRNA2 AND TMEM231 AND NOT-CLDN8	1	1	1
NOT-CHRNA2 AND LGR4 AND NOT-IL20RA	0.952381	1	0.909091
NOT-CHRNA2 AND WLS AND NOT-CLDN8	1	1	1
NOT-CHRNA2 AND SDC1 AND NOT-CADM2	0.952381	1	0.909091
NOT-CHRNA2 AND SDC1 AND NOT-CADM2	0.952381	1	0.909091
COMPLEX-LDLRAD3/PMEL/SLC23A2	1	1	1
COMPLEX-MUC12/LDLRAD3/PMEL	1	1	1
COMPLEX-MLANA/CDH20/FREM2	1	1	1
COMPLEX-MLANA/NOX1/FREM2	1	1	1
COMPLEX-ZACN/PMEL/RNF144A	1	1	1
COMPLEX-GJA3/PMEL/RNF144A	1	1	1
COMPLEX-ACSL6/PMEL/RNF144A	1	1	1
COMPLEX-EFNB2/PMEL/RNF144A	1	1	1
COMPLEX-HTRA2/PMEL/SLC2A1	1	1	1
COMPLEX-SEMA4B/PMEL/RNF144A	1	1	1
COMPLEX-SLC6A15/PMEL/RNF144A	1	1	1
COMPLEX-GPR158/PMEL/RNF144A	1	1	1
COMPLEX-MLANA/OR10J1/FREM2	1	1	1
COMPLEX-LDLRAD3/PMEL/CD163	1	1	1
COMPLEX-MLANA/SCARA5/FREM2	1	1	1
COMPLEX-ABCA12/HTRA2/PMEL	1	1	1
COMPLEX-MLANA/FREM2/ATP13A5	1	1	1
COMPLEX-TNFRSF12A/PMEL/RNF144A	1	1	1
COMPLEX-MLANA/FREM2/NPFFR1	1	1	1
COMPLEX-MLANA/FREM2/CCKBR	1	1	1
COMPLEX-CDH18/LDLRAD3/PMEL	1	1	1
COMPLEX-MLANA/TMEM235/FREM2	1	1	1

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
FAP AND SLC44A3 AND NOT-CDH8	0.842105	1	0.727273
FAP AND VNN2 AND NOT-PTPRT	0.842105	1	0.727273
FAP AND LTB AND NOT-ADAM20	0.842105	1	0.727273
FAP AND VNN2 AND NOT-GRM3	0.842105	1	0.727273
FAP AND IL2RG AND NOT-GRM3	0.842105	1	0.727273
SST AND IFNAR2 AND NOT-PTPRH	0.842105	1	0.727273
FAP AND LTB AND NOT-DSCAM	0.842105	1	0.727273
FAP AND VNN2 AND NOT-KCNJ9	0.842105	1	0.727273
FAP AND LTB AND NOT-SLC6A15	0.842105	1	0.727273
FAP AND IL2RG AND NOT-MDGA2	0.842105	1	0.727273
FAP AND LTB AND NOT-KCNA1	0.842105	1	0.727273
FAP AND VNN2 AND NOT-KCNA1	0.842105	1	0.727273
FAP AND VNN2 AND NOT-CNGA4	0.842105	1	0.727273
FAP AND VNN2 AND NOT-ADAM20	0.842105	1	0.727273
FAP AND VNN2 AND NOT-ADCY10	0.842105	1	0.727273
STEAP1 AND NOT-LRP4 AND GRM8	0.833333	0.769231	0.909091
THY1 AND VNN2 AND GJB4	0.842105	1	0.727273
THY1 AND VNN2 AND PTPRT	0.842105	1	0.727273
THY1 AND VNN2 AND NOT-ADAM20	0.842105	1	0.727273
THY1 AND LRIG3 AND NOT-XG	0.842105	1	0.727273
THY1 AND VNN2 AND GRM3	0.842105	1	0.727273
THY1 AND VNN2 AND KCNA1	0.842105	1	0.727273
FAP AND SLC44A3 AND NOT-ACSL6	0.842105	1	0.727273
NTM AND NOT-MAL AND CLDN1	0.842105	1	0.727273
THY1 AND NOT-MAL AND SLC44A3	0.857143	0.9	0.818182
NTM AND NOT-MAL AND ERBB3	0.842105	1	0.727273
NTM AND NOT-MAL AND TNFRSF10A	0.842105	1	0.727273
THY1 AND NOT-MAL AND TMEM30B	0.9	1	0.818182
NOX4 AND NOT-SLC47A1 AND CLDN1	0.9	1	0.818182
THY1 AND NOT-MAL AND IL2RG	0.9	1	0.818182
FAP AND NOT-GJC1 AND PTPRH	0.842105	1	0.727273
NOX4 AND NOT-MAL AND ERBB3	0.9	1	0.818182
THY1 AND NOT-MAL AND LY75	0.9	1	0.818182
FAP AND LTB AND NOT-LRFN5	0.842105	1	0.727273
GPR68 AND NOT-DUOX1 AND CLDN1	0.869565	0.833333	0.909091
FAP AND NOT-SLC47A1 AND PTPRH	0.842105	1	0.727273
ABCC3 AND CLDN1 AND NOT-SLC47A1	0.842105	1	0.727273
FAP AND SLC44A3 AND NOT-MAL	0.842105	1	0.727273
FAP AND SLC44A3 AND NOT-SCN3B	0.842105	1	0.727273
NOX4 AND NOT-MAL AND TNFRSF10A	0.842105	1	0.727273
FAP AND VNN2 AND NOT-LRFN5	0.842105	1	0.727273
NALCN AND NOT-LRP4 AND SDC1	0.857143	0.9	0.818182
THY1 AND VNN2 AND SLC44A3	0.842105	1	0.727273
FAP AND SLC44A3 AND NOT-CELSR1	0.842105	1	0.727273
FAP AND SLC44A3 AND NOT-CLEC2B	0.842105	1	0.727273
FAP AND SLC44A3 AND NOT-GPR1	0.842105	1	0.727273
FAP AND NOT-SLC47A1 AND SLC44A3	0.842105	1	0.727273
FAP AND SLC44A3 AND NOT-LYPD6B	0.842105	1	0.727273
FAP AND SLC44A3 AND NOT-PTPRZ1	0.842105	1	0.727273
THY1 AND SLC44A3 AND NOT-SLC47A1	0.857143	0.9	0.818182
THY1 AND VNN2 AND NOT-RTN3	0.842105	1	0.727273
THY1 AND VNN2 AND PCDH7	0.842105	1	0.727273
FAP AND VNN2 AND NOT-CD300LG	0.842105	1	0.727273
FAP AND LTB AND NOT-IL1RL2	0.842105	1	0.727273
FAP AND NOT-SLC47A1 AND SLC5A4	0.842105	1	0.727273
FAP AND NOT-SLC47A1 AND ABCC9	0.842105	1	0.727273
FAP AND IGSF6 AND NOT-LRFN5	0.842105	1	0.727273
FAP AND IL1R2 AND NOT-TSPAN14	0.842105	1	0.727273
FAP AND IL2RG AND NOT-MAL	0.842105	1	0.727273
FAP AND NOT-SLC47A1 AND GRIK3	0.842105	1	0.727273
FAP AND TMEM30B AND NOT-GPR1	0.842105	1	0.727273
FAP AND NOT-SLC47A1 AND PAQR9	0.842105	1	0.727273
NTM AND NOT-CELSR2 AND EPHB2	0.842105	1	0.727273
THY1 AND VNN2 AND NOT-GJC1	0.842105	1	0.727273
FAP AND SLC44A3 AND NOT-KCNA5	0.842105	1	0.727273
THY1 AND NOT-MAL AND ERBB3	0.818182	0.818182	0.818182
FAP AND ERBB3 AND NOT-MAL	0.842105	1	0.727273
FAP AND NOT-SLC47A1 AND CLDN1	0.842105	1	0.727273
FAP AND SLC44A3 AND NOT-NCAM1	0.842105	1	0.727273
THY1 AND NOT-MAL AND TNFRSF10A	0.842105	1	0.727273
FAP AND NOT-SLC47A1 AND L1CAM	0.842105	1	0.727273
FAP AND ERBB3 AND NOT-CELSR2	0.842105	1	0.727273
FAP AND NOT-SLC47A1 AND ABCB5	0.842105	1	0.727273
FAP AND LTB AND NOT-ERBB4	0.842105	1	0.727273
FAP AND ERBB3 AND NOT-RTN3	0.842105	1	0.727273
FAP AND IL2RG AND NOT-L1CAM	0.842105	1	0.727273
FAP AND IL2RG AND NOT-CLDN1	0.842105	1	0.727273

Antigen Pair Logic	F1	Precision	Recall
COMPLEX-ATP8A1/LDLRAD3/PMEL	1	1	1
COMPLEX-CLCN1/LDLRAD3/PMEL	1	1	1
COMPLEX-TMPRSS11B/LDLRAD3/PMEL	1	1	1
COMPLEX-LDLRAD3/PMEL/OR2C3	1	1	1
COMPLEX-LDLRAD3/PMEL/TRPM8	1	1	1
COMPLEX-LDLRAD3/PMEL/SYT4	1	1	1
COMPLEX-LDLRAD3/PMEL/SLC6A6	1	1	1
COMPLEX-LDLRAD3/PMEL/PCDH15	1	1	1
COMPLEX-LDLRAD3/PMEL/SLC1A2	1	1	1
COMPLEX-LDLRAD3/OR1Q1/PMEL	1	1	1
COMPLEX-LDLRAD3/LRRC52/PMEL	1	1	1
COMPLEX-LDLRAD3/IL13/PMEL	1	1	1
COMPLEX-LDLRAD3/SLC6A18/PMEL	1	1	1
COMPLEX-LDLRAD3/NOX1/PMEL	1	1	1
COMPLEX-LDLRAD3/EMP3/PMEL	1	1	1
COMPLEX-LDLRAD3/DIO1/PMEL	1	1	1
COMPLEX-MLANA/GHR/FREM2	1	1	1
COMPLEX-LDLRAD3/F2R/PMEL	1	1	1
COMPLEX-LDLRAD3/KCNV1/PMEL	1	1	1
COMPLEX-SLCO6A1/LDLRAD3/PMEL	1	1	1
COMPLEX-ANTXR2/LDLRAD3/PMEL	1	1	1
COMPLEX-SLCO1B1/LDLRAD3/PMEL	1	1	1
COMPLEX-LDLRAD3/PMEL/CNTNAP4	1	1	1
COMPLEX-LDLRAD3/PMEL/GJA10	1	1	1
COMPLEX-LDLRAD3/PMEL/GLRA3	1	1	1
COMPLEX-LDLRAD3/PMEL/STEAP4	1	1	1
COMPLEX-LDLRAD3/PMEL/SHISA9	1	1	1
COMPLEX-LDLRAD3/TMPRSS11E/PMEL	1	1	1
COMPLEX-LDLRAD3/PMEL/CD58	1	1	1
COMPLEX-LDLRAD3/PMEL/GGTLC1	1	1	1
COMPLEX-LDLRAD3/PMEL/ADAM21	1	1	1
COMPLEX-LDLRAD3/PMEL/GPR137B	1	1	1
COMPLEX-LDLRAD3/CACNG6/PMEL	1	1	1
COMPLEX-LDLRAD3/TRPC7/PMEL	1	1	1
NOT-CHRNA2 AND TTYH2 AND LRIG3	1	1	1
NOT-CHRNA2 AND TTYH2 AND LRIG3	1	1	1
NOT-CHRNA2 AND TTYH2 AND FAT1	1	1	1
NOT-CHRNA2 AND TTYH2 AND PLVAP	1	1	1
NOT-CHRNA2 AND FAT1 AND NOT-ATP13A5	1	1	1
NOT-CHRNA2 AND TTYH2 AND PLVAP	1	1	1
NOT-CHRNA2 AND TTYH2 AND EFNB2	1	1	1
NOT-CHRNA2 AND TTYH2 AND GHR	1	1	1
NOT-CHRNA2 AND TTYH2 AND GHR	1	1	1
NOT-CHRNA2 AND TTYH2 AND NOT-SORL1	0.956522	0.916667	1
NOT-CHRNA2 AND NOT-CLDN10 AND LRIG3	1	1	1
GPR137B AND TTYH2 AND NOT-MPL	0.952381	1	0.909091
NOT-CHRNA2 AND TTYH2 AND TNFRSF12A	0.952381	1	0.909091
NOT-CHRNA2 AND TTYH2 AND CDH11	0.952381	1	0.909091
NOT-CLDN19 AND TTYH2 AND LRIG3	0.952381	1	0.909091
COMPLEX-CALHM3/SLC6A15/GPR137B	0.952381	1	0.909091
NOT-CHRNA2 AND TTYH2 AND ENPP1	0.952381	1	0.909091
NOT-CHRNA2 AND EMP3 AND FAT1	0.952381	1	0.909091
NOT-CHRNA2 AND EMP3 AND LRIG3	0.952381	1	0.909091
NOT-CLDN19 AND FAT1 AND NOT-ATP8B1	0.952381	1	0.909091
NOT-CLDN19 AND TTYH2 AND NOT-S1PR1	0.952381	1	0.909091
GPR137B AND NOT-CLDN10 AND FAT1	1	1	1
NOT-CHRNA2 AND FAT1 AND NOT-CLDN10	1	1	1
NOT-SLC10A2 AND TTYH2 AND LRIG3	0.952381	1	0.909091
NOT-CHRNA2 AND TTYH2 AND ATP6V0A4	0.952381	1	0.909091
NOT-CHRNA2 AND NOT-ATP13A5 AND LRIG3	0.952381	1	0.909091
NOT-CHRNA2 AND TTYH2 AND LRRC8E	0.952381	1	0.909091
NOT-CHRNA2 AND FAT1 AND NOT-CD207	0.952381	1	0.909091
NOT-CHRNA2 AND FAT1 AND NOT-PCDH8	1	1	1
NOT-CLDN19 AND LRIG3 AND NOT-CLDN10	1	1	1
NOT-CLDN19 AND VANGL1 AND NOT-ATP8B1	0.952381	1	0.909091
NOT-CHRNA2 AND FAT1 AND NOT-SLC1A6	1	1	1
GPR137B AND NOT-LCT AND FAT1	0.952381	1	0.909091
NOT-CHRNA2 AND NOT-ATP13A5 AND PLVAP	0.952381	1	0.909091
NOT-CHRNA2 AND FAT1 AND NOT-GHR	1	1	1
NOT-CHRNA2 AND FAT1 AND NOT-CEACAM7	1	1	1
NOT-CHRNA2 AND FAT1 AND NOT-GHR	1	1	1
NOT-CACNG6 AND FAT1 AND NOT-ATP8B1	0.952381	1	0.909091
NOT-CHRNA2 AND FAT1 AND NOT-AOC3	0.952381	1	0.909091
NOT-CHRNA2 AND FAT1 AND NOT-SLC30A10	0.952381	1	0.909091
NOT-CLDN19 AND LRIG3 AND NOT-ATP8B1	1	1	1
GPR137B AND NOT-P2RY4 AND FAT1	0.952381	1	0.909091
GPR137B AND NOT-ATP8B1 AND FAT1	1	1	1

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
THY1 AND NOT-MAL AND SDC1	0.9	1	0.818182
FAP AND ERBB3 AND NOT-PTPRZ1	0.842105	1	0.727273
FAP AND ERBB3 AND NOT-LYPD6B	0.842105	1	0.727273
FAP AND LTB AND NOT-IL20RA	0.842105	1	0.727273
THY1 AND VNN2 AND IL20RA	0.842105	1	0.727273
THY1 AND VNN2 AND ERBB3	0.842105	1	0.727273
FAP AND SLC44A3 AND NOT-FCRL5	0.842105	1	0.727273
FAP AND IL2RG AND NOT-ROR1	0.842105	1	0.727273
FAP AND LTB AND NOT-SDC1	0.842105	1	0.727273
FAP AND VNN2 AND NOT-IL20RA	0.842105	1	0.727273
FAP AND VNN2 AND NOT-SDC1	0.842105	1	0.727273
THY1 AND NOT-MAL AND FOLH1	0.818182	0.818182	0.818182
SST AND CLDN1 AND IL2RG	0.842105	1	0.727273
FAP AND CLEC7A AND NOT-CLDN1	0.842105	1	0.727273
SST AND CLEC7A AND NOT-GUCY2C	0.842105	1	0.727273
FAP AND FAM26F AND NOT-ERBB4	0.842105	1	0.727273
FAP AND FOLH1 AND NOT-MAL	0.842105	1	0.727273
FAP AND EPHB2 AND NOT-GPR1	0.842105	1	0.727273
SST AND CLDN1 AND PTGER2	0.842105	1	0.727273
FAP AND FOLH1 AND NOT-RTN3	0.842105	1	0.727273
FAP AND PTPRC AND NOT-ABCB5	0.842105	1	0.727273
FAP AND PTPRC AND NOT-IL20RA	0.842105	1	0.727273
STEAP2 AND CLDN1 AND NOT-CELSR1	0.869565	0.833333	0.909091
FAP AND IGSF6 AND NOT-SDC1	0.842105	1	0.727273
FAP AND ERBB3 AND NOT-PTPRC	0.842105	1	0.727273
FAP AND ERBB3 AND KCNJ3	0.842105	1	0.727273
SST AND CLEC7A AND CLDN1	0.842105	1	0.727273
SST AND CLDN1 AND FZD10	0.842105	1	0.727273
FAP AND FCRL5 AND NOT-NRP2	0.842105	1	0.727273
SYT13 AND CLDN1 AND ABCB5	0.956522	0.916667	1
THY1 AND NOT-CELSR2 AND EPHB2	0.842105	1	0.727273
FAP AND IGSF6 AND GUCY2C	0.842105	1	0.727273
FAP AND PTPRC AND GUCY2C	0.842105	1	0.727273
FAP AND PTPRC AND NOT-CLDN12	0.842105	1	0.727273
FAP AND FAM26F AND NOT-SSTR3	0.842105	1	0.727273
FAP AND FOLH1 AND NOT-PTPRZ1	0.842105	1	0.727273
SST AND CLDN1 AND CELSR1	0.842105	1	0.727273
SST AND CLDN1 AND SLC47A1	0.842105	1	0.727273
FAP AND KLRD1 AND NOT-CLDN1	0.842105	1	0.727273
SST AND CLDN1 AND SGCD	0.842105	1	0.727273
SST AND CLDN1 AND GPR68	0.842105	1	0.727273
NTM AND NOT-MAL AND LY75	0.842105	1	0.727273
NTM AND NOT-MAL AND IL2RG	0.842105	1	0.727273
NTM AND NOT-MAL AND SLC44A3	0.842105	1	0.727273
NTM AND NOT-MAL AND CDH1	0.842105	1	0.727273
KCNE4 AND LY75 AND NOT-MAL	0.842105	1	0.727273
KCNE4 AND SLC44A3 AND NOT-MAL	0.842105	1	0.727273
NTM AND NOT-MAL AND TMEM30B	0.842105	1	0.727273
NOX4 AND SLC44A3 AND NOT-MAL	0.9	1	0.818182
KCNE4 AND CDH1 AND NOT-MAL	0.842105	1	0.727273
KCNE4 AND IGSF6 AND NOT-TSPAN14	0.842105	1	0.727273
KCNE4 AND FAM26F AND NOT-TSPAN14	0.842105	1	0.727273
NTM AND NOT-MAL AND GPR160	0.842105	1	0.727273
KCNE4 AND ITGB8 AND NOT-VAMP2	0.842105	1	0.727273
KCNE4 AND LTB AND NOT-FLT3LG	0.842105	1	0.727273
KCNE4 AND FAM26F AND NOT-ITGAD	0.842105	1	0.727273
KCNE4 AND FAM26F AND NOT-NPY5R	0.842105	1	0.727273
NTM AND NOT-MAL AND OR2F2	0.842105	1	0.727273
KCNE4 AND IGSF6 AND NOT-FLT3LG	0.842105	1	0.727273
NOX4 AND NOT-MAL AND TMEM30B	0.857143	0.9	0.818182
KCNE4 AND IGSF6 AND NOT-ADCY7	0.842105	1	0.727273
PCDHB16 AND NOT-MAL AND IL2RG	0.9	1	0.818182
KCNE4 AND IGSF6 AND NOT-NPY5R	0.842105	1	0.727273
NOX4 AND NOT-MAL AND LY75	0.9	1	0.818182
NTM AND NOT-MAL AND CHRNB1	0.842105	1	0.727273
NTM AND NOT-RTN3 AND FAM26F	0.842105	1	0.727273
NOX4 AND NOT-SLC47A1 AND SLC44A3	0.9	1	0.818182
NOX4 AND NOT-SLC47A1 AND CHRNB1	0.857143	0.9	0.818182
NTM AND NOT-RTN3 AND TMEM30B	0.842105	1	0.727273
SYT13 AND IGSF6 AND PCDHB16	1	1	1
NOX4 AND NOT-SLC47A1 AND SLC5A4	0.857143	0.9	0.818182
MUC1 AND NOT-RTN3 AND IFNAR2	0.952381	1	0.909091
GRM8 AND ITGAM AND MUC1	0.842105	1	0.727273
OR5I1 AND GPC3 AND IFNAR2	0.857143	0.9	0.818182
OTOF AND MUC1 AND CLEC2B	0.833333	0.769231	0.909091
THY1 AND IFNAR2 AND OTOF	0.9	1	0.818182
FAP AND OTOF AND IFNAR2	0.842105	1	0.727273

Antigen Pair Logic	F1	Precision	Recall
NOT-CLDN19 AND LRIG3 AND NOT-ATP8B1	1	1	1
GPR137B AND NOT-AOC3 AND FAT1	0.952381	1	0.909091
COMPLEX-NPFFR1/GPR137B/SLC23A2	0.916667	0.846154	1
NOT-CHRNA2 AND TTYH2 AND VANGL1	0.916667	0.846154	1
GPR137B AND TTYH2 AND NOT-UPK3A	0.916667	0.846154	1
NOT-CHRNA2 AND FAT1 AND NOT-ATP8B1	1	1	1
NOT-CHRNA2 AND FAT1 AND NOT-ATP6V0A4	1	1	1
NOT-GJD2 AND LRIG3 AND GPR19	1	1	1
GPR137B AND NOT-SLC1A6 AND FAT1	1	1	1
NOT-SLC10A2 AND LRIG3 AND GPR19	1	1	1
NOT-CHRNA2 AND FAT1 AND NOT-LCT	0.952381	1	0.909091
NOT-CHRNA2 AND TTYH2 AND AOC3	0.909091	0.909091	0.909091
NOT-CHRNA2 AND TTYH2 AND ATP8B1	0.909091	0.909091	0.909091
NOT-CHRNA2 AND TTYH2 AND NOT-SEMA4D	0.909091	0.909091	0.909091
NOT-CHRNA2 AND FAT1 AND NOT-SHISA9	0.952381	1	0.909091
NOT-CHRNA2 AND FAT1 AND NOT-SLC13A2	0.952381	1	0.909091
GPR137B AND NOT-LCT AND PLVAP	0.952381	1	0.909091
NOT-CHRNA2 AND FAT1 AND NOT-SLC13A2	0.952381	1	0.909091
NOT-CHRNA2 AND FAT1 AND NOT-PIRT	1	1	1
GPR137B AND NOT-SLC13A2 AND FAT1	0.952381	1	0.909091
NOT-CHRNA2 AND FAT1 AND NOT-TMPRSS11E	1	1	1
GPR137B AND NOT-GHR AND FAT1	0.952381	1	0.909091
NOT-OR10J1 AND LRIG3 AND GPR19	1	1	1
NOT-CHRNA2 AND GPR19 AND NOT-KCNJ6	1	1	1
GPR137B AND GPR19 AND NOT-NPFFR1	1	1	1
NOT-CHRNA2 AND GPR19 AND NOT-SCN1A	1	1	1
NOT-CHRNA2 AND GPR19 AND NOT-SYT4	1	1	1
NOT-CHRNA2 AND GPR19 AND SLC30A1	1	1	1
GPR19 AND LRIG3 AND NOT-OR1Q1	0.952381	1	0.909091
GPR19 AND LRIG3 AND NOT-KCNK12	0.952381	1	0.909091
NOT-CHRNA2 AND ZACN AND GPR19	1	1	1
NOT-CLDN19 AND GPR19 AND ZACN	1	1	1
NOT-CHRNA2 AND GPR19 AND LRIG3	1	1	1
NOT-CHRNA2 AND GPR19 AND SEMA4B	1	1	1
NOT-CHRNA2 AND GPR19 AND SLC22A18	1	1	1
GPR19 AND LRIG3 AND NOT-CYP4A11	0.952381	1	0.909091
GPR137B AND GPR19 AND NOT-NMUR2	1	1	1
NOT-NOX1 AND GPR19 AND LRIG3	0.952381	1	0.909091
GPR137B AND GPR19 AND SLC22A5	0.952381	1	0.909091
NOT-CLDN19 AND GPR19 AND LRIG3	1	1	1
NOT-CLDN19 AND LRIG3 AND NOT-PCDH8	1	1	1
NOT-CHRNA2 AND LRIG3 AND NOT-PCDH8	1	1	1
NOT-CHRNA2 AND FAT1 AND NOT-P2RY4	0.952381	1	0.909091
COMPLEX-GPR19/SLC6A18/GPR137B	0.956522	0.916667	1
COMPLEX-SLC5A8/GPR19/GPR137B	1	1	1
COMPLEX-CALHM3/GPR19/GPR137B	1	1	1
COMPLEX-GPR19/GPR137B/TAS1R1	0.916667	0.846154	1
COMPLEX-SLC6A15/CACNG6/GPR137B	0.9	1	0.818182
COMPLEX-SLC6A15/GPR137B/LPPR3	0.9	1	0.818182
COMPLEX-GRIK4/SLC6A15/GPR137B	0.9	1	0.818182
COMPLEX-NPHS1/SLC6A15/GPR137B	0.9	1	0.818182
COMPLEX-SLC6A15/CACNG7/GPR137B	0.9	1	0.818182
COMPLEX-NPBWR1/SLC6A15/GPR137B	0.9	1	0.818182
NOT-CHRNA2 AND TTYH2 AND GPNMB	1	1	1
NOT-CLDN19 AND GPNMB AND TTYH2	1	1	1
NOT-CHRNA2 AND TTYH2 AND CSPG4	1	1	1
NOT-CHRNA2 AND TTYH2 AND VCAM1	1	1	1
NOT-CHRNA2 AND TTYH2 AND VCAM1	1	1	1
NOT-CHRNA2 AND GPNMB AND NOT-ATP13A5	1	1	1
NOT-CHRNA2 AND SDC1 AND NOT-ATP13A5	1	1	1
NOT-CHRNA2 AND FAT1 AND NOT-CLDN8	1	1	1
NOT-CHRNA2 AND TTYH2 AND STEAP2	1	1	1
NOT-CHRNA2 AND TTYH2 AND STEAP2	1	1	1
NOT-CHRNA2 AND TTYH2 AND SDC1	0.956522	0.916667	1
NOT-CHRNA2 AND TTYH2 AND SDC1	0.956522	0.916667	1
NOT-CHRNA2 AND TTYH2 AND ERBB2	0.956522	0.916667	1
NOT-CHRNA2 AND TTYH2 AND TPBG	1	1	1
NOT-CHRNA2 AND TTYH2 AND TPBG	1	1	1
NOT-CHRNA2 AND VCAM1 AND NOT-ATP13A5	1	1	1
NOT-CHRNA2 AND TTYH2 AND CLDN1	0.956522	0.916667	1
NOT-CHRNA2 AND TTYH2 AND ROR1	0.956522	0.916667	1
NOT-CHRNA2 AND TTYH2 AND ERBB2	0.956522	0.916667	1
GPR137B AND NOT-ATP13A5 AND SDC1	0.956522	0.916667	1
NOT-CLDN19 AND TTYH2 AND TPBG	0.952381	1	0.909091
NOT-CHRNA2 AND TTYH2 AND FAP	0.952381	1	0.909091
NOT-CHRNA2 AND TTYH2 AND TNC	0.952381	1	0.909091
NOT-CHRNA2 AND TTYH2 AND ERBB3	0.952381	1	0.909091

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
FAP AND CDH18 AND IFNAR2	0.842105	1	0.727273
THY1 AND USH2A AND IFNAR2	0.9	1	0.818182
FAP AND CRB1 AND IFNAR2	0.842105	1	0.727273
TNFRSF12A AND IFNAR2 AND ABCB5	0.9	1	0.818182
CRB1 AND NOT-RAET1E AND LRIG3	0.842105	1	0.727273
THY1 AND IFNAR2 AND CNTNAP4	0.9	1	0.818182
THY1 AND IFNAR2 AND GRM8	0.9	1	0.818182
THY1 AND IFNAR2 AND GRIN2B	0.9	1	0.818182
THY1 AND IFNAR2 AND DCC	0.9	1	0.818182
GRM8 AND NOT-RAET1E AND LRIG3	0.842105	1	0.727273
GJB4 AND NOT-RAET1E AND CLDN1	0.842105	1	0.727273
SST AND IFNAR2 AND NOT-EPHB2	0.842105	1	0.727273
SST AND CLDN1 AND IFNAR2	0.842105	1	0.727273
FAP AND ERBB3 AND NOT-OTOF	0.842105	1	0.727273
SST AND CLDN1 AND NOT-LCT	0.842105	1	0.727273
FAP AND ERBB3 AND NOT-IFNAR2	0.842105	1	0.727273
FAP AND ERBB3 AND NOT-LCT	0.842105	1	0.727273
SST AND CLDN1 AND GRIN2B	0.842105	1	0.727273
SST AND CLDN1 AND NOT-GPR158	0.842105	1	0.727273
SST AND CLDN1 AND KCNA1	0.842105	1	0.727273
SST AND CLDN1 AND ABCG5	0.842105	1	0.727273
SST AND CLDN1 AND CRB1	0.842105	1	0.727273
SST AND CLDN1 AND NOT-GABRB2	0.842105	1	0.727273
SST AND CLDN1 AND NOT-USH2A	0.842105	1	0.727273
SST AND CLDN1 AND GRM8	0.842105	1	0.727273
SST AND CLDN1 AND CD207	0.842105	1	0.727273
CRB1 AND NOT-RAET1E AND ROR1	0.842105	1	0.727273
FAP AND EPHB2 AND NOT-THBD	0.842105	1	0.727273
SST AND CLDN1 AND NOT-SCN2A	0.842105	1	0.727273
SST AND CLDN1 AND NOT-MMP24	0.842105	1	0.727273
SST AND CLDN1 AND NOT-GJB4	0.842105	1	0.727273
SST AND CLDN1 AND CSMD2	0.842105	1	0.727273
GJB4 AND NOT-RAET1E AND SDC1	0.842105	1	0.727273
THY1 AND IFNAR2 AND NOT-SDC1	0.9	1	0.818182
THY1 AND IFNAR2 AND ABCB5	0.9	1	0.818182
FAP AND GPR158 AND NOT-CLDN1	0.842105	1	0.727273
THY1 AND IFNAR2 AND NOT-L1CAM	0.9	1	0.818182
MUC1 AND NOT-RTN3 AND CLEC2B	0.833333	0.769231	0.909091
MUC1 AND ABCB5 AND IFNAR2	0.842105	1	0.727273
EPHB2 AND IFNAR2 AND GPC3	0.857143	0.9	0.818182
MUC1 AND CSMD2 AND CLDN1	0.833333	0.769231	0.909091
TNFRSF12A AND GJB4 AND IFNAR2	0.9	1	0.818182
TNFRSF12A AND LCT AND IFNAR2	0.9	1	0.818182
TNFRSF12A AND OTOF AND IFNAR2	0.9	1	0.818182
TNFRSF12A AND GRIN2B AND IFNAR2	0.9	1	0.818182
TNFRSF12A AND SLC6A15 AND IFNAR2	0.842105	1	0.727273
TNFRSF12A AND FCAMR AND IFNAR2	0.9	1	0.818182
TNFRSF12A AND OR7C1 AND IFNAR2	0.9	1	0.818182
TNFRSF12A AND IFNAR2 AND SLC28A1	0.9	1	0.818182
TNFRSF12A AND SLC22A6 AND IFNAR2	0.9	1	0.818182
TNFRSF12A AND SHISA9 AND IFNAR2	0.9	1	0.818182
TNFRSF12A AND CDH18 AND IFNAR2	0.842105	1	0.727273
TNFRSF12A AND CRB1 AND IFNAR2	0.9	1	0.818182
TNFRSF12A AND IFNAR2 AND CACNA1S	0.842105	1	0.727273
TNFRSF12A AND CNTNAP4 AND IFNAR2	0.9	1	0.818182
TNFRSF12A AND CHRND AND IFNAR2	0.9	1	0.818182
TNFRSF12A AND GRM8 AND IFNAR2	0.9	1	0.818182
TNFRSF12A AND CSMD2 AND IFNAR2	0.842105	1	0.727273
TNFRSF12A AND IFNAR2 AND SLC30A10	0.9	1	0.818182
TNFRSF12A AND GRM8 AND LRIG3	0.818182	0.818182	0.818182
TNFRSF12A AND IFNAR2 AND LRIG3	0.9	1	0.818182
TNFRSF12A AND GPR6 AND IFNAR2	0.842105	1	0.727273
TNFRSF12A AND IFNAR2 AND NOT-LRRN4	0.842105	1	0.727273
TNFRSF12A AND GPR158 AND IFNAR2	0.9	1	0.818182
TNFRSF12A AND IFNAR2 AND SLC39A2	0.9	1	0.818182
TNFRSF12A AND USH2A AND IFNAR2	0.9	1	0.818182
TNFRSF12A AND IFNAR2 AND CLDN19	0.9	1	0.818182
TNFRSF12A AND IFNAR2 AND CDH8	0.9	1	0.818182
TNFRSF12A AND IFNAR2 AND GPRC6A	0.9	1	0.818182
TNFRSF12A AND IFNAR2 AND GABRG2	0.9	1	0.818182
TNFRSF12A AND CHRNA1 AND IFNAR2	0.842105	1	0.727273
TNFRSF12A AND IFNAR2 AND CD1B	0.9	1	0.818182
TNFRSF12A AND CSMD2 AND LRIG3	0.857143	0.9	0.818182
ABCB5 AND NOT-RAET1E AND CLDN1	0.842105	1	0.727273
STEAP1 AND ABCB5 AND CLDN1	0.833333	0.769231	0.909091
STEAP2 AND ABCB5 AND CLDN1	0.869565	0.833333	0.909091
STEAP2 AND CLDN1 AND NOT-SDC1	0.869565	0.833333	0.909091

Antigen Pair Logic	F1	Precision	Recall
GPR137B AND NOT-ATP13A5 AND ERBB2	0.952381	1	0.909091
NOT-CHRNA2 AND TTYH2 AND CD276	0.952381	1	0.909091
NOT-CACNG6 AND TTYH2 AND GPNMB	0.952381	1	0.909091
NOT-CHRNA2 AND TTYH2 AND IL13RA2	0.952381	1	0.909091
NOT-CHRNA2 AND TTYH2 AND CD34	0.952381	1	0.909091
NOT-CHRNA2 AND TTYH2 AND AXL	0.952381	1	0.909091
NOT-CHRNA2 AND TTYH2 AND L1CAM	0.952381	1	0.909091
NOT-CHRNA2 AND TTYH2 AND STEAP1	0.952381	1	0.909091
GPR137B AND NOT-CLDN8 AND FAT1	1	1	1
NOT-CHRNA2 AND NOT-CLDN10 AND SDC1	1	1	1
GPR137B AND NOT-CLDN10 AND ERBB3	1	1	1
NOT-CLDN19 AND GPNMB AND NOT-CLDN10	1	1	1
NOT-SLC10A2 AND TTYH2 AND GPNMB	0.952381	1	0.909091
NOT-CHRNA2 AND FAT1 AND NOT-VTCN1	0.952381	1	0.909091
NOT-CHRNA2 AND LRIG3 AND NOT-EPCAM	0.952381	1	0.909091
GPR137B AND NOT-CLDN10 AND ERBB2	0.952381	1	0.909091
NOT-CHRNA2 AND NOT-CLDN10 AND ERBB2	0.952381	1	0.909091
NOT-CLDN19 AND LRIG3 AND NOT-EPCAM	0.952381	1	0.909091
NOT-CLDN19 AND TTYH2 AND VCAM1	0.952381	1	0.909091
NOT-CHRNA2 AND FAT1 AND NOT-ERBB2	1	1	1
NOT-CHRNA2 AND FAT1 AND NOT-ERBB2	1	1	1
GPR137B AND NOT-CLDN8 AND LRIG3	1	1	1
GPR137B AND NOT-AOC3 AND ERBB3	0.952381	1	0.909091
NOT-CLDN19 AND GPNMB AND NOT-SLC30A10	0.952381	1	0.909091
NOT-CLDN19 AND GPNMB AND NOT-PCDH8	1	1	1
NOT-CHRNA2 AND FAT1 AND NOT-IL20RA	0.952381	1	0.909091
NOT-CLDN19 AND GPNMB AND NOT-ATP8B1	1	1	1
NOT-CHRNA2 AND SDC1 AND NOT-PCDH8	1	1	1
NOT-CHRNA2 AND ENG AND NOT-GHR	1	1	1
GPR137B AND NOT-IL20RA AND FAT1	0.952381	1	0.909091
NOT-CLDN19 AND GPNMB AND GPR19	1	1	1
GPR137B AND NOT-P2RY4 AND ERBB3	0.952381	1	0.909091
NOT-CHRNA2 AND NOT-CLDN10 AND GPNMB	1	1	1
NOT-CLDN19 AND ENG AND NOT-GHR	0.952381	1	0.909091
NOT-CLDN19 AND GPNMB AND NOT-SLC1A6	1	1	1
GPR137B AND NOT-ATP8B1 AND ERBB2	0.952381	1	0.909091
NOT-NOX1 AND GPNMB AND GPR19	1	1	1
NOT-CHRNA2 AND ENG AND NOT-SLC13A2	0.952381	1	0.909091
NOT-CHRNA2 AND NOT-CLDN10 AND VCAM1	1	1	1
NOT-CLDN19 AND STEAP1 AND NOT-ATP8B1	0.952381	1	0.909091
NOT-CLDN19 AND ENG AND NOT-ATP8B1	1	1	1
NOT-CHRNA2 AND FAT1 AND NOT-EPCAM	0.952381	1	0.909091
GPR137B AND NOT-LCT AND ERBB3	0.952381	1	0.909091
NOT-CLDN19 AND GPNMB AND NOT-TMPRSS11E	1	1	1
GPR137B AND NOT-GHR AND ERBB3	0.952381	1	0.909091
NOT-CHRNA2 AND GPNMB AND NOT-CD207	0.952381	1	0.909091
NOT-CHRNA2 AND SDC1 AND NOT-CD207	0.952381	1	0.909091
GPR137B AND NOT-CLDN19 AND L1CAM	0.952381	1	0.909091
NOT-CHRNA2 AND GPNMB AND NOT-PCDH8	1	1	1
NOT-CHRNA2 AND PLVAP AND NOT-CLDN8	1	1	1
NOT-CLDN19 AND ENG AND NOT-NOX1	1	1	1
NOT-CLDN19 AND GPNMB AND NOT-CD207	0.952381	1	0.909091
GPR137B AND NOT-NOX1 AND L1CAM	0.952381	1	0.909091
NOT-CHRNA2 AND VCAM1 AND NOT-PCDH8	1	1	1
NOT-CACNG6 AND SDC1 AND NOT-PCDH8	1	1	1
GPR137B AND NOT-ATP8B1 AND ERBB3	1	1	1
GPR137B AND NOT-CLDN19 AND CSPG4	0.952381	1	0.909091
NOT-CHRNA2 AND ITGAV AND NOT-PCDH8	0.916667	0.846154	1
NOT-CHRNA2 AND ERBB2 AND NOT-ATP13A5	0.916667	0.846154	1
NOT-CHRNA2 AND NOT-CLDN10 AND ITGAV	0.916667	0.846154	1
GPR137B AND NOT-SLC1A6 AND ERBB3	1	1	1
NOT-CLDN19 AND VCAM1 AND NOT-PCDH8	1	1	1
NOT-CHRNA2 AND NOT-CLDN10 AND TPBG	0.956522	0.916667	1
GPR137B AND NOT-EPCAM AND FAT1	0.952381	1	0.909091
NOT-CACNG6 AND GPNMB AND NOT-CLDN10	1	1	1
NOT-CHRNA2 AND LRIG3 AND NOT-VTCN1	0.952381	1	0.909091
NOT-CLDN19 AND ERBB2 AND NOT-ATP8B1	0.952381	1	0.909091
NOT-CHRNA2 AND SDC1 AND NOT-SLC1A6	1	1	1
NOT-CHRNA2 AND FAT1 AND NOT-EGFR	0.952381	1	0.909091
GPR137B AND NOT-CLDN10 AND SDC1	0.956522	0.916667	1
NOT-CHRNA2 AND GPNMB AND NOT-SLC1A6	1	1	1
NOT-CLDN19 AND GPNMB AND NOT-ATP6V0A4	1	1	1
NOT-CLDN19 AND GPNMB AND NOT-LCT	0.952381	1	0.909091
NOT-CLDN19 AND GPNMB AND NOT-GRM6	1	1	1
NOT-CHRNA2 AND EMP3 AND SDC1	0.909091	0.909091	0.909091
NOT-SLC5A5 AND ZNRF3 AND NOT-CADM2	1	1	1
NOT-SLC5A5 AND TYRO3 AND NOT-CADM2	1	1	1

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
EPHB2 AND CLDN1 AND NOT-ABCB5	0.818182	0.818182	0.818182
OTOF AND IFNAR2 AND GPC3	0.818182	0.818182	0.818182
GRIN2B AND GPC3 AND IFNAR2	0.9	1	0.818182
USH2A AND GPC3 AND IFNAR2	0.857143	0.9	0.818182
FCAMR AND GPC3 AND IFNAR2	0.818182	0.818182	0.818182
GRM8 AND GPC3 AND IFNAR2	0.818182	0.818182	0.818182
LCT AND IFNAR2 AND GPC3	0.818182	0.818182	0.818182
CSMD2 AND IFNAR2 AND GPC3	0.818182	0.818182	0.818182
MUC1 AND CNTNAP4 AND IFNAR2	0.842105	1	0.727273
LCT AND MUC1 AND IFNAR2	0.869565	0.833333	0.909091
MUC1 AND GRIN2B AND IFNAR2	0.909091	0.909091	0.909091
MUC1 AND CSMD2 AND IFNAR2	0.909091	0.909091	0.909091
Renal Cancer			
CDH6 AND NOT-HAS1 AND TYROBP	0.857143	1	0.75
CDH6 AND NOT-ACPP AND LITAF	0.857143	1	0.75
CDH6 AND NOT-CORIN AND ITGAM	0.857143	1	0.75
CDH6 AND CD63 AND GJB6	0.857143	1	0.75
CDH6 AND NOT-SLC24A4 AND OR51M1	0.857143	1	0.75
CDH6 AND NOT-SLC24A4 AND NTRK1	0.857143	1	0.75
CDH6 AND NOT-COLQ AND KCNS1	0.857143	1	0.75
CDH6 AND JTB AND TRAT1	0.857143	1	0.75
CDH6 AND KCNS1 AND NOT-VIPR1	0.857143	1	0.75
CDH6 AND NOT-DPP6 AND DSC2	0.857143	1	0.75
CDH6 AND NOT-HAS1 AND C5AR1	0.857143	1	0.75
CDH6 AND NOT-KCNK2 AND ZDHHC5	0.857143	1	0.75
CDH6 AND ATP1B1 AND CLPTM1	0.857143	1	0.75
CDH6 AND NOT-CORIN AND NTRK1	0.857143	1	0.75
CDH6 AND NOT-NRG4 AND TAAR8	0.857143	1	0.75
COMPLEX-CDH6/ATRAID/TPM1	0.857143	1	0.75
CDH6 AND NOT-NRG4 AND LRRC32	0.857143	1	0.75
CDH6 AND SLC25A5 AND CAV3	0.857143	1	0.75
CDH6 AND NOT-FGFR3 AND NOT-FZD6	0.857143	1	0.75
CDH6 AND NOT-SCNN1B AND GPR82	0.857143	1	0.75
COMPLEX-CDH6/CLCN3/ATRAID	0.857143	1	0.75
CDH6 AND NOT-NRG4 AND KCNS1	0.857143	1	0.75
CDH6 AND CD53 AND NOT-KRT5	0.857143	1	0.75
COMPLEX-CDH6/ATRAID/PTPRJ	0.857143	1	0.75
COMPLEX-CDH6/ATRAID/DST	0.857143	1	0.75
CDH6 AND NOT-PTPRZ1 AND NOT-MARVELD2	0.857143	1	0.75
CDH6 AND NOT-NRG4 AND C5AR1	0.857143	1	0.75
CDH6 AND JTB AND CD3D	0.857143	1	0.75
CDH6 AND NOT-AKAP6 AND VIPR1	0.857143	1	0.75
CDH6 AND NOT-TPM1 AND NOT-WNT4	0.857143	1	0.75
CDH6 AND NOT-TM9SF2 AND CD63	0.857143	1	0.75
COMPLEX-CDH6/IL6ST/ATRAID	0.857143	1	0.75
CDH6 AND NOT-STX2 AND NOT-PTPRZ1	0.857143	1	0.75
CDH6 AND CD63 AND ZDHHC5	0.857143	1	0.75
CDH6 AND OR51M1 AND NOT-OR2B2	0.857143	1	0.75
CDH6 AND TYROBP AND NOT-WNT4	0.857143	1	0.75
CDH6 AND NOT-NRG4 AND ITGA2	0.857143	1	0.75
CDH6 AND NOT-NRG4 AND DCHS2	0.857143	1	0.75
CDH6 AND NOT-SCNN1B AND GRM2	0.857143	1	0.75
CDH6 AND OR51M1 AND NOT-OR51B4	0.857143	1	0.75
CDH6 AND NOT-RHBDL2 AND LITAF	0.857143	1	0.75
CDH6 AND NOT-RHBDL2 AND KCNS1	0.857143	1	0.75
CDH6 AND NOT-SGCG AND PTPRU	0.857143	1	0.75
CDH6 AND NOT-ESYT3 AND DCHS2	0.857143	1	0.75
COMPLEX-CDH6/ATRAID/PTPRG	0.857143	1	0.75
CDH6 AND NOT-RET AND NOT-SCNN1B	0.857143	1	0.75
CDH6 AND NOT-PRRT2 AND CLPTM1	0.857143	1	0.75
COMPLEX-CDH6/LRRC8B/RET	0.857143	1	0.75
CDH16 AND NOT-SCNN1B AND OR51M1	0.857143	1	0.75
COMPLEX-CDH6/ATRAID/ETNK1	0.857143	1	0.75
COMPLEX-CDH6/ATRAID/GPRC5A	0.857143	1	0.75
CDH6 AND NOT-COLQ AND CLPTM1	0.857143	1	0.75
COMPLEX-CDH6/ATP1B3/ATRAID	0.857143	1	0.75
CDH16 AND NOT-SCNN1B AND SLC5A5	0.857143	1	0.75
CDH6 AND SLC25A5 AND OPN5	0.857143	1	0.75
COMPLEX-CDH6/CADM1/ATRAID	0.857143	1	0.75
CDH16 AND NOT-RET AND NOT-SCNN1B	0.857143	1	0.75
CDH6 AND NOT-DPP6 AND CLPTM1	0.857143	1	0.75
CDH6 AND ATP1B1 AND ADAM15	0.857143	1	0.75
CDH6 AND ATP1B1 AND KLRB1	0.857143	1	0.75
CDH6 AND NOT-TPM1 AND NOT-KCNK2	0.857143	1	0.75
CDH6 AND CD63 AND ADAM15	0.857143	1	0.75
CDH6 AND ATP1B1 AND MEP1A	0.857143	1	0.75
CDH6 AND NOT-KCNJ12 AND TYROBP	0.857143	1	0.75

Antigen Pair Logic	F1	Precision	Recall
NOT-SLC5A5 AND LGR4 AND NOT-CADM2	1	1	1
NOT-SLC5A5 AND IGDCC4 AND NOT-CADM2	1	1	1
NOT-SLC5A5 AND PCDHB14 AND NOT-CADM2	1	1	1
NOT-SLC5A5 AND SORT1 AND NOT-CADM2	1	1	1
NOT-GIPR AND SORT1 AND NOT-ATP1A2	1	1	1
NOT-TM4SF5 AND NOT-CADM2 AND TYRO3	1	1	1
NOT-OR2S2 AND NOT-ATP1A2 AND PCDHB14	1	1	1
NOT-SLC5A5 AND FZD7 AND NOT-CADM2	1	1	1
NOT-OR2S2 AND NOT-ATP1A2 AND GPM6B	1	1	1
NOT-OR2S2 AND NOT-ATP1A2 AND NLGN1	1	1	1
NOT-SLC5A5 AND TSPAN11 AND NOT-CADM2	1	1	1
NOT-GIPR AND GPR161 AND NOT-ATP1A2	1	1	1
NOT-CD300LB AND PAQR6 AND NOT-CADM2	0.956522	0.916667	1
NOT-SLC34A3 AND PCDHB14 AND NOT-CADM2	1	1	1
NOT-TM4SF5 AND NOT-CADM2 AND ZNRF3	1	1	1
NOT-SLC34A3 AND LGR4 AND NOT-CADM2	1	1	1
NOT-TM4SF5 AND NOT-CADM2 AND APLNR	1	1	1
NOT-SLC34A3 AND IGDCC4 AND NOT-CADM2	0.956522	0.916667	1
NOT-CD300LB AND PAQR6 AND NOT-CADM2	0.956522	0.916667	1
NOT-TM4SF5 AND NOT-CADM2 AND TIE1	0.956522	0.916667	1
NOT-TM4SF5 AND NOT-ATP1A2 AND TYRO3	1	1	1
NOT-TM4SF5 AND NOT-CADM2 AND PCDHB14	1	1	1
NOT-TM4SF5 AND NOT-CADM2 AND LGR4	1	1	1
CLCN5 AND NOT-CD300LB AND NOT-CADM2	1	1	1
NOT-SLC5A5 AND FZD7 AND NOT-TSPAN8	0.952381	1	0.909091
NOT-GIPR AND BEST1 AND LGR4	0.952381	1	0.909091
NOT-SLC5A5 AND BEST1 AND LGR4	0.952381	1	0.909091
NOT-SLC5A5 AND ZNRF3 AND NOT-ATP1A2	0.952381	1	0.909091
NOT-TM4SF5 AND NOT-TSPAN8 AND PCDHB14	0.952381	1	0.909091
NOT-TAAR8 AND LGR4 AND NOT-TSPAN8	0.952381	1	0.909091
NOT-SLC5A5 AND LGR4 AND NOT-TSPAN8	0.952381	1	0.909091
NOT-TAAR8 AND SORT1 AND NOT-TSPAN8	0.952381	1	0.909091
NOT-SLC5A5 AND LGR4 AND NOT-ATP1A2	0.952381	1	0.909091
NOT-SLC5A5 AND SORT1 AND NOT-TSPAN8	0.952381	1	0.909091
NOT-SLC5A5 AND TSPAN11 AND NOT-TSPAN8	0.952381	1	0.909091
COMPLEX-ITPR3/GPR143/OR2S2	0.952381	1	0.909091
NOT-OR2S2 AND NOT-ATP1A2 AND PRRG1	0.952381	1	0.909091
NOT-SLC5A5 AND TYRO3 AND NOT-NTRK3	0.952381	1	0.909091
NOT-SLC5A5 AND TYRO3 AND NOT-ATP1A2	0.952381	1	0.909091
NOT-OR2S2 AND NOT-NTRK3 AND TYRO3	0.952381	1	0.909091
NOT-OR2S2 AND NOT-ATP1A2 AND KCNN2	0.952381	1	0.909091
NOT-OR2S2 AND NOT-ATP1A2 AND SLC25A4	0.952381	1	0.909091
NOT-OR2S2 AND NOT-ATP1A2 AND PLP1	0.952381	1	0.909091
NOT-CD300LB AND TGFB3 AND NOT-CADM2	0.952381	1	0.909091
NOT-OR2S2 AND NOT-ATP1A2 AND APOLD1	0.952381	1	0.909091
TRPV1 AND NOT-CADM2 AND NOT-CD300LB	0.952381	1	0.909091
NOT-CD300LB AND TSPAN11 AND NOT-TSPAN8	0.952381	1	0.909091
NOT-GIPR AND GPR161 AND NOT-TSPAN8	0.952381	1	0.909091
NOT-SLC5A5 AND TIE1 AND NOT-ATP1A2	0.952381	1	0.909091
NOT-OR51I2 AND TSPAN11 AND NOT-TSPAN8	0.952381	1	0.909091
NOT-SLC34A3 AND PCDHB14 AND NOT-TSPAN8	0.952381	1	0.909091
NOT-SLC5A5 AND PCDHB14 AND NOT-ATP1A2	0.952381	1	0.909091
NOT-CD300LB AND LGR4 AND NOT-TSPAN8	0.952381	1	0.909091
NOT-TM4SF5 AND NOT-TSPAN8 AND LGR4	0.952381	1	0.909091
NOT-SLC5A5 AND SORT1 AND NOT-ATP1A2	0.952381	1	0.909091
NOT-TAAR8 AND PCDHB14 AND NOT-TSPAN8	0.952381	1	0.909091
NOT-GIPR AND LGR4 AND NOT-TSPAN8	0.952381	1	0.909091
NOT-SLC5A5 AND PCDHB14 AND NOT-TSPAN8	0.952381	1	0.909091
NOT-CD300LB AND ATP9B AND NOT-CADM2	0.952381	1	0.909091
PTPRS AND NOT-CADM2 AND NOT-CD300LB	0.952381	1	0.909091
NOT-CD300LB AND TGFBR1 AND NOT-CADM2	0.952381	1	0.909091
NOT-SLC34A3 AND LGR4 AND NOT-TSPAN8	0.952381	1	0.909091
NOT-SLC5A5 AND NOT-ATP1A2 AND NTN4	0.956522	0.916667	1
NOT-SLC5A5 AND APCDD1 AND NOT-ATP1A2	0.956522	0.916667	1
NOT-OR2S2 AND TSPAN11 AND NOT-TSPAN8	0.952381	1	0.909091
NOT-OR2S2 AND NOT-ATP1A2 AND VANGL2	1	1	1
NOT-SLC5A5 AND FZD4 AND NOT-ATP1A2	1	1	1
NOT-SLC5A5 AND ZNRF3 AND NOT-TSPAN8	0.952381	1	0.909091
NOT-SLC5A5 AND PDGFRA AND NOT-ATP1A2	1	1	1
NOT-SLC5A5 AND TIE1 AND NOT-TSPAN8	0.952381	1	0.909091
NOT-TAAR8 AND TSPAN11 AND NOT-FNDC5	1	1	1
NOT-SLC5A5 AND PERP AND NOT-ATP1A2	1	1	1
NOT-SLC5A5 AND WNT5A AND NOT-TSPAN8	0.952381	1	0.909091
NOT-CD300LB AND FAM57A AND NOT-CADM2	0.952381	1	0.909091
NOT-OR2S2 AND NOT-CADM2 AND PCDHB14	1	1	1
NOT-SLC5A5 AND OSMR AND NOT-CADM2	1	1	1
NOT-CD300LB AND PTGDR2 AND NOT-TSPAN8	0.952381	1	0.909091

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
CDH6 AND OR51M1 AND NOT-SLC6A14	0.857143	1	0.75
CDH6 AND CD63 AND PRLHR	0.857143	1	0.75
CDH6 AND NOT-AKAP6 AND NOT-ATP1B3	0.857143	1	0.75
COMPLEX-CDH6/LRRC8B/ATRAID	0.857143	1	0.75
COMPLEX-CDH6/HAS1/PTPRJ	0.857143	1	0.75
CDH6 AND CCR5 AND NOT-SLC6A14	0.857143	1	0.75
CDH6 AND NOT-KCNK2 AND ITGAM	0.857143	1	0.75
CDH6 AND NOT-COLQ AND NOT-FZD6	0.857143	1	0.75
CDH6 AND CD63 AND SLC27A1	0.857143	1	0.75
CDH6 AND NOT-NPY2R AND CLPTM1	0.857143	1	0.75
CDH6 AND C5AR1 AND ATP1B1	0.857143	1	0.75
CDH6 AND NOT-PTPRZ1 AND NOT-PSEN1	0.857143	1	0.75
CDH6 AND NOT-MEGF10 AND LRRC32	0.857143	1	0.75
CDH6 AND NOT-AKAP6 AND CATSPER3	0.857143	1	0.75
CDH6 AND SLC25A5 AND OR8D1	0.857143	1	0.75
CDH6 AND NOT-HAS1 AND CHRNA2	0.857143	1	0.75
CDH6 AND NOT-TAS2R14 AND CALHM3	0.857143	1	0.75
KCNJ16 AND ATRAID AND NOT-CLDN16	0.857143	1	0.75
CDH6 AND NOT-HTR5A AND KCNS1	0.857143	1	0.75
CDH6 AND NOT-ESYT3 AND CATSPER3	0.857143	1	0.75
CDH6 AND NOT-SLC4A8 AND CLPTM1	0.857143	1	0.75
CDH6 AND NOT-ADCYAP1R1 AND CATSPER3	0.857143	1	0.75
CDH6 AND NOT-MEGF10 AND TMEM8A	0.857143	1	0.75
CDH6 AND NOT-MEGF10 AND HRH1	0.857143	1	0.75
CDH6 AND NOT-MEGF10 AND C5AR1	0.857143	1	0.75
CDH6 AND ATP1B1 AND SLC12A9	0.857143	1	0.75
CDH6 AND NOT-P2RX3 AND C5AR1	0.857143	1	0.75
CDH6 AND NOT-PTPRZ1 AND HLA-DRB1	0.857143	1	0.75
CDH6 AND NOT-KCNQ5 AND ABCA12	0.857143	1	0.75
CDH6 AND NOT-FLVCR1 AND ATP1B1	0.857143	1	0.75
COMPLEX-CDH6/GRIN2A/ATRAID	0.857143	1	0.75
CDH6 AND NOT-SHH AND ABCA12	0.857143	1	0.75
CDH6 AND NOT-NRG4 AND ROS1	0.857143	1	0.75
CDH6 AND NOT-SLCO1A2 AND NOT-ATP1B3	0.857143	1	0.75
CDH6 AND NOT-SLC9B1 AND C5AR1	0.857143	1	0.75
CDH6 AND NOT-XG AND CATSPER3	0.857143	1	0.75
CDH6 AND CD63 AND CATSPER3	0.857143	1	0.75
CDH6 AND OR51M1 AND NOT-DSC1	0.857143	1	0.75
CDH6 AND NOT-CD207 AND DSC2	0.857143	1	0.75
CDH6 AND NOT-MEGF10 AND NOT-MCOLN2	0.857143	1	0.75
CDH6 AND NOT-HAS1 AND ABCA12	0.857143	1	0.75
CDH6 AND NOT-HAS1 AND CALHM3	0.857143	1	0.75
CDH6 AND NOT-SGCG AND CATSPER3	0.857143	1	0.75
CDH6 AND ATP1B1 AND BEST3	0.857143	1	0.75
CDH6 AND ABCA12 AND NOT-RHBDL2	0.857143	1	0.75
CDH6 AND NOT-MEGF10 AND SLC5A5	0.857143	1	0.75
CDH6 AND NOT-SHH AND CALHM3	0.857143	1	0.75
CDH6 AND NOT-NRG4 AND C8B	0.857143	1	0.75
CDH6 AND NOT-OXTR AND NOT-COLQ	0.857143	1	0.75
CDH6 AND NOT-TAS2R14 AND NOX1	0.857143	1	0.75
CDH6 AND ATP1B1 AND PCDHA6	0.857143	1	0.75
CDH6 AND NOT-KIAA0319 AND VIPR1	0.857143	1	0.75
CDH6 AND ATP1B1 AND SELP	0.857143	1	0.75
CDH6 AND NOT-CD207 AND TAAR8	0.857143	1	0.75
CDH6 AND ABCA12 AND NOT-HRH4	0.857143	1	0.75
CDH6 AND NOT-ADCYAP1R1 AND ABCA12	0.857143	1	0.75
CDH6 AND NOT-DPP6 AND ABCA12	0.857143	1	0.75
CDH6 AND NOT-COLQ AND CALHM3	0.857143	1	0.75
CDH6 AND NOT-SLC4A8 AND C5AR1	0.857143	1	0.75
CDH6 AND NOT-MEGF10 AND ZDHHC5	0.857143	1	0.75
CDH6 AND NOT-COLQ AND ABCA12	0.857143	1	0.75
COMPLEX-CDH6/NFASC/ATRAID	0.857143	1	0.75
CDH6 AND ATP1B1 AND CALHM1	0.857143	1	0.75
CDH6 AND NOT-MEGF10 AND TMIGD2	0.857143	1	0.75
CDH6 AND NOT-HTR5A AND OR51M1	0.857143	1	0.75
CDH6 AND VIPR1 AND NOT-OR8D1	0.857143	1	0.75
CDH6 AND NOT-CORIN AND SLC12A9	0.857143	1	0.75
CDH6 AND NOX1 AND NOT-NPY2R	0.857143	1	0.75
CDH6 AND NOT-MLC1 AND VIPR1	0.857143	1	0.75
CDH6 AND NOT-SLC20A2 AND ABCA12	0.857143	1	0.75
CDH6 AND NOT-MEGF10 AND CLPTM1	0.857143	1	0.75
CDH6 AND NOT-NRG4 AND ABCA12	0.857143	1	0.75
CDH6 AND NOT-MEGF10 AND TYROBP	0.857143	1	0.75
CDH6 AND C8B AND CD63	0.857143	1	0.75
CDH6 AND NOT-NRG4 AND CATSPER3	0.857143	1	0.75
CDH6 AND NOT-CORIN AND HLA-DRB1	0.857143	1	0.75
CDH6 AND NOT-SYT8 AND CATSPER3	0.857143	1	0.75

Antigen Pair Logic	F1	Precision	Recall
NOT-CD300LB AND B4GALT1 AND NOT-CADM2	0.952381	1	0.909091
NOT-SLC5A5 AND NOT-CADM2 AND GJA1	1	1	1
NOT-CD300LB AND CYP51A1 AND NOT-CADM2	0.952381	1	0.909091
NOT-SLC5A5 AND NOT-ATP1A2 AND GJA1	0.956522	0.916667	1
NOT-OR2S2 AND NOT-ATP1A2 AND SORT1	1	1	1
NOT-SLC5A5 AND IGDCC4 AND NOT-ATP1A2	1	1	1
NOT-CD300LB AND PTGDR2 AND NOT-CADM2	1	1	1
NOT-SLC5A5 AND ANO1 AND NOT-TSPAN8	0.952381	1	0.909091
NOT-OR2S2 AND TSPAN11 AND NOT-CADM2	1	1	1
NOT-OR2S2 AND NOT-CADM2 AND IGDCC4	1	1	1
NOT-SLC5A5 AND TIE1 AND NOT-CADM2	1	1	1
NOT-CD300LB AND TGFBR1 AND NOT-ATP1A2	0.952381	1	0.909091
NOT-CD300LB AND PTGDR2 AND NOT-CADM2	1	1	1
NOT-SLC5A5 AND IGDCC4 AND NOT-TSPAN8	0.952381	1	0.909091
NOT-TM4SF5 AND NOT-CADM2 AND ANO1	1	1	1
NOT-TM4SF5 AND NOT-TSPAN8 AND ZNRF3	0.952381	1	0.909091
NOT-GPR21 AND PCDHB14 AND NOT-TSPAN8	0.952381	1	0.909091
NOT-TAAR8 AND TSPAN11 AND NOT-ATP1A2	1	1	1
NOT-TM4SF5 AND NOT-TSPAN8 AND FZD7	0.952381	1	0.909091
NOT-SLC34A3 AND TSPAN11 AND NOT-CADM2	0.952381	1	0.909091
NOT-GPR21 AND TSPAN11 AND NOT-NTRK3	0.952381	1	0.909091
NOT-SLC5A5 AND GPNMB AND NOT-KCNK5	1	1	1
NOT-SLC5A5 AND GPNMB AND NOT-CFTR	1	1	1
NOT-SLC5A5 AND GPNMB AND NOT-CADM2	1	1	1
NOT-TM4SF5 AND GPNMB AND NOT-KCNK5	1	1	1
NOT-SLC5A5 AND GPNMB AND NOT-TSPAN8	0.952381	1	0.909091
NOT-SLC5A5 AND GPNMB AND NOT-SYT8	0.952381	1	0.909091
NOT-SLC34A3 AND LGR4 AND NOT-CLDN8	1	1	1
NOT-CD300LB AND GPNMB AND NOT-TSPAN8	0.952381	1	0.909091
NOT-TM4SF5 AND GPNMB AND NOT-CFTR	1	1	1
NOT-OR2S2 AND GPNMB AND NOT-KCNK5	1	1	1
NOT-SLC5A5 AND LGR4 AND NOT-CLDN8	1	1	1
NOT-OR2S2 AND TSPAN11 AND NOT-CLDN8	1	1	1
NOT-TM4SF5 AND NOT-CADM2 AND SDC1	1	1	1
NOT-CD300LB AND PCYT1A AND NOT-CADM2	0.952381	1	0.909091
NOT-OR2S2 AND NOT-NTRK3 AND SDC1	0.952381	1	0.909091
NOT-TAAR8 AND GPNMB AND NOT-TSPAN8	0.952381	1	0.909091
NOT-TM4SF5 AND NOT-CADM2 AND GPNMB	1	1	1
NOT-SLC5A5 AND GPNMB AND NOT-ATP1A2	0.952381	1	0.909091
NOT-OR2S2 AND NOT-NTRK3 AND GPNMB	0.952381	1	0.909091
NOT-GIPR AND GPNMB AND NOT-TSPAN8	0.952381	1	0.909091
NOT-GIPR AND GPNMB AND NOT-KCNK5	1	1	1
NOT-TM4SF5 AND GPNMB AND NOT-SYT8	0.952381	1	0.909091
NOT-SLC5A5 AND GPNMB AND NOT-NTRK3	0.952381	1	0.909091
NOT-TAAR8 AND TSPAN11 AND NOT-CLDN8	1	1	1
NOT-OR2S2 AND GPNMB AND NOT-ATP1A2	0.952381	1	0.909091
NOT-OR2S2 AND GPNMB AND NOT-ATP1A2	0.952381	1	0.909091
NOT-TM4SF5 AND GPNMB AND NOT-TSPAN8	0.952381	1	0.909091
NOT-SLC5A5 AND GPNMB AND PTPRJ	0.952381	1	0.909091
NOT-GIPR AND GPNMB AND NOT-NPY1R	0.952381	1	0.909091
NOT-GIPR AND TSPAN11 AND NOT-EPCAM	0.952381	1	0.909091
NOT-OR2S2 AND NOT-ATP1A2 AND L1CAM	0.952381	1	0.909091
NOT-TM4SF5 AND GPNMB AND NOT-NPY1R	0.952381	1	0.909091
NOT-GIPR AND WNT5A AND NOT-EPCAM	0.952381	1	0.909091
NOT-OR2S2 AND GPNMB AND PTPRJ	0.952381	1	0.909091
NOT-SLC5A5 AND GPNMB AND NOT-NPY1R	0.952381	1	0.909091
NOT-SLC5A5 AND GPNMB AND NOT-NRXN1	0.952381	1	0.909091
NOT-GIPR AND GPNMB AND BEST1	0.952381	1	0.909091
NOT-SLC5A5 AND GPNMB AND NOT-CNGA1	0.952381	1	0.909091
NOT-TM4SF5 AND LGR4 AND NOT-EPCAM	0.952381	1	0.909091
NOT-SLC34A3 AND LGR4 AND NOT-EPCAM	0.952381	1	0.909091
NOT-TAAR8 AND LGR4 AND NOT-EPCAM	0.952381	1	0.909091
NOT-SLC5A5 AND LGR4 AND NOT-EPCAM	0.952381	1	0.909091
NOT-OR2S2 AND GPNMB AND NOT-CFTR	1	1	1
NOT-SLC5A5 AND GPNMB AND NOT-GPRC5D	0.952381	1	0.909091
NOT-SLC5A5 AND GPNMB AND NOT-ACSL1	0.952381	1	0.909091
NOT-GIPR AND GPNMB AND NOT-CFTR	1	1	1
NOT-SLC5A5 AND TSPAN11 AND NOT-CLDN8	1	1	1
NOT-SLC34A3 AND PCDHB14 AND NOT-CLDN8	1	1	1
NOT-GPR21 AND TSPAN11 AND NOT-CLDN8	1	1	1
NOT-OR2S2 AND SDC1 AND NOT-TSPAN8	0.952381	1	0.909091
NOT-SLC5A5 AND SDC1 AND NOT-TSPAN8	0.952381	1	0.909091
NOT-OR2S2 AND NOT-ATP1A2 AND THY1	0.952381	1	0.909091
NOT-OR2S2 AND GPNMB AND NOT-NPR3	1	1	1
NOT-SLC5A5 AND PCDHB14 AND NOT-CLDN8	1	1	1
NOT-SLC5A5 AND FZD7 AND NOT-CLDN8	1	1	1
NOT-SLC5A5 AND GPNMB AND NOT-NPR3	1	1	1

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
CDH6 AND NOT-CDH11 AND SDCBP	0.857143	1	0.75
CDH6 AND NOT-CORIN AND C8B	0.857143	1	0.75
CDH6 AND CD63 AND STAB1	0.857143	1	0.75
CDH6 AND NOT-OPALIN AND OR51M1	0.857143	1	0.75
CDH6 AND NOT-MEGF10 AND NOT-ACPP	0.857143	1	0.75
CDH6 AND NOT-DPP6 AND CALHM3	0.857143	1	0.75
CDH6 AND ATP1B1 AND OR3A1	0.857143	1	0.75
CDH6 AND ATP1B1 AND OR7C1	0.857143	1	0.75
CDH6 AND NOT-MEGF10 AND ATP1B1	0.857143	1	0.75
COMPLEX-CDH6/ATRAID/PPAPDC1B	0.857143	1	0.75
CDH6 AND NOT-ACPP AND NOT-SLC4A8	0.857143	1	0.75
CDH6 AND NOT-NRG4 AND HLA-DRB1	0.857143	1	0.75
CDH6 AND NOT-MEGF10 AND NOT-HAS3	0.857143	1	0.75
CDH6 AND NOT-NPY2R AND CATSPER3	0.857143	1	0.75
CDH6 AND NOT-ATP1B3 AND NOT-CRB1	0.857143	1	0.75
COMPLEX-CDH6/CDH11/ATRAID	0.857143	1	0.75
CDH6 AND ABCB5 AND NOT-OR2B2	0.857143	1	0.75
CDH6 AND NOT-LRP4 AND CD276	0.857143	1	0.75
KCNJ16 AND NOT-COLQ AND CD276	0.857143	1	0.75
CDH6 AND NOT-LGR5 AND OR51M1	0.857143	1	0.75
CDH6 AND ABCB5 AND NOT-SLC26A3	0.857143	1	0.75
CDH6 AND NOT-LRRTM2 AND NOT-CD276	0.857143	1	0.75
CDH6 AND ATP1B1 AND CD19	0.857143	1	0.75
CDH6 AND NOT-LYPD6B AND GPNMB	0.857143	1	0.75
CDH6 AND NOT-KCNJ12 AND CD276	0.857143	1	0.75
CDH6 AND ABCB5 AND NOT-HTR1A	0.857143	1	0.75
CDH6 AND NOT-TAS2R14 AND ABCB5	0.857143	1	0.75
CDH6 AND NOT-HAS1 AND GPNMB	0.857143	1	0.75
CDH6 AND NOT-GPR52 AND ABCB5	0.857143	1	0.75
CDH6 AND NOT-PTPRZ1 AND CD276	0.857143	1	0.75
CDH6 AND ABCB5 AND NOT-SCIMP	0.857143	1	0.75
CDH6 AND ABCB5 AND NOT-TACR3	0.857143	1	0.75
CDH16 AND NOT-SCNN1B AND CD276	0.857143	1	0.75
CDH6 AND NOT-COLQ AND CD276	0.857143	1	0.75
CDH6 AND NOT-PMEL AND DSC2	0.857143	1	0.75
CDH6 AND ABCB5 AND NOT-HRH4	0.857143	1	0.75
CDH6 AND JTB AND FCRL2	0.857143	1	0.75
CDH6 AND NOT-FGFR3 AND GPNMB	0.857143	1	0.75
CDH6 AND NOT-ATP1B3 AND NOT-ERBB2	0.857143	1	0.75
CDH6 AND ATP1B1 AND ABCB5	0.857143	1	0.75
CDH6 AND NOT-SHH AND ABCB5	0.857143	1	0.75
CDH6 AND NOT-NRG4 AND ERBB2	0.857143	1	0.75
CDH6 AND NOT-LYPD6B AND ERBB2	0.857143	1	0.75
CDH6 AND NOT-ATP1B3 AND NOT-ABCB5	0.857143	1	0.75
SLC3A1 AND CD276 AND NOT-GDPD2	0.857143	1	0.75
CDH6 AND NOT-COLQ AND PTK7	0.857143	1	0.75
SLC3A1 AND CD276 AND NOT-SLC35G1	0.857143	1	0.75
CDH6 AND NOT-NCAM2 AND CD276	0.857143	1	0.75
CDH6 AND ABCB5 AND NOT-NPY2R	0.857143	1	0.75
CDH6 AND NOT-COLQ AND GPNMB	0.857143	1	0.75
SLC3A1 AND CD276 AND NOT-ATP1B3	0.857143	1	0.75
SLC3A1 AND CD276 AND NOT-SLC26A3	0.857143	1	0.75
CDH6 AND NOT-GPR63 AND ABCB5	0.857143	1	0.75
CDH6 AND ABCB5 AND NOT-WNT4	0.857143	1	0.75
CDH6 AND NOT-ATP1B3 AND NOT-CD276	0.857143	1	0.75
SLC3A1 AND CD276 AND NOT-RHBDL2	0.857143	1	0.75
CDH6 AND NOT-DPP6 AND ERBB2	0.857143	1	0.75
CDH6 AND ABCB5 AND NOT-DUOX1	0.857143	1	0.75
KCNJ16 AND NOT-SLC6A14 AND CD276	0.857143	1	0.75
CDH6 AND ATP1B1 AND ERBB2	0.857143	1	0.75
CDH6 AND NOT-SLC20A2 AND GPNMB	0.857143	1	0.75
CDH6 AND NOT-PMEL AND ITGAM	0.857143	1	0.75
CDH6 AND NOT-SYT8 AND ERBB2	0.857143	1	0.75
CDH6 AND NOT-REEP2 AND CD276	0.857143	1	0.75
CDH6 AND NOT-ADCYAP1R1 AND CD276	0.857143	1	0.75
SLC3A1 AND CD276 AND NOT-DSC2	0.857143	1	0.75
KCNJ16 AND NOT-MUC15 AND CD276	0.857143	1	0.75
CDH6 AND NOT-HAS1 AND CD276	0.857143	1	0.75
SLC3A1 AND CD276 AND NOT-SMPD3	0.857143	1	0.75
CDH6 AND NOT-TAS2R16 AND CD276	0.857143	1	0.75
SLC3A1 AND CD276 AND NOT-PTPRF	0.857143	1	0.75
CDH6 AND NOT-PRRT2 AND ABCB5	0.857143	1	0.75
CDH6 AND NOT-COLQ AND ERBB2	0.857143	1	0.75
CDH6 AND NOT-DPP6 AND NOT-CD276	0.857143	1	0.75
CDH6 AND NOT-AKAP6 AND ABCB5	0.857143	1	0.75
CDH6 AND NOT-NRG4 AND PTK7	0.857143	1	0.75
CDH6 AND JTB AND ERBB2	0.857143	1	0.75

Antigen Pair Logic	F1	Precision	Recall
NOT-OR2S2 AND GPNMB AND NOT-TSPAN8	0.952381	1	0.909091
NOT-OR2S2 AND GPNMB AND NOT-SGCG	0.952381	1	0.909091
NOT-SLC5A5 AND SDC1 AND NOT-CADM2	1	1	1
NOT-SLC5A5 AND GPNMB AND NOT-EDAR	1	1	1
NOT-GIPR AND GPNMB AND NOT-CADM2	1	1	1
NOT-GIPR AND GPNMB AND NOT-ATP1A2	1	1	1
NOT-TM4SF5 AND GPNMB AND NOT-EDAR	1	1	1
NOT-SLC5A5 AND VCAM1 AND NOT-TSPAN8	0.952381	1	0.909091
NOT-OR2S2 AND NOT-ATP1A2 AND STEAP1	0.952381	1	0.909091
COMPLEX-SGCD/TPBG/TM4SF5	0.952381	1	0.909091
NOT-SLC5A5 AND VCAM1 AND NOT-CADM2	1	1	1
NOT-GIPR AND TSPAN11 AND NOT-CLDN8	1	1	1
NOT-OR2S2 AND NOT-NTRK3 AND VCAM1	0.952381	1	0.909091
NOT-CD300LB AND PCYT1A AND NOT-ATP1A2	0.952381	1	0.909091
NOT-SLC5A5 AND SDC1 AND NOT-NTRK3	0.952381	1	0.909091
NOT-TM4SF5 AND NOT-CADM2 AND TPBG	0.956522	0.916667	1
NOT-TM4SF5 AND NOT-ATP1A2 AND SDC1	1	1	1
NOT-TM4SF5 AND GPNMB AND GPR3	0.952381	1	0.909091
NOT-SLC5A5 AND TNFRSF19 AND NOT-CLDN8	1	1	1
NOT-SLC34A3 AND GPNMB AND NOT-KCNK5	1	1	1
NOT-TM4SF5 AND GPNMB AND NOT-SLC6A2	0.952381	1	0.909091
NOT-SLC5A5 AND GPNMB AND NOT-SLC6A2	0.952381	1	0.909091
NOT-SLC5A5 AND VCAM1 AND NOT-NTRK3	0.952381	1	0.909091
NOT-TM4SF5 AND LGR4 AND NOT-CLDN4	0.952381	1	0.909091
NOT-CD300LB AND GPNMB AND PTPRJ	0.952381	1	0.909091
NOT-TM4SF5 AND GPNMB AND NOT-ATP1A2	0.952381	1	0.909091
NOT-TM4SF5 AND GPNMB AND NOT-NTRK3	0.952381	1	0.909091
NOT-TM4SF5 AND NOT-NTRK3 AND SDC1	0.952381	1	0.909091
NOT-SLC5A5 AND GPNMB AND NOT-CYP4F12	0.952381	1	0.909091
NOT-GIPR AND LGR4 AND NOT-EPCAM	0.952381	1	0.909091
NOT-GIPR AND GPNMB AND NOT-CNGA1	1	1	1
NOT-SLC5A5 AND GPNMB AND NOT-SCNN1G	1	1	1
NOT-OR2S2 AND GPNMB AND NOT-NPY1R	0.952381	1	0.909091
NOT-OR2S2 AND GPNMB AND NOT-CYP4F12	0.952381	1	0.909091
NOT-GIPR AND PCDHB14 AND NOT-EPCAM	0.952381	1	0.909091
NOT-TM4SF5 AND TSPAN11 AND NOT-EPCAM	0.952381	1	0.909091
NOT-OR51I2 AND TNFRSF19 AND NOT-CLDN8	0.952381	1	0.909091
NOT-TM4SF5 AND GPNMB AND NOT-CNGA1	0.952381	1	0.909091
NOT-SLC5A5 AND WNT5A AND NOT-EPCAM	0.952381	1	0.909091
NOT-TM4SF5 AND SDC1 AND NOT-KIAA1324	1	1	1
ITM2B AND NOT-TMPRSS2 AND SDC1	0.956522	0.916667	1
NOT-SLC5A5 AND TSPAN11 AND NOT-EPCAM	0.952381	1	0.909091
NOT-TM4SF5 AND GPNMB AND NOT-NRXN1	0.952381	1	0.909091
NOT-SLC34A3 AND ANO1 AND NOT-EPCAM	0.952381	1	0.909091
COMPLEX-LDLRAD3/PMEL/CDH1	1	1	1
COMPLEX-MERTK/PMEL/RNF144A	1	1	1
COMPLEX-MLANA/FREM2/GPRC5C	1	1	1
COMPLEX-MLANA/FREM2/TMEM100	1	1	1
COMPLEX-PMEL/NIPA2/ST3GAL5	1	1	1
COMPLEX-LDLRAD3/MARCH1/PMEL	1	1	1
COMPLEX-MLANA/FREM2/NPY5R	1	1	1
COMPLEX-MLANA/FREM2/NPR3	1	1	1
COMPLEX-HTRA2/PMEL/NIPA2	1	1	1
COMPLEX-MLANA/HRH2/FREM2	1	1	1
COMPLEX-LDLRAD3/PMP22/PMEL	1	1	1
COMPLEX-IL31RA/MLANA/FREM2	1	1	1
COMPLEX-LDLRAD3/ATP2A3/PMEL	1	1	1
COMPLEX-LDLRAD3/ASPH/PMEL	1	1	1
COMPLEX-LDLRAD3/FAM26F/PMEL	1	1	1
COMPLEX-LDLRAD3/BCAM/PMEL	1	1	1
COMPLEX-LDLRAD3/SHISA6/PMEL	1	1	1
COMPLEX-MLANA/FREM2/ZP2	1	1	1
COMPLEX-MLANA/FREM2/AQP10	1	1	1
COMPLEX-LDLRAD3/SI/PMEL	1	1	1
COMPLEX-LDLRAD3/PMEL/TAS1R2	1	1	1
COMPLEX-LDLRAD3/PMEL/TMPRSS2	1	1	1
COMPLEX-LDLRAD3/PMEL/VAMP7	1	1	1
COMPLEX-LDLRAD3/PMEL/SLC26A10	1	1	1
COMPLEX-HCST/LDLRAD3/PMEL	1	1	1
COMPLEX-LDLRAD3/SCN4A/PMEL	1	1	1
COMPLEX-VAMP5/LDLRAD3/PMEL	1	1	1
COMPLEX-LDLRAD3/ESYT2/PMEL	1	1	1
COMPLEX-LDLRAD3/C2orf83/PMEL	1	1	1
COMPLEX-LDLRAD3/KCNH2/PMEL	1	1	1
COMPLEX-SLC27A4/PMEL/RNF144A	1	1	1
COMPLEX-PMEL/CD81/RNF144A	1	1	1
COMPLEX-LDLRAD3/HFE/PMEL	1	1	1

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
CDH6 AND NOT-PMEL AND ZDHHC5	0.857143	1	0.75
COMPLEX-CDH6/EPHB2/LRRC8B	0.857143	1	0.75
CDH6 AND NOT-KCNJ12 AND GPNMB	0.857143	1	0.75
CDH6 AND NOT-FGFR3 AND CD276	0.857143	1	0.75
CDH6 AND NOT-CORIN AND NOT-CD276	0.857143	1	0.75
CDH6 AND NOT-PMEL AND CLPTM1	0.857143	1	0.75
CDH6 AND NOT-ADCYAP1R1 AND ABCB5	0.857143	1	0.75
CDH6 AND NOT-PMEL AND ITGA6	0.857143	1	0.75
CDH6 AND NOT-PMEL AND NOT-FZD6	0.857143	1	0.75
CDH6 AND NOT-SYT8 AND GPNMB	0.857143	1	0.75
CDH6 AND NOT-KCNK2 AND GPNMB	0.857143	1	0.75
CDH6 AND ATP1B1 AND PTK7	0.857143	1	0.75
CDH6 AND NOT-SLC7A5 AND SDCBP	0.857143	1	0.75
CDH6 AND CD63 AND MUC4	0.857143	1	0.75
CDH6 AND ATP1B1 AND CD22	0.857143	1	0.75
CDH6 AND NOT-SLC7A5 AND SLC25A5	0.857143	1	0.75
CDH6 AND CD63 AND CD19	0.857143	1	0.75
CDH6 AND NOT-ADCYAP1R1 AND GPNMB	0.857143	1	0.75
KCNJ16 AND NOT-KRT5 AND CD276	0.857143	1	0.75
KCNJ16 AND NOT-SCNN1B AND CD276	0.857143	1	0.75
CDH6 AND CD63 AND RAET1E	0.857143	1	0.75
KCNJ16 AND NOT-MUC15 AND SDC1	0.857143	1	0.75
SLC3A1 AND CD276 AND GPR63	0.857143	1	0.75
SLC3A1 AND CD276 AND EPHB6	0.857143	1	0.75
SLC3A1 AND CD276 AND NOT-DPP6	0.857143	1	0.75
SLC3A1 AND CD276 AND LY6D	0.857143	1	0.75
SLC3A1 AND CD276 AND XG	0.857143	1	0.75
SLC3A1 AND CD276 AND MUC15	0.857143	1	0.75
CDH6 AND NOT-PMEL AND ATP1B1	0.857143	1	0.75
CDH6 AND NOT-PMEL AND CAV1	0.857143	1	0.75
CDH6 AND NOT-OR1G1 AND PTK7	0.857143	1	0.75
CDH6 AND NOT-PMEL AND ITGA2	0.857143	1	0.75
CDH6 AND NOT-PMEL AND LRRC32	0.857143	1	0.75
CDH6 AND ATP1B1 AND CD79A	0.857143	1	0.75
CDH6 AND NOT-MEGF10 AND CD276	0.857143	1	0.75
CDH6 AND NOT-SLCO1A2 AND ABCB5	0.857143	1	0.75
CDH6 AND ABCB5 AND NOT-KCNA1	0.857143	1	0.75
CDH6 AND ABCB5 AND NOT-GPR135	0.857143	1	0.75
CDH6 AND NOT-PMEL AND CALHM3	0.857143	1	0.75
CDH6 AND NOT-SLC9B1 AND ABCB5	0.857143	1	0.75
CDH6 AND NOT-MOG AND ABCB5	0.857143	1	0.75
CDH6 AND ABCA12 AND NOT-PMEL	0.857143	1	0.75
SLC3A1 AND CD276 AND NOT-CHRND	0.857143	1	0.75
CDH6 AND NOT-HTR5A AND CD276	0.857143	1	0.75
CDH6 AND NOT-SLC4A8 AND ABCB5	0.857143	1	0.75
CDH6 AND ABCB5 AND NOT-CATSPERD	0.857143	1	0.75
CDH6 AND ABCB5 AND NOT-P2RY4	0.857143	1	0.75
CDH6 AND NOT-GPR26 AND ABCB5	0.857143	1	0.75
COMPLEX-CDH6/GPNMB/MEGF10	0.857143	1	0.75
COMPLEX-CDH6/ERBB2/MEGF10	0.857143	1	0.75
CDH6 AND NOT-HTR5A AND GPNMB	0.857143	1	0.75
SLC3A1 AND CD276 AND DSC1	0.857143	1	0.75
SLC3A1 AND CD276 AND CRB1	0.857143	1	0.75
CDH6 AND NOT-PVRL1 AND GPNMB	0.857143	1	0.75
SLC3A1 AND CD276 AND NOT-CABP7	0.818182	0.9	0.75
SLC3A1 AND CD276 AND NOT-HTR5A	0.818182	0.9	0.75
SLC3A1 AND CD276 AND MEGF10	0.818182	0.9	0.75
SLC3A1 AND CD276 AND NOT-OR2L2	0.818182	0.9	0.75
KCNJ16 AND NOT-CRB1 AND CD276	0.818182	0.9	0.75
KCNJ16 AND NOT-MEGF10 AND CD276	0.857143	1	0.75
KCNJ16 AND NOT-HTR5A AND CD276	0.818182	0.9	0.75
CDH6 AND NOT-CHRND AND PTK7	0.8	1	0.666667
KCNJ16 AND NOT-OR2L2 AND CD276	0.8	1	0.666667
CDH6 AND NOT-CHRND AND ERBB2	0.8	1	0.666667
CDH6 AND NOT-CHRND AND ABCB5	0.8	1	0.666667
CDH6 AND NOT-CABP7 AND CD276	0.8	1	0.666667
CDH6 AND NOT-AGTR2 AND ERBB2	0.8	1	0.666667
CLDN2 AND ATRAID AND NOT-GABRA2	0.8	1	0.666667
CDH6 AND NOT-CHRND AND NOT-SDC1	0.8	1	0.666667
CDH6 AND NOT-HTR5A AND ERBB2	0.8	1	0.666667
KCNJ16 AND NOT-CHRND AND CD276	0.8	1	0.666667
SLC3A1 AND CD276 AND SLC13A5	0.8	1	0.666667
CDH6 AND NOT-CALN1 AND CD276	0.8	1	0.666667
CDH6 AND NOT-CHRND AND CD276	0.8	1	0.666667
VCAM1 AND ABCA12 AND NOT-SLC24A4	0.8	1	0.666667
CDH6 AND NOT-MEGF10 AND ABCA12	0.857143	1	0.75
CDH6 AND NOT-MEGF10 AND NOX1	0.857143	1	0.75

Antigen Pair Logic	F1	Precision	Recall
COMPLEX-ABCC9/MLANA/FREM2	1	1	1
COMPLEX-LDLRAD3/GRIA3/PMEL	1	1	1
COMPLEX-LDLRAD3/GPM6B/PMEL	1	1	1
COMPLEX-LDLRAD3/GP5/PMEL	1	1	1
COMPLEX-LDLRAD3/OR1A2/PMEL	1	1	1
COMPLEX-LDLRAD3/TAS2R50/PMEL	1	1	1
COMPLEX-LDLRAD3/TAS2R39/PMEL	1	1	1
COMPLEX-LDLRAD3/TPSG1/PMEL	1	1	1
COMPLEX-LDLRAD3/FRRS1L/PMEL	1	1	1
COMPLEX-LDLRAD3/BACE1/PMEL	1	1	1
COMPLEX-LDLRAD3/ACSL1/PMEL	1	1	1
COMPLEX-LDLRAD3/F3/PMEL	1	1	1
COMPLEX-LDLRAD3/CADM4/PMEL	1	1	1
COMPLEX-LDLRAD3/MARVELD2/PMEL	1	1	1
COMPLEX-LDLRAD3/CTSZ/PMEL	1	1	1
COMPLEX-LDLRAD3/IHH/PMEL	1	1	1
COMPLEX-ATP1A1/PMEL/RNF144A	1	1	1
COMPLEX-GPR176/PMEL/RNF144A	1	1	1
COMPLEX-SLCO2B1/PMEL/RNF144A	1	1	1
COMPLEX-DCBLD2/PMEL/RNF144A	1	1	1
COMPLEX-LDLRAD3/KCNA3/PMEL	1	1	1
COMPLEX-HRH1/PMEL/RNF144A	1	1	1
COMPLEX-SLCO4C1/PMEL/RNF144A	1	1	1
COMPLEX-LRP5/PMEL/RNF144A	1	1	1
COMPLEX-MLANA/TMEM235/SLC2A1	1	1	1
COMPLEX-MLANA/HTR1E/SLC2A1	1	1	1
COMPLEX-MLANA/NMUR2/SLC2A1	1	1	1
COMPLEX-MLANA/ABCA12/OR10J1	1	1	1
COMPLEX-MLANA/ABCA12/TMEM235	1	1	1
COMPLEX-MLANA/ABCA12/GHR	1	1	1
COMPLEX-MLANA/ABCA12/SCARA5	1	1	1
COMPLEX-MLANA/ABCA12/CCKBR	1	1	1
COMPLEX-MLANA/SLC2A1/CCKBR	1	1	1
COMPLEX-MLANA/ABCA12/NOX1	1	1	1
COMPLEX-MLANA/ABCA12/ATP13A5	1	1	1
COMPLEX-MLANA/NPFFR1/SLC2A1	1	1	1
COMPLEX-MLANA/OR10J1/SLC2A1	1	1	1
COMPLEX-MLANA/GHR/SLC2A1	1	1	1
COMPLEX-MLANA/ABCA12/NPFFR1	1	1	1
NOT-CHRNA2 AND TTYH2 AND KDR	1	1	1
COMPLEX-MLANA/SCARA5/SLC2A1	1	1	1
COMPLEX-MLANA/ATP13A5/SLC2A1	1	1	1
COMPLEX-MLANA/NOX1/SLC2A1	1	1	1
NOT-CHRNA2 AND TTYH2 AND KDR	1	1	1
COMPLEX-MLANA/CDH20/SLC2A1	1	1	1
COMPLEX-MLANA/ABCA12/NMUR2	1	1	1
COMPLEX-MLANA/ABCA12/CDH20	1	1	1
COMPLEX-MLANA/ABCA12/HTR1E	1	1	1
MLANA AND NOT-NOX1 AND NOT-SYT4	1	1	1
MLANA AND NOT-NOX1 AND EFNB2	1	1	1
MLANA AND NOT-OR10J1 AND NOT-LRIG3	1	1	1
MLANA AND NOT-NOX1 AND NOT-OR10J1	1	1	1
MLANA AND NOT-NOX1 AND LRIG3	1	1	1
MLANA AND NOT-NOX1 AND NOT-KCNA1	1	1	1
COMPLEX-MLANA/ABCA12/SLC28A1	1	1	1
COMPLEX-MLANA/ABCA12/SLC22A13	1	1	1
COMPLEX-MLANA/ABCA12/SLC22A6	1	1	1
COMPLEX-MLANA/ABCA12/CACNG8	1	1	1
COMPLEX-MLANA/ABCA12/LRFN2	1	1	1
COMPLEX-MLANA/ABCA12/SLC4A5	1	1	1
COMPLEX-MLANA/ABCA12/SLC10A1	1	1	1
COMPLEX-MLANA/ABCA12/GPR50	1	1	1
COMPLEX-MLANA/ABCA12/GPR137B	1	1	1
COMPLEX-MLANA/ABCA12/CACNA1S	1	1	1
COMPLEX-MLANA/ABCA12/GJA10	1	1	1
COMPLEX-MLANA/ABCA12/UPK3A	1	1	1
COMPLEX-MLANA/ABCA12/SLC22A5	1	1	1
COMPLEX-HRH3/MLANA/SLC2A1	1	1	1
COMPLEX-CLDN16/MLANA/SLC2A1	1	1	1
COMPLEX-CHRNA4/MLANA/SLC2A1	1	1	1
COMPLEX-HCN4/MLANA/SLC2A1	1	1	1
COMPLEX-CLCN1/MLANA/SLC2A1	1	1	1
COMPLEX-MLANA/GHSR/SLC2A1	1	1	1
COMPLEX-MLANA/TSPAN16/SLC2A1	1	1	1
COMPLEX-MLANA/GRIN2B/SLC2A1	1	1	1
COMPLEX-MRGPRX2/MLANA/SLC2A1	1	1	1
COMPLEX-MLANA/EPHA10/SLC2A1	1	1	1

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
CDH6 AND NOT-OPALIN AND ABCA12	0.857143	1	0.75
CDH6 AND CHRNA2 AND NOT-CRB1	0.857143	1	0.75
CDH6 AND NOT-SCN1A AND CATSPER3	0.857143	1	0.75
CDH6 AND NOT-SLCO1A2 AND CATSPER3	0.857143	1	0.75
CDH6 AND NOT-OPALIN AND CATSPER3	0.857143	1	0.75
CDH6 AND NOT-CD207 AND SLC12A9	0.857143	1	0.75
CDH6 AND C8B AND NOT-DSC1	0.857143	1	0.75
CDH6 AND NOT-MEGF10 AND CATSPER3	0.857143	1	0.75
CDH6 AND NOT-SLC9B1 AND CALHM3	0.857143	1	0.75
CDH6 AND NOT-MOG AND CHRNA2	0.857143	1	0.75
CDH6 AND NOT-CHRND AND NOT-OXTR	0.857143	1	0.75
CDH6 AND NOT-CD207 AND ABCA12	0.857143	1	0.75
CDH6 AND NOT-KCNA1 AND CATSPER3	0.857143	1	0.75
CDH6 AND NOT-SLCO1A2 AND CHRNA2	0.857143	1	0.75
CDH6 AND NOT-SLC4A8 AND CALHM3	0.857143	1	0.75
CDH6 AND NOT-SLC4A8 AND CHRNA2	0.857143	1	0.75
CDH6 AND NOT-SCN1A AND ABCA12	0.857143	1	0.75
CDH6 AND NOX1 AND NOT-MOG	0.857143	1	0.75
CDH6 AND NOT-CD207 AND C8B	0.857143	1	0.75
COMPLEX-CDH6/CALHM3/MEGF10	0.857143	1	0.75
COMPLEX-CDH6/DSC1/MEGF10	0.857143	1	0.75
CDH6 AND NOT-SLCO1A2 AND CALHM3	0.857143	1	0.75
CDH6 AND HLA-DRB1 AND NOT-CD207	0.857143	1	0.75
CDH6 AND NOT-KCNA1 AND CALHM3	0.857143	1	0.75
NOT-SCARA5 AND CAV2 AND CD74	0.869565	0.909091	0.833333
KCNJ16 AND NOT-SCN1A AND ABCA12	0.818182	0.9	0.75
KCNJ16 AND NOT-SCN1A AND NOT-CATSPER3	0.818182	0.9	0.75
CDH6 AND NOT-CHRND AND ABCA12	0.8	1	0.666667
CDH6 AND CATSPERD AND NOT-CALHM3	0.8	1	0.666667
CDH6 AND NOT-CHRND AND CHRNA2	0.8	1	0.666667
CDH6 AND NOT-CHRND AND CATSPER3	0.8	1	0.666667
CDH6 AND NOT-CHRND AND VANGL1	0.8	1	0.666667
KCNJ16 AND HLA-DRB1 AND NOT-CD207	0.818182	0.9	0.75
TSPAN4 AND NOT-SCARA5 AND CD74	0.846154	0.785714	0.916667
TSPAN4 AND NOT-SCARA5 AND VANGL1	0.857143	1	0.75
ABCA1 AND NOT-SCARA5 AND VANGL1	0.8	1	0.666667
GGTLC1 AND NOT-SCARA5 AND ATRAID	0.818182	0.9	0.75
KCNJ16 AND HLA-DRB1 AND NOT-DSC1	0.857143	1	0.75
NOT-SCARA5 AND NOT-SLC4A8 AND ABCA3	0.909091	1	0.833333
SLC3A1 AND CATSPER3 AND NOT-MEGF10	0.8	1	0.666667
KCNJ16 AND NOT-DSC1 AND C8B	0.818182	0.9	0.75
SLC3A1 AND CD276 AND NOT-GPA33	0.857143	1	0.75
SLC3A1 AND CD276 AND NOT-MUC4	0.857143	1	0.75
CDH6 AND NOT-LGR5 AND ERBB2	0.857143	1	0.75
CDH6 AND NOT-LGR5 AND GPNMB	0.857143	1	0.75
CDH6 AND NOT-LGR5 AND CD276	0.857143	1	0.75
CDH6 AND NOT-PMEL AND GPNMB	0.857143	1	0.75
SLC3A1 AND CD276 AND NOT-SDC1	0.857143	1	0.75
SLC3A1 AND CD276 AND NOT-PMEL	0.818182	0.9	0.75
CDH6 AND NOT-PMEL AND ERBB2	0.8	1	0.666667
KCNJ16 AND NOT-LGR5 AND CD276	0.8	1	0.666667
CLDN2 AND NOT-HHIP AND CD276	0.8	1	0.666667
CLDN2 AND ATRAID AND CD276	0.8	1	0.666667
CLDN2 AND CD276 AND CD63	0.8	1	0.666667
CLDN2 AND GPNMB AND NOT-PLA2R1	0.8	1	0.666667
CLDN2 AND PTK7 AND NOT-NPY2R	0.8	1	0.666667
NOT-SLITRK6 AND ERBB2 AND NOT-LGR5	0.8	1	0.666667
CLDN2 AND CD276 AND NOT-LRRTM2	0.8	1	0.666667
CLDN2 AND CD276 AND NOT-PTPRZ1	0.8	1	0.666667
CLDN2 AND PTK7 AND NOT-DPP6	0.8	1	0.666667
NOT-IL11RA AND ATRAID AND NOT-MUC4	0.8	1	0.666667
VCAM1 AND ATP1B1 AND NOT-CD276	0.857143	1	0.75
CLDN2 AND ITGAM AND NOT-PMEL	0.8	1	0.666667
VCAM1 AND ATP1B1 AND ABCB5	0.857143	1	0.75
CLDN2 AND ABCA12 AND NOT-SCN1A	0.8	1	0.666667
VCAM1 AND ABCA12 AND NOT-MOG	0.818182	0.9	0.75
VCAM1 AND ABCA12 AND NOT-CHRND	0.818182	0.9	0.75
VCAM1 AND ABCA12 AND NOT-CRB1	0.818182	0.9	0.75
VCAM1 AND ABCA12 AND NOT-CD207	0.857143	1	0.75
VCAM1 AND ABCA12 AND NOT-OPALIN	0.8	1	0.666667
VCAM1 AND CALHM3 AND NOT-P2RY4	0.818182	0.9	0.75
VCAM1 AND CALHM3 AND NOT-GPR135	0.857143	1	0.75
VCAM1 AND CALHM3 AND NOT-SLC9B1	0.8	1	0.666667
VCAM1 AND CALHM3 AND NOT-CD207	0.857143	1	0.75
CLDN2 AND CALHM3 AND NOT-DSC1	0.8	1	0.666667
VCAM1 AND CALHM3 AND NOT-CATSPERD	0.857143	1	0.75
VCAM1 AND NOX1 AND NOT-CD207	0.8	1	0.666667

Antigen Pair Logic	F1	Precision	Recall
COMPLEX-MLANA/AMHR2/SLC2A1	1	1	1
COMPLEX-MLANA/ABCA12/HTR6	1	1	1
COMPLEX-MLANA/OR2L2/SLC2A1	1	1	1
COMPLEX-MLANA/ABCA12/SLC6A18	1	1	1
COMPLEX-MLANA/OR1C1/SLC2A1	1	1	1
COMPLEX-GPR156/MLANA/SLC2A1	1	1	1
COMPLEX-LRRN4/MLANA/SLC2A1	1	1	1
COMPLEX-SLC5A8/MLANA/SLC2A1	1	1	1
COMPLEX-MLANA/CALHM1/SLC2A1	1	1	1
COMPLEX-MRGPRF/MLANA/SLC2A1	1	1	1
COMPLEX-LRIG3/MLANA/SLC2A1	1	1	1
COMPLEX-FGF6/MLANA/SLC2A1	1	1	1
COMPLEX-CHRNG/MLANA/SLC2A1	1	1	1
COMPLEX-CHRND/MLANA/SLC2A1	1	1	1
COMPLEX-CYP4A11/MLANA/SLC2A1	1	1	1
COMPLEX-CNGA4/MLANA/SLC2A1	1	1	1
COMPLEX-SLC22A12/MLANA/SLC2A1	1	1	1
COMPLEX-SLC5A11/MLANA/SLC2A1	1	1	1
COMPLEX-MLANA/ABCA12/GPR135	1	1	1
COMPLEX-CHRNA2/MLANA/SLC2A1	1	1	1
COMPLEX-MLANA/ABCA12/GABBR2	1	1	1
COMPLEX-CACNG2/MLANA/SLC2A1	1	1	1
COMPLEX-MLANA/ABCA12/IGDCC3	1	1	1
COMPLEX-MLANA/ABCA12/PCDH8	1	1	1
COMPLEX-MLANA/ABCA12/KCNH1	1	1	1
COMPLEX-MLANA/ABCA12/CACNG6	1	1	1
COMPLEX-LRIG3/MLANA/ABCA12	1	1	1
COMPLEX-MLANA/SLC2A1/UMOD	1	1	1
COMPLEX-MLANA/SLC2A1/CALN1	1	1	1
COMPLEX-MLANA/SLC2A1/CASR	1	1	1
COMPLEX-MLANA/SLC2A1/CCKAR	1	1	1
COMPLEX-MLANA/ABCA12/CCKAR	1	1	1
COMPLEX-MLANA/SLC2A1/OTOF	1	1	1
COMPLEX-MLANA/ABCA12/CASR	1	1	1
COMPLEX-MLANA/ABCA12/CALN1	1	1	1
COMPLEX-GPR156/MLANA/ABCA12	1	1	1
COMPLEX-LRRN4/MLANA/ABCA12	1	1	1
COMPLEX-SLC5A8/MLANA/ABCA12	1	1	1
COMPLEX-MLANA/ABCA12/UMOD	1	1	1
COMPLEX-CHRNG/MLANA/ABCA12	1	1	1
COMPLEX-MLANA/ABCA12/OR52D1	1	1	1
COMPLEX-MLANA/ABCA12/SLC10A2	1	1	1
COMPLEX-MLANA/ABCA12/SLC1A6	1	1	1
COMPLEX-MLANA/ABCA12/ABCG8	1	1	1
COMPLEX-MLANA/ABCA12/CACNG7	1	1	1
COMPLEX-CLDN16/MLANA/ABCA12	1	1	1
COMPLEX-MLANA/ABCA12/GJD2	1	1	1
NOT-SLC5A5 AND GPNMB AND NOT-CLDN8	1	1	1
NOT-GIPR AND GPNMB AND NOT-CLDN8	1	1	1
NOT-TM4SF5 AND GPNMB AND NOT-CLDN8	1	1	1
NOT-GIPR AND GPNMB AND NOT-EPCAM	0.952381	1	0.909091
NOT-SLC5A5 AND GPNMB AND NOT-VTCN1	0.952381	1	0.909091
NOT-TM4SF5 AND GPNMB AND NOT-VTCN1	0.952381	1	0.909091
NOT-SLC5A5 AND GPNMB AND NOT-ERBB2	1	1	1
NOT-OR2S2 AND GPNMB AND NOT-CLDN8	1	1	1
NOT-SLC34A3 AND GPNMB AND NOT-CLDN8	1	1	1
NOT-SLC5A5 AND GPNMB AND NOT-IL20RA	0.952381	1	0.909091
NOT-GIPR AND GPNMB AND NOT-VTCN1	0.952381	1	0.909091
NOT-TAAR8 AND GPNMB AND NOT-EPCAM	0.952381	1	0.909091
NOT-TM4SF5 AND SDC1 AND NOT-EPCAM	0.952381	1	0.909091
NOT-SLC5A5 AND GPNMB AND NOT-EPCAM	0.952381	1	0.909091
NOT-TM4SF5 AND GPNMB AND NOT-EPCAM	0.952381	1	0.909091
NOT-OR51M1 AND GPNMB AND NOT-ERBB2	0.952381	1	0.909091
NOT-TM4SF5 AND GPNMB AND NOT-ERBB2	1	1	1
NOT-TM4SF5 AND GPNMB AND NOT-ERBB2	1	1	1
NOT-TMEM150B AND GPNMB AND NOT-ERBB2	1	1	1
NOT-TAAR8 AND GPNMB AND NOT-CLDN8	1	1	1
NOT-GIPR AND GPNMB AND NOT-IL20RA	0.952381	1	0.909091
NOT-TM4SF5 AND GPNMB AND NOT-CLDN4	0.952381	1	0.909091
NOT-OR2S2 AND GPNMB AND NOT-VTCN1	0.952381	1	0.909091
NOT-OR2S2 AND SDC1 AND NOT-EPCAM	0.952381	1	0.909091
NOT-CD300LB AND GPNMB AND NOT-EPCAM	0.952381	1	0.909091
NOT-TM4SF5 AND GPNMB AND NOT-IL20RA	0.952381	1	0.909091
NOT-CD300LB AND GPNMB AND NOT-CLDN8	1	1	1
NOT-GIPR AND GPNMB AND NOT-ERBB2	1	1	1
NOT-GIPR AND GPNMB AND NOT-ERBB2	1	1	1
NOT-SLC5A5 AND GPNMB AND NOT-CEACAM6	1	1	1

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
VCAM1 AND OXTR AND NOT-CHRND	0.818182	0.9	0.75
VCAM1 AND CHRNA2 AND NOT-MOG	0.818182	0.9	0.75
VCAM1 AND NOT-MEGF10 AND CATSPER3	0.818182	0.9	0.75
VCAM1 AND NOX1 AND NOT-CATSPERD	0.8	1	0.666667
VCAM1 AND CATSPER3 AND NOT-CHRND	0.857143	1	0.75
VCAM1 AND CHRNA2 AND NOT-CHRND	0.818182	0.9	0.75
NOT-SCARA5 AND AXL AND NOT-SLC4A8	0.916667	0.916667	0.916667
VCAM1 AND CHRNA2 AND NOT-CRB1	0.818182	0.9	0.75
CLDN2 AND HLA-DRB1 AND NOT-DSC1	0.8	1	0.666667
NOT-SCARA5 AND AXL AND CD74	0.916667	0.916667	0.916667
VCAM1 AND CATSPER3 AND NOT-DSC1	0.857143	1	0.75
VCAM1 AND CATSPER3 AND NOT-CD207	0.857143	1	0.75
VCAM1 AND CATSPER3 AND NOT-SCN1A	0.8	1	0.666667
VCAM1 AND CATSPER3 AND NOT-CRB1	0.857143	1	0.75
VCAM1 AND CATSPER3 AND NOT-KCNA1	0.818182	0.9	0.75
VCAM1 AND NOT-SCARA5 AND ERBB2	0.857143	1	0.75
VCAM1 AND NOT-SCARA5 AND SDC1	0.8	1	0.666667
VCAM1 AND ABCB5 AND NOT-MOG	0.818182	0.9	0.75
CLDN2 AND PTK7 AND NOT-DSC1	0.8	1	0.666667
VCAM1 AND ABCA12 AND NOT-PMEL	0.818182	0.9	0.75
VCAM1 AND ABCB5 AND NOT-GPR135	0.857143	1	0.75
VCAM1 AND ABCB5 AND NOT-CRB1	0.818182	0.9	0.75
VCAM1 AND ABCB5 AND NOT-P2RY4	0.818182	0.9	0.75
VCAM1 AND ABCB5 AND NOT-CHRND	0.818182	0.9	0.75
VCAM1 AND CALHM3 AND NOT-PMEL	0.818182	0.9	0.75
VCAM1 AND ABCB5 AND NOT-SLC9B1	0.8	1	0.666667
VCAM1 AND ABCB5 AND NOT-CATSPERD	0.857143	1	0.75
VCAM1 AND NOT-CHRND AND NOT-CD276	0.818182	0.9	0.75
VCAM1 AND SDC1 AND NOT-MEGF10	0.8	1	0.666667
VCAM1 AND NOT-CD276 AND NOT-GPR6	0.8	1	0.666667
VCAM1 AND NOT-CHRND AND PTK7	0.818182	0.9	0.75
VCAM1 AND NOT-CD276 AND NOT-OR2L2	0.8	1	0.666667
NOT-SCARA5 AND AXL AND ERBB2	0.88	0.846154	0.916667
NOT-SCARA5 AND AXL AND SDC1	0.869565	0.909091	0.833333
NOT-LGR5 AND CD276 AND NOT-STEAP4	0.857143	1	0.75
CLDN2 AND NOT-LGR5 AND CD276	0.8	1	0.666667
VCAM1 AND NOT-LGR5 AND NOT-CD276	0.8	1	0.666667
VCAM1 AND SDC1 AND NOT-CD276	0.8	1	0.666667
VCAM1 AND NOT-CD276 AND NOT-GPA33	0.857143	1	0.75
NOT-LGR5 AND CD276 AND NOT-CLDN11	0.818182	0.9	0.75
NOT-CEACAM5 AND CD276 AND NOT-LGR5	0.818182	0.9	0.75
NOT-LGR5 AND EGFR AND ERBB2	0.8	1	0.666667
NOT-LGR5 AND EGFR AND GPNMB	0.8	1	0.666667
TNFRSF12A AND NOT-SCARA5 AND CD74	0.818182	0.9	0.75
GGTLC1 AND CALHM3 AND NOT-SLC9B1	0.818182	0.9	0.75
TNFRSF12A AND CATSPER3 AND NOT-SLCO1A2	0.8	1	0.666667
Colon Cancer			
BIRC5 AND TSPAN8 AND NOT-SEMA6D	1	1	1
HEPH AND CLDN1 AND NOT-CD200R1	1	1	1
DPEP1 AND TMEM123 AND ATP2C2	1	1	1
DPEP1 AND TMEM123 AND HEPH	1	1	1
MUC13 AND EDNRA AND NOT-NPY5R	1	1	1
MUC13 AND EDNRA AND NOT-NGFR	1	1	1
HEPH AND CLDN1 AND NOT-CATSPER2	1	1	1
MUC13 AND EDNRA AND NOT-MUSK	1	1	1
HEPH AND CLDN1 AND NOT-ADCY9	1	1	1
MUC13 AND EDNRA AND NOT-MPZ	1	1	1
MUC13 AND EDNRA AND NOT-MMP16	1	1	1
HEPH AND CLDN1 AND NOT-GPR146	1	1	1
DPEP1 AND TMEM123 AND NOT-GLP2R	1	1	1
MUC13 AND EDNRA AND NOT-MC4R	1	1	1
DPEP1 AND TMEM123 AND NOT-CD7	1	1	1
HEPH AND CLDN1 AND NOT-FZD10	1	1	1
MUC13 AND EDNRA AND NOT-ERVFRD-1	1	1	1
HEPH AND CLDN1 AND NOT-BTN2A1	1	1	1
DPEP1 AND TMEM123 AND NOT-XPR1	1	1	1
DPEP1 AND NOT-SLC22A7 AND NOT-SLC52A3	1	1	1
COMPLEX-SLC22A7/OR10A5/DPEP1	1	1	1
HEPH AND CLDN1 AND NOT-RAMP2	1	1	1
DPEP1 AND NOT-SLC22A7 AND NOT-ADRA1B	1	1	1
MUC13 AND EDNRA AND NOT-PCDHB8	1	1	1
DPEP1 AND NOT-SLC22A7 AND MCOLN2	1	1	1
MUC13 AND EDNRA AND NOT-PCDHGA11	1	1	1
HEPH AND CLDN1 AND F3	1	1	1
DPEP1 AND NOT-SLC22A7 AND NOT-FCGRT	1	1	1
HEPH AND CLDN1 AND NOT-PIANP	1	1	1
MUC13 AND EDNRA AND NOT-DCHS2	1	1	1

Antigen Pair Logic	F1	Precision	Recall
NOT-TM4SF5 AND GPNMB AND NOT-CEACAM6	1	1	1
NOT-SLC34A3 AND GPNMB AND NOT-LGR5	0.952381	1	0.909091
NOT-SLC34A3 AND GPNMB AND NOT-IL20RA	0.952381	1	0.909091
NOT-OR2S2 AND GPNMB AND NOT-IL20RA	0.952381	1	0.909091
NOT-SLC5A5 AND GPNMB AND NOT-EGFR	0.952381	1	0.909091
COMPLEX-CD300LB/EDNRB/CLDN6	0.9	1	0.818182
COMPLEX-EDNRB/SDC1/SLC14A2	0.9	1	0.818182
COMPLEX-GPNMB/EDNRB/PCDHGA9	0.9	1	0.818182
COMPLEX-EDNRB/SLC5A5/TPBG	0.9	1	0.818182
COMPLEX-GPNMB/EDNRB/GPR15	0.9	1	0.818182
COMPLEX-EDNRB/OR51I2/SDC1	0.9	1	0.818182
COMPLEX-GPNMB/EDNRB/GCGR	0.9	1	0.818182
COMPLEX-CD300C/EDNRB/SDC1	0.9	1	0.818182
EDNRB AND NOT-AQP10 AND GPNMB	0.9	1	0.818182
COMPLEX-GPNMB/EDNRB/TACSTD2	0.9	1	0.818182
COMPLEX-GPNMB/EDNRB/LRRC4B	0.9	1	0.818182
NOT-SLC5A5 AND GPNMB AND NOT-FOLR1	0.9	1	0.818182
COMPLEX-GPNMB/EDNRB/SLC14A2	0.9	1	0.818182
COMPLEX-GPNMB/EDNRB/HCRTR1	0.9	1	0.818182
COMPLEX-FCRL1/EDNRB/TM4SF5	0.9	1	0.818182
EDNRB AND NOT-SSTR1 AND NOT-AQP10	0.9	1	0.818182
COMPLEX-CD300C/EDNRB/SSTR1	0.9	1	0.818182
COMPLEX-GPNMB/EDNRB/FLRT3	0.9	1	0.818182
COMPLEX-GPNMB/EDNRB/ATP1A1	0.9	1	0.818182
COMPLEX-CHRM4/EDNRB/STEAP2	0.9	1	0.818182
COMPLEX-EDNRB/STEAP2/OR2S2	0.9	1	0.818182
COMPLEX-EDNRB/STEAP2/OR4D1	0.9	1	0.818182
COMPLEX-CD300LB/EDNRB/STEAP2	0.9	1	0.818182
COMPLEX-CD300C/EDNRB/STEAP2	0.9	1	0.818182
COMPLEX-CLDN23/EDNRB/SLC5A5	0.9	1	0.818182
COMPLEX-FCRL1/EDNRB/GIPR	0.9	1	0.818182
COMPLEX-GPNMB/EDNRB/OSMR	0.9	1	0.818182
COMPLEX-GPNMB/CD300C/EDNRB	0.9	1	0.818182
COMPLEX-GPNMB/EDNRB/IHH	0.9	1	0.818182
COMPLEX-EDNRB/OR4D1/VCAM1	0.9	1	0.818182
COMPLEX-EDNRB/RNF43/SLC5A5	0.9	1	0.818182
COMPLEX-EDNRB/SDC1/TAAR8	0.9	1	0.818182
COMPLEX-GPNMB/TSPAN9/EDNRB	0.9	1	0.818182
COMPLEX-GPNMB/CHRM4/EDNRB	0.9	1	0.818182
EDNRB AND VCAM1 AND NOT-AQP10	0.9	1	0.818182
COMPLEX-EDNRB/ITGAV/TAAR8	0.9	1	0.818182
COMPLEX-EDNRB/OR51I2/VCAM1	0.9	1	0.818182
COMPLEX-EDNRB/GIPR/FCRL2	0.9	1	0.818182
COMPLEX-EDNRB/TPBG/TM4SF5	0.9	1	0.818182
COMPLEX-GPNMB/CD300LB/EDNRB	0.9	1	0.818182
NOT-GIPR AND GPNMB AND NOT-CLDN4	0.9	1	0.818182
COMPLEX-EDNRB/OR2S2/VCAM1	0.9	1	0.818182
COMPLEX-EDNRB/SDC1/TM4SF5	0.9	1	0.818182
COMPLEX-EDNRB/SLC5A5/VCAM1	0.9	1	0.818182
NOT-SLC5A5 AND GPNMB AND NOT-CLDN4	0.9	1	0.818182
COMPLEX-EDNRB/STEAP2/TAAR8	0.9	1	0.818182
COMPLEX-FCRL1/SLC34A3/EDNRB	0.9	1	0.818182
COMPLEX-EDNRB/GPR15/SDC1	0.9	1	0.818182
COMPLEX-GPNMB/EDNRB/FGFR3	0.9	1	0.818182
COMPLEX-EDNRB/STEAP2/SLC5A5	0.9	1	0.818182
NOT-OR2S2 AND GPNMB AND NOT-FOLR1	0.9	1	0.818182
COMPLEX-GPNMB/CHRM5/EDNRB	0.9	1	0.818182
COMPLEX-EDNRB/SDC1/SLC5A5	0.9	1	0.818182
COMPLEX-GPNMB/EDNRB/ANO1	0.9	1	0.818182
COMPLEX-SLC34A3/EDNRB/FCRL2	0.9	1	0.818182
NOT-TAAR8 AND GPNMB AND NOT-VTCN1	0.952381	1	0.909091
NOT-OR2S2 AND GPNMB AND NOT-EPCAM	0.952381	1	0.909091
NOT-SLC5A5 AND SDC1 AND NOT-EPCAM	0.952381	1	0.909091
NOT-CLDN19 AND GPNMB AND NOT-CLDN8	1	1	1
NOT-CHRNA2 AND GPNMB AND NOT-CLDN8	1	1	1
NOT-CHRNA2 AND SDC1 AND NOT-CLDN8	1	1	1
GPR137B AND NOT-CLDN8 AND ERBB3	1	1	1
GPR137B AND NOT-CLDN8 AND ERBB3	1	1	1
GPR137B AND NOT-CLDN8 AND ERBB2	0.952381	1	0.909091
NOT-CLDN19 AND GPNMB AND NOT-LGR5	0.952381	1	0.909091
GPR137B AND NOT-CLDN8 AND ERBB2	0.952381	1	0.909091
NOT-CLDN19 AND GPNMB AND NOT-IL20RA	0.952381	1	0.909091
NOT-CLDN19 AND GPNMB AND NOT-VTCN1	0.952381	1	0.909091
NOT-CHRNA2 AND SDC1 AND NOT-VTCN1	0.952381	1	0.909091
NOT-CLDN19 AND GPNMB AND NOT-ERBB2	1	1	1
NOT-CLDN19 AND GPNMB AND NOT-ERBB2	1	1	1
NOT-CHRNA2 AND VCAM1 AND NOT-CLDN8	1	1	1

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
MUC13 AND EDNRA AND NOT-EQTN	1	1	1
HEPH AND CLDN1 AND NOT-ADRB2	1	1	1
MUC13 AND EDNRA AND SLC38A2	1	1	1
HEPH AND CLDN1 AND CYBA	1	1	1
HEPH AND CLDN1 AND NOT-CTLA4	1	1	1
MUC13 AND EDNRA AND NOT-ADAM22	1	1	1
HEPH AND CLDN1 AND NOT-CD300LG	1	1	1
MUC13 AND EDNRA AND NOT-PKD1	1	1	1
DPEP1 AND NOT-SLC22A7 AND NOT-CNST	1	1	1
MUC13 AND EDNRA AND NOT-C14orf180	1	1	1
MUC13 AND EDNRA AND NOT-LAG3	1	1	1
MUC13 AND EDNRA AND NOT-HCRTR1	1	1	1
DPEP1 AND TMEM123 AND NOT-TNFRSF25	1	1	1
DPEP1 AND NOT-ZP2 AND TM4SF5	1	1	1
DPEP1 AND TMEM123 AND NOT-SLC4A4	1	1	1
MUC13 AND EDNRA AND NOT-CD274	1	1	1
MUC13 AND EDNRA AND NOT-GRIA3	1	1	1
DPEP1 AND TMEM123 AND NOT-UNC5C	1	1	1
DPEP1 AND TMEM123 AND NOT-ITGA8	1	1	1
MUC13 AND EDNRA AND NOT-SLC9A9	1	1	1
MUC13 AND EDNRA AND NOT-GPR25	1	1	1
MUC13 AND EDNRA AND NOT-SLC9C2	1	1	1
MUC13 AND EDNRA AND NOT-IZUMO1	1	1	1
MUC13 AND EDNRA AND NOT-GPR17	1	1	1
MUC13 AND EDNRA AND NOT-NPBWR2	1	1	1
MUC13 AND EDNRA AND NOT-OR8D2	1	1	1
DPEP1 AND TMEM123 AND NOT-COL25A1	1	1	1
DPEP1 AND TMEM123 AND NOT-LRRC8C	1	1	1
MUC13 AND EDNRA AND NOT-GLP1R	1	1	1
MUC13 AND EDNRA AND NOT-SIGLEC9	1	1	1
DPEP1 AND NOT-ZP2 AND DEGS1	1	1	1
DPEP1 AND NOT-ZP2 AND GALR3	1	1	1
DPEP1 AND TMEM123 AND NOT-COLQ	1	1	1
DPEP1 AND TMEM123 AND NOT-ADAM23	1	1	1
MUC13 AND EDNRA AND NOT-KCNIP3	1	1	1
DPEP1 AND TMEM123 AND NOT-SYT12	1	1	1
MUC13 AND EDNRA AND NOT-AOC2	1	1	1
HEPH AND CLDN1 AND NOT-NAALAD2	1	1	1
DPEP1 AND NOT-ZP2 AND TNFSF15	1	1	1
MUC13 AND EDNRA AND NOT-KCNS2	1	1	1
MUC13 AND EDNRA AND NOT-KCNK3	1	1	1
MUC13 AND EDNRA AND NOT-KCNJ14	1	1	1
MUC13 AND EDNRA AND NOT-KCNJ4	1	1	1
MUC13 AND EDNRA AND NOT-KCNC3	1	1	1
MUC13 AND EDNRA AND NOT-KCNA3	1	1	1
DPEP1 AND NOT-ZP2 AND ATP2C2	1	1	1
MUC13 AND EDNRA AND NOT-ITGA9	1	1	1
MUC13 AND EDNRA AND NOT-ITGA2B	1	1	1
MUC13 AND EDNRA AND NOT-INSRR	1	1	1
MUC13 AND EDNRA AND NOT-AQP4	1	1	1
DPEP1 AND NOT-ZP2 AND HEPH	1	1	1
MUC13 AND EDNRA AND NOT-IL5RA	1	1	1
DPEP1 AND TMEM123 AND NOT-CACNA1H	1	1	1
MUC13 AND EDNRA AND NOT-PAQR9	1	1	1
DPEP1 AND TMEM123 AND NOT-VNN2	1	1	1
MUC13 AND EDNRA AND NOT-HTR3A	1	1	1
MUC13 AND EDNRA AND NOT-APLP1	1	1	1
DPEP1 AND NOT-ZP2 AND ACVR1	1	1	1
LY75 AND THY1 AND NOT-CLDN10	1	1	1
DPEP1 AND NOT-SPAM1 AND NOT-FLVCR2	1	1	1
DPEP1 AND NOT-SPAM1 AND NOT-IL17RB	1	1	1
BIRC5 AND IFITM1 AND NOT-FUT3	1	1	1
DPEP1 AND NOT-SLC13A3 AND FXYD6	1	1	1
DPEP1 AND NOT-PCDHGB6 AND CDH11	1	1	1
DPEP1 AND SLC39A10 AND NOT-NMBR	1	1	1
TGFBI AND NOT-LPAR1 AND STEAP2	1	1	1
DPEP1 AND NOT-SPAM1 AND NOT-GDPD2	1	1	1
DPEP1 AND NOT-CSMD3 AND JPH1	1	1	1
DPEP1 AND NOT-NMBR AND PCDH18	1	1	1
DPEP1 AND FXYD5 AND NOT-NMBR	1	1	1
DPEP1 AND NOT-SPAM1 AND NOT-CDHR5	1	1	1
DPEP1 AND NOT-CD1B AND NOT-SLC7A13	1	1	1
DPEP1 AND NOT-SPAM1 AND OLR1	1	1	1
DPEP1 AND NOT-MMP16 AND CD93	1	1	1
DPEP1 AND NOT-NMBR AND NINJ2	1	1	1
DPEP1 AND NOT-SPAM1 AND ABCC1	1	1	1
DPEP1 AND NOT-SPAM1 AND TACSTD2	1	1	1

Antigen Pair Logic	F1	Precision	Recall
GPR137B AND NOT-IL20RA AND ERBB3	0.952381	1	0.909091
GPR137B AND NOT-CLDN8 AND L1CAM	0.952381	1	0.909091
NOT-CHRNA2 AND GPNMB AND NOT-VTCN1	0.952381	1	0.909091
NOT-CACNG6 AND SDC1 AND NOT-EPCAM	0.952381	1	0.909091
NOT-CHRNA2 AND SDC1 AND NOT-EPCAM	0.952381	1	0.909091
NOT-CLDN19 AND GPNMB AND NOT-EPCAM	0.952381	1	0.909091
GPR137B AND NOT-ERBB2 AND ERBB3	1	1	1
GPR137B AND NOT-ERBB2 AND ERBB3	1	1	1
NOT-CHRNA2 AND GPNMB AND NOT-ERBB2	1	1	1
NOT-CLDN19 AND GPNMB AND NOT-EGFR	1	1	1
GPR137B AND NOT-CLDN8 AND SDC1	0.956522	0.916667	1
COMPLEX-CHRNA2/ITGAV/L1CAM	0.952381	1	0.909091
GPR137B AND NOT-CLDN8 AND SDC1	0.956522	0.916667	1
NOT-CLDN19 AND GPNMB AND NOT-CEACAM6	1	1	1
NOT-CHRNA2 AND GPNMB AND NOT-IL20RA	0.952381	1	0.909091
NOT-CHRNA2 AND ENG AND NOT-ERBB2	1	1	1
NOT-CHRNA2 AND ITGAV AND NOT-CLDN8	0.916667	0.846154	1
NOT-CHRNA2 AND ERBB2 AND NOT-CLDN8	0.916667	0.846154	1
GPR137B AND NOT-EPCAM AND ERBB3	0.952381	1	0.909091
NOT-CHRNA2 AND VCAM1 AND NOT-VTCN1	0.952381	1	0.909091
GPR137B AND NOT-EGFR AND ERBB3	0.952381	1	0.909091
COMPLEX-CHRNA2/ABCB5/VCAM1	1	1	1
NOT-CHRNA2 AND GPNMB AND NOT-EPCAM	0.952381	1	0.909091
NOT-CACNG6 AND GPNMB AND NOT-EPCAM	0.952381	1	0.909091
NOT-CLDN19 AND GPNMB AND NOT-CLDN1	1	1	1
NOT-GJD2 AND GPNMB AND NOT-EPCAM	0.952381	1	0.909091
NOT-CHRNA2 AND SDC1 AND NOT-IL20RA	0.952381	1	0.909091
GPR137B AND NOT-CLDN8 AND IL13RA2	0.952381	1	0.909091
NOT-CHRNA2 AND SDC1 AND NOT-CEACAM6	1	1	1
GPR137B AND NOT-CLDN8 AND THY1	0.952381	1	0.909091
NOT-CHRNA2 AND VCAM1 AND NOT-IL20RA	0.952381	1	0.909091
GPR137B AND NOT-IL20RA AND SDC1	0.909091	0.909091	0.909091
NOT-CACNG6 AND GPNMB AND NOT-CLDN8	1	1	1
NOT-GJD2 AND GPNMB AND NOT-CLDN8	1	1	1
NOT-CLDN19 AND VCAM1 AND NOT-CLDN8	1	1	1
GPR137B AND NOT-LGR5 AND ERBB3	0.952381	1	0.909091
GPR137B AND NOT-CLDN8 AND AXL	0.952381	1	0.909091
NOT-CHRNA2 AND SDC1 AND NOT-EGFR	0.952381	1	0.909091
NOT-CACNG6 AND GPNMB AND NOT-CLDN4	0.952381	1	0.909091
NOT-CACNG6 AND GPNMB AND NOT-VTCN1	0.952381	1	0.909091
NOT-CHRNA2 AND SDC1 AND NOT-FOLR1	0.9	1	0.818182
COMPLEX-GPNMB/EDNRB/GRIK4	0.9	1	0.818182
COMPLEX-EDNRB/GPR137B/CLDN8	0.9	1	0.818182
GPR137B AND NOT-CLDN8 AND TNC	0.9	1	0.818182
COMPLEX-GPNMB/EDNRB/GPR26	0.9	1	0.818182
COMPLEX-EDNRB/SLC10A2/FCRL2	0.9	1	0.818182
GPR137B AND NOT-IL20RA AND ERBB2	0.9	1	0.818182
COMPLEX-CHRNA2/L1CAM/VCAM1	0.9	1	0.818182
COMPLEX-CHRNA2/EDNRB/MST1R	0.9	1	0.818182
COMPLEX-FCRL1/EDNRB/SLC10A2	0.9	1	0.818182
COMPLEX-OR1Q1/EDNRB/SDC1	0.9	1	0.818182
COMPLEX-EDNRB/STEAP2/SLC10A2	0.9	1	0.818182
COMPLEX-GPNMB/EDNRB/LRFN2	0.9	1	0.818182
COMPLEX-CLDN19/EDNRB/RNF43	0.9	1	0.818182
NOT-CLDN19 AND GPNMB AND NOT-FOLR1	0.9	1	0.818182
COMPLEX-EDNRB/STEAP2/CACNG6	0.9	1	0.818182
GPR137B AND NOT-CLDN8 AND FAP	0.9	1	0.818182
COMPLEX-EDNRB/RNF43/GPR137B	0.9	1	0.818182
COMPLEX-GPNMB/EDNRB/VN1R2	0.9	1	0.818182
COMPLEX-GPNMB/EDNRB/OR3A1	0.9	1	0.818182
NOT-CHRNA2 AND STEAP1 AND NOT-SSTR1	0.9	1	0.818182
COMPLEX-GPNMB/EDNRB/CACNG6	0.9	1	0.818182
NOT-CHRNA2 AND STEAP1 AND NOT-ERBB2	0.9	1	0.818182
COMPLEX-CLDN19/EDNRB/FCRL2	0.9	1	0.818182
COMPLEX-EDNRB/GPR137B/MUC16	0.9	1	0.818182
COMPLEX-CHRNA2/EDNRB/STEAP2	0.9	1	0.818182
COMPLEX-CHRNA2/EDNRB/CLDN1	0.9	1	0.818182
COMPLEX-GPNMB/EDNRB/ABCG8	0.9	1	0.818182
COMPLEX-EDNRB/STEAP2/NOX1	0.9	1	0.818182
COMPLEX-FCRL1/CLDN19/EDNRB	0.9	1	0.818182
COMPLEX-CHRNA2/EDNRB/RNF43	0.9	1	0.818182
EDNRB AND NOT-CHRNA2 AND IL11RA	0.9	1	0.818182
COMPLEX-EDNRB/SLAMF7/GPR137B	0.9	1	0.818182
COMPLEX-CHRNA2/CLDN23/EDNRB	0.9	1	0.818182
COMPLEX-EDNRB/GPR137B/CLDN1	0.9	1	0.818182
EDNRB AND GPR137B AND IL11RA	0.9	1	0.818182
COMPLEX-EDNRB/SDC1/ABCG8	0.9	1	0.818182

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
DPEP1 AND NOT-SPAM1 AND OR7A10	1	1	1
DPEP1 AND NOT-NMBR AND ITGA4	1	1	1
DPEP1 AND NOT-SLC13A1 AND CDH7	1	1	1
DPEP1 AND LRP8 AND NOT-NMBR	1	1	1
DPEP1 AND NOT-SPAM1 AND NOT-SLC2A9	1	1	1
DPEP1 AND SLC52A2 AND NOT-NMBR	1	1	1
IFI6 AND GJB3 AND NOT-MAGEA11	1	1	1
DPEP1 AND NOT-SPAM1 AND NOT-SLC15A1	1	1	1
DPEP1 AND NOT-SPAM1 AND NOT-SLC8A1	1	1	1
DPEP1 AND NOT-SPAM1 AND NOT-SLC5A1	1	1	1
DPEP1 AND NOT-SPAM1 AND NOT-SLC3A1	1	1	1
DPEP1 AND NOT-SPAM1 AND SLC1A4	1	1	1
DPEP1 AND NOT-SPAM1 AND NOT-SORT1	1	1	1
DPEP1 AND NOT-GABRD AND CYBB	1	1	1
DPEP1 AND NOT-PCDHGB6 AND MSMO1	1	1	1
IFI6 AND GJB3 AND NOT-FLOT2	1	1	1
DPEP1 AND NOT-PCDHGB6 AND TNFRSF12A	1	1	1
DPEP1 AND NOT-CSMD3 AND SLC1A4	1	1	1
IFI6 AND GJB3 AND NOT-NCAM1	1	1	1
DPEP1 AND NOT-SPAM1 AND NOT-ABCD4	1	1	1
DPEP1 AND NOT-SLC13A1 AND ABCA8	1	1	1
IFI6 AND GJB3 AND NOT-VTCN1	1	1	1
DPEP1 AND NOT-SPAM1 AND PTPRM	1	1	1
IFI6 AND GJB3 AND NOT-MUC16	1	1	1
DPEP1 AND NOT-SPAM1 AND NOT-PTPRF	1	1	1
DPEP1 AND NOT-PCDHGB6 AND S1PR1	1	1	1
DPEP1 AND NOT-SPAM1 AND JPH1	1	1	1
DPEP1 AND NOT-SPAM1 AND KCNMA1	1	1	1
LY75 AND THY1 AND NOT-CALN1	1	1	1
DPEP1 AND PLP2 AND NOT-CDH22	1	1	1
DPEP1 AND NOT-CDH7 AND CD44	1	1	1
DPEP1 AND NOT-SPAM1 AND NOT-AQP11	1	1	1
LY75 AND THY1 AND NOT-SLC12A5	1	1	1
DPEP1 AND NOT-CLDN20 AND KCNE3	1	1	1
DPEP1 AND NOX1 AND NOT-SIGLEC9	1	1	1
DPEP1 AND NOT-EPHA10 AND SLC22A3	1	1	1
DPEP1 AND NOT-KCNA10 AND KCNE3	1	1	1
DPEP1 AND NOT-TAS2R39 AND CEACAM7	1	1	1
DPEP1 AND KCNE3 AND NOT-GRIN2C	1	1	1
DPEP1 AND NOT-SPAM1 AND MYOF	1	1	1
COMPLEX-DPEP1/SERINC5/SPAM1	1	1	1
DPEP1 AND KCNE3 AND NOT-GPR12	1	1	1
DPEP1 AND NOT-SPAM1 AND NOT-TSPAN12	1	1	1
DPEP1 AND KCNE3 AND NOT-OR10J1	1	1	1
DPEP1 AND NOX1 AND NOT-TAS2R39	1	1	1
DPEP1 AND NOT-SPAM1 AND ABCA1	1	1	1
IFI6 AND CLDN3 AND NOT-IL18RAP	1	1	1
DPEP1 AND NOT-SPAM1 AND NOT-DSC2	1	1	1
DPEP1 AND NOT-DSC1 AND KCNE3	1	1	1
DPEP1 AND NOT-ABCG5 AND SLC22A3	1	1	1
DPEP1 AND NOT-SLC2A2 AND SLC22A3	1	1	1
DPEP1 AND NOT-OR1C1 AND NOT-SIGLEC9	1	1	1
DPEP1 AND KCNE3 AND NOT-SLC30A10	1	1	1
DPEP1 AND KCNE3 AND NOT-CD36	1	1	1
LY75 AND THY1 AND NOT-STEAP4	1	1	1
DPEP1 AND MMD AND NOT-NMBR	1	1	1
DPEP1 AND NOT-SPAM1 AND NOT-AQP7	1	1	1
IFI6 AND BIRC5 AND NOT-CLEC1A	1	1	1
IFI6 AND MUC13 AND NOT-CLEC1A	1	1	1
DPEP1 AND NOT-SIGLEC9 AND CEACAM7	1	1	1
DPEP1 AND NOT-ATP1B2 AND NOT-CSMD3	1	1	1
IFI6 AND GPA33 AND NOT-FCER2	1	1	1
DPEP1 AND NOT-GABBR2 AND KCNE3	1	1	1
BIRC5 AND IFITM1 AND NOT-SLC13A2	1	1	1
DPEP1 AND NOT-GRM2 AND NOT-SPAM1	1	1	1
IFI6 AND EPHB2 AND NOT-FCER2	1	1	1
DPEP1 AND KCNE3 AND NOT-FCAMR	1	1	1
DPEP1 AND NOT-OR2C3 AND KCNE3	1	1	1
DPEP1 AND KCNE3 AND NOT-TRPM3	1	1	1
DPEP1 AND NOT-SPAM1 AND NOT-APLP2	1	1	1
DPEP1 AND NOT-SIGLEC9 AND CD93	1	1	1
DPEP1 AND NOT-SPAM1 AND NOT-HMOX1	1	1	1
LY75 AND THY1 AND NOT-SLC1A6	1	1	1
DPEP1 AND NOT-SPAM1 AND NOT-TRHDE	1	1	1
DPEP1 AND NOT-SPAM1 AND TGFBR3	1	1	1
IFI6 AND SLC26A3 AND NOT-TNFRSF13C	1	1	1
DPEP1 AND NOT-SPAM1 AND TRPC1	1	1	1

Antigen Pair Logic	F1	Precision	Recall
COMPLEX-EDNRB/SSTR5/GPR137B	0.9	1	0.818182
COMPLEX-GPNMB/SLC22A12/EDNRB	0.9	1	0.818182
COMPLEX-EDNRB/GPR137B/CLDN2	0.9	1	0.818182
COMPLEX-CLDN23/CLDN19/EDNRB	0.9	1	0.818182
COMPLEX-CLDN19/EDNRB/STEAP2	0.9	1	0.818182
COMPLEX-EDNRB/GPR137B/TPBG	0.9	1	0.818182
COMPLEX-EDNRB/SDC1/GPR137B	0.9	1	0.818182
COMPLEX-MLANA/SLC2A1/SSTR1	1	1	1
COMPLEX-MLANA/SLC2A1/LGR5	1	1	1
COMPLEX-MLANA/ABCA12/SSTR1	1	1	1
COMPLEX-MLANA/EPCAM/SLC2A1	1	1	1
COMPLEX-MLANA/ABCA12/EPCAM	1	1	1
COMPLEX-MLANA/ABCA12/LGR5	1	1	1
COMPLEX-MLANA/SLC2A1/SSTR5	1	1	1
COMPLEX-CLDN3/MLANA/ABCA12	1	1	1
COMPLEX-MLANA/ABCA12/VTCN1	1	1	1
COMPLEX-MLANA/ABCA12/ULBP3	1	1	1
COMPLEX-EGFR/MLANA/ABCA12	1	1	1
COMPLEX-ERBB2/MLANA/ABCA12	1	1	1
COMPLEX-DPEP1/MLANA/ABCA12	1	1	1
COMPLEX-MLANA/SLC2A1/CLDN1	1	1	1
COMPLEX-DPEP1/MLANA/SLC2A1	1	1	1
COMPLEX-ERBB2/MLANA/SLC2A1	1	1	1
COMPLEX-MLANA/SLC2A1/SSTR3	1	1	1
COMPLEX-MLANA/ABCA12/SSTR5	1	1	1
COMPLEX-CLDN3/MLANA/SLC2A1	1	1	1
COMPLEX-MLANA/ABCA12/CLDN1	1	1	1
COMPLEX-MLANA/ABCA12/SSTR3	1	1	1
COMPLEX-DNAJB8/MLANA/ABCA12	1	1	1
COMPLEX-DNAJB8/MLANA/SLC2A1	1	1	1
COMPLEX-MLANA/ABCA12/ULBP1	1	1	1
COMPLEX-MLANA/SLC2A1/ULBP3	1	1	1
COMPLEX-MLANA/SLC2A1/VTCN1	1	1	1
COMPLEX-MLANA/SLC2A1/ULBP1	1	1	1
COMPLEX-EGFR/MLANA/SLC2A1	1	1	1
COMPLEX-MLANA/SLC2A1/CLDN8	1	1	1
MLANA AND GPR137B AND IL11RA	1	1	1
MLANA AND NOT-CLDN1 AND GPR137B	1	1	1
MLANA AND GPR137B AND NOT-ERBB2	1	1	1
MLANA AND NOT-CHRNA2 AND IL11RA	1	1	1
COMPLEX-CLDN7/MLANA/ABCA12	1	1	1
COMPLEX-MLANA/STEAP2/NOX1	1	1	1
COMPLEX-TNFRSF13C/MLANA/ABCA12	1	1	1
MLANA AND NOT-CHRNA2 AND NOT-CLDN1	1	1	1
COMPLEX-MLANA/PTGER4/TPBG	1	1	1
COMPLEX-TNFRSF13C/MLANA/SLC2A1	1	1	1
COMPLEX-MLANA/ABCA12/CLDN8	1	1	1
MLANA AND NOT-CHRNA2 AND NOT-ERBB2	1	1	1
COMPLEX-MLANA/OR10J1/ULBP1	1	1	1
COMPLEX-SLC34A2/MLANA/SLC2A1	1	1	1
COMPLEX-SLC34A2/MLANA/ABCA12	1	1	1
COMPLEX-CLDN7/MLANA/SLC2A1	1	1	1
COMPLEX-MLANA/SLC2A1/SST	1	1	1
COMPLEX-MLANA/ABCA12/ENPP3	0.952381	1	0.909091
COMPLEX-MLANA/ABCA12/CD34	0.952381	1	0.909091
COMPLEX-MLANA/ENPP3/SLC2A1	0.952381	1	0.909091
COMPLEX-CLDN19/PMEL/VCAM1	0.952381	1	0.909091
COMPLEX-MLANA/SLC2A1/CD34	0.952381	1	0.909091
COMPLEX-MLANA/LCT/TPBG	0.952381	1	0.909091
COMPLEX-GPA33/MLANA/ABCA12	0.952381	1	0.909091
COMPLEX-MLANA/ABCA12/CEACAM6	0.952381	1	0.909091
COMPLEX-GPA33/MLANA/SLC2A1	0.952381	1	0.909091
COMPLEX-MLANA/SLC2A1/CLDN5	0.952381	1	0.909091
COMPLEX-MLANA/ABCA12/CLDN5	0.952381	1	0.909091
COMPLEX-MLANA/MOK/SLC2A1	1	1	1
COMPLEX-MLANA/ABCA12/MOK	1	1	1
COMPLEX-MLANA/TRPM4/SLC2A1	0.952381	1	0.909091
COMPLEX-MLANA/BIRC5/SLC2A1	0.952381	1	0.909091
COMPLEX-MLANA/ABCA12/IL11RA	1	1	1
COMPLEX-ENG/MLANA/SLC2A1	1	1	1
COMPLEX-ENG/MLANA/ABCA12	1	1	1
COMPLEX-MLANA/IL11RA/SLC2A1	1	1	1
COMPLEX-RAET1E/MLANA/SLC2A1	0.952381	1	0.909091
COMPLEX-MLANA/SLC2A1/SSTR4	1	1	1
COMPLEX-MLANA/ABCA12/TRPM4	0.952381	1	0.909091
COMPLEX-MLANA/ABCA12/BIRC5	0.952381	1	0.909091
COMPLEX-RAET1E/MLANA/ABCA12	0.952381	1	0.909091

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
DPEP1 AND NOT-P2RX6 AND NOT-LCT	1	1	1
DPEP1 AND NOT-GPC3 AND NOT-DISP1	1	1	1
DPEP1 AND NOT-GPC3 AND NOT-ADRA1B	1	1	1
DPEP1 AND NOT-GPC3 AND NOT-SLC5A1	1	1	1
DPEP1 AND NOT-GPC3 AND NOT-UNC5D	1	1	1
DPEP1 AND NOT-GPC3 AND NOT-SLC34A3	1	1	1
DPEP1 AND NOT-SLC5A12 AND NOT-GPC3	1	1	1
DPEP1 AND NOT-GPC3 AND NOT-TMEM25	1	1	1
DPEP1 AND NOT-TRHDE AND NOT-GPC3	1	1	1
DPEP1 AND NOT-GPC3 AND NOT-SLC51A	1	1	1
DPEP1 AND NOT-GPC3 AND NOT-ENPEP	1	1	1
GUCY2C AND NOT-ABHD6 AND NOT-GPC3	1	1	1
SLC26A3 AND THY1 AND NOT-GPC3	1	1	1
DPEP1 AND NOT-GPC3 AND EPHB4	1	1	1
DPEP1 AND NOT-GPC3 AND FGFR1	1	1	1
DPEP1 AND NOT-GPC3 AND GJB3	1	1	1
DPEP1 AND NOT-KCNG2 AND NOT-GPC3	1	1	1
JPH1 AND CLDN1 AND NOT-GPC3	1	1	1
MUC13 AND SLC39A10 AND NOT-MUC1	1	1	1
DPEP1 AND NOT-GPC3 AND ITGA4	1	1	1
DPEP1 AND NOT-GPC3 AND PCDH17	1	1	1
MUC13 AND CDH3 AND NOT-MUC1	1	1	1
DPEP1 AND NOT-GPC3 AND KCNE3	1	1	1
DPEP1 AND NOT-GPC3 AND CDH5	1	1	1
DPEP1 AND NOT-GPC3 AND NOT-SLC51A	1	1	1
DPEP1 AND NOT-GPC3 AND NOT-EMB	1	1	1
DPEP1 AND NOT-GPC3 AND NOT-SLC51B	1	1	1
DPEP1 AND NOT-GPC3 AND CALCRL	1	1	1
DPEP1 AND NOT-GPC3 AND PLP2	1	1	1
DPEP1 AND NOT-GPC3 AND MMD	1	1	1
DPEP1 AND NOT-GPC3 AND FXYD5	1	1	1
DPEP1 AND NOT-GPC3 AND KCNJ2	1	1	1
KCNE3 AND THY1 AND NOT-GPC3	1	1	1
DPEP1 AND NOT-GPC3 AND PIGU	1	1	1
MUC13 AND SLC39A10 AND NOT-MSLN	1	1	1
DPEP1 AND NOT-GPC3 AND CDH3	1	1	1
MEP1A AND THY1 AND NOT-ERBB2	1	1	1
XK AND CLDN1 AND NOT-GPC3	1	1	1
GUCY2C AND EDNRA AND NOT-GPC3	1	1	1
MEP1A AND THY1 AND NOT-GPC3	1	1	1
DPEP1 AND NOT-GPC3 AND NOT-SLC22A24	1	1	1
DPEP1 AND NOT-GPC3 AND NOT-SCAMP5	1	1	1
DPEP1 AND NOT-GPC3 AND CFTR	1	1	1
GUCY2C AND NOT-CEACAM1 AND NOT-GPC3	1	1	1
DPEP1 AND NOT-GPC3 AND LRP8	1	1	1
DPEP1 AND NOT-GPC3 AND CKAP4	1	1	1
DPEP1 AND NOT-GPC3 AND ADCY7	1	1	1
DPEP1 AND NOT-ADRA1B AND ERBB2	1	1	1
DPEP1 AND NOT-GPC3 AND KCNE3	1	1	1
DPEP1 AND NOT-GPC3 AND NOT-ABCC2	1	1	1
MEP1A AND CLDN1 AND NOT-ERBB2	1	1	1
DPEP1 AND NOT-GPC3 AND KCNN4	1	1	1
DPEP1 AND NOT-GPC3 AND CDH3	1	1	1
GUCY2C AND SLC39A10 AND NOT-GPC3	1	1	1
DPEP1 AND NOT-GPC3 AND NOT-PMP22	1	1	1
DPEP1 AND NOT-GPC3 AND APCDD1	1	1	1
ATP10B AND THY1 AND NOT-GPC3	1	1	1
MEP1A AND THY1 AND NOT-MUC1	1	1	1
DPEP1 AND NOT-GPC3 AND ADORA2B	1	1	1
DPEP1 AND NOT-GPC3 AND CKAP4	1	1	1
DPEP1 AND NOT-GPC3 AND NOT-PRLR	1	1	1
DPEP1 AND NOT-GPC3 AND NOT-ADAM10	1	1	1
DPEP1 AND NOT-GPC3 AND SEMA4C	1	1	1
MUC13 AND NOT-FLVCR2 AND NOT-MUC1	1	1	1
DPEP1 AND NOT-GPC3 AND NOT-SLC52A1	1	1	1
DPEP1 AND NOT-GPC3 AND EDAR	1	1	1
DPEP1 AND TMEM123 AND NOT-GPC3	1	1	1
GPA33 AND SLC39A10 AND NOT-GPC3	1	1	1
GUCY2C AND NOT-FLVCR2 AND NOT-MUC1	1	1	1
DPEP1 AND NOT-GPC3 AND NOT-NAALADL1	1	1	1
DPEP1 AND NOT-GPC3 AND TSPAN2	1	1	1
GPA33 AND SLC39A10 AND NOT-MUC1	1	1	1
DPEP1 AND NOT-GPC3 AND NOT-ATP6V0A1	1	1	1
DPEP1 AND NOT-GPC3 AND NOT-ABCB1	1	1	1
DPEP1 AND NOT-GPC3 AND NOT-SLC46A1	1	1	1
DPEP1 AND NOT-GPC3 AND NOT-AQP1	1	1	1
DPEP1 AND NOT-GPC3 AND PLXNC1	1	1	1

Antigen Pair Logic	F1	Precision	Recall
COMPLEX-MLANA/ABCA12/SSTR4	1	1	1
COMPLEX-CLDN19/MLANA/STEAP2	1	1	1
COMPLEX-MLANA/GPR137B/CLDN2	1	1	1
COMPLEX-MLANA/STEAP2/ABCG8	1	1	1
COMPLEX-ERBB3/MLANA/GPR137B	1	1	1
COMPLEX-MLANA/SSTR4/GPR137B	1	1	1
COMPLEX-MLANA/ABCA12/ANXA1	1	1	1
COMPLEX-MLANA/SLC2A1/CLDN9	1	1	1
COMPLEX-MLANA/GPR137B/CLDN9	1	1	1
COMPLEX-MLANA/STEAP2/CACNG6	1	1	1
COMPLEX-AFP/MLANA/SLC2A1	0.952381	1	0.909091
COMPLEX-MLANA/ABCA12/CLDN6	1	1	1
COMPLEX-MLANA/ABCA12/CLDN9	1	1	1
COMPLEX-MLANA/ABCA12/GUCY2C	0.952381	1	0.909091
COMPLEX-AFP/MLANA/ABCA12	0.952381	1	0.909091
COMPLEX-FCRL1/MLANA/GJD2	1	1	1
COMPLEX-MLANA/ABCA12/MST1R	1	1	1
COMPLEX-MLANA/ABCA12/CLDN2	1	1	1
COMPLEX-CLDN19/MLANA/RNF43	1	1	1
COMPLEX-MLANA/MST1R/SLC2A1	1	1	1
COMPLEX-MLANA/ANXA1/SLC2A1	1	1	1
COMPLEX-MLANA/SLC2A1/CLDN6	1	1	1
COMPLEX-STEAP1/PMEL/RNF144A	1	1	1
COMPLEX-MLANA/FREM2/EPCAM	1	1	1
COMPLEX-LDLRAD3/ANXA1/PMEL	1	1	1
COMPLEX-LDLRAD3/MUC4/PMEL	1	1	1
COMPLEX-LDLRAD3/RNF43/PMEL	1	1	1
COMPLEX-LDLRAD3/PMEL/BMPR1B	1	1	1
COMPLEX-BIRC5/PMEL/RNF144A	1	1	1
COMPLEX-FAP/PMEL/RNF144A	1	1	1
COMPLEX-CLDN4/LDLRAD3/PMEL	1	1	1
COMPLEX-LDLRAD3/EPHA3/PMEL	1	1	1
COMPLEX-LDLRAD3/MUC1/PMEL	1	1	1
COMPLEX-LDLRAD3/PMEL/CD33	1	1	1
COMPLEX-LDLRAD3/PMEL/CD38	1	1	1
COMPLEX-CLDN3/LDLRAD3/PMEL	1	1	1
COMPLEX-MOK/PMEL/NIPA2	1	1	1
COMPLEX-LDLRAD3/IL13RA1/PMEL	1	1	1
COMPLEX-MLANA/FREM2/LGR5	1	1	1
COMPLEX-GPA33/LDLRAD3/PMEL	1	1	1
COMPLEX-HHLA2/LDLRAD3/PMEL	1	1	1
COMPLEX-LDLRAD3/TRPM4/PMEL	1	1	1
COMPLEX-MLANA/FREM2/SSTR1	1	1	1
COMPLEX-L1CAM/PMEL/RNF144A	1	1	1
COMPLEX-FCRL1/PMEL/RNF144A	1	1	1
COMPLEX-LDLRAD3/GUCY2C/PMEL	1	1	1
COMPLEX-HTRA2/TNC/PMEL	1	1	1
COMPLEX-HTRA2/PMEL/LGR5	1	1	1
COMPLEX-HTRA2/PMEL/VTCN1	1	1	1
COMPLEX-HTRA2/PMEL/SSTR1	1	1	1
COMPLEX-HTRA2/PMEL/CD34	1	1	1
COMPLEX-HTRA2/PMEL/CLDN1	1	1	1
COMPLEX-ERBB2/HTRA2/PMEL	1	1	1
COMPLEX-CLDN7/HTRA2/PMEL	1	1	1
COMPLEX-HTRA2/EPCAM/PMEL	1	1	1
COMPLEX-HTRA2/ENPP3/PMEL	1	1	1
COMPLEX-ERBB2/MLANA/FREM2	1	1	1
COMPLEX-MLANA/FREM2/ULBP1	1	1	1
COMPLEX-MLANA/FREM2/SSTR3	1	1	1
COMPLEX-DPEP1/MLANA/FREM2	1	1	1
COMPLEX-MLANA/FREM2/SSTR5	1	1	1
COMPLEX-MLANA/FREM2/ULBP3	1	1	1
COMPLEX-MLANA/FREM2/VTCN1	1	1	1
COMPLEX-EGFR/MLANA/FREM2	1	1	1
COMPLEX-MLANA/FREM2/CLDN1	1	1	1
COMPLEX-CLDN3/MLANA/FREM2	1	1	1
COMPLEX-DNAJB8/MLANA/FREM2	1	1	1
COMPLEX-CLDN4/PMEL/NIPA2	1	1	1
COMPLEX-CEACAM5/PMEL/NIPA2	1	1	1
COMPLEX-FOLR1/PMEL/MARVELD1	1	1	1
COMPLEX-EGFR/PMEL/NIPA2	1	1	1
COMPLEX-SLC34A2/MLANA/FREM2	1	1	1
COMPLEX-CD160/PMEL/NIPA2	1	1	1
COMPLEX-HHLA2/PMEL/NIPA2	1	1	1
COMPLEX-AFP/PMEL/NIPA2	1	1	1
COMPLEX-CDH3/MLANA/SSTR1	1	1	1
COMPLEX-DDX3X/PMEL/NIPA2	1	1	1

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
DPEP1 AND NOT-GPC3 AND NOT-SLC27A4	1	1	1
DPEP1 AND NOT-GPC3 AND NOT-AQP3	1	1	1
DPEP1 AND NOT-GPC3 AND NOT-KCNG1	1	1	1
DPEP1 AND NOT-KCNJ13 AND NOT-GPC3	1	1	1
DPEP1 AND NOT-GPC3 AND NOT-PTPRH	1	1	1
TSPAN8 AND THY1 AND NOT-GPC3	1	1	1
DPEP1 AND NOT-GPC3 AND PCDH7	1	1	1
NOT-LIFR AND THY1 AND MUC1	1	1	1
DPEP1 AND NOT-GPC3 AND NOT-KLRB1	1	1	1
DPEP1 AND NOT-GPC3 AND NOT-SLC2A13	1	1	1
DPEP1 AND NOT-ATP6V0A1 AND ERBB2	1	1	1
MUC13 AND EDNRA AND NOT-GPC3	1	1	1
DPEP1 AND NOT-GPC3 AND NOT-ATP6V0A1	1	1	1
JPH1 AND CLDN1 AND NOT-GPC3	1	1	1
DPEP1 AND NOT-GPC3 AND PDPN	1	1	1
GUCY2C AND CDH3 AND NOT-MUC1	1	1	1
DPEP1 AND NOT-GPC3 AND NOT-SLC19A3	1	1	1
DPEP1 AND NOT-GPC3 AND JPH1	1	1	1
MEP1A AND SLC7A5 AND NOT-MUC1	1	1	1
DPEP1 AND NOT-GPC3 AND NOT-CD7	1	1	1
BIRC5 AND EPCAM AND NOT-SLC41A2	1	1	1
BIRC5 AND NOT-MUC15 AND CLDN1	1	1	1
GUCY2C AND SLC7A5 AND NOT-AMN	1	1	1
MUC13 AND SLC7A5 AND NOT-KCNG3	1	1	1
GUCY2C AND SLC7A5 AND NOT-PDCD1LG2	1	1	1
GUCY2C AND SLC7A5 AND NOT-ATP8B4	1	1	1
GUCY2C AND SLC7A5 AND NOT-CATSPERB	1	1	1
GUCY2C AND SLC7A5 AND NOT-SLC30A2	1	1	1
GUCY2C AND SLC7A5 AND NOT-UPK1B	1	1	1
GPA33 AND SLC7A5 AND NOT-C1QTNF1	1	1	1
GUCY2C AND SLC7A5 AND NOT-TMPRSS2	1	1	1
MUC13 AND CLDN1 AND LY75	1	1	1
GUCY2C AND THY1 AND NOT-KCNG3	1	1	1
DPEP1 AND NOT-EVC AND TNC	1	1	1
DPEP1 AND IFITM1 AND LGR5	1	1	1
GUCY2C AND SLC7A5 AND NOT-VAMP2	1	1	1
DPEP1 AND SLC7A5 AND NOT-KCNE1	1	1	1
GUCY2C AND SLC7A5 AND NOT-SLCO5A1	1	1	1
GUCY2C AND SLC7A5 AND NOT-COLQ	1	1	1
GUCY2C AND SLC7A5 AND NOT-CD99L2	1	1	1
GUCY2C AND SLC7A5 AND NOT-OR1G1	1	1	1
GUCY2C AND SLC7A5 AND NOT-CACNA1H	1	1	1
GUCY2C AND SLC7A5 AND NOT-VNN2	1	1	1
GUCY2C AND SLC7A5 AND NOT-IL18RAP	1	1	1
DPEP1 AND NOT-MAGEA1 AND SLC52A2	1	1	1
GUCY2C AND SLC7A5 AND NOT-ADAM23	1	1	1
DPEP1 AND NOT-CLDN18 AND SLC52A2	1	1	1
GUCY2C AND SLC7A5 AND NOT-TNFRSF25	1	1	1
GUCY2C AND SLC7A5 AND NOT-UNC5C	1	1	1
GUCY2C AND SLC7A5 AND NOT-CAV3	1	1	1
GUCY2C AND SLC7A5 AND NOT-ABCC11	1	1	1
GUCY2C AND SLC7A5 AND NOT-ITGA8	1	1	1
GUCY2C AND SLC7A5 AND NOT-RELT	1	1	1
GUCY2C AND SLC7A5 AND NOT-GALR3	1	1	1
GUCY2C AND SLC7A5 AND NOT-COL25A1	1	1	1
GUCY2C AND SLC7A5 AND NOT-LRRC8C	1	1	1
DPEP1 AND TMED1 AND NOT-IL11RA	1	1	1
DPEP1 AND SLC7A5 AND NOT-TEK	1	1	1
DPEP1 AND NOT-SLC26A7 AND TNC	1	1	1
DPEP1 AND NOT-IL12RB2 AND PTK7	1	1	1
DPEP1 AND NOT-CD79B AND CLEC2B	1	1	1
GUCY2C AND SLC7A5 AND NOT-SCTR	1	1	1
GUCY2C AND SLC7A5 AND NOT-SCN4B	1	1	1
DPEP1 AND SLC7A5 AND NOT-LIFR	1	1	1
GUCY2C AND SLC7A5 AND NOT-SLC5A7	1	1	1
DPEP1 AND IFITM1 AND NOT-CD79B	1	1	1
DPEP1 AND NOT-CD79B AND RYK	1	1	1
MUC13 AND THY1 AND NOT-IL1R2	1	1	1
DPEP1 AND NOT-SLC6A9 AND TNC	1	1	1
GUCY2C AND SLC7A5 AND NOT-TMEM8A	1	1	1
DPEP1 AND KCNJ2 AND NOT-CD79B	1	1	1
GUCY2C AND SLC7A5 AND NOT-PTPRN	1	1	1
GUCY2C AND SLC7A5 AND NOT-PTPRG	1	1	1
DPEP1 AND ANXA1 AND NOT-VSTM4	1	1	1
GUCY2C AND SLC7A5 AND NOT-PTPRC	1	1	1
GUCY2C AND SLC7A5 AND NOT-LRRC4	1	1	1
GUCY2C AND SLC7A5 AND NOT-CDH24	1	1	1

Antigen Pair Logic	F1	Precision	Recall
COMPLEX-MLANA/SLC27A1/EPCAM	1	1	1
COMPLEX-RAET1E/PMEL/NIPA2	1	1	1
COMPLEX-MSLN/PMEL/NIPA2	1	1	1
COMPLEX-CLDN7/PMEL/NIPA2	1	1	1
COMPLEX-GPA33/PMEL/NIPA2	1	1	1
COMPLEX-SLC34A2/PMEL/NIPA2	1	1	1
COMPLEX-CD52/PMEL/NIPA2	1	1	1
COMPLEX-MLANA/LGR5/NRP1	1	1	1
COMPLEX-ANXA1/PMEL/MARVELD1	1	1	1
COMPLEX-CNIH3/MLANA/LGR5	1	1	1
COMPLEX-L1CAM/PMEL/MARVELD1	1	1	1
COMPLEX-CEACAM6/PMEL/MARVELD1	1	1	1
COMPLEX-MLANA/SSTR1/NRP1	1	1	1
COMPLEX-PMEL/MARVELD1/CLDN8	1	1	1
COMPLEX-CNIH3/MLANA/SSTR1	1	1	1
COMPLEX-FOLR1/HTRA2/PMEL	1	1	1
COMPLEX-HTRA2/IL11RA/PMEL	1	1	1
COMPLEX-HTRA2/L1CAM/PMEL	1	1	1
COMPLEX-HTRA2/CEACAM6/PMEL	1	1	1
COMPLEX-MLANA/SLC20A2/LGR5	1	1	1
COMPLEX-HTRA2/PMEL/BMPR1B	1	1	1
COMPLEX-HTRA2/PMEL/SST	1	1	1
COMPLEX-HTRA2/PMEL/TYR	1	1	1
COMPLEX-MLANA/MACF1/SSTR1	1	1	1
COMPLEX-HSPA5/PMEL/MARVELD1	1	1	1
COMPLEX-MLANA/PMEL/MARVELD1	1	1	1
COMPLEX-MAGEA1/PMEL/NIPA2	1	1	1
COMPLEX-B4GALNT1/PMEL/NIPA2	1	1	1
COMPLEX-KDR/PMEL/NIPA2	1	1	1
COMPLEX-CDH3/MLANA/PMEL	1	1	1
COMPLEX-MLANA/FREM2/PMEL	0.952381	1	0.909091
COMPLEX-CNIH3/MLANA/PMEL	0.952381	1	0.909091
MLANA AND NOT-SLC5A5 AND KDR	1	1	1
MLANA AND NOT-OR2S2 AND KDR	1	1	1
MLANA AND NOT-CD300LB AND KDR	1	1	1
COMPLEX-MLANA/PMEL/NRP1	0.952381	1	0.909091
MLANA AND NOT-IHH AND KDR	1	1	1
COMPLEX-MLANA/CLEC1A/PMEL	0.952381	1	0.909091
MLANA AND NOT-SLC34A3 AND KDR	1	1	1
COMPLEX-MLANA/SLC27A1/PMEL	0.952381	1	0.909091
COMPLEX-MLANA/SLCO4A1/PMEL	0.952381	1	0.909091
COMPLEX-MLANA/MACF1/PMEL	0.952381	1	0.909091
COMPLEX-MLANA/PMEL/SLC20A2	0.952381	1	0.909091
MLANA AND NOT-TM4SF5 AND KDR	1	1	1
MLANA AND NOT-GIPR AND KDR	1	1	1
COMPLEX-MLANA/KCNN4/PMEL	0.952381	1	0.909091
COMPLEX-MLANA/MACF1/B4GALNT1	1	1	1
COMPLEX-MLANA/B4GALNT1/SLCO4A1	1	1	1
COMPLEX-MLANA/B4GALNT1/SLC20A2	1	1	1
COMPLEX-MLANA/B4GALNT1/PLXDC1	0.956522	0.916667	1
COMPLEX-MLANA/PLXDC1/PMEL	0.909091	0.909091	0.909091
COMPLEX-MLANA/GLG1/KDR	0.9	1	0.818182
COMPLEX-MLANA/KCNN4/KDR	0.9	1	0.818182
COMPLEX-MLANA/SLCO4A1/KDR	0.9	1	0.818182
COMPLEX-MLANA/MACF1/KDR	0.9	1	0.818182
COMPLEX-CNIH3/MLANA/KDR	0.9	1	0.818182
COMPLEX-MLANA/KDR/SLC20A2	0.9	1	0.818182
COMPLEX-CDH3/MLANA/KDR	0.9	1	0.818182
COMPLEX-MLANA/FREM2/KDR	0.9	1	0.818182
COMPLEX-MLANA/B4GALNT1/APLP2	0.88	0.785714	1
COMPLEX-MLANA/KDR/PLXDC1	0.857143	0.9	0.818182
COMPLEX-MLANA/APLP2/PMEL	0.833333	0.769231	0.909091
NOT-OR2S2 AND KDR AND NOT-B4GALNT1	0.842105	1	0.727273
NOT-GIPR AND KDR AND NOT-B4GALNT1	0.842105	1	0.727273
MLANA AND NOT-NOX1 AND KDR	1	1	1
MLANA AND NOT-GJD2 AND KDR	1	1	1
MLANA AND NOT-CACNG6 AND KDR	1	1	1
COMPLEX-MLANA/ABCA12/PMEL	0.952381	1	0.909091
COMPLEX-MLANA/PMEL/SLC2A1	0.952381	1	0.909091
MLANA AND NOT-CLDN19 AND KDR	1	1	1
MLANA AND GPR137B AND KDR	1	1	1
MLANA AND NOT-CHRNA2 AND KDR	1	1	1
MLANA AND NOT-SLC10A2 AND KDR	1	1	1
COMPLEX-MLANA/B4GALNT1/SLC2A1	1	1	1
COMPLEX-MLANA/KDR/SLC2A1	0.9	1	0.818182
COMPLEX-MLANA/ABCA12/KDR	0.9	1	0.818182
COMPLEX-MLANA/FOLR1/TPBG	1	1	1

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
GUCY2C AND SLC7A5 AND NOT-SPN	1	1	1
GUCY2C AND SLC7A5 AND NOT-SHH	1	1	1
GUCY2C AND SLC7A5 AND NOT-SIGLEC1	1	1	1
BIRC5 AND NOT-ADCY7 AND CLDN1	1	1	1
DPEP1 AND NOT-IL12RB2 AND STEAP2	1	1	1
GUCY2C AND SLC7A5 AND NOT-SLCO2A1	1	1	1
GUCY2C AND SLC7A5 AND NOT-SLC18A2	1	1	1
GUCY2C AND SLC7A5 AND NOT-SLC15A2	1	1	1
DPEP1 AND NOT-IL12RB2 AND ANXA1	1	1	1
GUCY2C AND SLC7A5 AND NOT-SLC6A8	1	1	1
GUCY2C AND SLC7A5 AND NOT-SLC6A2	1	1	1
GUCY2C AND SLC7A5 AND NOT-BMP2	1	1	1
DPEP1 AND SLC7A5 AND NOT-OR10A4	1	1	1
GUCY2C AND SLC7A5 AND NOT-SLC1A1	1	1	1
DPEP1 AND NOT-IL12RB2 AND IL20RA	1	1	1
GUCY2C AND SLC7A5 AND NOT-SLC13A3	1	1	1
DPEP1 AND GUCY2C AND NOT-SLC1A1	1	1	1
DPEP1 AND NOT-SST AND NOT-SUSD2	1	1	1
DPEP1 AND ITGAV AND NOT-PCDH9	1	1	1
DPEP1 AND ITGAV AND NOT-SLC16A2	1	1	1
MUC13 AND CLDN1 AND NOT-SLC38A6	1	1	1
DPEP1 AND OAS1 AND NOT-ABHD6	1	1	1
DPEP1 AND CEACAM6 AND NOT-ABHD6	1	1	1
DPEP1 AND ITGAV AND NOT-SMO	1	1	1
MUC13 AND CLDN1 AND NOT-CRHR2	1	1	1
DPEP1 AND CLDN4 AND NOT-ABHD6	1	1	1
DPEP1 AND OAS1 AND NOT-SLC5A1	1	1	1
MUC13 AND CLDN1 AND NOT-SLC26A9	1	1	1
DPEP1 AND ITGAV AND NOT-TIE1	1	1	1
DPEP1 AND NOT-SLC15A1 AND OAS1	1	1	1
XK AND CLDN1 AND NOT-CD160	1	1	1
DPEP1 AND ITGAV AND NOT-TRPC5	1	1	1
DPEP1 AND ITGAV AND NOT-SLC8A2	1	1	1
DPEP1 AND ITGAV AND NOT-TRPV1	1	1	1
DPEP1 AND ITGAV AND NOT-SLC4A1	1	1	1
MUC13 AND CLDN1 AND NOT-PCDHGB7	1	1	1
DPEP1 AND NOT-SLC13A1 AND PCDHA9	1	1	1
CDH17 AND THY1 AND NOT-PTPRT	1	1	1
IFI6 AND EPHB2 AND NOT-MMP24	1	1	1
CDH17 AND THY1 AND NOT-CLDN16	1	1	1
IFI6 AND EPHB2 AND NOT-CHRNB4	1	1	1
CDH17 AND THY1 AND NOT-CDH18	1	1	1
IFI6 AND EPHB2 AND NOT-NKAIN4	1	1	1
IFI6 AND EPHB2 AND NOT-SLCO6A1	1	1	1
IFI6 AND EPHB2 AND NOT-SLC32A1	1	1	1
IFI6 AND EPHB2 AND NOT-LRRN4	1	1	1
IFI6 AND EPHB2 AND NOT-KCNV2	1	1	1
CLDN7 AND TGFBI AND NOT-PAQR8	1	1	1
IFI6 AND EPHB2 AND NOT-DSC1	1	1	1
IFI6 AND EPHB2 AND NOT-AGTR2	1	1	1
IFI6 AND EPHB2 AND NOT-S1PR1	1	1	1
DPEP1 AND NOT-EPHA8 AND NOT-KCNK12	1	1	1
CDH17 AND THY1 AND NOT-ADAM29	1	1	1
CDH17 AND THY1 AND NOT-HRH3	1	1	1
IFI6 AND EPHB2 AND NOT-TMEFF2	1	1	1
CDH17 AND THY1 AND NOT-CHRND	1	1	1
IFI6 AND CLDN7 AND NOT-MLANA	1	1	1
NOX1 AND THY1 AND NOT-SLC22A12	1	1	1
NOX1 AND THY1 AND NOT-CSMD2	1	1	1
NOX1 AND THY1 AND NOT-CHRND	1	1	1
MUC13 AND TGFBI AND CDH11	1	1	1
NOX1 AND THY1 AND NOT-HRH3	1	1	1
NOX1 AND THY1 AND NOT-PTPRT	1	1	1
NOX1 AND THY1 AND NOT-ADAM29	1	1	1
NOX1 AND THY1 AND NOT-CLDN16	1	1	1
NOX1 AND THY1 AND NOT-CDH18	1	1	1
CDH17 AND THY1 AND NOT-MRGPRX2	1	1	1
CDH17 AND THY1 AND NOT-SLC22A12	1	1	1
CDH17 AND THY1 AND NOT-CSMD2	1	1	1
IFI6 AND MUC13 AND NOT-MLANA	1	1	1
IFI6 AND EPHB2 AND NOT-ATP8A1	1	1	1
IFI6 AND EPHB2 AND NOT-SYT11	1	1	1
DPEP1 AND NOT-KIRREL2 AND CD93	1	1	1
NOX1 AND THY1 AND NOT-ANTXR2	1	1	1
DPEP1 AND NOT-KCNK10 AND NOX1	1	1	1
IFI6 AND EPHB2 AND NOT-KCNJ9	1	1	1
DPEP1 AND NOX1 AND NOT-AQP6	1	1	1

Antigen Pair Logic	F1	Precision	Recall
COMPLEX-CLDN7/MLANA/TPBG	1	1	1
COMPLEX-MLANA/CEACAM6/TPBG	0.952381	1	0.909091
COMPLEX-MSLN/MLANA/TPBG	0.916667	0.846154	1
COMPLEX-MLANA/TPBG/CLDN1	0.916667	0.846154	1
COMPLEX-MLANA/FLOT2/TPBG	0.916667	0.846154	1
COMPLEX-ERBB2/MLANA/TPBG	0.916667	0.846154	1
COMPLEX-DPEP1/MLANA/TPBG	0.916667	0.846154	1
COMPLEX-CSPG4/MLANA/TPBG	0.916667	0.846154	1
COMPLEX-CR2/MLANA/TPBG	0.916667	0.846154	1
COMPLEX-MLANA/TPBG/CLDN8	0.916667	0.846154	1
COMPLEX-MLANA/SLC39A6/TPBG	0.916667	0.846154	1
COMPLEX-MLANA/TPBG/ULBP2	0.916667	0.846154	1
COMPLEX-MLANA/TPBG/TYR	0.916667	0.846154	1
COMPLEX-MLANA/SSTR3/TPBG	0.916667	0.846154	1
COMPLEX-SLC34A2/MLANA/TPBG	0.916667	0.846154	1
COMPLEX-MLANA/MUC4/TPBG	0.916667	0.846154	1
COMPLEX-MLANA/FOLR2/TPBG	0.916667	0.846154	1
COMPLEX-MLANA/IGF1R/TPBG	0.916667	0.846154	1
COMPLEX-MLANA/ITGAV/TPBG	0.916667	0.846154	1
COMPLEX-MLANA/MAGEA4/TPBG	0.916667	0.846154	1
COMPLEX-MLANA/TPBG/MUC16	0.916667	0.846154	1
COMPLEX-MLANA/TPBG/CLDN9	0.916667	0.846154	1
COMPLEX-MLANA/TPBG/CLDN6	0.916667	0.846154	1
COMPLEX-MLANA/TPBG/CLDN12	0.916667	0.846154	1
COMPLEX-MLANA/TPBG/PROM1	0.916667	0.846154	1
COMPLEX-MLANA/TPBG/LGR5	0.916667	0.846154	1
COMPLEX-MLANA/TPBG/CD276	0.916667	0.846154	1
COMPLEX-MLANA/TPBG/ULBP1	0.916667	0.846154	1
COMPLEX-MLANA/TPBG/PSCA	0.916667	0.846154	1
COMPLEX-MLANA/TPBG/VTCN1	0.916667	0.846154	1
COMPLEX-MLANA/TPBG/ULBP3	0.916667	0.846154	1
COMPLEX-MLANA/SSTR4/TPBG	0.916667	0.846154	1
COMPLEX-MLANA/SSTR1/TPBG	0.916667	0.846154	1
COMPLEX-MLANA/SDC1/TPBG	0.916667	0.846154	1
COMPLEX-MLANA/OAS1/TPBG	0.916667	0.846154	1
COMPLEX-MLANA/EPCAM/TPBG	0.916667	0.846154	1
COMPLEX-MLANA/SST/TPBG	0.916667	0.846154	1
COMPLEX-MLANA/SSTR5/TPBG	0.916667	0.846154	1
COMPLEX-MLANA/MUC13/TPBG	0.916667	0.846154	1
COMPLEX-MLANA/TPBG/CLDN2	0.916667	0.846154	1
COMPLEX-MLANA/TPBG/TNFSF11	0.916667	0.846154	1
COMPLEX-MLANA/TPBG/SLC7A5	0.916667	0.846154	1
COMPLEX-MLANA/IL11RA/TPBG	0.916667	0.846154	1
COMPLEX-MLANA/SSTR2/TPBG	0.916667	0.846154	1
COMPLEX-MLANA/SLAMF7/TPBG	0.916667	0.846154	1
COMPLEX-MLANA/MUC1/TPBG	0.916667	0.846154	1
COMPLEX-SPON2/MLANA/TPBG	0.916667	0.846154	1
COMPLEX-MLANA/MST1R/TPBG	0.916667	0.846154	1
COMPLEX-MLANA/TNC/TPBG	0.916667	0.846154	1
COMPLEX-ALDH1A1/MLANA/TPBG	0.916667	0.846154	1
COMPLEX-EPHA3/MLANA/TPBG	0.916667	0.846154	1
COMPLEX-EGFR/MLANA/TPBG	0.916667	0.846154	1
COMPLEX-DDX3X/MLANA/TPBG	0.916667	0.846154	1
COMPLEX-CBX3/MLANA/TPBG	0.916667	0.846154	1
COMPLEX-MLANA/TPBG/CD22	0.916667	0.846154	1
COMPLEX-MLANA/TPBG/TNFRSF8	0.916667	0.846154	1
COMPLEX-TNFRSF13C/MLANA/TPBG	0.916667	0.846154	1
COMPLEX-CLDN3/MLANA/TPBG	0.916667	0.846154	1
COMPLEX-MLANA/IL3RA/TPBG	0.916667	0.846154	1
COMPLEX-MLANA/ANXA1/TPBG	0.916667	0.846154	1
COMPLEX-MLANA/GAGE1/TPBG	0.916667	0.846154	1
COMPLEX-ERBB4/MLANA/TPBG	0.916667	0.846154	1
COMPLEX-ENG/MLANA/TPBG	0.916667	0.846154	1
COMPLEX-CLDN23/MLANA/TPBG	0.916667	0.846154	1
COMPLEX-DNAJB8/MLANA/TPBG	0.916667	0.846154	1
COMPLEX-AFP/MLANA/TPBG	0.869565	0.833333	0.909091
COMPLEX-MLANA/SEMA5B/TPBG	0.869565	0.833333	0.909091
COMPLEX-MLANA/CLDN5/TPBG	0.869565	0.833333	0.909091
COMPLEX-MLANA/PCYT1A/TPBG	0.869565	0.833333	0.909091
COMPLEX-MLANA/ENPP3/TPBG	0.869565	0.833333	0.909091
COMPLEX-MLANA/TPBG/WT1	0.869565	0.833333	0.909091
COMPLEX-MLANA/TRPM4/TPBG	0.869565	0.833333	0.909091
COMPLEX-MLANA/GUCY2C/TPBG	0.869565	0.833333	0.909091
COMPLEX-RAET1E/MLANA/TPBG	0.869565	0.833333	0.909091
COMPLEX-GPA33/MLANA/TPBG	0.869565	0.833333	0.909091
COMPLEX-MLANA/TPBG/CD70	0.869565	0.833333	0.909091
COMPLEX-MLANA/STEAP2/TPBG	0.869565	0.833333	0.909091

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
IFI6 AND EPHB2 AND NOT-MAG	1	1	1
DPEP1 AND NOX1 AND NOT-HTR6	1	1	1
DPEP1 AND NOX1 AND NOT-VN1R2	1	1	1
DPEP1 AND NOX1 AND NOT-GRM4	1	1	1
IFI6 AND IL20RA AND NOT-SLCO6A1	1	1	1
CDH17 AND TGFBI AND NOT-PCYT1A	1	1	1
IFI6 AND EPHB2 AND NOT-MIP	1	1	1
IFI6 AND GUCY2C AND NOT-SLCO6A1	1	1	1
IFI6 AND EPHB2 AND NOT-NMBR	1	1	1
DPEP1 AND NOX1 AND NOT-GABRA5	1	1	1
IFI6 AND EPHB2 AND NOT-SLCO6A1	1	1	1
DPEP1 AND IFI6 AND NOT-SLCO6A1	1	1	1
DPEP1 AND NOT-HTR3C AND NOT-SHISA9	1	1	1
DPEP1 AND NOX1 AND NOT-RXFP3	1	1	1
IFI6 AND EPHB2 AND NOT-KCNH1	1	1	1
IFI6 AND EPHB2 AND NOT-GABRD	1	1	1
CDH17 AND TGFBI AND NOT-ENPP3	1	1	1
IFI6 AND EPHB2 AND NOT-GHRHR	1	1	1
DPEP1 AND NOT-SLC1A6 AND FXYD6	1	1	1
IFI6 AND EPHB2 AND NOT-GPR78	1	1	1
CDH17 AND TGFBI AND CLDN1	1	1	1
IFI6 AND EPHB2 AND NOT-GNRHR	1	1	1
IFI6 AND EPHB2 AND NOT-GRIK1	1	1	1
DPEP1 AND NOX1 AND NOT-CCKBR	1	1	1
IFI6 AND EPHB2 AND NOT-GRIK4	1	1	1
DPEP1 AND NOT-SLC10A2 AND NOX1	1	1	1
DPEP1 AND NOX1 AND NOT-SLC9A1	1	1	1
IFI6 AND EPHB2 AND NOT-GRIN2C	1	1	1
CDH17 AND TGFBI AND THY1	1	1	1
IFI6 AND EPHB2 AND NOT-KCNA10	1	1	1
DPEP1 AND NOX1 AND NOT-PTGER4	1	1	1
DPEP1 AND NOX1 AND NOT-AJAP1	1	1	1
NOX1 AND THY1 AND NOT-MRGPRX2	1	1	1
NOX1 AND THY1 AND NOT-CALHM3	1	1	1
DPEP1 AND NOT-SLC28A1 AND NKAIN2	1	1	1
NOX1 AND THY1 AND NOT-PORCN	1	1	1
NOX1 AND THY1 AND NOT-ABCG4	1	1	1
NOX1 AND THY1 AND NOT-SCN1A	1	1	1
NOX1 AND THY1 AND NOT-CACNG8	1	1	1
DPEP1 AND NOT-SLC28A1 AND NOT-ATP4B	1	1	1
NOX1 AND THY1 AND NOT-JPH3	1	1	1
NOX1 AND THY1 AND NOT-ATP8B2	1	1	1
DPEP1 AND NOT-SLC13A1 AND KCNK12	1	1	1
NOX1 AND THY1 AND NOT-SLC12A9	1	1	1
NOX1 AND THY1 AND NOT-MRAP	1	1	1
NOX1 AND THY1 AND NOT-PCDHAC2	1	1	1
NOX1 AND THY1 AND NOT-ENPP1	1	1	1
NOX1 AND THY1 AND NOT-KCNK9	1	1	1
NOX1 AND THY1 AND NOT-KCNK4	1	1	1
NOX1 AND THY1 AND NOT-OXTR	1	1	1
NOX1 AND THY1 AND NOT-OPRK1	1	1	1
NOX1 AND THY1 AND NOT-GPR135	1	1	1
NOX1 AND THY1 AND NOT-SLC6A7	1	1	1
NOX1 AND THY1 AND NOT-ASTN1	1	1	1
NOX1 AND THY1 AND NOT-SLC6A13	1	1	1
NOX1 AND THY1 AND NOT-OTOF	1	1	1
NOX1 AND THY1 AND NOT-CD1A	1	1	1
DPEP1 AND NOT-ATP4B AND NOT-ABCG5	1	1	1
DPEP1 AND NOT-SLC13A1 AND CD72	1	1	1
IFI6 AND CLDN3 AND NOT-FLOT2	1	1	1
DPEP1 AND NOT-SLC1A6 AND STEAP2	1	1	1
DPEP1 AND NOT-SLC6A11 AND STEAP2	1	1	1
DPEP1 AND NOT-SLC13A1 AND STEAP2	1	1	1
DPEP1 AND NOT-SPAM1 AND STEAP2	1	1	1
DPEP1 AND STEAP2 AND NOT-DGKE	1	1	1
DPEP1 AND NOT-CNTNAP4 AND STEAP2	1	1	1
DPEP1 AND STEAP2 AND NOT-SLC22A6	1	1	1
DPEP1 AND ITGAV AND NOT-KCNK12	1	1	1
DPEP1 AND ITGAV AND NOT-CACNG2	1	1	1
DPEP1 AND NOT-CACNG2 AND TNC	1	1	1
DPEP1 AND NOT-KCNH4 AND STEAP2	1	1	1
DPEP1 AND NOT-KCNH4 AND ANXA1	1	1	1
DPEP1 AND NOT-KCNH4 AND PTK7	1	1	1
CLDN7 AND SLC7A5 AND NOT-TMPRSS11B	1	1	1
DPEP1 AND STEAP2 AND NOT-MPL	1	1	1
CLDN7 AND SLC7A5 AND NOT-PORCN	1	1	1
DPEP1 AND NOT-CHRND AND MST1R	1	1	1

Antigen Pair Logic	F1	Precision	Recall
COMPLEX-MLANA/TPBG/CD34	0.869565	0.833333	0.909091
COMPLEX-MLANA/BIRC5/TPBG	0.869565	0.833333	0.909091
COMPLEX-MLANA/L1CAM/TPBG	0.869565	0.833333	0.909091
COMPLEX-MLANA/CLDN18/TPBG	0.818182	0.818182	0.818182
COMPLEX-MLANA/CLDN11/TPBG	0.818182	0.818182	0.818182
COMPLEX-MLANA/GPC3/TPBG	0.818182	0.818182	0.818182
COMPLEX-MLANA/AXL/TPBG	0.818182	0.818182	0.818182
COMPLEX-MLANA/B4GALNT1/TPBG	0.916667	0.846154	1
COMPLEX-MLANA/PMEL/TPBG	0.869565	0.833333	0.909091
COMPLEX-MLANA/KDR/TPBG	0.818182	0.818182	0.818182
NOT-SLC34A2 AND GPNMB AND ERBB2	0.952381	1	0.909091
NOT-VTCN1 AND GPNMB AND ERBB2	0.842105	1	0.727273
EDNRB AND GPNMB AND ERBB2	0.842105	1	0.727273
NOT-TNFRSF13C AND GPNMB AND NOT-ERBB2	0.9	1	0.818182
NOT-AFP AND GPNMB AND NOT-ERBB2	0.952381	1	0.909091
CLDN23 AND GPNMB AND NOT-ERBB2	0.842105	1	0.727273
NOT-CLDN6 AND GPNMB AND ERBB2	0.952381	1	0.909091
GPNMB AND NOT-MUC4 AND ERBB2	0.952381	1	0.909091
GPNMB AND FLOT2 AND NOT-ERBB2	0.9	1	0.818182
TNC AND NOT-ERBB2 AND GPNMB	0.842105	1	0.727273
GPNMB AND OAS1 AND ERBB2	0.842105	1	0.727273
GPNMB AND NOT-HHLA2 AND ERBB2	0.9	1	0.818182
GPNMB AND NOT-SST AND ERBB2	0.909091	0.909091	0.909091
ENG AND GPNMB AND NOT-ERBB2	0.857143	0.9	0.818182
Neuroblastoma			
SLC10A4 AND TMEM97 AND NOT-AGPAT3	0.994012	1	0.988095
SLC10A4 AND NOT-NTM AND NOT-HTR1B	0.994012	1	0.988095
COMPLEX-SLC10A4/SLC6A8/SLC15A2	0.994012	1	0.988095
COMPLEX-SLC10A4/SLC6A1/SLC15A2	0.994012	1	0.988095
SLC10A4 AND NOT-CD99L2 AND NOT-SLC5A1	0.994012	1	0.988095
SLC10A4 AND NOT-PAQR6 AND NOT-MCOLN2	0.994012	1	0.988095
COMPLEX-SLC10A4/CLCA4/SLC7A10	0.994012	1	0.988095
COMPLEX-SLC10A4/CLSTN2/SLC15A2	0.994012	1	0.988095
COMPLEX-IGSF11/SLC10A4/SLC7A10	0.994012	1	0.988095
SLC10A4 AND NOT-NTRK2 AND NOT-SLC5A1	0.994012	1	0.988095
COMPLEX-SLC10A4/JAM2/SLC15A2	0.994012	1	0.988095
COMPLEX-SLC10A4/SLC15A1/SLC15A2	0.994012	1	0.988095
SLC10A4 AND NOT-CD99L2 AND NOT-OR5I1	0.994012	1	0.988095
COMPLEX-SLC10A4/SLC15A2/VMP1	0.994012	1	0.988095
SLC10A4 AND NOT-PAQR6 AND NOT-TNFRSF14	0.994012	1	0.988095
SLC10A4 AND NOT-AQP4 AND NOT-CRHR2	0.994012	1	0.988095
SLC10A4 AND NOT-APLP1 AND NOT-PCDHAC1	0.994012	1	0.988095
COMPLEX-SLC10A4/PRRG1/SLC15A2	0.994012	1	0.988095
COMPLEX-SLC10A4/SLC15A2/SYP	0.994012	1	0.988095
COMPLEX-SLC10A4/PTPRZ1/SLC15A2	0.994012	1	0.988095
SLC10A4 AND NOT-AQP4 AND NOT-SLC5A1	0.994012	1	0.988095
SLC10A4 AND NOT-PAQR6 AND NOT-IL17RC	0.994012	1	0.988095
COMPLEX-SLC10A4/SLC22A23/SLC15A2	0.994012	1	0.988095
SLC10A4 AND NOT-SLC22A23 AND NOT-MRGPRX3	0.994012	1	0.988095
SLC10A4 AND NOT-PAQR6 AND NOT-MRGPRX3	0.994012	1	0.988095
COMPLEX-MRGPRX3/SLC10A4/ABCG2	0.994012	1	0.988095
COMPLEX-IGSF11/SLC10A4/TRPC6	0.994012	1	0.988095
SLC10A4 AND NOT-PAQR6 AND NOT-AGPAT3	0.994012	1	0.988095
COMPLEX-SLC10A4/TMEM100/SLC15A2	0.994012	1	0.988095
SLC10A4 AND NOT-SLC22A23 AND NOT-OR10H3	0.994012	1	0.988095
COMPLEX-SLC10A4/PLP1/SLC15A2	0.994012	1	0.988095
COMPLEX-SLC10A4/ATP1A2/SLC15A2	0.994012	1	0.988095
COMPLEX-SLC10A4/ATP1A2/SLC15A2	0.994012	1	0.988095
SLC10A4 AND NOT-SLC22A23 AND NOT-SLC5A1	0.994012	1	0.988095
SLC10A4 AND NOT-GPR62 AND NOT-ABCA4	0.994012	1	0.988095
COMPLEX-IGSF11/SLC10A4/NTSR1	0.994012	1	0.988095
COMPLEX-SLC10A4/NTSR1/STYK1	0.994012	1	0.988095
COMPLEX-SLC10A4/NTSR1/SCN2B	0.994012	1	0.988095
COMPLEX-SLC10A4/NTSR1/ABCG2	0.994012	1	0.988095
COMPLEX-CHRNB3/SLC10A4/SLC7A10	0.994012	1	0.988095
COMPLEX-MRGPRX3/IGSF11/SLC10A4	0.994012	1	0.988095
SLC10A4 AND NOT-GJC2 AND NOT-PHEX	0.994012	1	0.988095
COMPLEX-SLC10A4/PRRG1/SLC15A2	0.994012	1	0.988095
COMPLEX-SLC10A4/SLC15A2/ATRN	0.994012	1	0.988095
COMPLEX-CHRNB3/SLC10A4/CADM2	0.994012	1	0.988095
COMPLEX-SLC10A4/SHISA6/STYK1	0.994012	1	0.988095
COMPLEX-SLC10A4/SHISA6/SCN2B	0.994012	1	0.988095
COMPLEX-CX3CR1/SLC2A12/SLC10A4	0.994012	1	0.988095
SLC10A4 AND NOT-PAQR6 AND NOT-USP48	0.994012	1	0.988095
SLC10A4 AND NOT-PAQR6 AND NOT-SLC16A2	0.994012	1	0.988095
COMPLEX-GPR62/SLC10A4/STX3	0.994012	1	0.988095
COMPLEX-SLC2A12/SLC10A4/SLC7A8	0.994012	1	0.988095

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
DPEP1 AND ITGAV AND NOT-CHRNA4	1	1	1
DPEP1 AND GUCY2C AND NOT-SLC9A1	1	1	1
DPEP1 AND NOT-SLC10A2 AND GUCY2C	1	1	1
DPEP1 AND NOT-CD1B AND GUCY2C	1	1	1
DPEP1 AND NOT-ANPEP AND GPA33	1	1	1
BIRC5 AND GPA33 AND NOT-SLC9A1	1	1	1
IFI6 AND CEACAM6 AND LGR5	1	1	1
DPEP1 AND NOT-TMPRSS11B AND CLDN1	1	1	1
DPEP1 AND NOT-CLDN19 AND CLDN1	1	1	1
DPEP1 AND NOT-SV2C AND CLDN1	1	1	1
DPEP1 AND NOT-SLC12A5 AND STEAP2	1	1	1
DPEP1 AND NOT-OR52D1 AND STEAP2	1	1	1
CLDN7 AND THY1 AND NOT-SLCO6A1	1	1	1
IFI6 AND CLDN3 AND NOT-DDX3X	1	1	1
DPEP1 AND SLC7A5 AND NOT-KCNU1	1	1	1
DPEP1 AND NOT-PCDHB1 AND CD70	1	1	1
IFI6 AND CEACAM5 AND NOT-SSTR4	1	1	1
DPEP1 AND NOT-PCDHB1 AND PTK7	1	1	1
DPEP1 AND NOT-PCDHB1 AND IL20RA	1	1	1
DPEP1 AND NOT-PCDHB1 AND ANXA1	1	1	1
DPEP1 AND NOT-PCDHB1 AND STEAP2	1	1	1
DPEP1 AND NOT-GABBR2 AND CD276	1	1	1
BIRC5 AND NOX1 AND ITGAV	1	1	1
DPEP1 AND NOT-GABBR2 AND THY1	1	1	1
GUCY2C AND THY1 AND NOT-CHRND	1	1	1
DPEP1 AND NOT-SLC28A1 AND MAGEA4	1	1	1
IFI6 AND MUC13 AND NOT-SDC1	1	1	1
DPEP1 AND NOT-CD1B AND NOT-CD79B	1	1	1
MUC13 AND THY1 AND NOT-MUC12	1	1	1
IFI6 AND CEACAM5 AND SLC7A5	1	1	1
DPEP1 AND NOT-SLC9B1 AND STEAP2	1	1	1
DPEP1 AND STEAP2 AND NOT-NKAIN2	1	1	1
DPEP1 AND NOT-KCNU1 AND STEAP2	1	1	1
DPEP1 AND NOT-EPHA8 AND STEAP2	1	1	1
DPEP1 AND STEAP2 AND NOT-CRB1	1	1	1
DPEP1 AND STEAP2 AND NOT-GABRA1	1	1	1
DPEP1 AND STEAP2 AND NOT-GABRG3	1	1	1
DPEP1 AND STEAP2 AND NOT-GRIN2B	1	1	1
DPEP1 AND STEAP2 AND NOT-GUCY2F	1	1	1
DPEP1 AND NOT-SST AND NOT-CACNG2	1	1	1
DPEP1 AND NOT-HTR1D AND STEAP2	1	1	1
DPEP1 AND NOT-HTR2C AND STEAP2	1	1	1
DPEP1 AND NOT-GABRB1 AND CLDN1	1	1	1
DPEP1 AND NOT-GPR22 AND CLDN1	1	1	1
IFI6 AND ITGB6 AND NOT-EDNRB	1	1	1
DPEP1 AND NOT-TSHR AND STEAP2	1	1	1
IFI6 AND CLDN7 AND NOT-FCRL1	1	1	1
IFI6 AND MUC13 AND NOT-FCRL1	1	1	1
IFI6 AND BIRC5 AND NOT-CLDN9	1	1	1
IFI6 AND BIRC5 AND NOT-SLC7A5	1	1	1
IFI6 AND CEACAM5 AND NOT-CD79B	1	1	1
IFI6 AND MUC13 AND NOT-BIRC5	1	1	1
IFI6 AND BIRC5 AND NOT-CD180	1	1	1
IFI6 AND BIRC5 AND NOT-ITGB3	1	1	1
IFI6 AND BIRC5 AND NOT-IGF1R	1	1	1
IFI6 AND BIRC5 AND NOT-GAGE1	1	1	1
IFI6 AND STEAP1 AND EPCAM	1	1	1
IFI6 AND BIRC5 AND NOT-DKK1	1	1	1
DPEP1 AND SLC7A5 AND NOT-MIP	1	1	1
DPEP1 AND NOT-HTR1F AND CLDN1	1	1	1
IFI6 AND MUC13 AND NOT-ITGB6	1	1	1
IFI6 AND BIRC5 AND NOT-EDNRB	1	1	1
IFI6 AND BIRC5 AND NOT-DDX3X	1	1	1
DPEP1 AND ITGAV AND NOT-SLC22A16	1	1	1
IFI6 AND BIRC5 AND NOT-TNFRSF13C	1	1	1
IFI6 AND CLDN3 AND ITGAV	1	1	1
DPEP1 AND NOT-SV2C AND PTK7	1	1	1
DPEP1 AND NOT-SV2C AND IL20RA	1	1	1
DPEP1 AND NOT-SV2C AND ANXA1	1	1	1
DPEP1 AND NOT-SLC28A1 AND ITGAV	1	1	1
DPEP1 AND ITGAV AND NOT-ADAM7	1	1	1
DPEP1 AND ITGAV AND NOT-TAS1R1	1	1	1
DPEP1 AND NOT-SLC28A1 AND GPA33	1	1	1
DPEP1 AND NOT-TSHR AND ANXA1	1	1	1
DPEP1 AND NOT-TSHR AND IL20RA	1	1	1
DPEP1 AND NOT-TSHR AND PTK7	1	1	1
DPEP1 AND NOT-OPRK1 AND CLDN1	1	1	1

Antigen Pair Logic	F1	Precision	Recall
COMPLEX-SLC10A4/KCNJ10/LGR4	0.994012	1	0.988095
COMPLEX-SLC10A4/KCNJ10/CEACAM1	0.994012	1	0.988095
COMPLEX-SLC10A4/KCNJ10/TRPV4	0.994012	1	0.988095
SLC10A4 AND NOT-GPR62 AND NOT-HTR6	0.994012	1	0.988095
COMPLEX-SLC10A4/TMEM235/ABCG2	0.994012	1	0.988095
COMPLEX-SLC10A4/KCNJ10/VIPR1	0.994012	1	0.988095
SLC10A4 AND NOT-ACSL6 AND NOT-NPY5R	0.994012	1	0.988095
SLC10A4 AND NOT-NKAIN2 AND NOT-USP48	0.994012	1	0.988095
SLC10A4 AND NOT-ACSL6 AND NOT-DCHS2	0.994012	1	0.988095
COMPLEX-SLC32A1/SLC10A4/SLC15A2	0.994012	1	0.988095
SLC10A4 AND NOT-NKAIN2 AND NOT-LY6D	0.994012	1	0.988095
SLC10A4 AND NOT-NKAIN2 AND NOT-TNFRSF14	0.994012	1	0.988095
COMPLEX-SLC10A4/GABRA2/SLC15A2	0.994012	1	0.988095
SLC10A4 AND NOT-KCNJ10 AND NOT-SLC5A1	0.994012	1	0.988095
COMPLEX-SLC10A4/GABRB1/SLC15A2	0.994012	1	0.988095
COMPLEX-SLC10A4/GABRG1/SLC15A2	0.994012	1	0.988095
SLC10A4 AND NOT-SYPL1 AND NOT-KCNJ10	0.994012	1	0.988095
COMPLEX-SLC10A4/KCNJ10/AMN	0.994012	1	0.988095
SLC10A4 AND NOT-KCNJ10 AND NOT-FAM73B	0.994012	1	0.988095
SLC10A4 AND NOT-KCNJ10 AND NOT-ABCC3	0.994012	1	0.988095
COMPLEX-IGSF11/SLC10A4/GLRA3	0.994012	1	0.988095
SLC10A4 AND NOT-NKAIN2 AND NOT-ABCG1	0.994012	1	0.988095
SLC10A4 AND TUSC3 AND NOT-GLRA3	0.994012	1	0.988095
SLC10A4 AND NOT-CDH26 AND NOT-HTR1F	0.994012	1	0.988095
SLC10A4 AND NOT-KCNJ10 AND NOT-P2RY2	0.994012	1	0.988095
SLC10A4 AND NOT-KCNJ10 AND NOT-HFE	0.994012	1	0.988095
SLC10A4 AND NOT-NKAIN2 AND NOT-SLC52A1	0.994012	1	0.988095
SLC10A4 AND NOT-NTRK2 AND SLC38A1	0.994012	1	0.988095
SLC10A4 AND NOT-CD99L2 AND NOT-ADAM7	0.994012	1	0.988095
SLC10A4 AND NOT-KCNJ10 AND NOT-APCDD1	0.994012	1	0.988095
COMPLEX-NKAIN2/SLC10A4/TAS2R38	0.994012	1	0.988095
SLC10A4 AND NOT-NKAIN2 AND NOT-SLC46A2	0.994012	1	0.988095
COMPLEX-SLC10A4/FGFR2/KCNJ10	0.994012	1	0.988095
COMPLEX-SLC10A4/SLC7A10/SLC8A3	0.994012	1	0.988095
SLC10A4 AND NOT-CD99L2 AND NOT-HTR1F	0.994012	1	0.988095
SLC10A4 AND NOT-KCNJ10 AND NOT-OR10H3	0.994012	1	0.988095
SLC10A4 AND NOT-CD99L2 AND NOT-GRIN2B	0.994012	1	0.988095
SLC10A4 AND NOT-KCNJ10 AND NOT-MR1	0.994012	1	0.988095
COMPLEX-SLC10A4/SLC7A10/CALN1	0.994012	1	0.988095
SLC10A4 AND NOT-NKAIN2 AND NOT-RXFP1	0.994012	1	0.988095
SLC10A4 AND NOT-CD99L2 AND NOT-OR5P3	0.994012	1	0.988095
SLC10A4 AND NOT-KCNJ10 AND NOT-ITPR2	0.994012	1	0.988095
COMPLEX-SLC10A4/KCNJ10/LIM2	0.994012	1	0.988095
SLC10A4 AND NOT-KCNJ10 AND NOT-BCAM	0.994012	1	0.988095
SLC10A4 AND NOT-NKAIN2 AND NOT-SCNN1A	0.994012	1	0.988095
SLC10A4 AND NOT-KCNJ10 AND NOT-MGST2	0.994012	1	0.988095
COMPLEX-CACNG3/SLC10A4/SLC15A2	0.994012	1	0.988095
SLC10A4 AND NOT-NKAIN2 AND NOT-BRS3	0.994012	1	0.988095
COMPLEX-SLC10A4/OPCML/SLC15A2	0.994012	1	0.988095
SLC10A4 AND NOT-GRM1 AND NOT-OPN5	0.994012	1	0.988095
COMPLEX-IGSF11/SLC10A4/SLC2A1	0.994012	1	0.988095
COMPLEX-SLC10A4/GPR26/SLC15A2	0.994012	1	0.988095
COMPLEX-IGSF11/SLC10A4/GABRA1	0.994012	1	0.988095
SLC10A4 AND NOT-GPR62 AND NOT-SLC12A3	0.994012	1	0.988095
COMPLEX-MDGA2/SLC10A4/SLC15A2	0.994012	1	0.988095
COMPLEX-SLC10A4/HLA-DRB1/ABCG2	0.994012	1	0.988095
SLC10A4 AND NOT-GPR62 AND NOT-LRRC8E	0.994012	1	0.988095
SLC10A4 AND NOT-AQP4 AND NOT-ATP1B4	0.994012	1	0.988095
COMPLEX-SLC2A12/SLC10A4/GABRB1	0.994012	1	0.988095
COMPLEX-SLC10A4/SCN8A/ABCG2	0.994012	1	0.988095
COMPLEX-SLC10A4/SLC8A3/SLC15A2	0.994012	1	0.988095
COMPLEX-SLC2A12/SLC10A4/GPR26	0.994012	1	0.988095
SLC10A4 AND NOT-SSTR1 AND NOT-LHCGR	0.994012	1	0.988095
COMPLEX-BVES/SLC10A4/CLDN11	0.994012	1	0.988095
COMPLEX-SLC10A4/AQP4/ITGB3	0.994012	1	0.988095
SLC10A4 AND NOT-GPR62 AND NOT-EGFR	0.994012	1	0.988095
SLC10A4 AND NOT-CD99L2 AND NOT-ITGB3	0.994012	1	0.988095
COMPLEX-SLC10A4/FOLH1/SLC15A2	0.994012	1	0.988095
SLC10A4 AND NOT-SSTR1 AND NOT-RRH	0.994012	1	0.988095
SLC10A4 AND NOT-AGPAT3 AND NOT-ABCB5	0.994012	1	0.988095
COMPLEX-SLC27A5/SLC10A4/CLDN11	0.994012	1	0.988095
SLC10A4 AND NOT-SSTR1 AND NOT-OPN5	0.994012	1	0.988095
COMPLEX-SLC10A4/CLDN11/PCDH10	0.994012	1	0.988095
COMPLEX-SLC10A4/ALDH1A1/PAQR6	0.994012	1	0.988095
COMPLEX-CLDN4/SLC10A4/AQP4	0.994012	1	0.988095
SLC10A4 AND NOT-KCNJ16 AND NOT-EGFR	0.994012	1	0.988095
SLC10A4 AND NOT-AQP4 AND NOT-EGFR	0.994012	1	0.988095

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
DPEP1 AND NOT-ENPP1 AND CLDN1	1	1	1
DPEP1 AND NOT-CD1B AND NOT-GPC3	1	1	1
IFI6 AND CLDN3 AND NOT-GPC3	1	1	1
DPEP1 AND NOT-GPC3 AND NOT-CCR9	1	1	1
DPEP1 AND NOT-GPC3 AND NOT-SLC10A2	1	1	1
NOX1 AND THY1 AND NOT-GPC3	1	1	1
NOX1 AND THY1 AND NOT-ERBB2	1	1	1
IFI6 AND CLDN4 AND NOT-GPC3	1	1	1
IFI6 AND EPHB2 AND NOT-GPC3	1	1	1
IFI6 AND NOT-GPC3 AND ITGB6	1	1	1
DPEP1 AND NOX1 AND NOT-MUC1	1	1	1
IFI6 AND MUC13 AND NOT-MUC1	1	1	1
IFI6 AND CLDN7 AND NOT-MUC1	1	1	1
IFI6 AND MUC13 AND NOT-GPC3	1	1	1
IFI6 AND NOT-GPC3 AND BIRC5	1	1	1
IFI6 AND CLDN3 AND NOT-MUC1	1	1	1
DPEP1 AND NOT-GPC3 AND NOT-FLVCR1	1	1	1
CLDN7 AND TGFBI AND NOT-MUC1	1	1	1
IFI6 AND STEAP1 AND NOT-GPC3	1	1	1
IFI6 AND CLDN4 AND NOT-GPC3	1	1	1
DPEP1 AND NOT-GPC3 AND CEACAM7	1	1	1
DPEP1 AND NOT-GPC3 AND NOT-SLC31A1	1	1	1
DPEP1 AND NOT-GPC3 AND ALCAM	1	1	1
IFI6 AND NOT-GPC3 AND EPCAM	1	1	1
DPEP1 AND NOT-CLDN20 AND NOT-GPC3	1	1	1
IFI6 AND NOT-GPC3 AND BIRC5	1	1	1
DPEP1 AND NOT-SLC13A1 AND NOT-GPC3	1	1	1
DPEP1 AND NOT-SLC13A1 AND ERBB2	1	1	1
CDH17 AND CLDN1 AND NOT-MSLN	1	1	1
DPEP1 AND NOT-ENPP1 AND MUC1	1	1	1
DPEP1 AND NOT-GPC3 AND CEACAM7	1	1	1
DPEP1 AND IFI6 AND NOT-ERBB2	1	1	1
IFI6 AND MUC13 AND NOT-MSLN	1	1	1
DPEP1 AND NOT-SLC28A1 AND NOT-GPC3	1	1	1
DPEP1 AND NOT-GPC3 AND NOT-SLC16A5	1	1	1
DPEP1 AND NOT-GPC3 AND CD44	1	1	1
DPEP1 AND NOT-GPC3 AND NOT-SLC2A2	1	1	1
DPEP1 AND NOT-GPC3 AND NOT-ABCG5	1	1	1
DPEP1 AND NOT-ANPEP AND NOT-GPC3	1	1	1
DPEP1 AND NOT-GPC3 AND NOX1	1	1	1
DPEP1 AND NOT-GPC3 AND CD93	1	1	1
DPEP1 AND NOT-GPC3 AND NOT-MUC17	1	1	1
DPEP1 AND NOT-EPHA8 AND NOT-GPC3	1	1	1
DPEP1 AND NOT-LCT AND NOT-GPC3	1	1	1
DPEP1 AND NOT-GPC3 AND PTGIS	1	1	1
DPEP1 AND NOT-GPC3 AND NOT-GRIN2C	1	1	1
IFI6 AND MUC13 AND NOT-ERBB2	1	1	1
DPEP1 AND NOT-GPC3 AND NOT-CD1A	1	1	1
DPEP1 AND NOT-GPC3 AND NOT-SLC13A2	1	1	1
DPEP1 AND NOT-GPC3 AND FXYD6	1	1	1
DPEP1 AND NOT-GPC3 AND NOT-FLVCR1	1	1	1
DPEP1 AND NOT-GPC3 AND NOT-CLDN15	1	1	1
DPEP1 AND NOT-CHRND AND NOT-GPC3	1	1	1
CDH17 AND SLC7A5 AND NOT-GPC3	1	1	1
CDH17 AND SLC7A5 AND NOT-ERBB2	1	1	1
IFI6 AND STEAP1 AND NOT-GPC3	1	1	1
NOX1 AND THY1 AND NOT-MUC1	1	1	1
NOX1 AND SLC7A5 AND NOT-GPC3	1	1	1
NOX1 AND SLC7A5 AND NOT-ERBB2	1	1	1
IFI6 AND IL20RA AND NOT-MUC1	1	1	1
DPEP1 AND NOT-GPC3 AND NOT-ABCG5	1	1	1
DPEP1 AND NOT-ABCG5 AND ERBB2	1	1	1
IFI6 AND CLDN3 AND NOT-MUC1	1	1	1
IFI6 AND NOT-GPC3 AND EPCAM	1	1	1
IFI6 AND MUC13 AND NOT-GPC3	1	1	1
NOX1 AND CLDN1 AND NOT-GPC3	1	1	1
IFI6 AND GPA33 AND NOT-ERBB2	1	1	1
DPEP1 AND NOT-GPC3 AND NOT-PTPRR	1	1	1
DPEP1 AND NOT-SLC13A1 AND NOT-GPC3	1	1	1
DPEP1 AND NOT-GPC3 AND VANGL1	1	1	1
DPEP1 AND NOT-GPC3 AND CD163	1	1	1
DPEP1 AND NOT-GPC3 AND CD44	1	1	1
CDH17 AND THY1 AND NOT-ERBB2	1	1	1
CDH17 AND THY1 AND NOT-GPC3	1	1	1
IFI6 AND GPA33 AND NOT-ERBB2	1	1	1
DPEP1 AND NOT-SPAM1 AND MUC1	1	1	1
DPEP1 AND NOT-SLC13A1 AND MUC1	1	1	1

Antigen Pair Logic	F1	Precision	Recall
SLC10A4 AND NOT-AQP4 AND NOT-EGFR	0.994012	1	0.988095
COMPLEX-SLC10A4/SLC15A2/CLDN1	0.994012	1	0.988095
COMPLEX-CELSR2/SLC10A4/CLDN11	0.994012	1	0.988095
COMPLEX-SLC10A4/NCAM1/SLC15A2	0.994012	1	0.988095
SLC10A4 AND NOT-GPR62 AND NOT-EGFR	0.994012	1	0.988095
COMPLEX-SLC10A4/CLDN11/RET	0.994012	1	0.988095
COMPLEX-GPR62/SLC10A4/ITGB3	0.994012	1	0.988095
COMPLEX-SLC10A4/GPM6B/CLDN11	0.994012	1	0.988095
COMPLEX-CHRNB3/SLC10A4/CLDN11	0.994012	1	0.988095
COMPLEX-SLC10A4/SLC7A10/SSTR1	0.994012	1	0.988095
COMPLEX-SLC10A4/CLDN11/SLC15A2	0.994012	1	0.988095
COMPLEX-SLC10A4/SLC15A2/SSTR1	0.994012	1	0.988095
COMPLEX-SLC2A12/SLC10A4/SSTR1	0.994012	1	0.988095
COMPLEX-SLC2A12/SLC10A4/TPBG	0.994012	1	0.988095
COMPLEX-GJB6/SLC10A4/CLDN1	0.994012	1	0.988095
COMPLEX-IGSF11/SLC10A4/ERBB4	0.994012	1	0.988095
COMPLEX-SLC10A4/TNFRSF21/CLDN11	0.994012	1	0.988095
COMPLEX-SLC10A4/NCAM1/SLC7A10	0.994012	1	0.988095
COMPLEX-PTPRU/SLC10A4/SSTR1	0.994012	1	0.988095
SLC10A4 AND NOT-AQP4 AND NOT-TYR	0.994012	1	0.988095
SLC10A4 AND NOT-STYK1 AND NOT-CLDN11	0.994012	1	0.988095
SLC10A4 AND NOT-AQP4 AND NOT-DNAJB8	0.994012	1	0.988095
SLC10A4 AND NOT-PAQR6 AND NOT-TYR	0.994012	1	0.988095
SLC10A4 AND NOT-PAQR6 AND NOT-DNAJB8	0.994012	1	0.988095
SLC10A4 AND NOT-GPR62 AND NOT-TYR	0.994012	1	0.988095
SLC10A4 AND NOT-GPR62 AND NOT-DNAJB8	0.994012	1	0.988095
COMPLEX-SLC2A12/SLC10A4/NCAM1	0.994012	1	0.988095
COMPLEX-SLC10A4/SLC7A10/SSTR1	0.994012	1	0.988095
COMPLEX-CELSR2/SLC10A4/CLDN11	0.994012	1	0.988095
SLC10A4 AND NOT-SSTR1 AND NOT-CLEC4D	0.994012	1	0.988095
COMPLEX-NKAIN2/SLC10A4/CLDN18	0.994012	1	0.988095
COMPLEX-SLC10A4/TMEFF2/CLDN11	0.994012	1	0.988095
COMPLEX-CACNG3/SLC10A4/CLDN11	0.994012	1	0.988095
SLC10A4 AND NOT-SSTR1 AND NOT-GNRHR	0.994012	1	0.988095
COMPLEX-NKAIN2/SLC10A4/KDR	0.994012	1	0.988095
COMPLEX-SLC10A4/CLDN11/SLC2A1	0.994012	1	0.988095
SLC10A4 AND NOT-KCNJ10 AND NOT-DNAJB8	0.994012	1	0.988095
SLC10A4 AND NOT-KCNJ10 AND NOT-TYR	0.994012	1	0.988095
SLC10A4 AND NOT-SSTR1 AND NOT-GLRA3	0.994012	1	0.988095
COMPLEX-MDGA2/SLC10A4/CLDN11	0.994012	1	0.988095
COMPLEX-SLC10A4/CLDN11/PAQR8	0.994012	1	0.988095
SLC10A4 AND NOT-KCNJ10 AND NOT-MSLN	0.994012	1	0.988095
SLC10A4 AND NOT-KCNJ10 AND NOT-EGFR	0.994012	1	0.988095
SLC10A4 AND NOT-NKAIN2 AND NOT-MSLN	0.994012	1	0.988095
SLC10A4 AND NOT-SSTR1 AND NOT-SCN10A	0.994012	1	0.988095
COMPLEX-SLC10A4/KCNJ10/CLDN18	0.994012	1	0.988095
SLC10A4 AND NOT-SSTR1 AND NOT-SLC13A1	0.994012	1	0.988095
SLC10A4 AND NOT-NKAIN2 AND NOT-EGFR	0.994012	1	0.988095
SLC10A4 AND NOT-NKAIN2 AND NOT-MSLN	0.994012	1	0.988095
COMPLEX-SLC10A4/ITGB3/KCNJ10	0.994012	1	0.988095
COMPLEX-NKAIN2/SLC10A4/IL13RA1	0.994012	1	0.988095
COMPLEX-CACNG3/SLC10A4/CLDN11	0.994012	1	0.988095
SLC10A4 AND NOT-NKAIN2 AND NOT-DNAJB8	0.994012	1	0.988095
SLC10A4 AND NOT-SSTR1 AND NOT-MRGPRX2	0.994012	1	0.988095
COMPLEX-SLC10A4/GABRB1/CLDN11	0.994012	1	0.988095
SLC10A4 AND NOT-SSTR1 AND NOT-MRGPRX2	0.994012	1	0.988095
SLC10A4 AND NOT-SSTR1 AND NOT-GLRA3	0.994012	1	0.988095
COMPLEX-SLC10A4/KCNJ10/CLDN18	0.994012	1	0.988095
SLC10A4 AND NOT-NKAIN2 AND NOT-TYR	0.994012	1	0.988095
COMPLEX-SLC10A4/GPR26/CLDN11	0.994012	1	0.988095
COMPLEX-NKAIN2/SLC10A4/IL13RA1	0.994012	1	0.988095
COMPLEX-NKAIN2/SLC10A4/KDR	0.994012	1	0.988095
COMPLEX-CSPG5/SLC10A4/CLDN11	0.994012	1	0.988095
SLC10A4 AND NOT-SSTR1 AND NOT-GNRHR	0.994012	1	0.988095
COMPLEX-MDGA2/SLC10A4/CLDN11	0.994012	1	0.988095
SLC10A4 AND NOT-FOLH1 AND NOT-GLRA3	0.994012	1	0.988095
SLC10A4 AND NOT-GABRB2 AND NOT-EGFR	0.994012	1	0.988095
COMPLEX-SLC10A4/CLDN11/PAQR8	0.994012	1	0.988095
SLC10A4 AND NOT-SSTR1 AND NOT-CACNG6	0.994012	1	0.988095
COMPLEX-CSMD3/SLC10A4/CLDN11	0.994012	1	0.988095
COMPLEX-SLC10A4/KCNJ10/KDR	0.994012	1	0.988095
SLC10A4 AND NOT-SSTR1 AND NOT-ADAM7	0.994012	1	0.988095
SLC10A4 AND NOT-GABRB2 AND NOT-EGFR	0.994012	1	0.988095
SLC10A4 AND NOT-SSTR1 AND NOT-ADAM7	0.994012	1	0.988095
SLC10A4 AND NOT-SSTR1 AND SLC38A1	0.994012	1	0.988095
SLC10A4 AND NOT-GABRB2 AND NOT-TYR	0.994012	1	0.988095
COMPLEX-SLC10A4/KCNJ10/ULBP3	0.994012	1	0.988095

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
IFI6 AND CEACAM5 AND NOT-GPC3	1	1	1
DPEP1 AND NOT-SLC28A1 AND ERBB2	1	1	1
IFI6 AND GUCY2C AND NOT-ERBB2	1	1	1
DPEP1 AND NOT-SLC28A1 AND NOT-GPC3	1	1	1
DPEP1 AND NOT-GPC3 AND NOT-SLC31A1	1	1	1
CLDN3 AND TGFBI AND NOT-MUC1	1	1	1
TGFBI AND NOT-IL11RA AND MUC1	1	1	1
DPEP1 AND NOT-GPC3 AND NOT-SLC9A1	1	1	1
IFI6 AND IL20RA AND NOT-MUC1	1	1	1
DPEP1 AND NOT-LCT AND NOT-GPC3	1	1	1
CDH17 AND THY1 AND NOT-MUC1	1	1	1
IFI6 AND MUC13 AND NOT-MUC1	1	1	1
DPEP1 AND NOT-SLC28A1 AND MSLN	1	1	1
DPEP1 AND NOT-GPC3 AND CDH11	1	1	1
DPEP1 AND NOT-CHRND AND NOT-GPC3	1	1	1
IFI6 AND MUC13 AND NOT-ERBB2	1	1	1
DPEP1 AND NOT-SLC13A1 AND MSLN	1	1	1
DPEP1 AND NOT-CHRND AND ERBB2	1	1	1
IFI6 AND GPA33 AND NOT-MSLN	1	1	1
HEPH AND SLC39A10 AND CDH1	1	1	1
HEPH AND SLC39A10 AND KCNQ4	1	1	1
HEPH AND SLC39A10 AND HTR3B	1	1	1
HEPH AND SLC39A10 AND NOT-CDHR1	1	1	1
HEPH AND SLC39A10 AND NOT-CD7	1	1	1
HEPH AND SLC39A10 AND NOT-IGSF8	1	1	1
HEPH AND SLC39A10 AND GLP2R	1	1	1
HEPH AND SLC39A10 AND NOT-FADS2	1	1	1
HEPH AND SLC39A10 AND ABCG2	1	1	1
HEPH AND SLC39A10 AND AKAP6	1	1	1
LY75 AND JPH1 AND NOT-KCNG3	1	1	1
HEPH AND SLC39A10 AND NOT-CELSR1	1	1	1
HEPH AND SLC39A10 AND NOT-ATP2C2	1	1	1
NOT-LIFR AND CDH3 AND NOT-ATP13A4	1	1	1
NOT-SLC16A7 AND MMD AND ITGA2	1	1	1
XK AND TACSTD2 AND NOT-PDE6B	1	1	1
XK AND TACSTD2 AND ATRAID	1	1	1
HEPH AND SLC39A10 AND NOT-SYT12	1	1	1
HEPH AND SLC39A10 AND NOT-ACVR1	1	1	1
HEPH AND SLC39A10 AND ITGA8	1	1	1
HEPH AND SLC39A10 AND SYT7	1	1	1
HEPH AND SLC39A10 AND NOT-DEGS1	1	1	1
HEPH AND SLC39A10 AND CAV3	1	1	1
HEPH AND SLC39A10 AND UNC5C	1	1	1
HEPH AND SLC39A10 AND SLC4A4	1	1	1
HEPH AND SLC39A10 AND NOT-TNFRSF25	1	1	1
HEPH AND SLC39A10 AND NOT-ADAM23	1	1	1
HEPH AND SLC39A10 AND TNFRSF11A	1	1	1
HEPH AND SLC39A10 AND IL18RAP	1	1	1
HEPH AND SLC39A10 AND VNN2	1	1	1
HEPH AND SLC39A10 AND CACNA1H	1	1	1
HEPH AND SLC39A10 AND SIGLEC10	1	1	1
HEPH AND SLC39A10 AND NOT-RHOT2	1	1	1
HEPH AND SLC39A10 AND NOT-TSPAN18	1	1	1
HEPH AND SLC39A10 AND TM4SF5	1	1	1
HEPH AND SLC39A10 AND NOT-DUOXA1	1	1	1
XK AND TACSTD2 AND NOT-LIFR	1	1	1
XK AND TACSTD2 AND NOT-KCNJ12	1	1	1
XK AND TACSTD2 AND NOT-SLC13A4	1	1	1
XK AND TACSTD2 AND NOT-C1orf101	1	1	1
HEPH AND TACSTD2 AND NOT-LIFR	1	1	1
NOT-SLC16A7 AND TM4SF5 AND NOT-FLVCR2	1	1	1
JPH1 AND TM4SF5 AND NOT-FLVCR2	1	1	1
ATP10B AND NOT-FLVCR2 AND NOT-SEMA6A	1	1	1
TM4SF5 AND CDH3 AND NOT-P2RX6	1	1	1
TM4SF5 AND CDH3 AND NOT-KCNQ5	1	1	1
HEPH AND SLC39A10 AND NOT-CNNM4	1	1	1
HEPH AND SLC39A10 AND PRRG1	1	1	1
TM4SF5 AND CDH3 AND NOT-KCNS2	1	1	1
ATP10B AND AMIGO2 AND NOT-SLC36A1	1	1	1
NOT-SLC16A7 AND CALCRL AND IGSF9	1	1	1
NOT-SLC16A7 AND CALCRL AND CA12	1	1	1
ATP10B AND AMIGO2 AND NOT-ACVR1C	1	1	1
JPH1 AND NOT-AQP4 AND FXYD5	1	1	1
XK AND AMIGO2 AND NOT-ADCY5	1	1	1
TM4SF5 AND NOT-LIFR AND TACSTD2	1	1	1
TSPAN8 AND NOT-LIFR AND TACSTD2	1	1	1
MEP1A AND TACSTD2 AND NOT-LIFR	1	1	1

Antigen Pair Logic	F1	Precision	Recall
COMPLEX-SEMA4D/SLC10A4/CLDN11	0.994012	1	0.988095
SLC10A4 AND NOT-KCNJ10 AND NOT-EGFR	0.994012	1	0.988095
COMPLEX-SLC10A4/TMEM235/CLDN11	0.994012	1	0.988095
COMPLEX-SEMA4D/SLC10A4/CLDN11	0.994012	1	0.988095
SLC10A4 AND NOT-SSTR1 AND NOT-SLC13A1	0.994012	1	0.988095
COMPLEX-SLC10A4/NTSR2/SSTR1	0.994012	1	0.988095
SLC10A4 AND NOT-SSTR1 AND NOT-SCN10A	0.994012	1	0.988095
SLC10A4 AND NOT-KCNJ10 AND NOT-MSLN	0.994012	1	0.988095
COMPLEX-LYPD1/SLC10A4/CLDN11	0.994012	1	0.988095
SLC10A4 AND NOT-SSTR1 AND NOT-HTR6	0.988095	0.988095	0.988095
SLC10A4 AND NOT-SSTR1 AND NOT-SEMA4B	0.988095	0.988095	0.988095
SLC10A4 AND NOT-SSTR1 AND NOT-CNGA4	0.988095	0.988095	0.988095
SLC10A4 AND NOT-SSTR1 AND NOT-STEAP4	0.988095	0.988095	0.988095
SLC10A4 AND NOT-SSTR1 AND NOT-GPR6	0.988095	0.988095	0.988095
COMPLEX-SLC10A4/TMEM235/CEACAM6	0.988095	0.988095	0.988095
SLC10A4 AND NOT-SSTR1 AND NOT-RXFP3	0.988095	0.988095	0.988095
SLC10A4 AND NOT-SSTR1 AND NOT-OR2L2	0.988095	0.988095	0.988095
SLC10A4 AND NOT-DTNA AND NOT-TYR	0.988095	0.988095	0.988095
SLC10A4 AND NOT-SSTR1 AND NOT-GRIK1	0.988095	0.988095	0.988095
SLC10A4 AND NOT-SSTR1 AND NOT-ROS1	0.988095	0.988095	0.988095
SLC10A4 AND NOT-SSTR1 AND NOT-SCARA5	0.988095	0.988095	0.988095
SLC10A4 AND NOT-TMEM235 AND NOT-EGFR	0.988095	0.988095	0.988095
COMPLEX-SLC10A4/ULBP3/OPALIN	0.988095	0.988095	0.988095
COMPLEX-SLC10A4/TMEM235/IL13RA1	0.988095	0.988095	0.988095
SLC10A4 AND NOT-SSTR1 AND NOT-SCARA5	0.988095	0.988095	0.988095
COMPLEX-SLC10A4/TMEM235/SLC7A5	0.988095	0.988095	0.988095
COMPLEX-SLC10A4/HTR5A/SSTR1	0.988095	0.988095	0.988095
SLC10A4 AND NOT-SSTR1 AND NOT-LRRC8E	0.988095	0.988095	0.988095
SLC10A4 AND NOT-KCNA1 AND NOT-EGFR	0.988095	0.988095	0.988095
COMPLEX-SLC10A4/TMEM235/EPCAM	0.988095	0.988095	0.988095
COMPLEX-SLC10A4/GABRA5/CLDN11	0.988095	0.988095	0.988095
COMPLEX-SLC10A4/GABRA5/CLDN11	0.988095	0.988095	0.988095
SLC10A4 AND NOT-SSTR1 AND NOT-CNGA4	0.988095	0.988095	0.988095
SLC10A4 AND NOT-SSTR1 AND NOT-XCR1	0.988095	0.988095	0.988095
COMPLEX-SLC10A4/EPHA3/TMEM235	0.988095	0.988095	0.988095
SLC10A4 AND NOT-SSTR1 AND NOT-ATP1B4	0.988095	0.988095	0.988095
GPR19 AND PTK7 AND NOT-KCNA1	0.957576	0.975309	0.940476
GPR19 AND PTK7 AND NOT-OR5P3	0.946108	0.951807	0.940476
CACNA1B AND NOT-MRGPRX2 AND PTK7	0.939024	0.9625	0.916667
GPR19 AND PTK7 AND NOT-GNRHR	0.934911	0.929412	0.940476
GPR19 AND PTK7 AND NOT-OR2L2	0.934911	0.929412	0.940476
GPR19 AND PTK7 AND NOT-MRGPRX2	0.934911	0.929412	0.940476
KCNQ2 AND NOT-MRGPRX2 AND PTK7	0.932515	0.962025	0.904762
GPR19 AND PTK7 AND NOT-CACNA1S	0.929412	0.918605	0.940476
GPR19 AND PTK7 AND NOT-GABRG3	0.929412	0.918605	0.940476
CACNA1B AND NOT-CACNA1S AND PTK7	0.925	0.973684	0.880952
KCNQ2 AND NOT-CACNA1S AND PTK7	0.924051	0.986486	0.869048
GPR19 AND PTK7 AND NOT-SLC13A1	0.923977	0.908046	0.940476
GPR19 AND PTK7 AND NOT-LCT	0.923977	0.908046	0.940476
GPR19 AND PTK7 AND NOT-ADAM7	0.923977	0.908046	0.940476
GPR19 AND PTK7 AND NOT-CACNG7	0.923977	0.908046	0.940476
GPR19 AND PTK7 AND NOT-KCNA10	0.923077	0.917647	0.928571
CACNA1B AND NOT-ROS1 AND PTK7	0.922156	0.927711	0.916667
CACNA1B AND NOT-GNRHR AND PTK7	0.922156	0.927711	0.916667
CACNA1B AND NOT-OR5P3 AND PTK7	0.922156	0.927711	0.916667
NKAIN1 AND NOT-GGTLC1 AND PTK7	0.920245	0.949367	0.892857
GPR19 AND PTK7 AND NOT-TMPRSS11B	0.918605	0.897727	0.940476
GPR19 AND PTK7 AND NOT-ROS1	0.918605	0.897727	0.940476
GPR19 AND PTK7 AND NOT-TRPM1	0.918605	0.897727	0.940476
KCNQ2 AND PTK7 AND NOT-TRPM1	0.918239	0.973333	0.869048
KCNQ2 AND PTK7 AND NOT-CACNG7	0.918239	0.973333	0.869048
GPR19 AND PTK7 AND NOT-GJA10	0.917647	0.906977	0.928571
GPR19 AND PTK7 AND NOT-SLC22A11	0.917647	0.906977	0.928571
CACNA1B AND PTK7 AND NOT-CACNG7	0.916667	0.916667	0.916667
GPR19 AND PTK7 AND NOT-CLDN17	0.916667	0.916667	0.916667
CACNA1B AND NOT-SLC22A11 AND PTK7	0.915663	0.926829	0.904762
KCNQ2 AND NOT-OR5P3 AND PTK7	0.915663	0.926829	0.904762
ASTN1 AND NOT-MLC1 AND NOT-CLDN1	0.914286	0.879121	0.952381
GPR19 AND PTK7 AND NOT-GRIN2B	0.913295	0.88764	0.940476
GPR19 AND PTK7 AND NOT-SLC5A8	0.913295	0.88764	0.940476
GPR19 AND PTK7 AND NOT-TRHR	0.913295	0.88764	0.940476
GPR19 AND PTK7 AND NOT-MIP	0.913295	0.88764	0.940476
GPR19 AND PTK7 AND NOT-MOG	0.913295	0.88764	0.940476
KCNQ2 AND NOT-GGTLC1 AND PTK7	0.9125	0.960526	0.869048
KCNQ2 AND PTK7 AND NOT-SLC5A8	0.9125	0.960526	0.869048
GPR19 AND PTK7 AND NOT-UMODL1	0.912281	0.896552	0.928571
ASTN1 AND NOT-MLC1 AND NOT-WT1	0.911243	0.905882	0.916667

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
HEPH AND SLC39A10 AND NOT-ATP6V0A1	1	1	1
XK AND TACSTD2 AND NOT-CRHR2	1	1	1
XK AND TACSTD2 AND NOT-CNR1	1	1	1
XK AND TACSTD2 AND NOT-TLR6	1	1	1
COMPLEX-ATP6V0A1/EVA1A/HEPH	1	1	1
HEPH AND NOT-ATP6V0A1 AND WNT5A	1	1	1
HEPH AND NOT-ATP6V0A1 AND SLC3A2	1	1	1
HEPH AND NOT-ATP6V0A1 AND TM4SF1	1	1	1
IYD AND TACSTD2 AND NOT-LIFR	1	1	1
HEPH AND NOT-ATP6V0A1 AND NOT-BEST4	1	1	1
NOT-LIFR AND TACSTD2 AND TSPAN1	1	1	1
NOT-LIFR AND TACSTD2 AND TSPAN15	1	1	1
NOT-LIFR AND TACSTD2 AND PARM1	1	1	1
PVRL3 AND NOT-LIFR AND TACSTD2	1	1	1
COMPLEX-IL6R/EVA1A/HEPH	1	1	1
HEPH AND SLC39A10 AND ABCC11	1	1	1
HEPH AND SLC39A10 AND COL25A1	1	1	1
HEPH AND SLC39A10 AND NOT-IL20RB	1	1	1
HEPH AND SLC39A10 AND NOT-PTGER3	1	1	1
HEPH AND SLC39A10 AND NOT-NLGN2	1	1	1
HEPH AND SLC39A10 AND ISLR2	1	1	1
HEPH AND SLC39A10 AND LRRC4C	1	1	1
HEPH AND SLC39A10 AND PTPRG	1	1	1
HEPH AND SLC39A10 AND PTPRN	1	1	1
HEPH AND SLC39A10 AND NOT-JAM2	1	1	1
HEPH AND SLC39A10 AND NOT-LGR6	1	1	1
HEPH AND SLC39A10 AND RHCE	1	1	1
HEPH AND SLC39A10 AND SLC5A7	1	1	1
HEPH AND SLC39A10 AND RTN2	1	1	1
HEPH AND SLC39A10 AND NOT-SCN1B	1	1	1
HEPH AND SLC39A10 AND NOT-SCN4B	1	1	1
HEPH AND SLC39A10 AND SCTR	1	1	1
HEPH AND SLC39A10 AND SECTM1	1	1	1
HEPH AND SLC39A10 AND NOT-PERP	1	1	1
HEPH AND SLC39A10 AND DSCAML1	1	1	1
HEPH AND SLC39A10 AND CYSLTR2	1	1	1
HEPH AND SLC39A10 AND EVA1A	1	1	1
HEPH AND SLC39A10 AND NOT-SLC17A7	1	1	1
HEPH AND SLC39A10 AND NOT-GPR173	1	1	1
HEPH AND SLC39A10 AND TAS2R5	1	1	1
HEPH AND SLC39A10 AND SLC6A20	1	1	1
HEPH AND SLC39A10 AND NOT-TRPM7	1	1	1
NOT-SLC16A7 AND TGFBI AND JPH1	1	1	1
IFI6 AND MEP1A AND NOT-ADRB3	1	1	1
IFI6 AND B3GNT3 AND NOT-DUOX1	1	1	1
IFI6 AND MEP1A AND CD9	1	1	1
IFI6 AND NOT-CLEC2D AND LY75	1	1	1
NOX1 AND IFITM1 AND NOT-LRMP	1	1	1
IFI6 AND NOT-CLEC2D AND PAQR5	1	1	1
IFI6 AND SLC39A4 AND NOT-CLEC2D	1	1	1
IFI6 AND NOT-SLC16A7 AND TMEM47	1	1	1
IFI6 AND SLC5A1 AND NOT-TMEM47	1	1	1
IFI6 AND ATP10B AND ATP1B1	1	1	1
NOX1 AND AMIGO2 AND NOT-LRMP	1	1	1
IFI6 AND PPAP2C AND NOT-CLEC2D	1	1	1
NOX1 AND EDNRA AND NOT-LRMP	1	1	1
IFI6 AND SLC26A3 AND ATP1B1	1	1	1
IFI6 AND ST14 AND NOT-ADRB3	1	1	1
IFI6 AND NOT-ADRB3 AND SLC52A2	1	1	1
IFI6 AND NOT-CLEC2D AND CXCR1	1	1	1
IFI6 AND NOT-CLEC2D AND ATP6V0A2	1	1	1
IFI6 AND GJB3 AND NOT-IL1RL1	1	1	1
IFI6 AND MEP1A AND NOT-ISLR2	1	1	1
IFI6 AND NOT-SLC19A3 AND ADAM15	1	1	1
IFI6 AND SLC35G1 AND NOT-ISLR2	1	1	1
IFI6 AND NOT-SLC19A3 AND GAL3ST1	1	1	1
IFI6 AND PPAP2C AND NOT-IZUMO1	1	1	1
IFI6 AND DUOX2 AND NOT-GPR63	1	1	1
IFI6 AND SLC39A4 AND NOT-IZUMO1	1	1	1
IFI6 AND DUOX2 AND NOT-KCNH7	1	1	1
IFI6 AND DUOX2 AND NOT-SLC16A7	1	1	1
IFI6 AND NOT-CLEC2D AND CYBA	1	1	1
IFI6 AND NOT-ADRB3 AND SLC27A4	1	1	1
IFI6 AND DUOX2 AND NOT-KCNB2	1	1	1
IFI6 AND DUOX2 AND NOT-ACVR2B	1	1	1
IFI6 AND DUOX2 AND NOT-NRG2	1	1	1
IFI6 AND SLC35G1 AND NOT-ADRB3	1	1	1

Antigen Pair Logic	F1	Precision	Recall
GPR19 AND PTK7 AND NOT-CLRN1	0.911243	0.905882	0.916667
CACNA1B AND NOT-ADAM7 AND PTK7	0.911243	0.905882	0.916667
CACNA1B AND NOT-OR2L2 AND PTK7	0.911243	0.905882	0.916667
CACNA1B AND NOT-TMPRSS11B AND PTK7	0.911243	0.905882	0.916667
CACNA1B AND NOT-GRIN2B AND PTK7	0.911243	0.905882	0.916667
SLC6A15 AND NOT-KCNA1 AND PTK7	0.910256	0.986111	0.845238
KCNQ2 AND NOT-TMPRSS11B AND PTK7	0.91018	0.915663	0.904762
CACNA1B AND NOT-GGTLC1 AND PTK7	0.91018	0.915663	0.904762
KCNQ2 AND NOT-GNRHR AND PTK7	0.91018	0.915663	0.904762
KCNQ2 AND NOT-ROS1 AND PTK7	0.91018	0.915663	0.904762
CACNA1B AND NOT-CLRN1 AND PTK7	0.91018	0.915663	0.904762
KCNQ2 AND NOT-SLC22A11 AND PTK7	0.909091	0.925926	0.892857
GPR19 AND PTK7 AND NOT-SLC12A3	0.908046	0.877778	0.940476
GPR19 AND PTK7 AND NOT-SLC28A1	0.908046	0.877778	0.940476
KCNQ2 AND NOT-MOG AND PTK7	0.905882	0.895349	0.916667
CACNA1B AND NOT-GABRG3 AND PTK7	0.905882	0.895349	0.916667
CACNA1B AND NOT-MOG AND PTK7	0.905882	0.895349	0.916667
CACNA1B AND NOT-LCT AND PTK7	0.905882	0.895349	0.916667
KCNQ2 AND NOT-GPR6 AND PTK7	0.904762	0.904762	0.904762
KCNQ2 AND NOT-GRIN2B AND PTK7	0.904762	0.904762	0.904762
GPR19 AND PTK7 AND NOT-UPK3A	0.902857	0.868132	0.940476
GPR19 AND PTK7 AND NOT-CCR9	0.902857	0.868132	0.940476
KCNQ2 AND PTK7 AND NOT-UMODL1	0.901235	0.935897	0.869048
KCNQ2 AND PTK7 AND NOT-GPR61	0.901235	0.935897	0.869048
CACNA1B AND NOT-MIP AND PTK7	0.900585	0.885057	0.916667
CACNA1B AND NOT-SLC13A1 AND PTK7	0.900585	0.885057	0.916667
CACNA1B AND PTK7 AND NOT-SLC5A8	0.900585	0.885057	0.916667
CACNA1B AND NOT-TRPM1 AND PTK7	0.900585	0.885057	0.916667
CACNA1B AND NOT-SLC28A1 AND PTK7	0.900585	0.885057	0.916667
CACNA1B AND NOT-SLC12A3 AND PTK7	0.900585	0.885057	0.916667
KCNQ2 AND NOT-ADAM7 AND PTK7	0.899408	0.894118	0.904762
KCNQ2 AND NOT-KCNA10 AND PTK7	0.898204	0.903614	0.892857
KCNQ2 AND NOT-CLRN1 AND PTK7	0.898204	0.903614	0.892857
NKAIN1 AND NOT-CLDN8 AND NOT-CALHM1	0.89697	0.91358	0.880952
ASTN1 AND NOT-MLC1 AND NOT-TYR	0.901734	0.876404	0.928571
NKAIN1 AND NOT-CLRN1 AND PTK7	0.89441	0.935065	0.857143
CACNA1B AND NOT-CLDN17 AND PTK7	0.894118	0.883721	0.904762
KCNQ2 AND NOT-LCT AND PTK7	0.894118	0.883721	0.904762
KCNQ2 AND NOT-OR2L2 AND PTK7	0.894118	0.883721	0.904762
SCN2A AND PTK7 AND NOT-CACNA1S	0.893082	0.946667	0.845238
UNC5A AND PTK7 AND NOT-MRGPRX2	0.898734	0.959459	0.845238
SCN2A AND PTK7 AND NOT-CACNG7	0.893082	0.946667	0.845238
UNC5A AND PTK7 AND NOT-ROS1	0.904459	0.972603	0.845238
ASTN1 AND NOT-NKAIN2 AND NOT-CLDN8	0.893082	0.946667	0.845238
SCN2A AND PTK7 AND NOT-OR5P3	0.893082	0.946667	0.845238
COMPLEX-SSTR1/NKAIN1/TMPRSS11D	0.892857	0.892857	0.892857
GPR19 AND PTK7 AND NOT-GPR6	0.892655	0.849462	0.940476
GPR19 AND PTK7 AND NOT-HTR1F	0.892655	0.849462	0.940476
SLC6A15 AND PTK7 AND NOT-OR5P3	0.89172	0.958904	0.833333
SLC6A15 AND NOT-MRGPRX2 AND PTK7	0.89172	0.958904	0.833333
CACNA1B AND PTK7 AND NOT-UMODL1	0.888889	0.873563	0.904762
NOT-EGFR AND ATP7A AND NOT-BTN3A1	0.888889	0.873563	0.904762
KCNQ2 AND NOT-SLC13A1 AND PTK7	0.888889	0.873563	0.904762
CACNA1B AND PTK7 AND NOT-PCDH15	0.888889	0.873563	0.904762
KCNQ2 AND NOT-HTR1F AND PTK7	0.888889	0.873563	0.904762
KCNQ2 AND NOT-SLC12A3 AND PTK7	0.888889	0.873563	0.904762
GPR19 AND PTK7 AND NOT-OR1C1	0.88764	0.840426	0.940476
NOT-EGFR AND PTK7 AND EBP	0.938272	0.974359	0.904762
NOT-EGFR AND CD276 AND NOT-EMP3	0.934911	0.929412	0.940476
NOT-EGFR AND PTK7 AND NOT-EMP3	0.927711	0.939024	0.916667
NOT-EGFR AND PTK7 AND NOT-ROS1	0.934911	0.929412	0.940476
NOT-EGFR AND PTK7 AND NOT-DSC1	0.946108	0.951807	0.940476
NOT-EGFR AND PTK7 AND NOT-GGTLC1	0.934911	0.929412	0.940476
KCNQ2 AND PTK7 AND NOT-GUCY2C	0.924051	0.986486	0.869048
NOT-EGFR AND PTK7 AND NOT-BTN3A1	0.940476	0.940476	0.940476
NOT-EGFR AND PTK7 AND NOT-TMPRSS11B	0.951807	0.963415	0.940476
NOT-EGFR AND PTK7 AND NOT-LCT	0.946108	0.951807	0.940476
NOT-EGFR AND PTK7 AND NOT-CACNA1S	0.934911	0.929412	0.940476
CACNA1B AND PTK7 AND NOT-GUCY2C	0.916667	0.916667	0.916667
COMPLEX-EGFR/NKAIN1/CD33	0.914634	0.9375	0.892857
GPR19 AND PTK7 AND NOT-ITGB3	0.913295	0.88764	0.940476
NOT-EGFR AND PTK7 AND ATP13A5	0.91018	0.915663	0.904762
NOT-EGFR AND CD276 AND NOT-DSC1	0.909091	0.869565	0.952381
GPR19 AND PTK7 AND NOT-CLDN2	0.908046	0.877778	0.940476
GPR19 AND PTK7 AND NOT-GUCY2C	0.908046	0.877778	0.940476
KCNQ2 AND PTK7 AND NOT-CLDN2	0.906832	0.948052	0.869048
CACNA1B AND PTK7 AND NOT-CLDN2	0.905882	0.895349	0.916667

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
IFI6 AND SLC39A4 AND NOT-OR10H1	1	1	1
IFI6 AND ST14 AND NOT-ISLR2	1	1	1
IFI6 AND TNFRSF11A AND NOT-ADRB3	1	1	1
IFI6 AND ATP10B AND NOT-IL1RL1	1	1	1
IFI6 AND AKAP1 AND NOT-SLC19A3	1	1	1
IFI6 AND PTPRH AND NOT-INSRR	1	1	1
IFI6 AND B3GNT3 AND NOT-SLC38A4	1	1	1
IFI6 AND SLC44A3 AND NOT-SLC26A7	1	1	1
IFI6 AND GPRC5A AND NOT-INSRR	1	1	1
IFI6 AND SLC26A3 AND NOT-SLC26A7	1	1	1
IFI6 AND ATP10B AND NOT-SLC26A7	1	1	1
IFI6 AND CDCP1 AND NOT-INSRR	1	1	1
IFI6 AND OR2F1 AND NOT-SLC26A7	1	1	1
IFI6 AND NOT-SLC4A3 AND PIGU	1	1	1
IFI6 AND PAQR5 AND NOT-PTPRM	1	1	1
IFI6 AND NOT-ATP6V0A1 AND KCNMA1	1	1	1
IFI6 AND DUOX2 AND NOT-INSRR	1	1	1
IFI6 AND KCNK1 AND NOT-INSRR	1	1	1
IFI6 AND ITGA2 AND NOT-SLC26A7	1	1	1
IFI6 AND MEP1A AND NOT-SLC4A3	1	1	1
IFI6 AND PPAP2C AND NOT-RXFP1	1	1	1
IFI6 AND ST14 AND NOT-SLC4A3	1	1	1
IFI6 AND SLC39A4 AND NOT-PTPRM	1	1	1
IFI6 AND JPH1 AND NOT-ATP6V0A1	1	1	1
IFI6 AND CYBA AND NOT-CCR3	1	1	1
IFI6 AND KCNK1 AND NOT-GPR55	1	1	1
IFI6 AND SLC26A3 AND NOT-IL1RL1	1	1	1
IFI6 AND SLC44A3 AND NOT-IL1RL1	1	1	1
IFI6 AND NOT-CCR3 AND ATP6V0A2	1	1	1
IFI6 AND NOT-CCR3 AND CXCR1	1	1	1
IFI6 AND GPRC5A AND NOT-PEAR1	1	1	1
IFI6 AND MEP1A AND NOT-IL17RB	1	1	1
IFI6 AND PAQR5 AND NOT-CCR3	1	1	1
IFI6 AND LGR4 AND NOT-SLC38A4	1	1	1
IFI6 AND SLC39A4 AND NOT-CCR3	1	1	1
IFI6 AND DUOX2 AND NOT-GPR55	1	1	1
IFI6 AND PPAP2C AND NOT-CCR3	1	1	1
IFI6 AND SLC5A1 AND NOT-ATP6V0A1	1	1	1
IFI6 AND SLC5A1 AND NOT-SLC38A4	1	1	1
IFI6 AND JPH1 AND NOT-SLC38A4	1	1	1
IFI6 AND IL17RB AND NOT-SLC38A4	1	1	1
IFI6 AND DUOX2 AND NOT-VIPR2	1	1	1
IFI6 AND NOT-SLC19A3 AND FAT4	1	1	1
IFI6 AND NOT-SLC16A7 AND PAQR5	1	1	1
IFI6 AND SLC44A3 AND NOT-CHRNE	1	1	1
IFI6 AND ATP10B AND NOT-PCDH20	1	1	1
IFI6 AND SLC26A3 AND NOT-PCDH20	1	1	1
IFI6 AND DUOX2 AND NOT-TAS2R50	1	1	1
IFI6 AND CFTR AND NOT-CHRNE	1	1	1
IFI6 AND SYT13 AND NOT-NPY2R	1	1	1
IFI6 AND DUOX2 AND NOT-GLRA1	1	1	1
IFI6 AND SYT13 AND VIMP	1	1	1
IFI6 AND DUOX2 AND NOT-NLGN4Y	1	1	1
IFI6 AND MEP1A AND NOT-SLC8A1	1	1	1
IFI6 AND SYT13 AND NOT-WDR19	1	1	1
IFI6 AND DUOX2 AND NOT-GRM2	1	1	1
IFI6 AND DUOX2 AND NOT-HCRTR2	1	1	1
IFI6 AND SLC26A3 AND NOT-CHRNE	1	1	1
IFI6 AND ATP10B AND NOT-CHRNE	1	1	1
IFI6 AND OR2F1 AND NOT-CHRNE	1	1	1
IFI6 AND SYT13 AND NOT-GRIA2	1	1	1
IFI6 AND DUOX2 AND NOT-FGFR2	1	1	1
MEP1A AND SLC39A10 AND NOT-GPC3	1	1	1
MEP1A AND TACSTD2 AND NOT-GPC3	1	1	1
NOT-SLC16A7 AND SLC39A10 AND MUC1	1	1	1
XK AND CDH3 AND NOT-MUC1	1	1	1
MEP1A AND CDH3 AND NOT-MUC1	1	1	1
ATP10B AND NOT-FLVCR2 AND NOT-MUC1	1	1	1
MEP1A AND TACSTD2 AND NOT-GPC3	1	1	1
MEP1A AND SLC39A10 AND NOT-MUC1	1	1	1
MEP1A AND SLC39A10 AND NOT-ERBB2	1	1	1
NOT-LIFR AND CD81 AND MUC1	1	1	1
HEPH AND SLC39A10 AND NOT-ERBB2	1	1	1
IYD AND CDH3 AND NOT-MUC1	1	1	1
IYD AND SLC39A10 AND NOT-GPC3	1	1	1
IYD AND NOT-ABHD6 AND NOT-GPC3	1	1	1
HEPH AND SLC39A10 AND MSLN	1	1	1

Antigen Pair Logic	F1	Precision	Recall
CACNA1B AND NOT-ITGB3 AND PTK7	0.900585	0.885057	0.916667
NKAIN1 AND NOT-CLDN8 AND NOT-SSTR1	0.898204	0.903614	0.892857
NOT-EGFR AND PTK7 AND OR1C1	0.89697	0.91358	0.880952
KCNQ2 AND PTK7 AND NOT-MUC16	0.89441	0.935065	0.857143
KCNQ2 AND NOT-ITGB3 AND PTK7	0.894118	0.883721	0.904762
SCN2A AND PTK7 AND NOT-GUCY2C	0.893082	0.946667	0.845238
NKAIN1 AND NOT-CLDN8 AND NOT-GPA33	0.887574	0.882353	0.892857
CACNA1B AND PTK7 AND NOT-MUC16	0.883721	0.863636	0.904762
UNC5A AND PTK7 AND NOT-CLDN2	0.881988	0.922078	0.845238
UNC5A AND PTK7 AND NOT-GUCY2C	0.881988	0.922078	0.845238
NOT-EGFR AND BMPR1B AND NOT-BTN3A1	0.878049	0.9	0.857143
NOT-EGFR AND BMPR1B AND NOT-DSC1	0.878049	0.9	0.857143
NOT-EGFR AND BMPR1B AND NOT-EMP3	0.878049	0.9	0.857143
SCN2A AND PTK7 AND NOT-ITGB3	0.876543	0.910256	0.845238
SCN2A AND PTK7 AND NOT-CLDN2	0.871166	0.898734	0.845238
NKAIN1 AND NOT-CLDN8 AND NOT-CLDN18	0.867052	0.842697	0.892857
NOT-EGFR AND FLVCR1 AND NOT-CD37	0.866242	0.931507	0.809524
GPR85 AND PTK7 AND NOT-GUCY2C	0.866242	0.931507	0.809524
UNC5A AND PTK7 AND NOT-ITGB3	0.865854	0.8875	0.845238
SLC6A15 AND PTK7 AND NOT-GUCY2C	0.864198	0.897436	0.833333
SLC6A15 AND PTK7 AND NOT-CLDN2	0.864198	0.897436	0.833333
SCN2A AND PTK7 AND NOT-MUC16	0.864198	0.897436	0.833333
GPR85 AND PTK7 AND NOT-CLDN2	0.860759	0.918919	0.809524
UNC5A AND PTK7 AND NOT-MUC16	0.858896	0.886076	0.833333
NOT-EGFR AND ATP7A AND NOT-CD33	0.858757	0.817204	0.904762
NOT-EGFR AND PTK7 AND UMODL1	0.853503	0.917808	0.797619
NOT-EGFR AND PTK7 AND TRPM1	0.853503	0.917808	0.797619
SLC6A15 AND NOT-ITGB3 AND PTK7	0.848485	0.864198	0.833333
NOT-EGFR AND ATP7A AND NOT-CD37	0.847458	0.806452	0.892857
SLC4A8 AND NOT-MUC16 AND PTK7	0.847059	0.837209	0.857143
JPH3 AND NOT-ITGB3 AND PTK7	0.845238	0.845238	0.845238
GPR85 AND PTK7 AND NOT-ITGB3	0.84472	0.883117	0.809524
NOT-EGFR AND FLVCR1 AND NOT-CD33	0.84472	0.883117	0.809524
ASTN1 AND PTK7 AND NOT-CLDN2	0.843373	0.853659	0.833333
SYT4 AND PTK7 AND NOT-GUCY2C	0.841463	0.8625	0.821429
NOT-CLDN4 AND PTK7 AND EBP	0.840909	0.804348	0.880952
TMEFF2 AND PTK7 AND NOT-ITGB3	0.840237	0.835294	0.845238
L1CAM AND NOT-MOG AND NOT-PTK7	0.836158	0.795699	0.880952
NOT-RNF43 AND PTK7 AND EBP	0.840237	0.835294	0.845238
NKAIN1 AND NOT-CLDN8 AND NOT-ITGB3	0.833333	0.78125	0.892857
SYT4 AND PTK7 AND NOT-CLDN2	0.831325	0.841463	0.821429
SYT11 AND NOT-EGFR AND CD37	0.830409	0.816092	0.845238
ASTN1 AND BIRC5 AND NOT-ITGB3	0.828947	0.926471	0.75
COMPLEX-PTK7/NKAIN1/CLDN2	0.828729	0.773196	0.892857
ASTN1 AND PTK7 AND NOT-FOLH1	0.828402	0.823529	0.833333
SYT4 AND PTK7 AND NOT-MUC16	0.826347	0.831325	0.821429
NOT-EGFR AND PTK7 AND GPR61	0.825806	0.901408	0.761905
NKAIN1 AND NOT-CD34 AND NOT-SSTR1	0.825	0.868421	0.785714
NOT-EGFR AND LPPR3 AND NOT-CD33	0.825	0.868421	0.785714
SLC6A15 AND PTK7 AND NOT-MUC16	0.823529	0.813953	0.833333
ASTN1 AND PTK7 AND NOT-ITGB3	0.823529	0.813953	0.833333
ASTN1 AND PTK7 AND NOT-ITGB3	0.823529	0.813953	0.833333
NOT-EGFR AND PTK7 AND SLC5A8	0.823529	0.913043	0.75
NKAIN1 AND NOT-CLDN8 AND NOT-SLC34A2	0.820225	0.776596	0.869048
GPR158 AND PTK7 AND NOT-ITGB3	0.818792	0.938462	0.72619
TMEFF2 AND PTK7 AND NOT-CLDN2	0.818182	0.782609	0.857143
ASTN1 AND BIRC5 AND NOT-FOLH1	0.818182	0.9	0.75
CNIH2 AND NOT-MUC16 AND PTK7	0.833333	0.902778	0.77381
NOT-CLDN1 AND PTK7 AND NOT-GGTLC1	0.811429	0.78022	0.845238
NOT-CLDN4 AND PTK7 AND NOT-ATP13A5	0.811429	0.78022	0.845238
NOT-LRIG3 AND PTK7 AND NOT-ITGB3	0.810127	0.864865	0.761905
ASTN1 AND NOT-ERBB4 AND NOT-CLDN8	0.813333	0.924242	0.72619
GPR158 AND PTK7 AND NOT-CLDN2	0.805195	0.885714	0.738095
NOT-EGFR AND FLVCR1 AND NOT-CD79B	0.804734	0.8	0.809524
SLC6A15 AND NOT-EGFR AND CD37	0.802548	0.863014	0.75
NOT-EGFR AND ATP8B2 AND NOT-CD33	0.80226	0.763441	0.845238
NOT-CLDN1 AND PTK7 AND NOT-KCNA1	0.813559	0.774194	0.857143
SLC10A4 AND NOT-NKAIN2 AND NOT-SEMA4B	0.994012	1	0.988095
SLC10A4 AND NOT-NKAIN2 AND NOT-SCN10A	0.994012	1	0.988095
SLC10A4 AND NOT-NKAIN2 AND NOT-ATP1B4	0.994012	1	0.988095
SLC10A4 AND NOT-TMEM235 AND NOT-SLC9B1	0.994012	1	0.988095
COMPLEX-SEMA4D/CSMD3/SLC10A4	0.994012	1	0.988095
SLC10A4 AND NOT-DTNA AND NOT-SLC13A1	0.994012	1	0.988095
SLC10A4 AND NOT-NKAIN2 AND NOT-ROS1	0.994012	1	0.988095
SLC10A4 AND NOT-KCNJ10 AND NOT-SLC9B1	0.994012	1	0.988095
SLC10A4 AND NOT-NKAIN2 AND NOT-XCR1	0.994012	1	0.988095
SLC10A4 AND NOT-MLC1 AND NOT-GLRA3	0.994012	1	0.988095

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
MEP1A AND SLC39A10 AND NOT-MSLN	1	1	1
ATP10B AND AMIGO2 AND NOT-ERBB2	1	1	1
LY75 AND EDNRA AND NOT-GPC3	1	1	1
NOT-GPC3 AND CDH3 AND NOT-ATP13A4	1	1	1
KCNE3 AND EDNRA AND NOT-GPC3	1	1	1
XK AND EDNRA AND NOT-GPC3	1	1	1
NOT-SLC16A7 AND EDNRA AND NOT-GPC3	1	1	1
NOT-LIFR AND CD81 AND ERBB2	1	1	1
SLC26A3 AND TACSTD2 AND NOT-GPC3	1	1	1
HEPH AND EDNRA AND NOT-GPC3	1	1	1
HEPH AND NOT-FLVCR2 AND NOT-MUC1	1	1	1
SLC26A3 AND SLC39A10 AND NOT-GPC3	1	1	1
HEPH AND NOT-ABHD6 AND NOT-GPC3	1	1	1
MEP1A AND AMIGO2 AND NOT-ERBB2	1	1	1
NOT-GPC3 AND CDH3 AND NOT-DUOX1	1	1	1
HEPH AND TACSTD2 AND NOT-GPC3	1	1	1
NOT-GPC3 AND CDH3 AND NOT-GPR87	1	1	1
SLC26A3 AND EDNRA AND NOT-GPC3	1	1	1
ATP10B AND SLC39A10 AND NOT-ERBB2	1	1	1
SLC26A3 AND SLC39A10 AND NOT-ERBB2	1	1	1
NOT-SLC16A7 AND SLC39A10 AND NOT-GPC3	1	1	1
HEPH AND NOT-CEACAM1 AND NOT-GPC3	1	1	1
MEP1A AND TACSTD2 AND NOT-MUC1	1	1	1
HEPH AND TACSTD2 AND NOT-GPC3	1	1	1
NOT-GPC3 AND CDH3 AND NOT-GPR87	1	1	1
NOT-GPC3 AND CDH3 AND NOT-DUOX1	1	1	1
MEP1A AND NOT-CEACAM1 AND NOT-GPC3	1	1	1
SLC26A3 AND TACSTD2 AND NOT-GPC3	1	1	1
JPH1 AND CDH3 AND NOT-GPC3	1	1	1
NOT-GPC3 AND CDH3 AND NOT-CLCA2	1	1	1
NOT-GPC3 AND CDH3 AND NOT-PVRL4	1	1	1
NOT-FAT2 AND CDH3 AND NOT-GPC3	1	1	1
SLC26A3 AND NOT-ABHD6 AND NOT-GPC3	1	1	1
NOT-GPC3 AND CDH3 AND NOT-PTPRZ1	1	1	1
PVRL3 AND CDH3 AND NOT-GPC3	1	1	1
NOT-GPC3 AND CDH3 AND NOT-MUC15	1	1	1
PVRL3 AND CDH3 AND NOT-GPC3	1	1	1
JPH1 AND CDH3 AND NOT-GPC3	1	1	1
ITM2C AND TACSTD2 AND NOT-GPC3	1	1	1
NOT-GPC3 AND CDH3 AND NOT-LTB4R	1	1	1
NOT-GPC3 AND CDH3 AND NOT-KRT5	1	1	1
ATP10B AND SLC39A10 AND NOT-GPC3	1	1	1
HEPH AND SLCO4A1 AND NOT-ERBB2	1	1	1
SLC26A3 AND NOT-CEACAM1 AND NOT-GPC3	1	1	1
MEP1A AND NOT-ABHD6 AND NOT-GPC3	1	1	1
XK AND CDH3 AND NOT-GPC3	1	1	1
NOT-LIFR AND EDNRA AND MUC1	1	1	1
NOT-FAT2 AND CDH3 AND NOT-GPC3	1	1	1
JPH1 AND IFITM1 AND MUC1	1	1	1
SLC26A3 AND SLCO4A1 AND NOT-ERBB2	1	1	1
IYD AND TACSTD2 AND NOT-GPC3	1	1	1
HEPH AND CDH3 AND NOT-MUC1	1	1	1
NOT-SLC16A7 AND CDH3 AND NOT-GPC3	1	1	1
MEP1A AND SLCO4A1 AND NOT-ERBB2	1	1	1
XK AND CDH3 AND NOT-GPC3	1	1	1
NOT-SLC16A7 AND SLCO4A1 AND MUC1	1	1	1
NOT-GPC3 AND CDH3 AND NOT-DUOXA1	1	1	1
NOT-GPC3 AND CDH3 AND NOT-DUOXA1	1	1	1
NOT-GPC3 AND CDH3 AND NOT-LTB4R	1	1	1
HEPH AND TACSTD2 AND NOT-ERBB2	1	1	1
NOT-GPC3 AND CDH3 AND NOT-MUC15	1	1	1
NOT-SLC16A7 AND CDH3 AND NOT-GPC3	1	1	1
NOT-ZP2 AND JPH1 AND ERBB2	0.909091	0.833333	1
CCRL2 AND NOT-FLVCR2 AND MUC1	0.909091	0.833333	1
TSPAN8 AND EDNRA AND NOT-GPC3	1	1	1
IYD AND SLC39A10 AND NOT-ERBB2	0.909091	0.833333	1
NOT-SLC16A7 AND WNT5A AND MUC1	1	1	1
IYD AND NOT-FLVCR2 AND NOT-GPC3	0.909091	0.833333	1
NOT-GPC3 AND CDH3 AND NOT-AQP3	1	1	1
NOT-SLC16A7 AND EDNRA AND MUC1	1	1	1
JPH1 AND TMEM97 AND NOT-GPC3	1	1	1
JPH1 AND NOT-KCNB1 AND ERBB2	0.909091	0.833333	1
TM4SF5 AND CDH3 AND NOT-GPC3	1	1	1
MEP1A AND SLC22A3 AND NOT-GPC3	1	1	1
HEPH AND NOT-FLVCR2 AND NOT-ERBB2	0.909091	0.833333	1
MEP1A AND AMIGO2 AND NOT-MSLN	0.909091	0.833333	1
JPH1 AND TACSTD2 AND NOT-GPC3	0.909091	0.833333	1

Antigen Pair Logic	F1	Precision	Recall
SLC10A4 AND NOT-DTNA AND NOT-SLC13A1	0.994012	1	0.988095
SLC10A4 AND NOT-NKAIN2 AND NOT-MRAP	0.994012	1	0.988095
SLC10A4 AND NOT-NKAIN2 AND NOT-ROS1	0.994012	1	0.988095
SLC10A4 AND NOT-NKAIN2 AND NOT-ADAM7	0.994012	1	0.988095
SLC10A4 AND NOT-KCNJ10 AND NOT-SEMA4B	0.994012	1	0.988095
SLC10A4 AND NOT-MLC1 AND NOT-SLC9B1	0.994012	1	0.988095
SLC10A4 AND NOT-OPALIN AND NOT-SLC9B1	0.994012	1	0.988095
SLC10A4 AND NOT-NKAIN2 AND NOT-LRRC8E	0.994012	1	0.988095
SLC10A4 AND NOT-KCNJ10 AND NOT-ATP1B4	0.994012	1	0.988095
SLC10A4 AND NOT-GABRB2 AND NOT-GLRA3	0.994012	1	0.988095
SLC10A4 AND NOT-KCNJ10 AND NOT-XCR1	0.994012	1	0.988095
SLC10A4 AND NOT-HCN2 AND NOT-GLRA3	0.994012	1	0.988095
COMPLEX-SLC10A4/HTR5A/KCNJ10	0.994012	1	0.988095
SLC10A4 AND NOT-KCNJ10 AND NOT-GPR6	0.994012	1	0.988095
SLC10A4 AND NOT-KCNJ10 AND NOT-MRAP	0.994012	1	0.988095
SLC10A4 AND NOT-GABRB2 AND NOT-SLC13A1	0.994012	1	0.988095
SLC10A4 AND NOT-MOG AND NOT-GLRA3	0.994012	1	0.988095
SLC10A4 AND NOT-MEGF10 AND NOT-GLRA3	0.994012	1	0.988095
SLC10A4 AND NOT-NKAIN2 AND NOT-STEAP4	0.994012	1	0.988095
SLC10A4 AND NOT-OPALIN AND NOT-SLC9B1	0.994012	1	0.988095
SLC10A4 AND NOT-NKAIN2 AND NOT-SLC12A3	0.994012	1	0.988095
SLC10A4 AND NOT-NKAIN2 AND NOT-SLC13A1	0.994012	1	0.988095
SLC10A4 AND NOT-NKAIN2 AND NOT-NMBR	0.994012	1	0.988095
COMPLEX-NKAIN2/SLC10A4/GUCY2F	0.994012	1	0.988095
SLC10A4 AND NOT-ACSL6 AND NOT-SLC13A1	0.994012	1	0.988095
SLC10A4 AND NOT-KCNJ10 AND NOT-MRGPRX2	0.994012	1	0.988095
SLC10A4 AND NOT-TTYH2 AND NOT-GLRA3	0.994012	1	0.988095
SLC10A4 AND NOT-NKAIN2 AND NOT-SLC28A1	0.994012	1	0.988095
SLC10A4 AND NOT-KCNJ10 AND NOT-ROS1	0.994012	1	0.988095
COMPLEX-CHRNA4/SLC10A4/KCNJ10	0.994012	1	0.988095
SLC10A4 AND NOT-GABRB2 AND NOT-GNRHR	0.994012	1	0.988095
SLC10A4 AND NOT-KCNJ10 AND NOT-OR2L2	0.994012	1	0.988095
SLC10A4 AND NOT-NKAIN2 AND SLC38A1	0.994012	1	0.988095
SLC10A4 AND NOT-NKAIN2 AND NOT-HTR6	0.994012	1	0.988095
SLC10A4 AND NOT-TTYH2 AND NOT-SLC9B1	0.994012	1	0.988095
SLC10A4 AND NOT-KCNJ10 AND NOT-SLC13A1	0.994012	1	0.988095
SLC10A4 AND NOT-NKAIN2 AND NOT-SLC9B1	0.994012	1	0.988095
SLC10A4 AND NOT-GABRB2 AND NOT-MRGPRX2	0.994012	1	0.988095
COMPLEX-SLC10A4/NTSR2/OPALIN	0.994012	1	0.988095
COMPLEX-SLC10A4/GRM7/KCNJ6	0.994012	1	0.988095
SLC10A4 AND NOT-MOG AND NOT-GLRA3	0.994012	1	0.988095
SLC10A4 AND NOT-ACSL6 AND NOT-GLRA3	0.994012	1	0.988095
SLC10A4 AND NOT-ACSL6 AND NOT-SLC13A1	0.994012	1	0.988095
COMPLEX-SLC10A4/KCNJ10/SLC22A11	0.994012	1	0.988095
SLC10A4 AND NOT-GABRB2 AND SLC38A1	0.994012	1	0.988095
SLC10A4 AND NOT-MOG AND NOT-SLC9B1	0.994012	1	0.988095
SLC10A4 AND NOT-KCNJ10 AND NOT-ROS1	0.994012	1	0.988095
SLC10A4 AND NOT-TMEM235 AND NOT-SLC9B1	0.994012	1	0.988095
SLC10A4 AND NOT-GABRB2 AND NOT-ADAM7	0.994012	1	0.988095
COMPLEX-SLC10A4/GRIN2C/KCNJ10	0.994012	1	0.988095
SLC10A4 AND NOT-NKAIN2 AND NOT-CASR	0.994012	1	0.988095
SLC10A4 AND NOT-NKAIN2 AND NOT-GLRA3	0.994012	1	0.988095
COMPLEX-CHRNA4/SLC10A4/KCNJ10	0.994012	1	0.988095
SLC10A4 AND NOT-NKAIN2 AND SLC38A1	0.994012	1	0.988095
SLC10A4 AND NOT-KCNJ10 AND NOT-MRGPRX2	0.994012	1	0.988095
SLC10A4 AND NOT-KCNJ10 AND NOT-SLC9B1	0.994012	1	0.988095
SLC10A4 AND NOT-SEMA4D AND NOT-HTR1F	0.994012	1	0.988095
SLC10A4 AND NOT-KCNJ10 AND NOT-GNRHR	0.994012	1	0.988095
COMPLEX-SLC10A4/CRB2/KCNJ10	0.994012	1	0.988095
SLC10A4 AND NOT-KCNJ10 AND NOT-SEMA4B	0.994012	1	0.988095
SLC10A4 AND NOT-KCNJ10 AND NOT-GPR6	0.994012	1	0.988095
SLC10A4 AND NOT-KCNJ10 AND NOT-HTR6	0.994012	1	0.988095
SLC10A4 AND NOT-NKAIN2 AND NOT-SLC13A1	0.994012	1	0.988095
SLC10A4 AND NOT-KCNJ10 AND NOT-SCN10A	0.994012	1	0.988095
SLC10A4 AND NOT-NKAIN2 AND NOT-SLC12A3	0.994012	1	0.988095
COMPLEX-SLC10A4/KCNJ10/OPRM1	0.994012	1	0.988095
SLC10A4 AND NOT-SEMA4D AND NOT-HTR1F	0.994012	1	0.988095
SLC10A4 AND NOT-DTNA AND NOT-GLRA3	0.994012	1	0.988095
SLC10A4 AND NOT-KCNJ10 AND NOT-RXFP3	0.994012	1	0.988095
SLC10A4 AND NOT-NKAIN2 AND NOT-ATP1B4	0.994012	1	0.988095
SLC10A4 AND NOT-KCNJ10 AND SLC38A1	0.994012	1	0.988095
SLC10A4 AND NOT-KCNJ10 AND NOT-CASR	0.994012	1	0.988095
SLC10A4 AND NOT-KCNJ10 AND NOT-CACNG6	0.994012	1	0.988095
SLC10A4 AND NOT-KCNJ10 AND SLC38A1	0.994012	1	0.988095
COMPLEX-SLC10A4/KCNJ10/SLC6A6	0.994012	1	0.988095
SLC10A4 AND NOT-MEGF10 AND NOT-SLC9B1	0.994012	1	0.988095
SLC10A4 AND NOT-TTYH2 AND NOT-GLRA3	0.994012	1	0.988095

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
NOT-SLC16A7 AND TMEM97 AND NOT-MUC1	1	1	1
ITM2C AND CDH3 AND NOT-GPC3	1	1	1
ATP10B AND AMIGO2 AND NOT-GPC3	0.909091	0.833333	1
NOT-ZP2 AND PRRG1 AND MUC1	1	1	1
NOT-SLC7A2 AND EDNRA AND ERBB2	0.909091	0.833333	1
COMPLEX-ERBB2/ATP10B/FLVCR2	0.909091	0.833333	1
HEPH AND NOT-SLC22A23 AND MUC1	0.909091	0.833333	1
SLC26A3 AND NOT-FLVCR2 AND ERBB2	0.909091	0.833333	1
IFI6 AND NOX1 AND NOT-CNR1	1	1	1
IFI6 AND NOX1 AND NOT-ADAM15	1	1	1
IFI6 AND NOX1 AND NOT-S1PR4	1	1	1
IFI6 AND NOX1 AND NOT-IL17RC	1	1	1
IFI6 AND NOX1 AND NOT-SLITRK6	1	1	1
IFI6 AND NOX1 AND NOT-JAM3	1	1	1
IFI6 AND NOX1 AND NOT-FCRL4	1	1	1
IFI6 AND NOX1 AND NOT-KCNH6	1	1	1
IFI6 AND NOX1 AND NOT-SLC19A3	1	1	1
IFI6 AND NOX1 AND NOT-IGFLR1	1	1	1
IFI6 AND NOX1 AND NOT-TGFBR2	1	1	1
IFI6 AND NOX1 AND NOT-SYP	1	1	1
IFI6 AND NOX1 AND NOT-SLC22A1	1	1	1
IFI6 AND NOX1 AND NOT-DNER	1	1	1
IFI6 AND SLC26A3 AND NOT-KCNU1	1	1	1
IFI6 AND SLC26A3 AND NOT-GABRA1	1	1	1
IFI6 AND SLC26A3 AND NOT-ADAM2	1	1	1
IFI6 AND SLC26A3 AND NOT-LAMP5	1	1	1
IFI6 AND SLC26A3 AND NOT-IL17RA	1	1	1
IFI6 AND SLC26A3 AND NOT-CRB1	1	1	1
IFI6 AND SLC26A3 AND NOT-LHFPL5	1	1	1
IFI6 AND SLC26A3 AND NOT-EPHA8	1	1	1
IFI6 AND SLC26A3 AND NOT-EMP3	1	1	1
IFI6 AND SLC26A3 AND NOT-EPHA2	1	1	1
IFI6 AND SLC26A3 AND NOT-PPAPDC1A	1	1	1
IFI6 AND SLC26A3 AND NOT-DRD5	1	1	1
IFI6 AND SLC26A3 AND NOT-SLC30A8	1	1	1
IFI6 AND SLC26A3 AND NOT-PAQR7	1	1	1
IFI6 AND SLC26A3 AND NOT-NKAIN2	1	1	1
IFI6 AND NOX1 AND NOT-NRXN2	1	1	1
IFI6 AND SLC26A3 AND NOT-SLC9B1	1	1	1
IFI6 AND SLC26A3 AND NOT-MUC17	1	1	1
IFI6 AND SLC26A3 AND NOT-ANO4	1	1	1
IFI6 AND SLC26A3 AND NOT-BTN3A1	1	1	1
IFI6 AND SLC26A3 AND NOT-GPR83	1	1	1
IFI6 AND SLC26A3 AND NOT-NRG3	1	1	1
IFI6 AND SLC26A3 AND NOT-SEMA4B	1	1	1
IFI6 AND SLC26A3 AND NOT-BTN3A3	1	1	1
IFI6 AND NOX1 AND NOT-KCNE2	1	1	1
IFI6 AND NOX1 AND NOT-LRIG2	1	1	1
IFI6 AND NOX1 AND NOT-CD27	1	1	1
IFI6 AND NOX1 AND NOT-SLC20A1	1	1	1
IFI6 AND CEACAM7 AND NOT-TAS2R3	1	1	1
IFI6 AND NOX1 AND NOT-TMEM63B	1	1	1
IFI6 AND NOX1 AND NOT-PAQR5	1	1	1
IFI6 AND NOX1 AND NOT-EQTN	1	1	1
IFI6 AND NOX1 AND SLC38A2	1	1	1
IFI6 AND NOX1 AND NOT-C11orf24	1	1	1
IFI6 AND NOX1 AND NOT-ADAM22	1	1	1
IFI6 AND NOX1 AND NOT-PDGFRB	1	1	1
IFI6 AND NOX1 AND NOT-PCDH12	1	1	1
IFI6 AND NOX1 AND NOT-NUCB2	1	1	1
IFI6 AND NOX1 AND NOT-NGFR	1	1	1
IFI6 AND NOX1 AND NOT-MPZ	1	1	1
IFI6 AND NOX1 AND NOT-MC4R	1	1	1
IFI6 AND NOX1 AND NOT-LRP1	1	1	1
IFI6 AND NOX1 AND NOT-C14orf180	1	1	1
IFI6 AND ATP10B AND NOT-GPC3	1	1	1
IFI6 AND ST14 AND NOT-GPC3	1	1	1
IFI6 AND MEP1A AND NOT-GPC3	1	1	1
IFI6 AND MEP1A AND NOT-ERBB2	1	1	1
IFI6 AND SLC39A4 AND NOT-MUC1	1	1	1
IFI6 AND SLC5A1 AND NOT-GPC3	1	1	1
NOX1 AND SLC39A10 AND NOT-GPC3	1	1	1
NOX1 AND AMIGO2 AND NOT-ERBB2	1	1	1
NOX1 AND NOT-CA12 AND NOT-GPC3	1	1	1
IFI6 AND NOT-GPC3 AND SLC27A4	1	1	1
IFI6 AND NOT-GPC3 AND PCDH18	1	1	1
IFI6 AND NOT-GPC3 AND PAQR5	1	1	1

Antigen Pair Logic	F1	Precision	Recall
SLC10A4 AND NOT-HCN2 AND NOT-GLRA3	0.994012	1	0.988095
SLC10A4 AND NOT-NKAIN2 AND NOT-GPR6	0.994012	1	0.988095
COMPLEX-SLC10A4/GRM3/KCNJ10	0.994012	1	0.988095
SLC10A4 AND NOT-NKAIN2 AND NOT-RXFP3	0.994012	1	0.988095
SLC10A4 AND NOT-SSTR1 AND NOT-TYR	0.994012	1	0.988095
SLC10A4 AND NOT-SSTR1 AND NOT-EGFR	0.994012	1	0.988095
COMPLEX-SLC10A4/ERBB4/CLDN11	0.994012	1	0.988095
COMPLEX-SLC10A4/CLDN11/CLDN1	0.994012	1	0.988095
SLC10A4 AND NOT-SSTR1 AND NOT-EGFR	0.994012	1	0.988095
COMPLEX-SLC10A4/CLDN11/SLC7A5	0.988095	0.988095	0.988095
SLC10A4 AND NOT-SSTR1 AND NOT-MSLN	0.988095	0.988095	0.988095
COMPLEX-SLC10A4/CLDN11/SLC7A5	0.988095	0.988095	0.988095
SLC10A4 AND NOT-SSTR1 AND NOT-MSLN	0.988095	0.988095	0.988095
SLC10A4 AND NOT-SSTR1 AND NOT-DNAJB8	0.988095	0.988095	0.988095
SLC10A4 AND NOT-CLDN11 AND NOT-EGFR	0.987952	1	0.97619
SLC10A4 AND NOT-CLDN11 AND NOT-MSLN	0.987952	1	0.97619
SLC10A4 AND CD276 AND NOT-CLDN18	0.987952	1	0.97619
SLC10A4 AND NOT-CLDN11 AND NOT-EGFR	0.987952	1	0.97619
SLC10A4 AND CD276 AND NOT-DNAJB8	0.987952	1	0.97619
SLC10A4 AND PTK7 AND NOT-TYR	0.987952	1	0.97619
SLC10A4 AND NOT-CLDN11 AND NOT-DNAJB8	0.987952	1	0.97619
SLC10A4 AND NOT-CLDN11 AND NOT-TYR	0.987952	1	0.97619
SLC10A4 AND CD276 AND NOT-TYR	0.987952	1	0.97619
SLC10A4 AND CD276 AND NOT-CLDN18	0.987952	1	0.97619
COMPLEX-SLC10A4/EPCAM/CLDN11	0.987952	1	0.97619
SLC10A4 AND CD276 AND NOT-ITGB3	0.987952	1	0.97619
SLC10A4 AND NOT-CLDN11 AND NOT-MSLN	0.987952	1	0.97619
SLC10A4 AND PTK7 AND NOT-ITGB3	0.987952	1	0.97619
SLC10A4 AND PTK7 AND NOT-CLDN18	0.987952	1	0.97619
SLC10A4 AND PTK7 AND NOT-ITGB3	0.987952	1	0.97619
SLC10A4 AND PTK7 AND NOT-CLDN18	0.987952	1	0.97619
COMPLEX-SLC10A4/LGR5/CD22	0.982249	0.976471	0.988095
COMPLEX-EGFR/SLC10A4/CD37	0.982249	0.976471	0.988095
COMPLEX-SLC10A4/CLDN11/PROM1	0.982249	0.976471	0.988095
COMPLEX-SLC10A4/CLDN11/CLDN8	0.982249	0.976471	0.988095
COMPLEX-CLDN7/SLC10A4/CLDN11	0.982249	0.976471	0.988095
COMPLEX-EGFR/SLC10A4/MUC1	0.982249	0.976471	0.988095
COMPLEX-EGFR/SLC10A4/SSTR2	0.982249	0.976471	0.988095
COMPLEX-EGFR/SLC10A4/ITGB3	0.982249	0.976471	0.988095
COMPLEX-SLC10A4/CLDN11/CD22	0.982249	0.976471	0.988095
COMPLEX-CLDN7/SLC10A4/LGR5	0.982249	0.976471	0.988095
COMPLEX-SLC10A4/LGR5/CD22	0.982249	0.976471	0.988095
COMPLEX-EGFR/SLC10A4/CD33	0.982249	0.976471	0.988095
COMPLEX-SLC10A4/CLDN11/LGR5	0.982249	0.976471	0.988095
COMPLEX-EGFR/SLC10A4/SLAMF7	0.982249	0.976471	0.988095
COMPLEX-SLC10A4/CLDN11/CD22	0.982249	0.976471	0.988095
COMPLEX-SLC10A4/ERBB3/CLDN11	0.982249	0.976471	0.988095
COMPLEX-EGFR/SLC10A4/ITGB3	0.982249	0.976471	0.988095
COMPLEX-EGFR/SLC10A4/CD37	0.982249	0.976471	0.988095
COMPLEX-MSLN/CLDN7/SLC10A4	0.982249	0.976471	0.988095
COMPLEX-EGFR/SLC10A4/CD33	0.982249	0.976471	0.988095
COMPLEX-EGFR/SLC10A4/ITGB6	0.982249	0.976471	0.988095
COMPLEX-EGFR/SLC10A4/ITGB6	0.982249	0.976471	0.988095
SLC10A4 AND PTK7 AND NOT-ABCB5	0.982036	0.987952	0.97619
SLC10A4 AND NOT-SSTR1 AND NOT-STEAP2	0.982036	0.987952	0.97619
SLC10A4 AND NOT-SSTR1 AND NOT-ENPP3	0.982036	0.987952	0.97619
SLC10A4 AND NOT-NCAM1 AND NOT-TYR	0.982036	0.987952	0.97619
SLC10A4 AND NOT-NCAM1 AND NOT-EGFR	0.982036	0.987952	0.97619
SLC10A4 AND CD276 AND NOT-ABCB5	0.982036	0.987952	0.97619
SLC10A4 AND NOT-SSTR1 AND NOT-PCYT1A	0.982036	0.987952	0.97619
SLC10A4 AND NOT-NCAM1 AND NOT-EGFR	0.982036	0.987952	0.97619
SLC10A4 AND NOT-SSTR1 AND NOT-HHLA2	0.982036	0.987952	0.97619
SLC10A4 AND CD276 AND NOT-ABCB5	0.982036	0.987952	0.97619
SLC10A4 AND NOT-SSTR1 AND NOT-IL20RA	0.982036	0.987952	0.97619
SLC10A4 AND PTK7 AND NOT-ULBP3	0.982036	0.987952	0.97619
SLC10A4 AND NOT-SSTR1 AND NOT-RNF43	0.982036	0.987952	0.97619
SLC10A4 AND NOT-SSTR1 AND NOT-STEAP2	0.982036	0.987952	0.97619
SLC10A4 AND CD276 AND NOT-MST1R	0.982036	0.987952	0.97619
SLC10A4 AND CD276 AND NOT-MST1R	0.982036	0.987952	0.97619
ASTN1 AND NOT-TRPC3 AND NOT-MLC1	0.939024	0.9625	0.916667
GPR19 AND NOT-DSC1 AND NOT-KCNA1	0.927711	0.939024	0.916667
ASTN1 AND NOT-MLC1 AND NOT-ENPP1	0.912281	0.896552	0.928571
ASTN1 AND SLC38A1 AND NOT-MLC1	0.911392	0.972973	0.857143
ASTN1 AND NOT-LRRC8E AND NOT-CSMD3	0.911243	0.905882	0.916667
ASTN1 AND NOT-NKAIN2 AND NOT-EMP3	0.910256	0.986111	0.845238
ASTN1 AND NOT-SCN10A AND NOT-MLC1	0.909091	0.869565	0.952381
ASTN1 AND NOT-MLC1 AND SLC4A8	0.904762	0.904762	0.904762

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
IFI6 AND SLC39A4 AND NOT-GPC3	1	1	1
IFI6 AND MEP1A AND NOT-GPC3	1	1	1
IFI6 AND NOT-GPC3 AND MCAM	1	1	1
IFI6 AND NOT-GPC3 AND TM4SF1	1	1	1
IFI6 AND NOT-LIFR AND MUC1	1	1	1
IFI6 AND NOT-ITGAL AND MUC1	1	1	1
IFI6 AND SLC5A1 AND NOT-ERBB2	1	1	1
IFI6 AND NOT-CNR1 AND MUC1	1	1	1
NOX1 AND NOT-CEACAM1 AND NOT-GPC3	1	1	1
TGFBI AND NOT-MAL AND MUC1	1	1	1
IFI6 AND ATP10B AND NOT-MSLN	1	1	1
CDH17 AND SLC39A10 AND NOT-MUC1	1	1	1
IFI6 AND NOT-GPC3 AND SGCB	1	1	1
IFI6 AND NOT-GPC3 AND JAG1	1	1	1
NOX1 AND NOT-FLVCR2 AND NOT-MUC1	1	1	1
MEP1A AND TGFBI AND NOT-MUC1	1	1	1
IFI6 AND SLC39A4 AND NOT-ERBB2	1	1	1
IFI6 AND SLC39A4 AND NOT-GPC3	1	1	1
IFI6 AND MEP1A AND NOT-MSLN	1	1	1
IFI6 AND CFTR AND NOT-ERBB2	1	1	1
CDH17 AND SLC39A10 AND NOT-ERBB2	1	1	1
CDH17 AND SLC39A10 AND NOT-GPC3	1	1	1
IFI6 AND SLC26A3 AND NOT-GPC3	1	1	1
TGFBI AND NOT-ABCA8 AND MUC1	1	1	1
IFI6 AND ATP10B AND NOT-MUC1	1	1	1
IFI6 AND NOT-GPC3 AND PDGFRB	1	1	1
IFI6 AND SLC5A1 AND NOT-GPC3	1	1	1
CDH17 AND TACSTD2 AND NOT-GPC3	1	1	1
IFI6 AND NOT-GPC3 AND SGCB	1	1	1
IFI6 AND NOT-TUSC3 AND MUC1	1	1	1
IFI6 AND NOT-GPC3 AND SLCO2A1	1	1	1
IFI6 AND NOT-LIFR AND ERBB2	1	1	1
IFI6 AND NOT-SLC7A2 AND ERBB2	1	1	1
IFI6 AND TNFRSF11A AND NOT-GPC3	1	1	1
IFI6 AND JPH1 AND NOT-GPC3	1	1	1
IFI6 AND NOT-GPC3 AND PARM1	1	1	1
IFI6 AND NOT-GPC3 AND TSPAN15	1	1	1
IFI6 AND NOT-NAALAD2 AND MUC1	1	1	1
IFI6 AND SLC26A3 AND NOT-ERBB2	1	1	1
CDH17 AND TACSTD2 AND NOT-GPC3	1	1	1
IFI6 AND NOT-GPC3 AND EDNRA	1	1	1
IFI6 AND NOT-GPC3 AND PARM1	1	1	1
IFI6 AND NOT-NAALAD2 AND ERBB2	1	1	1
IFI6 AND NOT-GPC3 AND B3GNT3	1	1	1
IFI6 AND CFTR AND NOT-ERBB2	1	1	1
IFI6 AND SLC39A4 AND NOT-ERBB2	1	1	1
IFI6 AND TNFRSF11A AND NOT-GPC3	1	1	1
CDH17 AND NOT-FLVCR2 AND NOT-MUC1	1	1	1
IFI6 AND NOT-GPC3 AND EPHB3	1	1	1
IFI6 AND SLC39A4 AND NOT-MSLN	1	1	1
IFI6 AND SLC26A3 AND NOT-MUC1	1	1	1
IFI6 AND NOT-GPC3 AND EDNRA	1	1	1
CDH17 AND SLC39A10 AND NOT-MSLN	1	1	1
IFI6 AND NOT-NAALAD2 AND ERBB2	1	1	1
MEP1A AND TGFBI AND NOT-MUC1	1	1	1
IFI6 AND NOT-GPC3 AND SLC52A3	1	1	1
IFI6 AND NOT-GPC3 AND FZD7	1	1	1
CDH17 AND NOT-CEACAM1 AND NOT-GPC3	1	1	1
IFI6 AND NOT-TRPC6 AND MUC1	1	1	1
IFI6 AND NOT-LIFR AND MUC1	1	1	1
IFI6 AND NOT-GPC3 AND HAS2	1	1	1
TGFBI AND NOT-LIFR AND MUC1	1	1	1
IFI6 AND NOT-GPC3 AND JAG1	1	1	1
IFI6 AND NOT-GPC3 AND GPRC5A	1	1	1
IFI6 AND NOT-GPC3 AND B3GNT3	1	1	1
IFI6 AND NOT-SLC7A2 AND ERBB2	1	1	1
IFI6 AND NOT-GPC3 AND EFNB1	1	1	1
IFI6 AND NOT-NAALAD2 AND MUC1	1	1	1
IFI6 AND SLC35G1 AND NOT-GPC3	1	1	1
IFI6 AND NOT-GPC3 AND C1orf210	1	1	1
IFI6 AND NOT-LIFR AND ERBB2	1	1	1
IFI6 AND PTPRH AND NOT-GPC3	1	1	1
IFI6 AND NOT-GPC3 AND SLC27A4	1	1	1
IFI6 AND SLC39A4 AND NOT-MUC1	1	1	1
IFI6 AND NOT-GPC3 AND BMP2	1	1	1
IFI6 AND NOT-TRPC6 AND MUC1	1	1	1
IFI6 AND NOT-GPC3 AND CDH3	1	1	1

Antigen Pair Logic	F1	Precision	Recall
ASTN1 AND NOT-MLC1 AND NOT-SLC13A5	0.902857	0.868132	0.940476
ASTN1 AND NOT-LRRC8E AND NOT-OMG	0.902439	0.925	0.880952
ASTN1 AND NOT-MLC1 AND NOT-ATP13A5	0.901734	0.876404	0.928571
GPR19 AND ASTN1 AND NOT-GABRD	0.901734	0.876404	0.928571
GPR19 AND ASTN1 AND NOT-MLC1	0.901734	0.876404	0.928571
ASTN1 AND NOT-GABRB2 AND NOT-MLC1	0.901235	0.935897	0.869048
ASTN1 AND NOT-GABRB2 AND NOT-MOG	0.901235	0.935897	0.869048
COMPLEX-KCNA1/KCNQ2/LRRTM4	0.901099	0.836735	0.97619
ASTN1 AND NOT-CSMD3 AND NOT-EMP3	0.9	0.947368	0.857143
COMPLEX-KCNQ2/SLC12A5/LRRTM4	0.898876	0.851064	0.952381
COMPLEX-GRM3/KCNQ2/LRRTM4	0.898876	0.851064	0.952381
ASTN1 AND NOT-NKAIN2 AND NOT-LRRC8E	0.898734	0.959459	0.845238
ASTN1 AND FXYD6 AND NOT-MLC1	0.898204	0.903614	0.892857
COMPLEX-KCNA2/KCNQ2/LRRTM4	0.896552	0.866667	0.928571
COMPLEX-GABRB2/KCNQ2/LRRTM4	0.896175	0.828283	0.97619
ASTN1 AND NOT-LRRC8E AND ATP7A	0.894118	0.883721	0.904762
ASTN1 AND NOT-NKAIN2 AND NOT-SLC28A1	0.893082	0.946667	0.845238
ASTN1 AND NOT-NKAIN2 AND NOT-GRIN2B	0.893082	0.946667	0.845238
ASTN1 AND NOT-NKAIN2 AND NOT-OR8B2	0.893082	0.946667	0.845238
ASTN1 AND NOT-NKAIN2 AND NOT-LCT	0.893082	0.946667	0.845238
ASTN1 AND NOT-NKAIN2 AND NOT-ADAM30	0.893082	0.946667	0.845238
COMPLEX-MOG/ASTN1/NKAIN1	0.892857	0.892857	0.892857
COMPLEX-SLC9B1/ASTN1/NKAIN1	0.898204	0.903614	0.892857
COMPLEX-KCNQ2/ATP2B2/LRRTM4	0.891304	0.82	0.97619
COMPLEX-MLC1/KCNQ2/LRRTM4	0.891304	0.82	0.97619
ASTN1 AND NOT-KCNA2 AND NOT-EMP3	0.890323	0.971831	0.821429
SCN2A AND NOT-MLC1 AND ASTN1	0.890244	0.9125	0.869048
ASTN1 AND NOT-MLC1 AND NOT-SCARA5	0.888889	0.923077	0.857143
ASTN1 AND NOT-MLC1 AND NOT-SLC22A11	0.88764	0.840426	0.940476
ASTN1 AND NOT-MLC1 AND NOT-TSPAN16	0.887574	0.882353	0.892857
ASTN1 AND NOT-SLCO1C1 AND NOT-EMP3	0.886364	0.847826	0.928571
GPR19 AND ASTN1 AND NOT-SLC39A12	0.886364	0.847826	0.928571
COMPLEX-CSMD3/KCNQ2/LRRTM4	0.886364	0.847826	0.928571
COMPLEX-NKAIN2/ASTN1/PHLDB2	0.893082	0.946667	0.845238
COMPLEX-KCNA2/SYT4/CACNA1B	0.886364	0.847826	0.928571
SCN2A AND NOT-KCNA1 AND NOT-DSC1	0.886228	0.891566	0.880952
ASTN1 AND NOT-MLC1 AND NOT-ATP4B	0.884848	0.901235	0.869048
COMPLEX-KCNA1/SYT4/CACNA1B	0.883978	0.824742	0.952381
COMPLEX-GABRB2/SYT4/CACNA1B	0.883978	0.824742	0.952381
ASTN1 AND NOT-MLC1 AND TMEFF2	0.883721	0.863636	0.904762
ASTN1 AND NOT-MLC1 AND NOT-CACNG1	0.883721	0.863636	0.904762
ASTN1 AND NOT-MLC1 AND NOT-GPRC6A	0.887574	0.882353	0.892857
COMPLEX-ATP2B2/SYT4/CACNA1B	0.882682	0.831579	0.940476
NKAIN1 AND NOT-KCNA7 AND NOT-SLC22A5	0.880952	0.880952	0.880952
ASTN1 AND NOT-MLC1 AND NOT-CD1B	0.880503	0.933333	0.833333
ASTN1 AND NOT-MLC1 AND NOT-VN1R2	0.913295	0.88764	0.940476
ASTN1 AND NOT-MLC1 AND NOT-ROS1	0.888889	0.833333	0.952381
ASTN1 AND NOT-MLC1 AND NOT-ROS1	0.888889	0.833333	0.952381
ASTN1 AND NOT-ADAM30 AND NOT-CSMD3	0.88	0.846154	0.916667
ASTN1 AND NOT-GABRB2 AND NOT-OPALIN	0.879518	0.890244	0.869048
ASTN1 AND NOT-KCNA2 AND NOT-MLC1	0.878981	0.945205	0.821429
ASTN1 AND NOT-MLC1 AND NOT-ADAM7	0.883978	0.824742	0.952381
ASTN1 AND NOT-MLC1 AND NOT-OR1C1	0.878613	0.853933	0.904762
ASTN1 AND NOT-MLC1 AND NOT-TMPRSS11E	0.878981	0.945205	0.821429
ASTN1 AND NOT-OPALIN AND SLC4A8	0.878613	0.853933	0.904762
ASTN1 AND NOT-MLC1 AND NOT-LCT	0.896552	0.866667	0.928571
ASTN1 AND NOT-CSMD3 AND NOT-SLC28A1	0.878049	0.9	0.857143
ASTN1 AND SLC38A1 AND NOT-OPALIN	0.878049	0.9	0.857143
ASTN1 AND NOT-MLC1 AND NOT-CD207	0.881356	0.83871	0.928571
GPR85 AND NOT-KCNA1 AND NOT-DSC1	0.877419	0.957746	0.809524
COMPLEX-ASTN1/ROS1/NKAIN1	0.877193	0.862069	0.892857
COMPLEX-KCNQ2/OMG/LRRTM4	0.877193	0.862069	0.892857
COMPLEX-GRM3/SYT4/CACNA1B	0.876404	0.829787	0.928571
ASTN1 AND NOT-CACNG6 AND NOT-MLC1	0.876404	0.829787	0.928571
ASTN1 AND NOT-LRRC8E AND NOT-SLCO1C1	0.876404	0.829787	0.928571
ASTN1 AND NOT-MLC1 AND NOT-CATSPER3	0.876404	0.829787	0.928571
SLC6A15 AND NOT-KCNA1 AND DSC1	0.875817	0.971014	0.797619
NKAIN1 AND NOT-MIP AND NOT-PHLDB2	0.87574	0.870588	0.880952
ASTN1 AND NOT-MLC1 AND NOT-CALHM1	0.902439	0.925	0.880952
ASTN1 AND NOT-MLC1 AND NOT-KCNU1	0.875	0.836957	0.916667
ASTN1 AND NOT-MOG AND CHRNB4	0.875	0.921053	0.833333
ASTN1 AND NOT-MLC1 AND NOT-ADAM30	0.874317	0.808081	0.952381
ASTN1 AND NOT-MLC1 AND NOT-PCDHB1	0.874317	0.808081	0.952381
ASTN1 AND NOT-MLC1 AND NOT-ADAM30	0.874317	0.808081	0.952381
ASTN1 AND NOT-MLC1 AND NOT-OR7C1	0.88764	0.840426	0.940476
ASTN1 AND NOT-MLC1 AND NOT-ADAM29	0.873563	0.844444	0.904762
COMPLEX-KCNA2/KCNQ2/CACNA1B	0.872928	0.814433	0.940476

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
IFI6 AND NOT-GPC3 AND MMP14	1	1	1
IFI6 AND NOT-DNER AND MUC1	1	1	1
IFI6 AND NOT-GPC3 AND SMAGP	1	1	1
IFI6 AND IFITM1 AND MUC1	1	1	1
IFI6 AND ATP10B AND NOT-ERBB2	1	1	1
NOX1 AND EDNRA AND NOT-ERBB2	1	1	1
NOX1 AND EDNRA AND NOT-GPC3	1	1	1
IFI6 AND SLC35G1 AND NOT-GPC3	1	1	1
CDH17 AND CDH3 AND NOT-MUC1	1	1	1
DPEP1 AND NOT-SST AND NOT-GPC3	1	1	1
DPEP1 AND NOT-GPC3 AND SLC7A5	1	1	1
GPA33 AND THY1 AND NOT-ERBB2	1	1	1
GUCY2C AND THY1 AND NOT-GPC3	1	1	1
GUCY2C AND THY1 AND NOT-ERBB2	1	1	1
MUC13 AND SLC7A5 AND NOT-GPC3	1	1	1
MUC13 AND SLC7A5 AND NOT-ERBB2	1	1	1
GPA33 AND SLC7A5 AND NOT-ERBB2	1	1	1
DPEP1 AND NOT-GPC3 AND ITGAV	1	1	1
GPA33 AND SLC7A5 AND NOT-GPC3	1	1	1
DPEP1 AND NOT-GPC3 AND NOT-CLDN7	1	1	1
MUC13 AND THY1 AND NOT-ERBB2	1	1	1
MUC13 AND SLC7A5 AND NOT-MUC1	1	1	1
CLDN3 AND THY1 AND NOT-GPC3	1	1	1
DPEP1 AND NOT-GPC3 AND SPON2	1	1	1
DPEP1 AND NOT-GPC3 AND NOT-PCYT1A	1	1	1
DPEP1 AND NOT-GPC3 AND NOT-ENPP3	1	1	1
GUCY2C AND SLC7A5 AND NOT-ERBB2	1	1	1
GUCY2C AND SLC7A5 AND NOT-GPC3	1	1	1
DPEP1 AND NOT-GPC3 AND THY1	1	1	1
GUCY2C AND THY1 AND NOT-MUC1	1	1	1
GPA33 AND THY1 AND NOT-GPC3	1	1	1
MUC13 AND THY1 AND NOT-GPC3	1	1	1
DPEP1 AND NOT-GPC3 AND CLDN1	1	1	1
DPEP1 AND NOT-GPC3 AND KDR	1	1	1
DPEP1 AND NOT-GPC3 AND CLDN1	1	1	1
DPEP1 AND NOT-GPC3 AND STEAP2	1	1	1
DPEP1 AND NOT-GPC3 AND NOT-GPA33	1	1	1
DPEP1 AND NOT-GPC3 AND NOT-TNFRSF13C	1	1	1
GPA33 AND SLC7A5 AND NOT-MUC1	1	1	1
DPEP1 AND NOT-GPC3 AND NOT-CD160	1	1	1
DPEP1 AND NOT-GPC3 AND SLC39A6	1	1	1
DPEP1 AND NOT-GPC3 AND STEAP1	1	1	1
DPEP1 AND BIRC5 AND NOT-MSLN	1	1	1
DPEP1 AND NOT-GPC3 AND PTK7	1	1	1
DPEP1 AND NOT-GPC3 AND THY1	1	1	1
CLDN3 AND SLC7A5 AND NOT-MUC1	1	1	1
DPEP1 AND NOT-GPC3 AND CD70	1	1	1
DPEP1 AND NOT-GPC3 AND PTK7	1	1	1
DPEP1 AND NOT-GPC3 AND IL20RA	1	1	1
DPEP1 AND NOT-GPC3 AND ANXA1	1	1	1
DPEP1 AND NOT-GPC3 AND STEAP2	1	1	1
DPEP1 AND NOT-GPC3 AND NOT-GPA33	1	1	1
DPEP1 AND NOT-GPC3 AND CD276	1	1	1
CLDN3 AND SLC7A5 AND NOT-GPC3	1	1	1
DPEP1 AND NOT-SST AND NOT-GPC3	1	1	1
GUCY2C AND SLC7A5 AND NOT-MUC1	1	1	1
DPEP1 AND NOT-GPC3 AND ITGAV	1	1	1
DPEP1 AND NOT-GPC3 AND NOT-TNFRSF13C	1	1	1
DPEP1 AND NOT-GPC3 AND NOT-CLDN7	1	1	1
DPEP1 AND NOT-GPC3 AND BIRC5	1	1	1
DPEP1 AND NOT-GPC3 AND ROR1	1	1	1
MUC13 AND THY1 AND NOT-MUC1	1	1	1
DPEP1 AND NOT-GPC3 AND BIRC5	1	1	1
DPEP1 AND NOT-GPC3 AND NOT-HHLA2	1	1	1
GPA33 AND THY1 AND NOT-MUC1	1	1	1
DPEP1 AND NOT-GPC3 AND NOT-IL13RA1	1	1	1
DPEP1 AND NOT-GPC3 AND PTK7	1	1	1
DPEP1 AND NOT-GPC3 AND ANXA1	1	1	1
DPEP1 AND NOT-GPC3 AND TNC	1	1	1
DPEP1 AND NOT-GPC3 AND SLC7A5	1	1	1
CLDN3 AND THY1 AND NOT-MUC1	1	1	1
DPEP1 AND CEACAM6 AND NOT-GPC3	1	1	1
DPEP1 AND NOT-GPC3 AND NOT-SLC34A2	1	1	1
DPEP1 AND CEACAM6 AND NOT-GPC3	1	1	1
CLDN7 AND THY1 AND NOT-GPC3	1	1	1
DPEP1 AND NOT-GPC3 AND MET	1	1	1
DPEP1 AND NOT-GPC3 AND NOT-GUCY2C	1	1	1

Antigen Pair Logic	F1	Precision	Recall
COMPLEX-KCNQ2/SCN1A/LRRTM4	0.872928	0.814433	0.940476
COMPLEX-KCNQ2/LRRTM4/OPALIN	0.87234	0.788462	0.97619
ASTN1 AND NOT-MLC1 AND NOT-HTR6	0.871508	0.821053	0.928571
COMPLEX-GABRB2/KCNQ2/CACNA1B	0.870968	0.794118	0.964286
COMPLEX-KCNA1/KCNQ2/CACNA1B	0.870968	0.794118	0.964286
COMPLEX-OMG/SYT4/CACNA1B	0.870056	0.827957	0.916667
ASTN1 AND NOT-MC2R AND NOT-MLC1	0.870056	0.827957	0.916667
COMPLEX-KCNQ2/SLCO1C1/LRRTM4	0.869565	0.8	0.952381
COMPLEX-KCNQ2/ATP2B2/CACNA1B	0.869565	0.8	0.952381
ASTN1 AND NOT-MLC1 AND CNTNAP2	0.869048	0.869048	0.869048
ASTN1 AND ATP7A AND NOT-EMP3	0.868571	0.835165	0.904762
NOT-EGFR AND PTK7 AND NOT-ITGB3	0.946108	0.951807	0.940476
NOT-EGFR AND PTK7 AND NOT-CD33	0.934911	0.929412	0.940476
NOT-EGFR AND CD276 AND NOT-CD33	0.91954	0.888889	0.952381
NOT-EGFR AND PTK7 AND CD37	0.825806	0.901408	0.761905
NOT-EGFR AND HSPA5 AND NOT-CD33	0.816754	0.728972	0.928571
NOT-EGFR AND PTK7 AND CD79B	0.807947	0.910448	0.72619
COMPLEX-SLC26A9/SLC10A4/B4GALNT1	0.982249	0.976471	0.988095
COMPLEX-SLC26A9/SLC10A4/B4GALNT1	0.982249	0.976471	0.988095
COMPLEX-SLC10A4/OPN5/B4GALNT1	0.982249	0.976471	0.988095
COMPLEX-SLC10A4/OPN5/B4GALNT1	0.982249	0.976471	0.988095
COMPLEX-SLC10A4/B4GALNT1/PCDHAC1	0.982249	0.976471	0.988095
COMPLEX-SLC10A4/B4GALNT1/ACE2	0.982249	0.976471	0.988095
COMPLEX-SLC10A4/B4GALNT1/CATSPERG	0.982249	0.976471	0.988095
COMPLEX-SLC10A4/B4GALNT1/CATSPERG	0.982249	0.976471	0.988095
COMPLEX-SLC10A4/B4GALNT1/ACE2	0.982249	0.976471	0.988095
COMPLEX-SLC10A4/B4GALNT1/PCDHAC1	0.982249	0.976471	0.988095
COMPLEX-SLC10A4/B4GALNT1/GPR143	0.988095	0.988095	0.988095
COMPLEX-SLC10A4/B4GALNT1/GPR143	0.988095	0.988095	0.988095
COMPLEX-SLC10A4/B4GALNT1/S1PR5	0.976471	0.965116	0.988095
COMPLEX-SLC10A4/B4GALNT1/CLCA2	0.976471	0.965116	0.988095
COMPLEX-SLC10A4/B4GALNT1/CDH26	0.976471	0.965116	0.988095
COMPLEX-CRHR2/SLC10A4/B4GALNT1	0.976471	0.965116	0.988095
COMPLEX-SLC10A4/B4GALNT1/GJB5	0.976471	0.965116	0.988095
COMPLEX-SLC10A4/B4GALNT1/GJB5	0.976471	0.965116	0.988095
COMPLEX-RRH/SLC10A4/B4GALNT1	0.976471	0.965116	0.988095
COMPLEX-SLC10A4/B4GALNT1/CLCA2	0.976471	0.965116	0.988095
COMPLEX-CRHR2/SLC10A4/B4GALNT1	0.976471	0.965116	0.988095
COMPLEX-RRH/SLC10A4/B4GALNT1	0.976471	0.965116	0.988095
COMPLEX-SLC10A4/B4GALNT1/CDH26	0.976471	0.965116	0.988095
COMPLEX-SLC10A4/B4GALNT1/S1PR5	0.976471	0.965116	0.988095
COMPLEX-SLC10A4/GABRA6/B4GALNT1	0.97619	0.97619	0.97619
COMPLEX-SLC10A4/GABRA6/B4GALNT1	0.97619	0.97619	0.97619
COMPLEX-OR5I1/SLC10A4/B4GALNT1	0.982249	0.976471	0.988095
COMPLEX-SLC10A4/B4GALNT1/XKRX	0.97619	0.97619	0.97619
COMPLEX-SLC10A4/B4GALNT1/TAS2R39	0.976471	0.965116	0.988095
COMPLEX-SLC10A4/B4GALNT1/GPR101	0.982249	0.976471	0.988095
COMPLEX-SLC10A4/B4GALNT1/TAS2R39	0.976471	0.965116	0.988095
COMPLEX-SLC10A4/B4GALNT1/GRPR	0.976471	0.965116	0.988095
COMPLEX-SLC10A4/B4GALNT1/MAS1	0.982249	0.976471	0.988095
COMPLEX-SLC10A4/B4GALNT1/GRPR	0.976471	0.965116	0.988095
COMPLEX-SLC10A4/B4GALNT1/OR51B2	0.982249	0.976471	0.988095
COMPLEX-OR5I1/SLC10A4/B4GALNT1	0.982249	0.976471	0.988095
COMPLEX-SLC10A4/B4GALNT1/HRH4	0.982249	0.976471	0.988095
COMPLEX-CNR2/SLC10A4/B4GALNT1	0.97619	0.97619	0.97619
COMPLEX-SLC10A4/B4GALNT1/XKRX	0.97619	0.97619	0.97619
COMPLEX-SLC10A4/B4GALNT1/GPR101	0.982249	0.976471	0.988095
COMPLEX-SLC10A4/B4GALNT1/OR51B2	0.982249	0.976471	0.988095
COMPLEX-SLC10A4/B4GALNT1/MAS1	0.982249	0.976471	0.988095
COMPLEX-SLC10A4/B4GALNT1/HRH4	0.982249	0.976471	0.988095
COMPLEX-CNR2/SLC10A4/B4GALNT1	0.97619	0.97619	0.97619
COMPLEX-SLC10A4/B4GALNT1/ERVFRD-1	0.982249	0.976471	0.988095
COMPLEX-SLC10A4/B4GALNT1/TAS2R41	0.982249	0.976471	0.988095
COMPLEX-SLC10A4/B4GALNT1/DCHS2	0.988095	0.988095	0.988095
COMPLEX-SLC10A4/B4GALNT1/HTR1B	0.982249	0.976471	0.988095
COMPLEX-SLC10A4/B4GALNT1/DCHS2	0.988095	0.988095	0.988095
COMPLEX-SLC10A4/B4GALNT1/OR7A17	0.976471	0.965116	0.988095
COMPLEX-SLC10A4/B4GALNT1/OR10A3	0.976471	0.965116	0.988095
COMPLEX-SLC10A4/B4GALNT1/HTR1B	0.982249	0.976471	0.988095
COMPLEX-SLC10A4/B4GALNT1/OR7A17	0.976471	0.965116	0.988095
COMPLEX-SLC10A4/B4GALNT1/TAS2R41	0.982249	0.976471	0.988095
COMPLEX-SLC10A4/B4GALNT1/DUOX1	0.97076	0.954023	0.988095
COMPLEX-SLC10A4/B4GALNT1/MEP1B	0.97076	0.954023	0.988095
COMPLEX-SLC10A4/B4GALNT1/IL12RB2	0.97076	0.954023	0.988095
COMPLEX-SLC10A4/B4GALNT1/HFE	0.97076	0.954023	0.988095
COMPLEX-SLC10A4/B4GALNT1/MEP1B	0.97076	0.954023	0.988095
COMPLEX-SLC10A4/B4GALNT1/CNGB3	0.97076	0.954023	0.988095

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
GPA33 AND SLC7A5 AND NOT-MSLN	1	1	1
DPEP1 AND NOT-TNFRSF13C AND ERBB2	1	1	1
MUC13 AND SLC7A5 AND NOT-MSLN	1	1	1
BIRC5 AND NOT-IL11RA AND MUC1	1	1	1
CLDN7 AND THY1 AND NOT-ERBB2	1	1	1
BIRC5 AND NOT-GPC3 AND CLDN1	1	1	1
DPEP1 AND NOT-MAGEA11 AND MUC1	1	1	1
BIRC5 AND NOT-GPC3 AND CLDN1	1	1	1
DPEP1 AND NOT-CLDN18 AND MUC1	1	1	1
DPEP1 AND NOT-MAGEA1 AND MUC1	1	1	1
BIRC5 AND NOT-IL11RA AND ERBB2	1	1	1
DPEP1 AND ITGAV AND ERBB2	0.909091	0.833333	1
DPEP1 AND SLC7A5 AND NOT-MUC1	0.909091	0.833333	1
DPEP1 AND ITGAV AND MUC1	0.909091	0.833333	1
DPEP1 AND ITGAV AND MUC1	0.909091	0.833333	1
GUCY2C AND ITGAV AND ERBB2	1	1	1
GUCY2C AND ITGAV AND NOT-MUC1	0.888889	1	0.8
CEACAM6 AND ITGAV AND NOT-MUC1	0.888889	1	0.8
MUC13 AND ITGAV AND NOT-MUC1	0.888889	1	0.8
DPEP1 AND NOT-SST AND BIRC5	1	1	1
GPA33 AND SLC7A5 AND NOT-ABCA5	1	1	1
BIRC5 AND CLDN3 AND NOT-CLDN23	1	1	1
GPA33 AND SLC7A5 AND NOT-GAGE1	1	1	1
GPA33 AND SLC7A5 AND NOT-IGF1R	1	1	1
GPA33 AND SLC7A5 AND NOT-IL13RA1	1	1	1
GPA33 AND SLC7A5 AND NOT-ITGB3	1	1	1
DPEP1 AND NOT-IGF1R AND PTK7	1	1	1
GPA33 AND SLC7A5 AND NOT-CD180	1	1	1
BIRC5 AND MUC13 AND NOT-CLDN23	1	1	1
GPA33 AND SLC7A5 AND NOT-MUC4	1	1	1
GPA33 AND SLC7A5 AND NOT-SEMA5B	1	1	1
GPA33 AND SLC7A5 AND NOT-TRPM4	1	1	1
GPA33 AND SLC7A5 AND NOT-TNFRSF17	1	1	1
GPA33 AND SLC7A5 AND NOT-SSTR4	1	1	1
GPA33 AND SLC7A5 AND NOT-ULBP3	1	1	1
MUC13 AND SLC7A5 AND NOT-SST	1	1	1
DPEP1 AND NOT-MAGEA11 AND MET	1	1	1
GPA33 AND SLC7A5 AND NOT-EDNRB	1	1	1
DPEP1 AND SLC7A5 AND NOT-KDR	1	1	1
DPEP1 AND ITGAV AND NOT-CLDN9	1	1	1
DPEP1 AND CEACAM6 AND SLC7A5	1	1	1
BIRC5 AND CLDN3 AND NOT-CLDN7	1	1	1
MUC13 AND SLC7A5 AND NOT-TNFRSF10A	1	1	1
MUC13 AND SLC7A5 AND NOT-CLDN6	1	1	1
DPEP1 AND NOT-SST AND NOT-TYR	1	1	1
MUC13 AND SLC7A5 AND NOT-CD22	1	1	1
BIRC5 AND GPA33 AND NOT-CLDN7	1	1	1
MUC13 AND SLC7A5 AND NOT-CD79B	1	1	1
DPEP1 AND NOT-SST AND NOT-MAGEA11	1	1	1
GPA33 AND SLC7A5 AND NOT-TNFRSF13C	1	1	1
GPA33 AND SLC7A5 AND NOT-CLDN7	1	1	1
BIRC5 AND GUCY2C AND NOT-CLDN7	1	1	1
GPA33 AND SLC7A5 AND NOT-DDX3X	1	1	1
BIRC5 AND MUC13 AND NOT-CLDN7	1	1	1
GPA33 AND SLC7A5 AND NOT-FCRL5	1	1	1
GPA33 AND SLC7A5 AND NOT-CLDN9	1	1	1
GPA33 AND SLC7A5 AND NOT-CD33	1	1	1
GPA33 AND SLC7A5 AND NOT-CD38	1	1	1
MUC13 AND CLDN1 AND ITGAV	1	1	1
DPEP1 AND ITGAV AND CLDN4	1	1	1
DPEP1 AND NOT-SST AND ANXA1	1	1	1
GUCY2C AND SLC7A5 AND NOT-CD37	1	1	1
DPEP1 AND CEACAM6 AND ITGAV	1	1	1
MUC13 AND CLDN1 AND NOT-CLDN3	1	1	1
MUC13 AND CLDN1 AND NOT-MAGEA11	1	1	1
DPEP1 AND ITGAV AND NOT-WT1	1	1	1
MUC13 AND CLDN1 AND NOT-ENPP3	1	1	1
GUCY2C AND THY1 AND NOT-FCRL1	1	1	1
GUCY2C AND THY1 AND NOT-CLDN23	1	1	1
DPEP1 AND ITGAV AND OAS1	1	1	1
DPEP1 AND THY1 AND NOT-IGF1R	1	1	1
MUC13 AND THY1 AND ITGAV	1	1	1
GPA33 AND THY1 AND ITGAV	1	1	1
DPEP1 AND CEACAM5 AND ANXA1	1	1	1
GPA33 AND SLC7A5 AND NOT-CD79A	1	1	1
CLDN3 AND THY1 AND NOT-CSPG4	1	1	1
CLDN3 AND THY1 AND NOT-ENG	1	1	1

Antigen Pair Logic	F1	Precision	Recall
COMPLEX-SLC10A4/B4GALNT1/KCNJ4	0.97076	0.954023	0.988095
COMPLEX-SLC10A4/B4GALNT1/KCNIP3	0.97076	0.954023	0.988095
COMPLEX-SLC10A4/B4GALNT1/CLEC12B	0.97076	0.954023	0.988095
COMPLEX-SLC10A4/B4GALNT1/CACNA1I	0.97076	0.954023	0.988095
COMPLEX-SLC10A4/B4GALNT1/CNGB3	0.97076	0.954023	0.988095
COMPLEX-SLC10A4/B4GALNT1/GPR32	0.97076	0.954023	0.988095
COMPLEX-SLC10A4/B4GALNT1/DUOX1	0.97076	0.954023	0.988095
COMPLEX-SLC10A4/B4GALNT1/KCNIP3	0.97076	0.954023	0.988095
COMPLEX-SLC10A4/B4GALNT1/IL12RB2	0.97076	0.954023	0.988095
COMPLEX-SLC10A4/B4GALNT1/CACNA1I	0.97076	0.954023	0.988095
COMPLEX-SLC10A4/B4GALNT1/KCNJ4	0.97076	0.954023	0.988095
COMPLEX-SLC10A4/B4GALNT1/HFE	0.97076	0.954023	0.988095
COMPLEX-SLC10A4/B4GALNT1/GLP1R	0.970414	0.964706	0.97619
COMPLEX-SLC10A4/B4GALNT1/GLP1R	0.970414	0.964706	0.97619
COMPLEX-SLC10A4/MS4A2/B4GALNT1	0.970414	0.964706	0.97619
COMPLEX-SLC10A4/MS4A2/B4GALNT1	0.970414	0.964706	0.97619
COMPLEX-SLC10A4/B4GALNT1/DUOXA1	0.97006	0.975904	0.964286
COMPLEX-SLC10A4/B4GALNT1/OR4D1	0.97006	0.975904	0.964286
COMPLEX-SLC10A4/B4GALNT1/OR51I1	0.97006	0.975904	0.964286
COMPLEX-SLC10A4/B4GALNT1/OR1J4	0.97006	0.975904	0.964286
COMPLEX-SLC10A4/B4GALNT1/SLC14A2	0.97006	0.975904	0.964286
COMPLEX-SLC10A4/B4GALNT1/FASLG	0.97006	0.975904	0.964286
COMPLEX-SLC10A4/B4GALNT1/KCNJ1	0.97006	0.975904	0.964286
COMPLEX-SLC10A4/B4GALNT1/SLC14A2	0.97006	0.975904	0.964286
COMPLEX-SLC10A4/B4GALNT1/DUOXA1	0.97006	0.975904	0.964286
COMPLEX-SLC10A4/B4GALNT1/HTR1F	0.994012	1	0.988095
COMPLEX-SLC10A4/B4GALNT1/HTR1F	0.994012	1	0.988095
COMPLEX-MRGPRX2/SLC10A4/B4GALNT1	0.976471	0.965116	0.988095
COMPLEX-SLC9B1/SLC10A4/B4GALNT1	0.976471	0.965116	0.988095
COMPLEX-SLC10A4/B4GALNT1/ADAM7	0.976471	0.965116	0.988095
COMPLEX-SLC9B1/SLC10A4/B4GALNT1	0.976471	0.965116	0.988095
COMPLEX-SLC10A4/B4GALNT1/GNRHR	0.976471	0.965116	0.988095
COMPLEX-SLC10A4/B4GALNT1/GNRHR	0.976471	0.965116	0.988095
COMPLEX-SLC10A4/B4GALNT1/ADAM7	0.976471	0.965116	0.988095
COMPLEX-MRGPRX2/SLC10A4/B4GALNT1	0.976471	0.965116	0.988095
COMPLEX-KCNU1/SLC10A4/B4GALNT1	0.97619	0.97619	0.97619
COMPLEX-SLC10A4/B4GALNT1/CACNG6	0.97619	0.97619	0.97619
COMPLEX-SLC10A4/GABRA1/B4GALNT1	0.97619	0.97619	0.97619
COMPLEX-SLC10A4/GABRA1/B4GALNT1	0.97619	0.97619	0.97619
COMPLEX-SLC10A4/GPRC6A/B4GALNT1	0.97619	0.97619	0.97619
COMPLEX-KCNU1/SLC10A4/B4GALNT1	0.97619	0.97619	0.97619
COMPLEX-SLC10A4/B4GALNT1/OR8B2	0.976471	0.965116	0.988095
COMPLEX-OR5P3/SLC10A4/B4GALNT1	0.982249	0.976471	0.988095
COMPLEX-SLC10A4/B4GALNT1/GRIN2B	0.988095	0.988095	0.988095
COMPLEX-OR5P3/SLC10A4/B4GALNT1	0.982249	0.976471	0.988095
COMPLEX-TMPRSS11B/SLC10A4/B4GALNT1	0.982249	0.976471	0.988095
COMPLEX-SLC10A4/B4GALNT1/GRM6	0.982249	0.976471	0.988095
COMPLEX-TMPRSS11B/SLC10A4/B4GALNT1	0.982249	0.976471	0.988095
COMPLEX-SLC10A4/B4GALNT1/LCT	0.976471	0.965116	0.988095
COMPLEX-SLC10A4/B4GALNT1/LCT	0.976471	0.965116	0.988095
COMPLEX-SLC10A4/B4GALNT1/OR8B2	0.976471	0.965116	0.988095
COMPLEX-SLC10A4/B4GALNT1/GRIN2B	0.988095	0.988095	0.988095
COMPLEX-SLC10A4/B4GALNT1/GLRA3	0.994012	1	0.988095
COMPLEX-SLC10A4/B4GALNT1/GLRA3	0.994012	1	0.988095
COMPLEX-SLC10A4/B4GALNT1/SCN10A	0.97076	0.954023	0.988095
COMPLEX-SLC10A4/CALHM1/B4GALNT1	0.97076	0.954023	0.988095
COMPLEX-ADAM30/SLC10A4/B4GALNT1	0.97076	0.954023	0.988095
COMPLEX-SLC10A4/GABRA5/B4GALNT1	0.97076	0.954023	0.988095
COMPLEX-SLC10A4/CALHM1/B4GALNT1	0.97076	0.954023	0.988095
COMPLEX-ADAM30/SLC10A4/B4GALNT1	0.97076	0.954023	0.988095
COMPLEX-SLC10A4/B4GALNT1/LRRC8E	0.97076	0.954023	0.988095
COMPLEX-SLC10A4/B4GALNT1/PCDHB1	0.97076	0.954023	0.988095
COMPLEX-SLC10A4/B4GALNT1/SCN1A	0.97076	0.954023	0.988095
COMPLEX-SLC10A4/B4GALNT1/SCN1A	0.97076	0.954023	0.988095
COMPLEX-SLC10A4/B4GALNT1/LRRC8E	0.97076	0.954023	0.988095
COMPLEX-SLC10A4/GABRA5/B4GALNT1	0.97076	0.954023	0.988095
COMPLEX-SLC10A4/B4GALNT1/SCN10A	0.97076	0.954023	0.988095
COMPLEX-SLC10A4/B4GALNT1/CD207	0.970414	0.964706	0.97619
COMPLEX-SLC10A4/B4GALNT1/SLC22A16	0.970414	0.964706	0.97619
COMPLEX-SLC10A4/B4GALNT1/SLC22A16	0.970414	0.964706	0.97619
COMPLEX-CLDN16/SLC10A4/B4GALNT1	0.97006	0.975904	0.964286
COMPLEX-SLC10A4/B4GALNT1/PCDH11X	0.97006	0.975904	0.964286
COMPLEX-HTR3C/SLC10A4/B4GALNT1	0.97006	0.975904	0.964286
COMPLEX-HTR3C/SLC10A4/B4GALNT1	0.97006	0.975904	0.964286
COMPLEX-SLC10A4/B4GALNT1/PCDH11X	0.97006	0.975904	0.964286
COMPLEX-CLDN16/SLC10A4/B4GALNT1	0.97006	0.975904	0.964286
COMPLEX-SLC10A4/B4GALNT1/SLC28A1	0.965116	0.943182	0.988095

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
CLDN3 AND THY1 AND NOT-MAGEA4	1	1	1
DPEP1 AND CEACAM6 AND THY1	1	1	1
CLDN3 AND THY1 AND NOT-CD72	1	1	1
DPEP1 AND BIRC5 AND NOT-TNFRSF8	1	1	1
DPEP1 AND NOT-ERBB4 AND PTK7	1	1	1
DPEP1 AND ANXA1 AND NOT-DLL3	1	1	1
GUCY2C AND SLC7A5 AND NOT-CSPG4	1	1	1
DPEP1 AND NOT-ERBB4 AND ANXA1	1	1	1
BIRC5 AND GPA33 AND ITGAV	1	1	1
GUCY2C AND SLC7A5 AND NOT-MAGEA4	1	1	1
BIRC5 AND GPA33 AND NOT-MST1R	1	1	1
DPEP1 AND SLC7A5 AND NOT-MAGEA4	1	1	1
MUC13 AND SLC7A5 AND NOT-FCRL2	1	1	1
GUCY2C AND THY1 AND NOT-ERBB4	1	1	1
DPEP1 AND ITGAV AND GUCY2C	1	1	1
CLDN3 AND SLC7A5 AND NOT-FCRL1	1	1	1
DPEP1 AND CEACAM6 AND NOT-CLDN7	1	1	1
BIRC5 AND MUC13 AND NOT-CLDN23	1	1	1
DPEP1 AND CEACAM6 AND SLC7A5	1	1	1
DPEP1 AND ITGAV AND OAS1	1	1	1
CLDN3 AND SLC7A5 AND NOT-ERBB4	1	1	1
GUCY2C AND SLC7A5 AND NOT-CLDN3	1	1	1
DPEP1 AND THY1 AND NOT-ERBB4	1	1	1
CLDN3 AND SLC7A5 AND NOT-ANXA1	1	1	1
DPEP1 AND NOT-MAGEA1 AND THY1	1	1	1
CLDN3 AND SLC7A5 AND NOT-HLA-DOB	1	1	1
DPEP1 AND NOT-CLDN18 AND THY1	1	1	1
BIRC5 AND EPCAM AND ITGAV	1	1	1
DPEP1 AND STEAP2 AND NOT-ULBP3	1	1	1
CLDN3 AND SLC7A5 AND NOT-PMEL	1	1	1
GPA33 AND SLC7A5 AND NOT-CLDN3	1	1	1
DPEP1 AND CEACAM6 AND BIRC5	1	1	1
MUC13 AND SLC7A5 AND NOT-SSX1	1	1	1
CLDN7 AND SLC7A5 AND NOT-PCYT1A	1	1	1
DPEP1 AND NOT-TNFRSF13C AND CD52	1	1	1
BIRC5 AND EPCAM AND NOT-CLDN7	1	1	1
DPEP1 AND ITGAV AND BIRC5	1	1	1
IFI6 AND NOX1 AND NOT-OTOF	1	1	1
IFI6 AND CEACAM7 AND NOT-SLC22A16	1	1	1
IFI6 AND CEACAM7 AND NOT-SLC12A3	1	1	1
IFI6 AND MUC12 AND NOT-GRIA4	1	1	1
IFI6 AND FUT3 AND NOT-AOC3	1	1	1
IFI6 AND MUC12 AND NOT-CHRND	1	1	1
IFI6 AND MUC12 AND NOT-GPR50	1	1	1
IFI6 AND CEACAM7 AND NOT-TMPRSS11D	1	1	1
IFI6 AND MUC12 AND NOT-RXFP3	1	1	1
IFI6 AND MUC12 AND NOT-CNGA4	1	1	1
IFI6 AND CEACAM7 AND NOT-GRIN1	1	1	1
IFI6 AND VANGL1 AND NOT-ADAM7	1	1	1
IFI6 AND MUC12 AND NOT-CRB2	1	1	1
IFI6 AND MUC12 AND NOT-SLC12A5	1	1	1
IFI6 AND FUT3 AND NOT-SLC12A5	1	1	1
IFI6 AND MUC12 AND NOT-SLC4A5	1	1	1
FUT3 AND NOT-CHRND AND CDH11	1	1	1
IFI6 AND MUC12 AND NOT-CACNG2	1	1	1
IFI6 AND VANGL1 AND NOT-ATP6V0A4	1	1	1
IFI6 AND CEACAM7 AND NOT-IFNAR2	1	1	1
IFI6 AND NOX1 AND NOT-IFNAR2	1	1	1
IFI6 AND MUC12 AND NOT-CRB1	1	1	1
IFI6 AND VANGL1 AND NOT-HTR1F	1	1	1
IFI6 AND MUC12 AND NOT-MLANA	1	1	1
IFI6 AND CEACAM7 AND NOT-KCNK9	1	1	1
IFI6 AND MUC12 AND NOT-SLC10A2	1	1	1
IFI6 AND CEACAM7 AND NOT-PTGER4	1	1	1
IFI6 AND NOX1 AND NOT-NETO1	1	1	1
IFI6 AND NOX1 AND NOT-MEGF10	1	1	1
IFI6 AND NOX1 AND NOT-KIRREL3	1	1	1
IFI6 AND NOX1 AND NOT-SLC22A16	1	1	1
IFI6 AND NOX1 AND NOT-ADAM21	1	1	1
IFI6 AND NOX1 AND NOT-CCKBR	1	1	1
IFI6 AND NOX1 AND NOT-CD1B	1	1	1
IFI6 AND NOX1 AND NOT-GUCY2F	1	1	1
IFI6 AND CEACAM7 AND NOT-SLC5A8	1	1	1
IFI6 AND CEACAM7 AND NOT-PIRT	1	1	1
IFI6 AND CEACAM7 AND NOT-GJA3	1	1	1
IFI6 AND MUC12 AND NOT-MLNR	1	1	1
NOX1 AND NOT-CD1B AND CDH11	1	1	1

Antigen Pair Logic	F1	Precision	Recall
COMPLEX-SLC10A4/B4GALNT1/ROS1	0.965116	0.943182	0.988095
COMPLEX-SLC10A4/B4GALNT1/CACNG8	0.965116	0.943182	0.988095
COMPLEX-CNGA4/SLC10A4/B4GALNT1	0.965116	0.943182	0.988095
COMPLEX-SLC10A4/B4GALNT1/HTR6	0.965116	0.943182	0.988095
COMPLEX-CNGA4/SLC10A4/B4GALNT1	0.965116	0.943182	0.988095
COMPLEX-SLC10A4/B4GALNT1/HTR6	0.965116	0.943182	0.988095
COMPLEX-SLC10A4/B4GALNT1/TRPM3	0.965116	0.943182	0.988095
COMPLEX-SLC10A4/B4GALNT1/ROS1	0.965116	0.943182	0.988095
COMPLEX-SLC10A4/B4GALNT1/SLC28A1	0.965116	0.943182	0.988095
COMPLEX-SLC10A4/B4GALNT1/GPR26	0.965116	0.943182	0.988095
COMPLEX-SLC10A4/B4GALNT1/AJAP1	0.964706	0.953488	0.97619
COMPLEX-SLC10A4/B4GALNT1/SLC1A6	0.964706	0.953488	0.97619
COMPLEX-SLC10A4/B4GALNT1/GJA10	0.964706	0.953488	0.97619
COMPLEX-SLC10A4/B4GALNT1/GJA10	0.964706	0.953488	0.97619
COMPLEX-SLC10A4/B4GALNT1/CLRN1	0.964286	0.964286	0.964286
COMPLEX-SLC10A4/B4GALNT1/DPP10	0.964286	0.964286	0.964286
COMPLEX-SLC10A4/B4GALNT1/CLRN1	0.964286	0.964286	0.964286
COMPLEX-SLC10A4/B4GALNT1/GRIN2C	0.963855	0.97561	0.952381
COMPLEX-SLC10A4/B4GALNT1/SLC12A3	0.959538	0.932584	0.988095
COMPLEX-SLC10A4/B4GALNT1/OR2L2	0.959538	0.932584	0.988095
COMPLEX-SLC10A4/B4GALNT1/SLC12A3	0.959538	0.932584	0.988095
COMPLEX-SLC10A4/B4GALNT1/OR2L2	0.959538	0.932584	0.988095
COMPLEX-SLC10A4/B4GALNT1/CLDN17	0.959064	0.942529	0.97619
COMPLEX-SLC10A4/B4GALNT1/CLDN17	0.959064	0.942529	0.97619
COMPLEX-SLC10A4/MLANA/B4GALNT1	0.95858	0.952941	0.964286
COMPLEX-SLC10A4/MLANA/B4GALNT1	0.95858	0.952941	0.964286
COMPLEX-SLC10A4/CATSPERD/B4GALNT1	0.958084	0.963855	0.952381
COMPLEX-SLC10A4/NFASC/B4GALNT1	0.958084	0.963855	0.952381
COMPLEX-SLC10A4/B4GALNT1/CRB2	0.954023	0.922222	0.988095
COMPLEX-SLC10A4/B4GALNT1/GPR6	0.954023	0.922222	0.988095
COMPLEX-SLC22A12/SLC10A4/B4GALNT1	0.954023	0.922222	0.988095
COMPLEX-SLC10A4/B4GALNT1/CACNA1S	0.954023	0.922222	0.988095
COMPLEX-SLC10A4/B4GALNT1/RXFP3	0.954023	0.922222	0.988095
COMPLEX-SLC10A4/B4GALNT1/GPR6	0.954023	0.922222	0.988095
COMPLEX-SLC22A12/SLC10A4/B4GALNT1	0.954023	0.922222	0.988095
COMPLEX-SLC10A4/B4GALNT1/OR2L13	0.954023	0.922222	0.988095
COMPLEX-SLC10A4/B4GALNT1/ATP7A	0.954023	0.922222	0.988095
COMPLEX-S1PR1/SLC10A4/B4GALNT1	0.954023	0.922222	0.988095
COMPLEX-SLC10A4/B4GALNT1/TYR	0.976471	0.965116	0.988095
COMPLEX-SLC10A4/B4GALNT1/ITGB3	0.976471	0.965116	0.988095
COMPLEX-SLC10A4/B4GALNT1/ITGB3	0.976471	0.965116	0.988095
COMPLEX-SLC10A4/B4GALNT1/ABCB5	0.97076	0.954023	0.988095
COMPLEX-SLC10A4/B4GALNT1/ABCB5	0.97076	0.954023	0.988095
COMPLEX-SLC10A4/B4GALNT1/CLDN18	0.97076	0.954023	0.988095
COMPLEX-SLC10A4/B4GALNT1/CLDN18	0.97076	0.954023	0.988095
COMPLEX-SLC10A4/B4GALNT1/ITGB6	0.97006	0.975904	0.964286
COMPLEX-SLC10A4/B4GALNT1/ITGB6	0.97006	0.975904	0.964286
COMPLEX-DNAJB8/SLC10A4/B4GALNT1	0.959538	0.932584	0.988095
COMPLEX-SLC10A4/B4GALNT1/PROM1	0.959538	0.932584	0.988095
COMPLEX-SLC10A4/B4GALNT1/IL20RA	0.95858	0.952941	0.964286
COMPLEX-SLC10A4/B4GALNT1/ULBP3	0.954023	0.922222	0.988095
COMPLEX-HHLA2/SLC10A4/B4GALNT1	0.953488	0.931818	0.97619
COMPLEX-SLC10A4/B4GALNT1/MST1R	0.948571	0.912088	0.988095
COMPLEX-SLC10A4/B4GALNT1/MST1R	0.948571	0.912088	0.988095
COMPLEX-SLC10A4/ERBB4/B4GALNT1	0.947368	0.931034	0.964286
COMPLEX-SLC10A4/ERBB4/B4GALNT1	0.947368	0.931034	0.964286
COMPLEX-SLC10A4/B4GALNT1/CD22	0.942529	0.911111	0.97619
COMPLEX-SLC10A4/B4GALNT1/RNF43	0.942529	0.911111	0.97619
COMPLEX-SLC10A4/B4GALNT1/RNF43	0.942529	0.911111	0.97619
COMPLEX-SLC10A4/B4GALNT1/ENPP3	0.942529	0.911111	0.97619
COMPLEX-SLC10A4/B4GALNT1/CD22	0.942529	0.911111	0.97619
COMPLEX-SLC10A4/B4GALNT1/ENPP3	0.942529	0.911111	0.97619
COMPLEX-SLC10A4/ENG/B4GALNT1	0.937143	0.901099	0.97619
COMPLEX-SLC10A4/B4GALNT1/ULBP1	0.936416	0.910112	0.964286
COMPLEX-SLC10A4/FOLR1/B4GALNT1	0.932515	0.962025	0.904762
COMPLEX-TNFRSF13C/SLC10A4/B4GALNT1	0.929412	0.918605	0.940476
COMPLEX-SLC10A4/B4GALNT1/CLDN6	0.928571	0.928571	0.928571
COMPLEX-SLC10A4/B4GALNT1/HSPA5	0.926554	0.88172	0.97619
COMPLEX-SLC10A4/B4GALNT1/CD34	0.924855	0.898876	0.952381
COMPLEX-SLC10A4/B4GALNT1/IL13RA1	0.922222	0.864583	0.988095
COMPLEX-SLC10A4/B4GALNT1/IL13RA1	0.922222	0.864583	0.988095
COMPLEX-SLC10A4/B4GALNT1/STEAP1	0.922156	0.927711	0.916667
COMPLEX-SLC10A4/B4GALNT1/CEACAM6	0.922156	0.927711	0.916667
COMPLEX-SLC10A4/ALK/B4GALNT1	0.91954	0.888889	0.952381
COMPLEX-SLC10A4/FLOT2/B4GALNT1	0.918605	0.897727	0.940476
COMPLEX-SLC10A4/B4GALNT1/STEAP2	0.914634	0.9375	0.892857
COMPLEX-SLC10A4/B4GALNT1/STEAP2	0.914634	0.9375	0.892857

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
IFI6 AND NOX1 AND NOT-CD1B	1	1	1
NOX1 AND NOT-CD1B AND SLC6A6	1	1	1
IFI6 AND MUC12 AND NOT-CLCN1	1	1	1
IFI6 AND CEACAM7 AND NOT-GUCY2F	1	1	1
IFI6 AND VANGL1 AND NOT-SLC22A25	1	1	1
IFI6 AND MUC12 AND NOT-GPR12	1	1	1
IFI6 AND FUT3 AND NOT-GPR12	1	1	1
IFI6 AND CEACAM7 AND NOT-PAQR7	1	1	1
IFI6 AND CEACAM7 AND NOT-GABRA5	1	1	1
IFI6 AND NOX1 AND NOT-GABRA5	1	1	1
IFI6 AND MUC12 AND NOT-DGKE	1	1	1
IFI6 AND VANGL1 AND NOT-TSHR	1	1	1
IFI6 AND VANGL1 AND NOT-MOG	1	1	1
CDH17 AND TGFBI AND NOT-CHRND	1	1	1
NOX1 AND TGFBI AND NOT-CHRND	1	1	1
IFI6 AND CEACAM7 AND NOT-IL13	1	1	1
IFI6 AND NOX1 AND NOT-IL13	1	1	1
IFI6 AND VANGL1 AND NOT-GRM4	1	1	1
NOX1 AND NOT-EPHA10 AND NOT-PCDHA6	1	1	1
NOX1 AND NOT-EPHA10 AND CD44	1	1	1
IFI6 AND CEACAM7 AND NOT-CD36	1	1	1
IFI6 AND NOT-SLCO6A1 AND ATP8B1	1	1	1
IFI6 AND NOX1 AND NOT-KCNK16	1	1	1
IFI6 AND NOT-STEAP4 AND VANGL1	1	1	1
IFI6 AND VANGL1 AND NOT-CACNG2	1	1	1
IFI6 AND VANGL1 AND NOT-GABRA5	1	1	1
IFI6 AND VANGL1 AND NOT-OR3A2	1	1	1
IFI6 AND VANGL1 AND NOT-SLC6A15	1	1	1
IFI6 AND VANGL1 AND NOT-AJAP1	1	1	1
IFI6 AND VANGL1 AND NOT-SLC4A5	1	1	1
IFI6 AND VANGL1 AND NOT-SCN2A	1	1	1
IFI6 AND MUC12 AND NOT-GPR26	1	1	1
IFI6 AND CEACAM7 AND NOT-MLC1	1	1	1
NOX1 AND TGFBI AND NOT-DISP2	1	1	1
MUC12 AND TGFBI AND NOT-SLC9A1	1	1	1
IFI6 AND NOT-CD36 AND NOT-PIRT	1	1	1
IFI6 AND CEACAM7 AND NOT-KCNK16	1	1	1
IFI6 AND MUC12 AND NOT-CDH22	1	1	1
IFI6 AND CEACAM7 AND NOT-GRIN2C	1	1	1
IFI6 AND CEACAM7 AND NOT-GPRC6A	1	1	1
NOX1 AND TGFBI AND NOT-MUC1	1	1	1
CDH17 AND TGFBI AND NOT-MUC1	1	1	1
IFI6 AND CEACAM7 AND NOT-MUC1	1	1	1
IFI6 AND MUC12 AND NOT-MUC1	1	1	1
CDH17 AND TGFBI AND NOT-MUC1	1	1	1
IFI6 AND NOT-MLC1 AND MUC1	1	1	1
IFI6 AND MUC12 AND NOT-MUC1	1	1	1
IFI6 AND MUC12 AND NOT-GPC3	1	1	1
IFI6 AND CEACAM7 AND NOT-GPC3	1	1	1
IFI6 AND MUC12 AND NOT-GPC3	1	1	1
IFI6 AND CEACAM7 AND NOT-GPC3	1	1	1
IFI6 AND CEACAM7 AND NOT-ERBB2	1	1	1
IFI6 AND NOT-KCNV2 AND MUC1	1	1	1
IFI6 AND NOT-KCNV2 AND MUC1	1	1	1
IFI6 AND NOT-GPC3 AND SLC1A5	1	1	1
IFI6 AND NOT-GPC3 AND CDH11	1	1	1
IFI6 AND NOT-GPC3 AND EFNB2	1	1	1
IFI6 AND NOT-GPC3 AND LRIG3	1	1	1
IFI6 AND NOT-GPC3 AND TNFRSF12A	1	1	1
NOX1 AND TGFBI AND NOT-MUC1	1	1	1
IFI6 AND NOX1 AND NOT-GPC3	1	1	1
IFI6 AND NOT-STEAP4 AND MUC1	1	1	1
IFI6 AND CEACAM7 AND NOT-ERBB2	1	1	1
IFI6 AND NOT-GPC3 AND ATP8B1	1	1	1
IFI6 AND MUC12 AND NOT-ERBB2	1	1	1
IFI6 AND CEACAM7 AND NOT-MSLN	1	1	1
IFI6 AND NOT-GPC3 AND VANGL1	1	1	1
CDH17 AND NOT-SLC9A1 AND ERBB2	1	1	1
IFI6 AND NOT-GPC3 AND VANGL1	1	1	1
CDH17 AND NOT-SLC9A1 AND NOT-GPC3	1	1	1
IFI6 AND NOT-GPR26 AND MUC1	1	1	1
IFI6 AND NOT-ATP8A2 AND MUC1	1	1	1
IFI6 AND NOT-SPAM1 AND MUC1	1	1	1
IFI6 AND NOT-MDGA2 AND MUC1	1	1	1
IFI6 AND NOT-MDGA2 AND MUC1	1	1	1
IFI6 AND NOT-GPR26 AND ERBB2	1	1	1
IFI6 AND NOT-GPR26 AND MUC1	1	1	1

Antigen Pair Logic	F1	Precision	Recall
COMPLEX-GPNMB/SLC10A4/B4GALNT1	0.913295	0.88764	0.940476
COMPLEX-SLC10A4/B4GALNT1/ANXA1	0.912281	0.896552	0.928571
COMPLEX-DDX3X/SLC10A4/B4GALNT1	0.907975	0.936709	0.880952
COMPLEX-SLC10A4/B4GALNT1/CLDN8	0.903614	0.914634	0.892857
COMPLEX-SLC10A4/B4GALNT1/THY1	0.902439	0.925	0.880952
COMPLEX-SPON2/SLC10A4/B4GALNT1	0.901734	0.876404	0.928571
COMPLEX-SLC10A4/B4GALNT1/IGF1R	0.900585	0.885057	0.916667
APLP1 AND CD276 AND NOT-B4GALNT1	0.900585	0.885057	0.916667
COMPLEX-SLC10A4/B4GALNT1/VCAM1	0.899408	0.894118	0.904762
COMPLEX-SLC10A4/B4GALNT1/MAGEA1	0.888889	0.873563	0.904762
COMPLEX-AFP/SLC10A4/B4GALNT1	0.882353	0.872093	0.892857
COMPLEX-CLDN23/SLC10A4/B4GALNT1	0.882353	0.872093	0.892857
COMPLEX-SLC10A4/B4GALNT1/TNFRSF8	0.878049	0.9	0.857143
COMPLEX-SLC10A4/B4GALNT1/SDC1	0.873418	0.932432	0.821429
COMPLEX-SLC10A4/B4GALNT1/ULBP2	0.872727	0.888889	0.857143
COMPLEX-SLC10A4/B4GALNT1/IL11RA	0.871166	0.898734	0.845238
COMPLEX-SLC10A4/B4GALNT1/IL11RA	0.871166	0.898734	0.845238
COMPLEX-SLC10A4/B4GALNT1/ROR1	0.86747	0.878049	0.857143
COMPLEX-SLC10A4/B4GALNT1/OAS1	0.866242	0.931507	0.809524
COMPLEX-SLC10A4/B4GALNT1/GPC3	0.865854	0.8875	0.845238
COMPLEX-SLC10A4/B4GALNT1/IL13RA2	0.862275	0.86747	0.857143
COMPLEX-SLC10A4/SLC39A6/B4GALNT1	0.860606	0.876543	0.845238
COMPLEX-SLC10A4/SLC39A6/B4GALNT1	0.860606	0.876543	0.845238
COMPLEX-SLC10A4/B4GALNT1/CD37	0.855422	0.865854	0.845238
COMPLEX-SLC10A4/B4GALNT1/GUCY2C	0.855422	0.865854	0.845238
COMPLEX-SLC10A4/EPHB2/B4GALNT1	0.848485	0.864198	0.833333
COMPLEX-CLDN3/SLC10A4/B4GALNT1	0.846626	0.873418	0.821429
COMPLEX-SLC10A4/B4GALNT1/PMEL	0.841463	0.8625	0.821429
COMPLEX-SLC10A4/B4GALNT1/SSTR5	0.841463	0.8625	0.821429
COMPLEX-SLC10A4/B4GALNT1/EPCAM	0.838323	0.843373	0.833333
COMPLEX-DLL3/SLC10A4/B4GALNT1	0.834356	0.860759	0.809524
COMPLEX-SLC10A4/B4GALNT1/TNFRSF10A	0.829268	0.85	0.809524
COMPLEX-SLC10A4/ABCA5/B4GALNT1	0.82716	0.858974	0.797619
COMPLEX-SLC10A4/B4GALNT1/CD180	0.82716	0.858974	0.797619
COMPLEX-SLC10A4/B4GALNT1/SSTR2	0.822785	0.878378	0.77381
SCN3B AND NOT-B4GALNT1 AND CD276	0.820513	0.888889	0.761905
COMPLEX-SLC10A4/B4GALNT1/IL2RA	0.819876	0.857143	0.785714
COMPLEX-SLC10A4/EPHA3/B4GALNT1	0.817073	0.8375	0.797619
COMPLEX-SLC10A4/ERBB2/B4GALNT1	0.815287	0.876712	0.761905
COMPLEX-SLC10A4/B4GALNT1/CD38	0.8125	0.855263	0.77381
COMPLEX-SLC10A4/B4GALNT1/CD38	0.8125	0.855263	0.77381
COMPLEX-MLC1/B4GALNT1/ASTN1	0.820513	0.720721	0.952381
Oligodendroglioma			
GRIA2 AND BMPR1A AND NOT-PTPRM	0.965517	1	0.933333
GRIA2 AND BMPR1A AND NOT-FLRT2	0.965517	1	0.933333
GRIA2 AND ANTXR1 AND NOT-RNF144A	0.928571	1	0.866667
GRIA2 AND BMPR1A AND NOT-RNF144A	0.965517	1	0.933333
GRIA2 AND ANTXR1 AND NOT-SLMAP	0.928571	1	0.866667
GRIA2 AND ANTXR1 AND NOT-ADIPOR1	0.928571	1	0.866667
GRIA2 AND ANTXR1 AND NOT-PTPRM	0.928571	1	0.866667
GRIA2 AND BMPR1A AND NOT-FAT4	0.965517	1	0.933333
GRIA2 AND ANTXR1 AND NOT-IL4R	0.928571	1	0.866667
NRCAM AND TSPAN11 AND NOT-SCAMP5	0.928571	1	0.866667
GRIA2 AND BMPR1A AND NOT-KIT	0.965517	1	0.933333
LRRTM2 AND BMPR1A AND NOT-PTPRM	0.903226	0.875	0.933333
LRRTM2 AND BMPR1A AND NOT-RNF144A	0.903226	0.875	0.933333
GRIA2 AND ANTXR1 AND NOT-ATRN	0.928571	1	0.866667
GRIA2 AND ANTXR1 AND NOT-KCNS3	0.928571	1	0.866667
GRIA2 AND ANTXR1 AND NOT-CNNM2	0.928571	1	0.866667
GRIA2 AND ANTXR1 AND NOT-TRPV2	0.928571	1	0.866667
GRIA2 AND ANTXR1 AND NOT-TAAR5	0.928571	1	0.866667
GRIA2 AND ANTXR1 AND NOT-SLC2A8	0.928571	1	0.866667
GRIA2 AND ANTXR1 AND NOT-EVI2B	0.928571	1	0.866667
GRIA2 AND ANTXR1 AND NOT-SCN4B	0.928571	1	0.866667
GRIA2 AND ANTXR1 AND NOT-FAT4	0.928571	1	0.866667
GRIA2 AND ANTXR1 AND NOT-BTN2A1	0.928571	1	0.866667
GRIA2 AND ANTXR1 AND NOT-TNFRSF14	0.928571	1	0.866667
GRIA2 AND ANTXR1 AND NOT-SLC7A2	0.928571	1	0.866667
GRIA2 AND ANTXR1 AND NOT-FCGR3B	0.928571	1	0.866667
GRIA2 AND ANTXR1 AND NOT-VAPA	0.928571	1	0.866667
GRIA2 AND ANTXR1 AND NOT-ABCB1	0.928571	1	0.866667
GRIA2 AND ANTXR1 AND NOT-FLRT2	0.928571	1	0.866667
GRIA2 AND ANTXR1 AND NOT-RHCG	0.928571	1	0.866667
GRIA2 AND ANTXR1 AND NOT-LGR4	0.928571	1	0.866667
GRIA2 AND ANTXR1 AND NOT-SLC17A5	0.928571	1	0.866667
GRIA2 AND ANTXR1 AND NOT-FZD6	0.928571	1	0.866667
GRIA2 AND ANTXR1 AND NOT-CLEC1A	0.928571	1	0.866667

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
IFI6 AND NOT-ATP8A2 AND MUC1	1	1	1
IFI6 AND MUC12 AND NOT-ERBB2	1	1	1
NOX1 AND NOT-SLC9A1 AND ERBB2	1	1	1
NOX1 AND NOT-SLC9A1 AND NOT-GPC3	1	1	1
IFI6 AND NOT-SPAM1 AND MUC1	1	1	1
IFI6 AND NOT-GPC3 AND NOT-SLC6A6	1	1	1
IFI6 AND NOT-OR2L2 AND MUC1	1	1	1
IFI6 AND NOT-GPC3 AND NOT-IFNAR2	1	1	1
IFI6 AND NOT-STEAP4 AND ERBB2	1	1	1
IFI6 AND NOT-OR3A2 AND MUC1	1	1	1
IFI6 AND MUC1 AND NOT-SLC4A5	1	1	1
IFI6 AND NOT-OR3A2 AND MUC1	1	1	1
IFI6 AND NOT-TSHR AND MUC1	1	1	1
IFI6 AND NOT-GABRA5 AND MUC1	1	1	1
IFI6 AND NOT-SLCO6A1 AND MUC1	1	1	1
IFI6 AND NOT-DSC1 AND MUC1	1	1	1
IFI6 AND NOT-SLC9B1 AND MUC1	1	1	1
IFI6 AND NOT-GPC3 AND NOT-SEMA4D	1	1	1
IFI6 AND NOT-GPC3 AND EPHA2	1	1	1
IFI6 AND NOT-GPC3 AND PROCR	1	1	1
IFI6 AND MUC1 AND NOT-SLC4A5	1	1	1
IFI6 AND NOT-GPC3 AND NOT-IFNAR2	1	1	1
IFI6 AND MUC1 AND NOT-AGTR2	1	1	1
IFI6 AND NOT-DTNA AND MUC1	1	1	1
IFI6 AND NOT-GPC3 AND NOT-SLC23A2	1	1	1
IFI6 AND NOT-STEAP4 AND ERBB2	1	1	1
IFI6 AND NOT-GPC3 AND NOT-STEAP4	1	1	1
IFI6 AND NOT-MOG AND MUC1	1	1	1
IFI6 AND NOT-DSC1 AND MUC1	1	1	1
IFI6 AND NOT-OR2L2 AND MUC1	1	1	1
IFI6 AND NOT-SCN1A AND MUC1	1	1	1
IFI6 AND NOT-TSHR AND MUC1	1	1	1
IFI6 AND NOT-SLC9B1 AND MUC1	1	1	1
IFI6 AND NOT-GABRA5 AND MUC1	1	1	1
IFI6 AND NOT-GPC3 AND NOT-SEMA4D	1	1	1
IFI6 AND NOT-DTNA AND MUC1	1	1	1
IFI6 AND NOT-SCN1A AND MUC1	1	1	1
IFI6 AND MUC1 AND NOT-SLC13A5	1	1	1
IFI6 AND MUC1 AND NOT-SLC13A5	1	1	1
IFI6 AND NOT-GPC3 AND NOT-STEAP4	1	1	1
IFI6 AND NOT-MOG AND MUC1	1	1	1
IFI6 AND NOT-GPC3 AND MRGPRF	1	1	1
IFI6 AND NOT-SLC1A2 AND MUC1	1	1	1
IFI6 AND NOT-HTR2C AND MUC1	1	1	1
IFI6 AND VANGL1 AND NOT-MUC1	0.909091	0.833333	1
IFI6 AND NOT-GPC3 AND CLDN19	0.909091	0.833333	1
IFI6 AND NOT-GPC3 AND NOT-P2RX3	0.909091	0.833333	1
IFI6 AND NOT-SLC13A1 AND MUC1	1	1	1
IFI6 AND NOT-PIRT AND MUC1	1	1	1
IFI6 AND NOT-GPC3 AND NOT-DGKE	0.909091	0.833333	1
IFI6 AND NOT-GPC3 AND NOT-MLC1	1	1	1
IFI6 AND NOT-GPC3 AND NOT-GRM4	0.909091	0.833333	1
IFI6 AND NOT-GPC3 AND NOT-KCNV2	0.909091	0.833333	1
IFI6 AND NOT-GPC3 AND NOT-BTN3A3	1	1	1
IFI6 AND NOT-GPC3 AND NOT-ANTXR2	1	1	1
IFI6 AND NOT-SLC1A2 AND NOT-GPC3	0.909091	0.833333	1
IFI6 AND NOT-GPC3 AND NOT-GABRD	0.909091	0.833333	1
IFI6 AND NOT-GPC3 AND NOT-GRM4	0.909091	0.833333	1
IFI6 AND NOT-GPC3 AND MUC1	1	1	1
IFI6 AND NOT-GPC3 AND ERBB2	1	1	1
IFI6 AND NOT-GPC3 AND MUC1	1	1	1
IFI6 AND NOT-GPC3 AND ERBB2	1	1	1
DPEP1 AND NOT-GPC3 AND MUC1	1	1	1
ITGAV AND NOT-GPC3 AND MUC1	1	1	1
ITGAV AND MUC1 AND NOT-ERBB2	0.909091	0.833333	1
NOT-GPC3 AND WNT5A AND MUC1	1	1	1
Ependymoma			
COMPLEX-CSPG5/UNC5D/FZD3	1	1	1
FZD3 AND CD44 AND NOT-SLC9A9	1	1	1
FZD3 AND CD44 AND NOT-DLL1	1	1	1
FZD3 AND CD44 AND NOT-CDH19	1	1	1
FZD3 AND CD44 AND NOT-TMEM150B	1	1	1
FZD3 AND CD44 AND NOT-GPR17	1	1	1
FZD3 AND CD44 AND NOT-UTS2R	1	1	1
FZD3 AND CD44 AND NOT-GPR4	1	1	1
FZD3 AND CD44 AND GPM6B	1	1	1
FZD3 AND CD44 AND NOT-GPR31	1	1	1

Antigen Pair Logic	F1	Precision	Recall
GRIA2 AND ANTXR1 AND NOT-EMCN	0.928571	1	0.866667
SCAMP5 AND TSPAN11 AND NOT-BDKRB2	0.903226	0.875	0.933333
GRIA2 AND ANTXR1 AND NOT-PRIMA1	0.928571	1	0.866667
GRIA2 AND BMPR1A AND NOT-SCN4B	0.965517	1	0.933333
GRIA2 AND BMPR1A AND NOT-CNNM2	0.965517	1	0.933333
GRIA2 AND BMPR1A AND NOT-IFITM2	0.928571	1	0.866667
GRIA2 AND BMPR1A AND NOT-SLMAP	0.965517	1	0.933333
SCAMP5 AND ANTXR1 AND NOT-SLC30A5	0.896552	0.928571	0.866667
SCAMP5 AND ANTXR1 AND NOT-FLRT2	0.896552	0.928571	0.866667
SCAMP5 AND ANTXR1 AND NOT-MYOF	0.896552	0.928571	0.866667
NRCAM AND SLC22A4 AND NOT-PTPRM	0.896552	0.928571	0.866667
NCAM2 AND ANTXR1 AND NOT-TSPAN8	0.928571	1	0.866667
LRRTM2 AND TSPAN11 AND NOT-SFRP1	0.903226	0.875	0.933333
GRIA2 AND BMPR1A AND BSG	0.965517	1	0.933333
GRIA2 AND ANTXR1 AND NOT-SLCO4A1	0.928571	1	0.866667
GRIA2 AND ANTXR1 AND NOT-PANX1	0.928571	1	0.866667
SCAMP5 AND ANTXR1 AND NOT-GPR160	0.896552	0.928571	0.866667
GRIA2 AND ANTXR1 AND NOT-AGTRAP	0.928571	1	0.866667
SCAMP5 AND ANTXR1 AND NOT-LSR	0.896552	0.928571	0.866667
GRIA2 AND ANTXR1 AND NOT-CHIC2	0.928571	1	0.866667
GRIA2 AND ANTXR1 AND NOT-CD4	0.928571	1	0.866667
GRIA2 AND ANTXR1 AND NOT-ABCD4	0.928571	1	0.866667
FAIM2 AND BMPR1A AND NOT-IFITM2	0.896552	0.928571	0.866667
GRIA2 AND VANGL2 AND NOT-TLR7	0.888889	1	0.8
GRIA2 AND VANGL2 AND NOT-SLMAP	0.888889	1	0.8
GPM6A AND CALCRL AND NOT-IGSF6	0.888889	1	0.8
GRIA2 AND BMP2 AND NOT-SLMAP	0.888889	1	0.8
GRIA2 AND VANGL2 AND NOT-VAPA	0.888889	1	0.8
SCAMP5 AND VANGL2 AND NOT-EVI2B	0.888889	1	0.8
GRIA2 AND VANGL2 AND NOT-HAVCR2	0.888889	1	0.8
SCAMP5 AND VANGL2 AND NOT-SLC30A5	0.888889	1	0.8
SCAMP5 AND VANGL2 AND NOT-PKD2L1	0.888889	1	0.8
GRIA2 AND CALCRL AND NOT-SLC24A1	0.888889	1	0.8
PTPRZ1 AND SIRPG AND NOT-ACPP	0.888889	1	0.8
GRIA2 AND CALCRL AND NOT-ATP2A2	0.888889	1	0.8
GPM6A AND CALCRL AND NOT-HAVCR2	0.888889	1	0.8
GRIA2 AND VANGL2 AND NOT-SLC24A1	0.888889	1	0.8
GRIA2 AND VANGL2 AND NOT-IL1R1	0.888889	1	0.8
GRIA2 AND VANGL2 AND NOT-LSR	0.888889	1	0.8
GRIA2 AND VANGL2 AND NOT-ROM1	0.888889	1	0.8
GRIA2 AND CALCRL AND NOT-KCNS3	0.888889	1	0.8
GRIA2 AND VANGL2 AND NOT-CD1C	0.888889	1	0.8
GRIA2 AND VANGL2 AND NOT-SLC17A5	0.888889	1	0.8
GRIA2 AND VANGL2 AND NOT-CLEC4E	0.888889	1	0.8
GRIA2 AND VANGL2 AND NOT-CHIC2	0.888889	1	0.8
GRIA2 AND VANGL2 AND NOT-ABCB1	0.888889	1	0.8
GRIA2 AND BMP2 AND NOT-SLC17A5	0.888889	1	0.8
GRIA2 AND CALCRL AND NOT-KIT	0.888889	1	0.8
SCAMP5 AND VANGL2 AND NOT-AGTRAP	0.888889	1	0.8
FAIM2 AND BMP2 AND NOT-ACSL1	0.888889	1	0.8
SCAMP5 AND VANGL2 AND NOT-CSF3R	0.888889	1	0.8
GRIA2 AND BEST3 AND NOT-SLMAP	0.967742	0.9375	1
GRIA2 AND BEST3 AND NOT-SCN4B	0.967742	0.9375	1
BEST3 AND LRRTM2 AND NOT-PLA2R1	0.967742	0.9375	1
BEST3 AND LRRTM2 AND NOT-SLMAP	0.9375	0.882353	1
GRIA2 AND TNFRSF10B AND NOT-FLRT2	0.928571	1	0.866667
SCAMP5 AND BEST3 AND NOT-SCN4B	0.9375	0.882353	1
GRIA2 AND BEST3 AND NOT-KIT	0.967742	0.9375	1
GRIA2 AND BEST3 AND NOT-VAPA	0.967742	0.9375	1
CDH10 AND BMPR1A AND NOT-PTPRM	0.965517	1	0.933333
GRIA2 AND BEST3 AND NOT-RNF144A	0.967742	0.9375	1
BEST3 AND LRRTM2 AND NOT-RNF144A	0.9375	0.882353	1
BEST3 AND ATP9A AND NOT-ABCC5	0.9375	0.882353	1
BEST3 AND ATP9A AND NOT-PLA2R1	0.967742	0.9375	1
GRIA2 AND BEST3 AND NOT-TRPV1	0.967742	0.9375	1
NCAM2 AND BEST3 AND NOT-TSPAN5	0.9375	0.882353	1
NCAM2 AND BEST3 AND NOT-SLMAP	0.9375	0.882353	1
BEST3 AND LRRTM2 AND NOT-CNNM4	0.9375	0.882353	1
NCAM2 AND BEST3 AND NOT-GPR52	0.9375	0.882353	1
NCAM2 AND BEST3 AND NOT-SCN4B	0.9375	0.882353	1
BEST3 AND LRRTM2 AND NOT-SLC26A10	0.9375	0.882353	1
NCAM2 AND BEST3 AND NOT-ABCC5	0.9375	0.882353	1
BEST3 AND LRRTM2 AND NOT-SCNN1G	0.9375	0.882353	1
GRIA2 AND TNFRSF10B AND NOT-IL4R	0.928571	1	0.866667
SCAMP5 AND BEST3 AND NOT-FLRT2	0.9375	0.882353	1
GRIA2 AND BEST3 AND NOT-FLRT2	0.967742	0.9375	1
NCAM2 AND BEST3 AND NOT-SLC5A6	0.9375	0.882353	1

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
FZD3 AND NOT-GLRB AND CD44	1	1	1
FZD3 AND CD44 AND NOT-HTRA2	1	1	1
FZD3 AND CD44 AND NOT-GLP1R	1	1	1
FZD3 AND CD44 AND NOT-KCNH5	1	1	1
FZD3 AND CD44 AND NOT-CACNG5	1	1	1
FZD3 AND CD44 AND NOT-GJB1	1	1	1
FZD3 AND CD44 AND NOT-GJA5	1	1	1
FZD3 AND CD44 AND NOT-EPHA6	1	1	1
FZD3 AND CD44 AND NOT-GPR32	1	1	1
FZD3 AND CD44 AND NOT-APOB	1	1	1
FZD3 AND CD44 AND NOT-GYPE	1	1	1
FZD3 AND CD44 AND NOT-ICAM3	1	1	1
FZD3 AND CD44 AND NOT-ANO9	1	1	1
FZD3 AND CD44 AND NOT-HTR7	1	1	1
FZD3 AND CD44 AND NOT-HTR3A	1	1	1
FZD3 AND CD44 AND NOT-HTR2B	1	1	1
FZD3 AND CD44 AND NOT-APLP2	1	1	1
FZD3 AND CD44 AND NOT-SLC2A8	1	1	1
FZD3 AND NOT-SLC25A4 AND CD44	1	1	1
FZD3 AND CD44 AND NOT-ANK1	1	1	1
FZD3 AND CD44 AND NOT-GPR39	1	1	1
FZD3 AND NOT-GPR37 AND CD44	1	1	1
FZD3 AND CD44 AND NOT-GIPR	1	1	1
FZD3 AND CD44 AND NOT-OR10H3	1	1	1
FZD3 AND CD44 AND NOT-SLC16A8	1	1	1
FZD3 AND NOT-FNDC5 AND CD44	1	1	1
FZD3 AND CD44 AND NOT-FLRT1	1	1	1
FZD3 AND CD44 AND NOT-FRRS1L	1	1	1
FZD3 AND CD44 AND NOT-KCNE4	1	1	1
FZD3 AND NOT-BACE1 AND CD44	1	1	1
FZD3 AND CD44 AND NOT-CLEC5A	1	1	1
FZD3 AND CD44 AND NOT-VSIG2	1	1	1
FZD3 AND CD44 AND NOT-FPR1	1	1	1
FZD3 AND CD44 AND NOT-LPAR3	1	1	1
FZD3 AND CD44 AND NOT-TNFRSF13B	1	1	1
FZD3 AND CD44 AND NOT-PLXND1	1	1	1
FZD3 AND CD44 AND NOT-NLGN4Y	1	1	1
FZD3 AND CD44 AND NOT-FGFR2	1	1	1
FZD3 AND CD44 AND NOT-CDHR3	1	1	1
FZD3 AND CD44 AND NOT-ZDHHC20	1	1	1
FZD3 AND CD44 AND NOT-FAM26E	1	1	1
FZD3 AND CD44 AND NOT-MCOLN2	1	1	1
FZD3 AND CD44 AND NOT-OR10A3	1	1	1
NOT-SPPL2A AND BMP2 AND DNAJC5	1	1	1
SLC7A11 AND NOT-FXYD5 AND CCR7	1	1	1
SLC7A11 AND NOT-FXYD5 AND ACE	1	1	1
NOT-SLC16A7 AND OR10H2 AND TMEM50B	1	1	1
NOT-SLC16A7 AND OR10H2 AND WNT5A	1	1	1
SLC7A11 AND NOT-FXYD5 AND QSOX2	1	1	1
SLC7A11 AND NOT-FXYD5 AND CLEC4C	1	1	1
SLC7A11 AND NOT-FXYD5 AND DSC3	1	1	1
SLC7A11 AND NOT-FXYD5 AND SCAMP5	1	1	1
SLC7A11 AND NOT-FXYD5 AND EGF	1	1	1
NOT-SLC16A7 AND OR10H2 AND TSPAN6	1	1	1
SLC7A11 AND NOT-FXYD5 AND EPHB1	1	1	1
SLC7A11 AND NOT-FXYD5 AND EVI2B	1	1	1
SLC7A11 AND NOT-FXYD5 AND DAGLB	1	1	1
SLC7A11 AND NOT-FXYD5 AND SLC35G1	1	1	1
SLC7A11 AND NOT-FXYD5 AND CD300LB	1	1	1
SLC7A11 AND NOT-FXYD5 AND GJD3	1	1	1
SLC7A11 AND NOT-FXYD5 AND NOT-ACVR1C	1	1	1
SLC7A11 AND NOT-FXYD5 AND SGCZ	1	1	1
NOT-SLC16A7 AND OR10H2 AND AMN	1	1	1
NOT-SLC16A7 AND OR10H2 AND NIPA2	1	1	1
SLC7A11 AND NOT-FXYD5 AND CR1	1	1	1
SLC7A11 AND NOT-FXYD5 AND MUC15	1	1	1
SLC7A11 AND NOT-FXYD5 AND OR10A5	1	1	1
SLC7A11 AND NOT-FXYD5 AND ADRA1D	1	1	1
SLC7A11 AND NOT-FXYD5 AND C1orf210	1	1	1
SLC7A11 AND NOT-FXYD5 AND PLB1	1	1	1
SLC7A11 AND NOT-FXYD5 AND BTLA	1	1	1
NOT-SLC16A7 AND OR10H2 AND ITSN1	1	1	1
SLC7A11 AND NOT-FXYD5 AND TENM4	1	1	1
SLC7A11 AND NOT-FXYD5 AND OR1A2	1	1	1
SLC7A11 AND NOT-FXYD5 AND OR2M4	1	1	1
SLC7A11 AND NOT-FXYD5 AND NOT-YIPF3	1	1	1
SLC7A11 AND NOT-FXYD5 AND OR8B8	1	1	1

Antigen Pair Logic	F1	Precision	Recall
BEST3 AND LRRTM2 AND NOT-ESYT3	0.9375	0.882353	1
NCAM2 AND BEST3 AND NOT-IL4R	0.9375	0.882353	1
NCAM2 AND BEST3 AND NOT-NTRK3	0.9375	0.882353	1
GRIA2 AND BEST3 AND NOT-PERP	0.933333	0.933333	0.933333
BEST3 AND ATP9A AND NOT-SCN4B	0.967742	0.9375	1
ASTN1 AND CYP4F2 AND NOT-SLC2A13	0.928571	1	0.866667
BEST3 AND LRRTM2 AND NOT-F10	0.9375	0.882353	1
BEST3 AND ATP9A AND NOT-CNNM4	0.9375	0.882353	1
GRIA2 AND BEST3 AND NOT-LGR4	0.967742	0.9375	1
NCAM2 AND BEST3 AND NOT-CNNM2	0.9375	0.882353	1
BEST3 AND LRRTM2 AND NOT-KIT	0.9375	0.882353	1
SCAMP5 AND BEST3 AND NOT-PERP	0.933333	0.933333	0.933333
NCAM2 AND BEST3 AND NOT-EFNB3	0.9375	0.882353	1
BEST3 AND ATP9A AND NOT-SCNN1G	0.9375	0.882353	1
NCAM2 AND BEST3 AND NOT-SLC12A6	0.9375	0.882353	1
BEST3 AND LRRTM2 AND NOT-AMICA1	0.9375	0.882353	1
GRIA2 AND BEST3 AND NOT-BTN2A1	0.967742	0.9375	1
LSAMP AND BEST3 AND NOT-ABCC5	0.9375	0.882353	1
NCAM2 AND BEST3 AND NOT-SLC2A13	0.967742	0.9375	1
BEST3 AND TM7SF2 AND NOT-PLA2R1	0.909091	0.833333	1
GRIA2 AND TNFRSF10B AND NOT-PTPRM	0.928571	1	0.866667
FAIM2 AND BEST3 AND NOT-SCN4B	0.909091	0.833333	1
GRIA2 AND TNFRSF10B AND NOT-GPR52	0.928571	1	0.866667
GRIA2 AND TNFRSF10B AND NOT-DIO2	0.928571	1	0.866667
FAIM2 AND BEST3 AND NOT-SLC2A13	0.909091	0.833333	1
FAIM2 AND BEST3 AND NOT-CLCN5	0.909091	0.833333	1
GRIA2 AND BMPR1A AND NOT-PROCR	0.965517	1	0.933333
GRIA2 AND BEST3 AND NOT-GPR52	0.967742	0.9375	1
GRIA2 AND BEST3 AND NOT-CLCN5	0.967742	0.9375	1
GRIA2 AND BEST3 AND NOT-IFITM2	0.933333	0.933333	0.933333
SCAMP5 AND BEST3 AND NOT-GPR52	0.909091	0.833333	1
SCAMP5 AND BEST3 AND NOT-CLCN5	0.9375	0.882353	1
BEST3 AND ATP9A AND NOT-FAT2	0.9375	0.882353	1
BEST3 AND LRRTM2 AND NOT-SYT2	0.933333	0.933333	0.933333
LRRC4C AND BEST3 AND NOT-GPR52	0.9375	0.882353	1
GRIA2 AND BEST3 AND NOT-CD1C	0.967742	0.9375	1
BEST3 AND ATP9A AND NOT-NRXN3	1	1	1
NCAM2 AND BEST3 AND NOT-LRFN1	0.903226	0.875	0.933333
ASTN1 AND GJB3 AND NOT-SLC2A13	0.933333	0.933333	0.933333
SCN2A AND TSPAN11 AND NOT-SLITRK6	0.967742	0.9375	1
GRIA2 AND ANTXR1 AND NOT-BST2	0.928571	1	0.866667
GRIA2 AND ANTXR1 AND NOT-PTPRT	0.928571	1	0.866667
GRIA2 AND ANTXR1 AND NOT-CD1B	0.928571	1	0.866667
LSAMP AND BEST3 AND NOT-SLC26A10	0.9375	0.882353	1
GRIA2 AND ANTXR1 AND NOT-PROCR	0.928571	1	0.866667
LSAMP AND BEST3 AND NOT-SCN4B	0.967742	0.9375	1
COMPLEX-CRB1/GRIK5/ADAM9	0.965517	1	0.933333
GRIA2 AND TNFRSF10B AND SERINC1	0.928571	1	0.866667
SCN2A AND TSPAN11 AND NOT-ITGA8	0.909091	0.833333	1
COMPLEX-CRB1/GRIK5/ITGB8	0.965517	1	0.933333
GRIA2 AND TNFRSF10B AND NOT-TNFRSF14	0.928571	1	0.866667
GRIA2 AND TNFRSF10B AND NOT-SLMAP	0.928571	1	0.866667
GRIA2 AND TNFRSF10B AND NOT-SLC7A2	0.928571	1	0.866667
GRIA2 AND TNFRSF10B AND NOT-ABCB1	0.928571	1	0.866667
OMG AND BMPR1A AND NOT-FLRT2	0.933333	0.933333	0.933333
GRIA2 AND TNFRSF10B AND NOT-CD1C	0.928571	1	0.866667
GRIA2 AND BMPR1A AND NOT-PTPRT	0.965517	1	0.933333
SCAMP5 AND ANTXR1 AND NOT-PROCR	0.896552	0.928571	0.866667
GRIA2 AND BEST3 AND NOT-TRPV2	0.933333	0.933333	0.933333
GRIA2 AND BEST3 AND NOT-IL4R	0.967742	0.9375	1
GRIA2 AND BEST3 AND NOT-CNNM2	0.967742	0.9375	1
SCAMP5 AND BEST3 AND NOT-TAAR5	0.909091	0.833333	1
COMPLEX-ATP10A/BCAN/ATRN	0.928571	1	0.866667
COMPLEX-SCAMP5/FCGR3B/BCAN	0.928571	1	0.866667
BCAN AND NOT-KCNJ8 AND SIRPG	0.928571	1	0.866667
BCAN AND NOT-TNFSF15 AND HAVCR2	0.928571	1	0.866667
BCAN AND NOT-TRPM7 AND SIRPG	0.928571	1	0.866667
COMPLEX-SCAMP5/AGTRAP/BCAN	0.928571	1	0.866667
BCAN AND NOT-LTB4R AND SLC22A4	0.928571	1	0.866667
BCAN AND NOT-CD2 AND SLC22A4	0.928571	1	0.866667
COMPLEX-LY9/ABCB1/BCAN	0.928571	1	0.866667
COMPLEX-ABCB1/BCAN/TLR5	0.928571	1	0.866667
BCAN AND NOT-KLRB1 AND SIRPG	0.928571	1	0.866667
COMPLEX-GOLM1/ABCB1/BCAN	0.928571	1	0.866667
COMPLEX-ATP10A/BCAN/RHOT2	0.928571	1	0.866667
COMPLEX-CLEC4E/ATP10A/BCAN	0.928571	1	0.866667
BCAN AND NOT-CYP4F12 AND HAVCR2	0.928571	1	0.866667

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
SLC7A11 AND NOT-FXYD5 AND OR10A3	1	1	1
SLC7A11 AND NOT-FXYD5 AND OR4D1	1	1	1
NOT-SLC16A7 AND OR10H2 AND RYK	1	1	1
SLC7A11 AND NOT-FXYD5 AND CACNG5	1	1	1
NOT-SLC16A7 AND OR10H2 AND SLC30A5	1	1	1
SLC7A11 AND NOT-FXYD5 AND TPSG1	1	1	1
SLC7A11 AND NOT-PLXND1 AND NOT-FXYD5	1	1	1
SLC16A2 AND NOT-SLC16A7 AND OR10H2	1	1	1
SLC7A11 AND NOT-FXYD5 AND FLT3	1	1	1
NOT-SLC16A7 AND OR10H2 AND BRCA1	1	1	1
SLC7A11 AND NOT-FXYD5 AND NCSTN	1	1	1
NOT-SLC16A7 AND OR10H2 AND NOT-SLC22A1	1	1	1
NOT-SLC16A7 AND OR10H2 AND SLC20A2	1	1	1
NOT-SLC16A7 AND OR10H2 AND BMP2	1	1	1
NOT-TNFSF10 AND GPR34 AND NOT-CLDN11	1	1	1
FZD3 AND NOT-CNTNAP1 AND NOT-CD180	1	1	1
FZD3 AND NOT-CNTNAP1 AND IL13RA1	1	1	1
FZD3 AND NOT-CNTNAP1 AND ABCB5	1	1	1
FZD3 AND NOT-CNTNAP1 AND NOT-EPHB2	1	1	1
FZD3 AND NOT-CNTNAP1 AND DDX3X	1	1	1
SLC7A11 AND WNT3 AND NOT-CD70	1	1	1
SLC7A11 AND WNT3 AND NOT-FCRL5	1	1	1
SLC7A11 AND WNT3 AND NOT-ULBP2	1	1	1
SLC7A11 AND WNT3 AND NOT-TPBG	1	1	1
FZD3 AND NOT-CNTNAP1 AND NOT-MAGEA1	1	1	1
FZD3 AND NOT-SLCO4A1 AND CR2	1	1	1
FZD3 AND NOT-SLCO4A1 AND CEACAM6	1	1	1
FZD3 AND NOT-SLCO4A1 AND MUC1	1	1	1
FZD3 AND NOT-SLCO4A1 AND ITGB6	1	1	1
FZD3 AND NOT-SLCO4A1 AND ITGB3	1	1	1
FZD3 AND NOT-SLCO4A1 AND IL13RA1	1	1	1
FZD3 AND NOT-SLCO4A1 AND ABCB5	1	1	1
FZD3 AND NOT-SLCO4A1 AND CTAG2	1	1	1
FZD3 AND NOT-SLCO4A1 AND ANXA1	1	1	1
FZD3 AND NOT-SLCO4A1 AND DDX3X	1	1	1
FZD3 AND NOT-SLCO4A1 AND RAET1E	1	1	1
FZD3 AND NOT-CNTNAP1 AND NOT-SDC1	1	1	1
FZD3 AND NOT-SLCO4A1 AND FCRL1	1	1	1
FZD3 AND NOT-SLCO4A1 AND CD160	1	1	1
FZD3 AND NOT-SLCO4A1 AND SLC34A2	1	1	1
FZD3 AND NOT-SLCO4A1 AND MSLN	1	1	1
FZD3 AND NOT-CNTNAP1 AND NOT-CD72	1	1	1
FZD3 AND NOT-CNTNAP1 AND CD33	1	1	1
FZD3 AND NOT-CNTNAP1 AND VTCN1	1	1	1
FZD3 AND NOT-CNTNAP1 AND NOT-SSTR5	1	1	1
FZD3 AND NOT-CNTNAP1 AND NOT-SSTR4	1	1	1
NOT-TRPM4 AND TSPAN11 AND APCDD1	1	1	1
SLC7A11 AND WNT3 AND NOT-THY1	1	1	1
FZD3 AND NOT-SLCO4A1 AND ENPP3	1	1	1
SLC7A11 AND WNT3 AND NOT-ERBB4	1	1	1
SLC7A11 AND WNT3 AND NOT-EPHB2	1	1	1
SLC7A11 AND NOT-ENG AND WNT3	1	1	1
SLC7A11 AND WNT3 AND NOT-DNAJB8	1	1	1
SLC7A11 AND WNT3 AND NOT-HHLA2	1	1	1
SLC7A11 AND WNT3 AND NOT-SLC34A2	1	1	1
SLC7A11 AND WNT3 AND NOT-MSLN	1	1	1
SLC7A11 AND WNT3 AND NOT-GPA33	1	1	1
SLC7A11 AND NOT-NINJ2 AND TNFRSF8	1	1	1
SLC7A11 AND NOT-NINJ2 AND MUC16	1	1	1
SLC7A11 AND NOT-NINJ2 AND ULBP1	1	1	1
SLC7A11 AND NOT-NINJ2 AND VTCN1	1	1	1
SLC7A11 AND NOT-NINJ2 AND ULBP3	1	1	1
SLC7A11 AND NOT-NINJ2 AND FCRL2	1	1	1
SLC7A11 AND NOT-NINJ2 AND CA9	1	1	1
SLC7A11 AND WNT3 AND NOT-FOLH1	1	1	1
SLC7A11 AND WNT3 AND NOT-SSTR5	1	1	1
SLC7A11 AND WNT3 AND NOT-SST	1	1	1
SLC7A11 AND WNT3 AND NOT-ENPP3	1	1	1
SLC7A11 AND NOT-IL13RA1 AND WNT3	1	1	1
SLC7A11 AND WNT3 AND NOT-IL3RA	1	1	1
SLC7A11 AND WNT3 AND NOT-GPC3	1	1	1
SLC7A11 AND WNT3 AND NOT-B4GALNT1	1	1	1
FZD3 AND NOT-SLCO4A1 AND OAS1	1	1	1
FZD3 AND NOT-SLCO4A1 AND IL20RA	1	1	1
SLC7A11 AND NOT-NINJ2 AND SSTR5	1	1	1
GRPR AND NOT-LTBR AND NOT-CD22	1	1	1
COMPLEX-CSPG5/SLC32A1/FZD3	1	1	1

Antigen Pair Logic	F1	Precision	Recall
COMPLEX-CCR7/ATP10A/BCAN	0.928571	1	0.866667
COMPLEX-F10/ABCB1/BCAN	0.928571	1	0.866667
COMPLEX-COL17A1/ATP10A/BCAN	0.928571	1	0.866667
COMPLEX-ATP10A/BCAN/CD1E	0.928571	1	0.866667
COMPLEX-FLRT3/ATP10A/BCAN	0.928571	1	0.866667
COMPLEX-FLRT3/BCAN/PPAP2A	0.928571	1	0.866667
BCAN AND NOT-EFNB3 AND NOT-SLC6A1	0.928571	1	0.866667
COMPLEX-ABCB1/BCAN/PPAP2A	0.928571	1	0.866667
BCAN AND NOT-TMPRSS2 AND SIRPG	0.928571	1	0.866667
COMPLEX-PLA2R1/ATP10A/BCAN	0.928571	1	0.866667
COMPLEX-EPHB4/ATP10A/BCAN	0.928571	1	0.866667
BCAN AND NOT-F10 AND SLC22A4	0.928571	1	0.866667
COMPLEX-GOLM1/ATP10A/BCAN	0.928571	1	0.866667
COMPLEX-F10/ATP10A/BCAN	0.928571	1	0.866667
COMPLEX-ATP10A/SCNN1G/BCAN	0.928571	1	0.866667
BCAN AND NOT-FAT2 AND SLC22A4	0.928571	1	0.866667
COMPLEX-SCAMP5/ABCB1/BCAN	0.928571	1	0.866667
BCAN AND NOT-ADRB1 AND HAVCR2	0.928571	1	0.866667
BCAN AND NOT-F10 AND CD84	0.928571	1	0.866667
COMPLEX-KLRB1/ABCB1/BCAN	0.928571	1	0.866667
BCAN AND NOT-CD27 AND CLEC4A	0.928571	1	0.866667
BCAN AND NOT-LTB4R AND CD84	0.928571	1	0.866667
COMPLEX-ABCB1/BCAN/TSPAN14	0.928571	1	0.866667
COMPLEX-ABCB1/BCAN/CD40	0.928571	1	0.866667
BCAN AND NOT-PTPRJ AND MME	0.928571	1	0.866667
COMPLEX-ATP10A/BCAN/PPAP2A	0.928571	1	0.866667
BCAN AND NOT-ADCY4 AND CD84	0.928571	1	0.866667
COMPLEX-ADRB1/ATP10A/BCAN	0.928571	1	0.866667
BCAN AND NOT-ANO9 AND HAVCR2	0.928571	1	0.866667
COMPLEX-ATP10A/BCAN/ITGA8	0.928571	1	0.866667
COMPLEX-ABCB1/BCAN/CD1C	0.928571	1	0.866667
COMPLEX-LTB/ATP10A/BCAN	0.928571	1	0.866667
COMPLEX-PCDH18/ATP10A/BCAN	0.928571	1	0.866667
COMPLEX-KCNS3/ATP10A/BCAN	0.928571	1	0.866667
BCAN AND NOT-SLC18A2 AND SIRPG	0.928571	1	0.866667
BCAN AND NOT-PCDH18 AND HAVCR2	0.928571	1	0.866667
COMPLEX-ATP10A/BCAN/TSPAN14	0.928571	1	0.866667
BCAN AND NOT-NCAM2 AND CLEC1A	0.928571	1	0.866667
COMPLEX-SCAMP5/BCAN/CD4	0.928571	1	0.866667
COMPLEX-FAT2/ATP10A/BCAN	0.928571	1	0.866667
COMPLEX-VAMP5/SCAMP5/BCAN	0.928571	1	0.866667
ASTN1 AND BEST3 AND NOT-IFITM2	0.933333	0.933333	0.933333
ASTN1 AND BEST3 AND NOT-FLRT2	0.967742	0.9375	1
ASTN1 AND BEST3 AND NOT-SLC2A13	1	1	1
ASTN1 AND BEST3 AND NOT-SCN4B	0.967742	0.9375	1
ASTN1 AND BEST3 AND NOT-SLC5A6	0.967742	0.9375	1
ASTN1 AND BEST3 AND NOT-MTUS1	0.967742	0.9375	1
ASTN1 AND BEST3 AND NOT-SLC31A2	0.967742	0.9375	1
ASTN1 AND BEST3 AND NOT-TM4SF1	0.933333	0.933333	0.933333
ASTN1 AND BEST3 AND NOT-ABCC5	0.967742	0.9375	1
OMG AND BEST3 AND NOT-SCN4B	0.967742	0.9375	1
ASTN1 AND BEST3 AND NOT-LPAR1	0.967742	0.9375	1
ASTN1 AND BEST3 AND NOT-TSPAN8	0.967742	0.9375	1
ASTN1 AND BEST3 AND NOT-SLC12A6	0.967742	0.9375	1
ASTN1 AND BEST3 AND NOT-EFNB3	0.967742	0.9375	1
ASTN1 AND BEST3 AND NOT-NRXN2	0.965517	1	0.933333
ASTN1 AND BEST3 AND NOT-ABCB1	0.967742	0.9375	1
OMG AND BEST3 AND NOT-SLC26A10	0.967742	0.9375	1
ASTN1 AND BEST3 AND NOT-EVI2B	0.967742	0.9375	1
ASTN1 AND BEST3 AND SERINC1	0.967742	0.9375	1
CDH10 AND BEST3 AND NOT-SCN4B	0.9375	0.882353	1
ASTN1 AND BEST3 AND NOT-CLEC4A	0.967742	0.9375	1
ASTN1 AND BEST3 AND NOT-ECSCR	0.967742	0.9375	1
ASTN1 AND BEST3 AND NOT-HAVCR2	0.967742	0.9375	1
ASTN1 AND BEST3 AND NOT-SLC30A5	0.967742	0.9375	1
ASTN1 AND BEST3 AND NOT-TGFBR2	0.967742	0.9375	1
ASTN1 AND BEST3 AND NOT-PKD2L1	0.967742	0.9375	1
ASTN1 AND BEST3 AND NOT-MYOF	0.967742	0.9375	1
ASTN1 AND BEST3 AND NOT-HLA-G	0.967742	0.9375	1
NCAM2 AND BEST3 AND NOT-JPH3	0.9375	0.882353	1
ASTN1 AND BEST3 AND NOT-SLC16A1	0.967742	0.9375	1
ASTN1 AND BEST3 AND NOT-CD84	0.967742	0.9375	1
ASTN1 AND BEST3 AND NOT-SCARB1	0.967742	0.9375	1
ASTN1 AND BEST3 AND NOT-SLC26A2	0.967742	0.9375	1
NCAM2 AND BEST3 AND NOT-PTPRT	0.9375	0.882353	1
ASTN1 AND BEST3 AND NOT-CD4	0.967742	0.9375	1
ASTN1 AND BEST3 AND NOT-CSF3R	0.933333	0.933333	0.933333

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
GRM6 AND ZDHHC17 AND NOT-GJA10	1	1	1
GRM6 AND ZDHHC17 AND NOT-CALN1	1	1	1
CADM1 AND NOT-SLC6A6 AND CCKAR	1	1	1
GRM6 AND ZDHHC17 AND NOT-TRPM8	1	1	1
FZD3 AND NOT-KCNK9 AND SLC43A1	1	1	1
GRM6 AND ZDHHC17 AND NOT-CACNA1S	1	1	1
GRM6 AND ZDHHC17 AND NOT-UNC5A	1	1	1
ZDHHC17 AND NOT-SEMA4D AND CALHM1	1	1	1
ZDHHC17 AND CALHM1 AND NOT-CACNG3	1	1	1
FZD3 AND NOT-KCNK9 AND PCDH11Y	1	1	1
ZDHHC17 AND CALHM1 AND NOT-SLC17A2	1	1	1
ZDHHC17 AND CALHM1 AND NOT-MUC12	1	1	1
ZDHHC17 AND CALHM1 AND NOT-HCN4	1	1	1
GRM6 AND ZDHHC17 AND NOT-CLSTN3	1	1	1
GRM6 AND ZDHHC17 AND NOT-SLC28A1	1	1	1
GRM6 AND ZDHHC17 AND NOT-TRPC3	1	1	1
GRM6 AND ZDHHC17 AND NOT-AJAP1	1	1	1
FZD3 AND NOT-KCNK9 AND TNFRSF10B	1	1	1
GRM6 AND ZDHHC17 AND NOT-RXFP3	1	1	1
FZD3 AND NOT-CNTNAP4 AND NOT-KCNK9	1	1	1
GRM6 AND ZDHHC17 AND NOT-PCDH15	1	1	1
FZD3 AND NOT-KCNK9 AND VAMP8	1	1	1
GRM6 AND ZDHHC17 AND NOT-GPR135	1	1	1
GRM6 AND ZDHHC17 AND NOT-ABCG5	1	1	1
GRM6 AND ZDHHC17 AND NOT-NPFFR1	1	1	1
FZD3 AND NOT-KCNK9 AND LAPTM5	1	1	1
FZD3 AND NOT-KCNK9 AND CACNA1S	1	1	1
ZDHHC17 AND CALHM1 AND NOT-OR4N4	1	1	1
ZDHHC17 AND CALHM1 AND NOT-GHR	1	1	1
COMPLEX-CSPG5/B4GALNT1/FZD3	1	1	1
GPM6B AND NOT-ANO4 AND CBX3	1	1	1
GRM6 AND ZDHHC17 AND NOT-CD70	1	1	1
ZDHHC17 AND CALHM1 AND NOT-CD160	1	1	1
ZDHHC17 AND CALHM1 AND NOT-CLDN7	1	1	1
ZDHHC17 AND CALHM1 AND NOT-CSPG4	1	1	1
ZDHHC17 AND CALHM1 AND NOT-CLEC14A	1	1	1
NOT-SLC6A6 AND SGCZ AND FCRL2	1	1	1
NOT-SLC6A6 AND SGCZ AND ABCB5	1	1	1
GPM6B AND NOT-ANO4 AND NOT-FOLH1	1	1	1
COL25A1 AND NOT-SLC6A6 AND ITGAV	1	1	1
COL25A1 AND NOT-SLC6A6 AND TNC	1	1	1
NOT-SLC16A7 AND GHRHR AND NOT-CLDN7	1	1	1
NOT-SLC6A6 AND SSTR3 AND EQTN	1	1	1
NOT-SLC6A6 AND EQTN AND NOT-EPCAM	1	1	1
NOT-SLC6A6 AND EQTN AND MSLN	1	1	1
NOT-SLC16A7 AND KCNH1 AND PROM1	1	1	1
NOT-SLC16A7 AND KCNH1 AND DDX3X	1	1	1
NOT-SLC16A7 AND KCNH1 AND CBX3	1	1	1
NOT-SLC6A6 AND MSLN AND NOT-CD151	1	1	1
GRM6 AND ZDHHC17 AND NOT-CD38	1	1	1
GRM6 AND ZDHHC17 AND NOT-CD180	1	1	1
GRM6 AND ZDHHC17 AND NOT-MST1R	1	1	1
GRM6 AND ZDHHC17 AND NOT-SEMA5B	1	1	1
GRM6 AND ZDHHC17 AND NOT-TRPM4	1	1	1
GRM6 AND ZDHHC17 AND NOT-MOK	1	1	1
GRM6 AND ZDHHC17 AND NOT-SSTR2	1	1	1
MEGF10 AND NOT-ADCY9 AND NOT-FOLH1	1	1	1
GRM6 AND ZDHHC17 AND NOT-WT1	1	1	1
MEGF10 AND NOT-ADCY9 AND NOT-ERBB3	1	1	1
MEGF10 AND NOT-ADCY9 AND DDX3X	1	1	1
GPM6B AND NOT-ANO4 AND MSLN	1	1	1
GRM6 AND ZDHHC17 AND NOT-ULBP2	1	1	1
GRM6 AND ZDHHC17 AND NOT-FCRL5	1	1	1
NOT-SLC6A6 AND MSLN AND ATP1A4	1	1	1
NOT-SLC6A6 AND MSLN AND BRS3	1	1	1
NOT-SLC6A6 AND MSLN AND SLC12A4	1	1	1
GPM6B AND NOT-ANO4 AND TNFRSF8	1	1	1
NOT-SLC6A6 AND MSLN AND STRA6	1	1	1
CDH23 AND NOT-SLC6A6 AND MSLN	1	1	1
NOT-SLC6A6 AND MSLN AND TRPV4	1	1	1
NOT-ATP11C AND CDH11 AND CBX3	1	1	1
NOT-SLC6A6 AND MSLN AND IL20RB	1	1	1
NOT-SLC6A6 AND MSLN AND P2RY13	1	1	1
NOT-SLC6A6 AND MSLN AND PMP22	1	1	1
NOT-SLC6A6 AND MSLN AND NTM	1	1	1
NOT-SLC6A6 AND MSLN AND TAS2R16	1	1	1
MEGF10 AND NOT-SEMA4D AND LGR5	1	1	1

Antigen Pair Logic	F1	Precision	Recall
ASTN1 AND BEST3 AND NOT-DIO2	1	1	1
ASTN1 AND BEST3 AND NOT-CX3CL1	0.967742	0.9375	1
ASTN1 AND BEST3 AND NOT-OSMR	0.933333	0.933333	0.933333
ASTN1 AND BEST3 AND NOT-PCDH19	0.967742	0.9375	1
ASTN1 AND BEST3 AND NOT-C1orf210	0.933333	0.933333	0.933333
ASTN1 AND BEST3 AND NOT-CXCR2	0.933333	0.933333	0.933333
BEST3 AND DISP2 AND NOT-AQP7	0.909091	0.833333	1
NCAM2 AND BEST3 AND NOT-ANTXR2	0.9375	0.882353	1
NCAM2 AND BEST3 AND NOT-CDH22	0.903226	0.875	0.933333
OMG AND BEST3 AND NOT-FLRT2	0.967742	0.9375	1
GRIA2 AND TNFRSF10B AND NOT-SLC30A10	0.928571	1	0.866667
GRIA2 AND TNFRSF10B AND NOT-SLC5A10	0.928571	1	0.866667
GRIA2 AND TNFRSF10B AND NOT-PTPRT	0.928571	1	0.866667
ASTN1 AND BEST3 AND NOT-TNFSF10	0.896552	0.928571	0.866667
GRIA2 AND BEST3 AND NOT-PTPRT	0.967742	0.9375	1
GRIA2 AND BEST3 AND NOT-SLC30A10	0.967742	0.9375	1
OMG AND BEST3 AND NOT-CLCN5	0.967742	0.9375	1
CDH10 AND BEST3 AND NOT-PTPRM	0.9375	0.882353	1
OMG AND BEST3 AND NOT-GPR52	0.967742	0.9375	1
COMPLEX-CRB1/GRIK5/CD82	0.9375	0.882353	1
GRIA2 AND TNFRSF10B AND NOT-PROCR	0.928571	1	0.866667
GRIA2 AND TNFRSF10B AND NOT-BTN3A1	0.928571	1	0.866667
GRIA2 AND TNFRSF10B AND NOT-BST2	0.928571	1	0.866667
COMPLEX-NKAIN4/GOLM1/THBD	0.888889	1	0.8
NKAIN4 AND NOT-PIK3IP1 AND P2RY13	0.888889	1	0.8
NKAIN4 AND NOT-OPCML AND HLA-G	0.888889	1	0.8
COMPLEX-ATP9A/NKAIN4/VANGL1	0.888889	1	0.8
NKAIN4 AND NOT-LAT AND P2RY13	0.888889	1	0.8
COMPLEX-NKAIN4/APOLD1/AOC3	0.888889	1	0.8
COMPLEX-NKAIN4/GOLM1/SMPD2	0.888889	1	0.8
ASTN1 AND TSPAN11 AND NOT-SLC5A11	0.888889	1	0.8
COMPLEX-NKAIN4/EMP2/DISP2	0.888889	1	0.8
ASTN1 AND TSPAN11 AND NOT-ANTXR2	0.888889	1	0.8
COMPLEX-NKAIN4/FLRT3/GHR	0.888889	1	0.8
COMPLEX-NKAIN4/TSPAN14/DISP2	0.888889	1	0.8
COMPLEX-ATP9A/NKAIN4/SLC39A8	0.888889	1	0.8
NKAIN4 AND NOT-FAIM2 AND SLC30A10	0.888889	1	0.8
ASTN1 AND TSPAN11 AND NOT-CNTNAP4	0.888889	1	0.8
COMPLEX-NKAIN4/GOLM1/LRRC8E	0.888889	1	0.8
NKAIN4 AND NOT-SMPD2 AND IL1RAP	0.888889	1	0.8
COMPLEX-NKAIN4/SLCO1B3/NTM	0.888889	1	0.8
COMPLEX-NKAIN4/SMPD2/VLDLR	0.888889	1	0.8
GRIA2 AND F2R AND NOT-PROCR	0.888889	1	0.8
COMPLEX-ATP10A/BCAN/LRRC8E	0.928571	1	0.866667
BCAN AND NOT-KCNC1 AND CD163L1	0.928571	1	0.866667
BCAN AND NOT-OPCML AND SLC24A1	0.928571	1	0.866667
BCAN AND NOT-GHR AND IGSF6	0.928571	1	0.866667
BCAN AND NOT-TGFBR2 AND NOT-ASTN1	0.928571	1	0.866667
COMPLEX-ATP10A/PTGIS/BCAN	0.928571	1	0.866667
COMPLEX-ASTN1/ATP10A/BCAN	0.928571	1	0.866667
COMPLEX-SCARA5/ABCB1/BCAN	0.928571	1	0.866667
BCAN AND NOT-GHR AND SLC22A4	0.928571	1	0.866667
COMPLEX-EPHA2/ATP10A/BCAN	0.928571	1	0.866667
COMPLEX-KCNC1/ABCB1/BCAN	0.928571	1	0.866667
COMPLEX-ABCB1/SCN2A/BCAN	0.928571	1	0.866667
BCAN AND NOT-SLC43A1 AND HAVCR2	0.928571	1	0.866667
BCAN AND NOT-NCAM2 AND CD1B	0.928571	1	0.866667
BCAN AND NOT-PLVAP AND HAVCR2	0.928571	1	0.866667
BCAN AND NOT-SELP AND SIRPG	0.928571	1	0.866667
COMPLEX-PIK3IP1/ATP10A/BCAN	0.928571	1	0.866667
BCAN AND NOT-SMPD2 AND CD84	0.928571	1	0.866667
BCAN AND NOT-RHOT2 AND NOT-ASTN1	0.928571	1	0.866667
BCAN AND NOT-CLEC4A AND NOT-ASTN1	0.928571	1	0.866667
BCAN AND NOT-SIRPG AND NOT-ASTN1	0.928571	1	0.866667
BCAN AND NOT-PIK3IP1 AND HAVCR2	0.928571	1	0.866667
BCAN AND NOT-GHR AND HAVCR2	0.928571	1	0.866667
COMPLEX-ABCB1/GPR158/BCAN	0.928571	1	0.866667
COMPLEX-ATP10A/BCAN/SLC43A1	0.928571	1	0.866667
COMPLEX-NCAM2/BCAN/LRRTM4	0.928571	1	0.866667
COMPLEX-GHR/ATP10A/BCAN	0.928571	1	0.866667
BCAN AND NOT-EMCN AND NOT-ASTN1	0.928571	1	0.866667
BCAN AND NOT-ABCD4 AND NOT-ASTN1	0.928571	1	0.866667
COMPLEX-LAT/ATP10A/BCAN	0.928571	1	0.866667
BCAN AND NOT-SLC24A2 AND CLEC4A	0.928571	1	0.866667
BCAN AND NOT-TMED1 AND NOT-ASTN1	0.928571	1	0.866667
COMPLEX-SCAMP5/BCAN/SLC39A8	0.928571	1	0.866667
BCAN AND NOT-SMPD2 AND CD1C	0.928571	1	0.866667

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
GRM8 AND NOT-SLC6A6 AND CBX3	1	1	1
NOT-SLC6A6 AND TAS1R1 AND SSTR3	1	1	1
NOT-SLC6A6 AND SLC9B1 AND NOT-EPCAM	1	1	1
NOT-SLC6A6 AND PCDHGC4 AND TNFSF11	1	1	1
GRM8 AND NOT-SLC6A6 AND DDX3X	1	1	1
NOT-SLC6A6 AND PCDHGC4 AND FCRL2	1	1	1
SLC7A5 AND NOT-SLC6A6 AND NPFFR1	1	1	1
NOT-SLC6A6 AND SLC9B1 AND ITGB3	1	1	1
GRM8 AND NOT-SLC6A6 AND SLC39A6	1	1	1
SLC7A5 AND NOT-SLC6A6 AND CHRNG	1	1	1
GRM8 AND NOT-SLC6A6 AND ITGAV	1	1	1
GRM8 AND NOT-SLC6A6 AND ITGB3	1	1	1
SLC7A5 AND NOT-SLC6A6 AND LAPTM5	1	1	1
NOT-SLC6A6 AND GPR78 AND AFP	1	1	1
NOT-SLC6A6 AND GPR78 AND CBX3	1	1	1
OTOF AND NOT-SLC6A6 AND CBX3	1	1	1
MEGF10 AND NOT-SLC6A6 AND SLC7A5	1	1	1
ABCB5 AND NOT-SLC6A6 AND UMODL1	1	1	1
NOT-SLC6A6 AND PCDHGC4 AND ITGB3	1	1	1
GRM8 AND NOT-SLC6A6 AND MOK	1	1	1
NOT-SLC6A6 AND MSLN AND CATSPER3	1	1	1
NOT-SLC6A6 AND CDH11 AND CLDN5	1	1	1
MEGF10 AND NOT-SLC6A6 AND DDX3X	1	1	1
NOT-SLC6A6 AND MSLN AND KCNJ6	1	1	1
MEGF10 AND NOT-SLC6A6 AND FAP	1	1	1
NOT-SLC6A6 AND UMODL1 AND HLA-DOB	1	1	1
NOT-SLC6A6 AND UMODL1 AND FAP	1	1	1
NOT-SLC6A6 AND UMODL1 AND ERBB4	1	1	1
NOT-SLC6A6 AND MSLN AND OXTR	1	1	1
NOT-SLC6A6 AND UMODL1 AND DDX3X	1	1	1
NOT-SLC6A6 AND UMODL1 AND CBX3	1	1	1
SLC7A5 AND NOT-SLC6A6 AND FLVCR1	1	1	1
MEGF10 AND NOT-SLC6A6 AND MOK	1	1	1
SLC7A5 AND NOT-SLC6A6 AND MLNR	1	1	1
SLC7A5 AND NOT-SLC6A6 AND OXTR	1	1	1
SLC7A5 AND NOT-SLC6A6 AND P2RX3	1	1	1
MEGF10 AND NOT-SLC6A6 AND TNFSF11	1	1	1
NOT-SLC6A6 AND TAS1R1 AND TNFSF11	1	1	1
MEGF10 AND NOT-SLC6A6 AND TNFRSF10A	1	1	1
NOT-SLC6A6 AND MSLN AND OR52D1	1	1	1
NOT-SLC6A6 AND LRRN4 AND IGF1R	1	1	1
NOT-SLC6A6 AND GHRHR AND ITGAV	1	1	1
GRM8 AND NOT-SLC6A6 AND MSLN	1	1	1
NOT-SLC6A6 AND KCNK9 AND TYR	1	1	1
NOT-SLC6A6 AND MLANA AND MUC13	1	1	1
NOT-SLC6A6 AND MSLN AND CSMD2	1	1	1
NOT-SLC6A6 AND ROS1 AND ABCB5	1	1	1
NOT-SLC6A6 AND CLDN17 AND MSLN	1	1	1
NOT-SLC6A6 AND CLDN17 AND IGF1R	1	1	1
NOT-SLC6A6 AND MLANA AND MSLN	1	1	1
NOT-SLC6A6 AND SLC12A3 AND SEMA5B	1	1	1
NOT-SLC6A6 AND CLDN17 AND ITGB3	1	1	1
NOT-SLC6A6 AND CLDN17 AND CD37	1	1	1
SLC22A14 AND NOT-SLC6A6 AND SLC39A6	1	1	1
NOT-SLC6A6 AND CLDN17 AND TNFSF11	1	1	1
NOT-SLC6A6 AND CLDN17 AND CLDN5	1	1	1
NOT-SLC6A6 AND CLDN17 AND NOT-TRPM4	1	1	1
NOT-SLC6A6 AND MSLN AND SLC13A5	1	1	1
NOT-SLC6A6 AND SLC12A3 AND TNFSF11	1	1	1
NOT-SLC6A6 AND MSLN AND SLC17A3	1	1	1
NOT-SLC6A6 AND PCDHGC4 AND FOLR1	1	1	1
NOT-SLC6A6 AND PCDHGC4 AND DKK1	1	1	1
NOT-SLC6A6 AND SSTR3 AND KCNK9	1	1	1
NOT-SLC6A6 AND SLC12A3 AND ITGAV	1	1	1
NOT-SLC6A6 AND PCDHGC4 AND ERBB4	1	1	1
MC2R AND NOT-CD52 AND UMODL1	1	1	1
NOT-SLC6A6 AND SLC12A3 AND ITGB3	1	1	1
NOT-SLC6A6 AND MSLN AND KCNK9	1	1	1
NOT-SLC6A6 AND PCDHGC4 AND DDX3X	1	1	1
NOT-SLC6A6 AND KCNK9 AND ERBB2	1	1	1
NOT-SLC6A6 AND KCNK9 AND CSPG4	1	1	1
ADCY10 AND NOT-CD52 AND CD33	1	1	1
NOT-SLC6A6 AND SSTR3 AND ABCB5	1	1	1
NOT-SLC6A6 AND SSTR3 AND CBX3	1	1	1
NOT-SLC6A6 AND SSTR3 AND DDX3X	1	1	1
NOT-SLC6A6 AND SSTR3 AND MSLN	1	1	1
NOT-SLC6A6 AND SSTR3 AND IGF1R	1	1	1

Antigen Pair Logic	F1	Precision	Recall
BCAN AND NOT-SELP AND HAVCR2	0.928571	1	0.866667
NCAM2 AND BEST3 AND NOT-ULBP2	0.967742	0.9375	1
BEST3 AND LRRTM2 AND NOT-TRPM4	0.9375	0.882353	1
GRIA2 AND EGFR AND NOT-PTPRT	0.903226	0.875	0.933333
GRIA2 AND TNFRSF10B AND NOT-ULBP2	0.928571	1	0.866667
GRIA2 AND EGFR AND NOT-SLC30A10	0.903226	0.875	0.933333
GRIA2 AND EGFR AND NOT-SLC5A10	0.903226	0.875	0.933333
GRIA2 AND BEST3 AND NOT-RAET1E	0.967742	0.9375	1
BEST3 AND ATP9A AND NOT-IL20RA	0.9375	0.882353	1
BEST3 AND ATP9A AND NOT-TRPM4	0.9375	0.882353	1
BEST3 AND ATP9A AND NOT-MST1R	0.9375	0.882353	1
GRIA2 AND BEST3 AND NOT-ULBP2	0.967742	0.9375	1
GRIA2 AND TNFRSF10B AND NOT-CLDN1	0.928571	1	0.866667
GRIA2 AND TNFRSF10B AND NOT-RAET1E	0.928571	1	0.866667
DLL3 AND NOT-SMPD2 AND CYP4F12	0.888889	1	0.8
ASTN1 AND VANGL2 AND NOT-IL13RA1	0.888889	1	0.8
DLL3 AND NOT-GHR AND SIRPG	0.888889	1	0.8
COMPLEX-NKAIN4/ITGAV/CA12	0.888889	1	0.8
COMPLEX-ITGAV/BCAN/PPAP2A	0.928571	1	0.866667
BCAN AND NOT-CLDN5 AND CLEC4A	0.928571	1	0.866667
BCAN AND NOT-CD37 AND CD84	0.928571	1	0.866667
BCAN AND NOT-NRXN2 AND VCAM1	0.928571	1	0.866667
COMPLEX-CLDN7/ATP10A/BCAN	0.928571	1	0.866667
BCAN AND NOT-MS4A1 AND SIRPG	0.928571	1	0.866667
COMPLEX-ATP10A/BCAN/CLDN5	0.928571	1	0.866667
COMPLEX-ATP10A/BCAN/CD34	0.928571	1	0.866667
COMPLEX-ERBB3/ATP10A/BCAN	0.928571	1	0.866667
BCAN AND NOT-AQP9 AND VCAM1	0.928571	1	0.866667
COMPLEX-ITGAV/ATP10A/BCAN	0.928571	1	0.866667
BCAN AND NOT-TRPM4 AND SLC22A4	0.928571	1	0.866667
COMPLEX-TRPM4/ATP10A/BCAN	0.928571	1	0.866667
BCAN AND NOT-RAET1E AND HAVCR2	0.928571	1	0.866667
BCAN AND NOT-CLDN1 AND SIRPG	0.928571	1	0.866667
COMPLEX-RAET1E/ATP10A/BCAN	0.928571	1	0.866667
BCAN AND NOT-TRPM4 AND NOT-LRP11	0.928571	1	0.866667
GRIA2 AND EGFR AND NOT-RAET1E	0.903226	0.875	0.933333
GRIA2 AND EGFR AND NOT-ULBP2	0.903226	0.875	0.933333
DLL3 AND NOT-MUC1 AND ATP10A	0.888889	1	0.8
DLL3 AND NOT-LSAMP AND CLDN5	0.888889	1	0.8
DLL3 AND NOT-CD34 AND FAIM2	0.888889	1	0.8
DLL3 AND NOT-ITGB6 AND GLDN	0.888889	1	0.8
DLL3 AND NOT-TRPM4 AND SIRPG	0.888889	1	0.8
DLL3 AND NOT-TRPM4 AND IL1RAP	0.888889	1	0.8
DLL3 AND NOT-CA12 AND CLDN5	0.888889	1	0.8
DLL3 AND NOT-CLDN5 AND FAIM2	0.888889	1	0.8
COMPLEX-DLL3/EPCAM/PTPRE	0.888889	1	0.8
DLL3 AND NOT-MS4A1 AND SIRPG	0.888889	1	0.8
DLL3 AND NOT-TRPM4 AND ATP10A	0.888889	1	0.8
DLL3 AND NOT-TRPM4 AND GLDN	0.888889	1	0.8
DLL3 AND NOT-IL20RA AND SIRPG	0.888889	1	0.8
DLL3 AND NOT-CLDN7 AND ATP10A	0.888889	1	0.8
DLL3 AND NOT-TRPM4 AND CD4	0.888889	1	0.8
DLL3 AND NOT-ITGB6 AND ATP10A	0.888889	1	0.8
DLL3 AND NOT-MUC1 AND GLDN	0.888889	1	0.8
NCAM2 AND EGFR AND NOT-CLDN1	0.888889	1	0.8
NCAM2 AND EGFR AND NOT-CD34	0.888889	1	0.8
DLL3 AND NOT-LYVE1 AND CLDN5	0.888889	1	0.8
NCAM2 AND EGFR AND NOT-IL13RA1	0.888889	1	0.8
DLL3 AND NOT-CD34 AND SCAMP5	0.888889	1	0.8
DLL3 AND NOT-RAET1E AND SCAMP5	0.888889	1	0.8
COMPLEX-DLL3/KLRF1/TRPM4	0.888889	1	0.8
NCAM2 AND EGFR AND NOT-ULBP2	0.888889	1	0.8
COMPLEX-DLL3/CXADR/TRPM4	0.888889	1	0.8
DLL3 AND NOT-CLDN5 AND ATP10A	0.857143	0.923077	0.8
DLL3 AND NOT-RAET1E AND GLDN	0.857143	0.923077	0.8
DLL3 AND NOT-TRPM4 AND CD84	0.888889	1	0.8
DLL3 AND NOT-CLDN5 AND NT5E	0.857143	0.923077	0.8
LSAMP AND EGFR AND NOT-CLDN1	0.857143	0.923077	0.8
COMPLEX-DLL3/ATP1B3/TRPM4	0.857143	0.923077	0.8
DLL3 AND NOT-LSAMP AND IL13RA1	0.857143	0.923077	0.8
DLL3 AND NOT-CLDN1 AND SIRPG	0.857143	0.923077	0.8
VANGL2 AND NOT-IL20RA AND CLDN5	0.857143	0.923077	0.8
DLL3 AND NOT-P2RX5 AND CD84	0.857143	0.923077	0.8
DLL3 AND NOT-CD34 AND CD84	0.857143	0.923077	0.8
DLL3 AND NOT-LSAMP AND CLDN1	0.857143	0.923077	0.8
DLL3 AND NOT-CD34 AND ATP10A	0.857143	0.923077	0.8
DLL3 AND NOT-FZD7 AND NOT-MST1R	0.846154	1	0.733333

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
NOT-SLC6A6 AND SSTR3 AND ITGAV	1	1	1
SLC7A5 AND NOT-SLC6A6 AND SLC39A6	1	1	1
GRM6 AND NOT-CD33 AND CBX3	1	1	1
MEGF10 AND NOT-SEMA4D AND CD44	1	1	1
NOT-SLC6A6 AND KCNK9 AND ABHD3	1	1	1
NOT-SLC6A6 AND KCNK9 AND HTR3C	1	1	1
NOT-SLC6A6 AND KCNK9 AND SLC31A1	1	1	1
NOT-SLC6A6 AND KCNK9 AND GJB4	1	1	1
NOT-SLC6A6 AND KCNK9 AND SLC22A12	1	1	1
NOT-SLC6A6 AND KCNK9 AND PROCR	1	1	1
GRM8 AND NOT-SLC6A6 AND GYPC	1	1	1
GRM8 AND NOT-SLC6A6 AND ERVW-1	1	1	1
GRM8 AND NOT-SLC6A6 AND SLC6A18	1	1	1
GRM8 AND NOT-SLC6A6 AND MIP	1	1	1
SLC22A14 AND NOT-SLC6A6 AND FLVCR1	1	1	1
GRM8 AND NOT-SLC6A6 AND OXTR	1	1	1
GRM8 AND NOT-SLC6A6 AND CHRNA9	1	1	1
GRM8 AND NOT-SLC6A6 AND TRPC7	1	1	1
GRM8 AND NOT-SLC6A6 AND PTGIR	1	1	1
GRM8 AND NOT-SLC6A6 AND SLC2A1	1	1	1
GRM8 AND NOT-SLC6A6 AND NOT-STEAP4	1	1	1
GRM8 AND NOT-SLC6A6 AND UMODL1	1	1	1
NOT-SLC6A6 AND KCNK9 AND NOT-SLC12A5	1	1	1
GRM8 AND NOT-SLC6A6 AND GPR156	1	1	1
NOT-SLC6A6 AND KCNK9 AND CD163	1	1	1
NOT-SLC6A6 AND KCNK9 AND KCNK16	1	1	1
NOT-SLC6A6 AND KCNK9 AND PCDH11Y	1	1	1
NOT-SLC6A6 AND SLC12A3 AND KCNK9	1	1	1
NOT-SLC6A6 AND KCNK9 AND NPFFR1	1	1	1
GRM8 AND NOT-SLC6A6 AND FLVCR1	1	1	1
NOT-SLC6A6 AND KCNK9 AND GJD2	1	1	1
GRM8 AND NOT-SLC6A6 AND OR8D1	1	1	1
NOT-SLC6A6 AND PCDHGC4 AND KCNK9	1	1	1
NOT-SLC6A6 AND KCNK9 AND MIP	1	1	1
NOT-SLC6A6 AND GPR78 AND KCNK9	1	1	1
NOT-SLC6A6 AND CLDN17 AND KCNK9	1	1	1
GRM8 AND NOT-SLC6A6 AND PCDH11Y	1	1	1
GRM8 AND NOT-SLC6A6 AND GPR50	1	1	1
NOT-SLC6A6 AND UMODL1 AND GPR19	1	1	1
NOT-SLC6A6 AND UMODL1 AND SLC32A1	1	1	1
NOT-SLC6A6 AND TAS1R1 AND NOT-STEAP4	1	1	1
NOT-SLC6A6 AND TAS1R1 AND GPR61	1	1	1
NOT-SLC6A6 AND TAS1R1 AND NOT-AOC3	1	1	1
NOT-SLC6A6 AND TAS1R1 AND CD44	1	1	1
NOT-SLC6A6 AND UMODL1 AND HCN4	1	1	1
NOT-SLC6A6 AND GPR78 AND SLC2A2	1	1	1
NOT-SLC6A6 AND GPR78 AND GJD2	1	1	1
NOT-SLC6A6 AND GPR78 AND MRAP	1	1	1
NOT-SLC6A6 AND UMODL1 AND CHRNG	1	1	1
NOT-SLC6A6 AND UMODL1 AND MRGPRX2	1	1	1
NOT-SLC6A6 AND GPR78 AND GABRA3	1	1	1
NOT-SLC6A6 AND UMODL1 AND GPR156	1	1	1
NOT-SLC6A6 AND TAS1R1 AND OXTR	1	1	1
NOT-SLC6A6 AND UMODL1 AND KCNV2	1	1	1
NOT-SLC6A6 AND UMODL1 AND HTR3C	1	1	1
NOT-SLC6A6 AND UMODL1 AND FGF6	1	1	1
NOT-SLC6A6 AND UMODL1 AND CNIH2	1	1	1
NOT-SLC6A6 AND UMODL1 AND OR1C1	1	1	1
NOT-SLC6A6 AND UMODL1 AND OR10J1	1	1	1
MEGF10 AND NOT-SLC6A6 AND MLANA	1	1	1
NOT-SLC6A6 AND PCDHGC4 AND MLANA	1	1	1
NOT-SLC6A6 AND MLANA AND P2RX4	1	1	1
NOT-SLC6A6 AND MLANA AND OXTR	1	1	1
NOT-SLC6A6 AND TAS1R1 AND UPK3A	1	1	1
NOT-SLC6A6 AND TAS1R1 AND KCNJ6	1	1	1
GRM8 AND NOT-SLC6A6 AND CD44	1	1	1
NOT-SLC6A6 AND SLC4A5 AND ZACN	1	1	1
NOT-SLC6A6 AND CLDN17 AND MMP24	1	1	1
NOT-SLC6A6 AND CLDN17 AND CABP7	1	1	1
NOT-SLC6A6 AND CLDN17 AND CNIH2	1	1	1
NOT-SLC6A6 AND CLDN17 AND GABRB1	1	1	1
NOT-SLC6A6 AND CLDN17 AND GPR22	1	1	1
NOT-SLC6A6 AND CLDN17 AND FLVCR1	1	1	1
NOT-SLC6A6 AND CLDN17 AND LCT	1	1	1
NOT-SLC6A6 AND CLDN17 AND OXTR	1	1	1
NOT-SLC6A6 AND CLDN17 AND CACNA1B	1	1	1
OTOF AND NOT-SLC6A6 AND CLDN17	1	1	1

Antigen Pair Logic	F1	Precision	Recall
COMPLEX-ATP9A/DLL3/MST1R	0.846154	1	0.733333
DLL3 AND NOT-IL3RA AND P2RY13	0.846154	1	0.733333
DLL3 AND NOT-MUC1 AND P2RY13	0.846154	1	0.733333
DLL3 AND NOT-TRPM4 AND P2RY13	0.846154	1	0.733333
DLL3 AND NOT-ITGB6 AND P2RY13	0.846154	1	0.733333
DLL3 AND NOT-CLDN7 AND P2RY13	0.846154	1	0.733333
DLL3 AND NOT-TNFRSF17 AND P2RY13	0.846154	1	0.733333
LSAMP AND EGFR AND NOT-IL13RA1	0.827586	0.857143	0.8
LSAMP AND EGFR AND NOT-RAET1E	0.827586	0.857143	0.8
LSAMP AND EGFR AND NOT-CD34	0.827586	0.857143	0.8
NRXN2 AND EGFR AND NOT-RAET1E	0.827586	0.857143	0.8
IFITM10 AND NOT-CLDN1 AND NOT-TRPM4	0.827586	0.857143	0.8
GRIA2 AND SSX1 AND ITGAV	0.846154	1	0.733333
NCAM2 AND CLDN1 AND NOT-EPCAM	0.823529	0.736842	0.933333
NLGN3 AND NOT-TRPM4 AND NOT-STEAP2	0.933333	0.933333	0.933333
VANGL2 AND NOT-PMEL AND CLDN5	0.827586	0.857143	0.8
COMPLEX-SEZ6/KCNQ2/BCAN	0.928571	1	0.866667
COMPLEX-SLC5A11/ASTN1/BCAN	0.928571	1	0.866667
BCAN AND NOT-KCNC1 AND SLC5A10	0.928571	1	0.866667
ASTN1 AND BEST3 AND NOT-RAET1E	0.967742	0.9375	1
ASTN1 AND BEST3 AND NOT-ULBP2	0.967742	0.9375	1
ASTN1 AND BEST3 AND NOT-CLDN5	0.967742	0.9375	1
ASTN1 AND BEST3 AND NOT-CD34	0.967742	0.9375	1
ASTN1 AND BEST3 AND NOT-IL13RA1	0.967742	0.9375	1
BEST3 AND NOT-AOC3 AND CLDN5	0.9375	0.882353	1
COMPLEX-NKAIN4/CLDN5/CD1A	0.888889	1	0.8
COMPLEX-CDH10/NKAIN4/CD34	0.888889	1	0.8
COMPLEX-NKAIN4/ITGAV/SCN2A	0.888889	1	0.8
COMPLEX-CDH10/NKAIN4/CLDN5	0.888889	1	0.8
ASTN1 AND EGFR AND NOT-GYPC	0.888889	1	0.8
ASTN1 AND EGFR AND NOT-PTPRT	0.888889	1	0.8
ASTN1 AND F2R AND NOT-CLDN1	0.888889	1	0.8
OMG AND BEST3 AND NOT-RAET1E	0.967742	0.9375	1
CDH10 AND BEST3 AND NOT-RAET1E	0.9375	0.882353	1
OMG AND BEST3 AND NOT-CLDN5	0.967742	0.9375	1
ASTN1 AND EGFR AND NOT-ANTXR2	0.888889	1	0.8
ASTN1 AND F2R AND NOT-IL13RA1	0.888889	1	0.8
ASTN1 AND EGFR AND NOT-SLC30A10	0.888889	1	0.8
GPR158 AND BEST3 AND NOT-RAET1E	0.9375	0.882353	1
OMG AND F2R AND NOT-RAET1E	0.888889	1	0.8
ASTN1 AND EGFR AND NOT-CNTNAP4	0.888889	1	0.8
ASTN1 AND TNFRSF10B AND NOT-IL13RA1	0.928571	1	0.866667
ASTN1 AND EGFR AND NOT-CD163	0.888889	1	0.8
BEST3 AND SCN2A AND NOT-CLDN1	0.967742	0.9375	1
COMPLEX-SEZ6/EGFR/KCNQ2	1	1	1
SCN2A AND EGFR AND NOT-ANTXR2	0.857143	0.923077	0.8
OMG AND EGFR AND NOT-SLC30A10	0.857143	0.923077	0.8
CDH10 AND EGFR AND NOT-SLC30A10	0.857143	0.923077	0.8
COMPLEX-DLL3/CD93/DISP2	0.857143	0.923077	0.8
CDH10 AND EGFR AND NOT-SLCO1B3	0.857143	0.923077	0.8
ASTN1 AND F2R AND NOT-VCAM1	0.888889	1	0.8
ASTN1 AND CXCL16 AND NOT-IL13RA1	0.875	0.823529	0.933333
COMPLEX-DLL3/CD93/OPCML	0.857143	0.923077	0.8
CDH10 AND EGFR AND NOT-SLC39A8	0.857143	0.923077	0.8
ASTN1 AND F2R AND ERBB3	0.888889	1	0.8
COMPLEX-NKAIN4/ERBB4/DISP2	0.846154	1	0.733333
ASTN1 AND EGFR AND NOT-SLC31A1	0.846154	1	0.733333
COMPLEX-NKAIN4/VCAM1/DISP2	0.846154	1	0.733333
COMPLEX-NKAIN4/SCN2A/PROM1	0.846154	1	0.733333
COMPLEX-NKAIN4/SCN2A/VCAM1	0.846154	1	0.733333
BEST3 AND NOT-SCARA5 AND CLDN5	0.9375	0.882353	1
ASTN1 AND TNFRSF10B AND NOT-CLDN1	0.928571	1	0.866667
CDH10 AND TNFRSF10B AND NOT-STEAP1	0.928571	1	0.866667
ASTN1 AND F2R AND CLDN5	0.888889	1	0.8
OMG AND TNFRSF10B AND RAET1E	0.83871	0.8125	0.866667
SCN2A AND EGFR AND NOT-SLC5A10	0.857143	0.923077	0.8
GPR158 AND F2R AND NOT-RAET1E	0.888889	1	0.8
OMG AND TNFRSF10B AND NOT-CD34	0.83871	0.8125	0.866667
CDH10 AND F2R AND NOT-STEAP1	0.888889	1	0.8
SCN2A AND EGFR AND NOT-CD1B	0.857143	0.923077	0.8
KCNC1 AND EGFR AND NOT-SLC5A10	0.827586	0.857143	0.8
KCNC1 AND EGFR AND NOT-SLC39A8	0.827586	0.857143	0.8
ASTN1 AND TNC AND NOT-CNTNAP4	0.827586	0.857143	0.8
ASTN1 AND TNC AND NOT-SLC5A11	0.827586	0.857143	0.8
CDH10 AND CXCL16 AND NOT-STEAP1	0.823529	0.736842	0.933333
ASTN1 AND CSPG4 AND NOT-SLC5A11	0.823529	0.736842	0.933333
ASTN1 AND F2R AND NOT-RAET1E	0.846154	1	0.733333

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
NOT-SLC6A6 AND TAS1R1 AND GUCY2D	1	1	1
NOT-SLC16A7 AND ABCB5 AND MSLN	1	1	1
SLC16A2 AND NOT-ENG AND CD160	1	1	1
SLC16A2 AND NOT-IL13RA1 AND SLC34A2	1	1	1
SLC16A2 AND NOT-IL13RA1 AND NOT-CD52	1	1	1
SLC16A2 AND NOT-IL13RA1 AND MSLN	1	1	1
SLC16A2 AND NOT-ENG AND CD38	1	1	1
SLC16A2 AND NOT-ENG AND MS4A1	1	1	1
SLC16A2 AND NOT-ENG AND NOT-CLDN12	1	1	1
SLC16A2 AND NOT-ENG AND TNFRSF10A	1	1	1
SLC16A2 AND NOT-ENG AND LGR5	1	1	1
SLC16A2 AND NOT-ENG AND VTCN1	1	1	1
SLC16A2 AND NOT-ENG AND CA9	1	1	1
SLC16A2 AND NOT-ENG AND NOT-VCAM1	1	1	1
SLC16A2 AND NOT-ENG AND TYR	1	1	1
SLC16A2 AND NOT-ENG AND SSX1	1	1	1
SLC16A2 AND NOT-ENG AND NOT-SDC1	1	1	1
SLC16A2 AND NOT-ENG AND MUC13	1	1	1
SLC16A2 AND NOT-ENG AND NOT-TRPM4	1	1	1
SLC16A2 AND NOT-ENG AND ENPP3	1	1	1
SLC16A2 AND NOT-ENG AND OAS1	1	1	1
SLC16A2 AND NOT-ENG AND CEACAM6	1	1	1
SLC16A2 AND NOT-ENG AND MUC1	1	1	1
SLC16A2 AND NOT-ENG AND NOT-EPCAM	1	1	1
SLC16A2 AND NOT-ENG AND ITGB6	1	1	1
SLC16A2 AND NOT-ENG AND ITGB3	1	1	1
SLC16A2 AND NOT-ENG AND ABCB5	1	1	1
SLC16A2 AND NOT-ENG AND CTAG2	1	1	1
SLC16A2 AND NOT-ENG AND AFP	1	1	1
SLC16A2 AND NOT-ENG AND DDX3X	1	1	1
SLC16A2 AND NOT-IL13RA1 AND CD160	1	1	1
SLC16A2 AND NOT-IL13RA1 AND FCRL1	1	1	1
SLC16A2 AND NOT-IL13RA1 AND DDX3X	1	1	1
SLC16A2 AND NOT-IL13RA1 AND VTCN1	1	1	1
SLC7A11 AND NOT-ENG AND FCRL1	1	1	1
SLC7A11 AND NOT-ENG AND CBX3	1	1	1
SLC16A2 AND NOT-IL13RA1 AND CD70	1	1	1
SLC16A2 AND NOT-IL13RA1 AND CD38	1	1	1
SLC16A2 AND NOT-IL13RA1 AND CD33	1	1	1
SLC16A2 AND NOT-IL13RA1 AND TNFRSF10A	1	1	1
SLC16A2 AND NOT-IL13RA1 AND LGR5	1	1	1
SLC16A2 AND NOT-IL13RA1 AND NOT-CD276	1	1	1
SLC16A2 AND NOT-IL13RA1 AND CA9	1	1	1
SLC16A2 AND NOT-IL13RA1 AND AFP	1	1	1
SLC16A2 AND NOT-IL13RA1 AND TYR	1	1	1
SLC16A2 AND NOT-IL13RA1 AND SSX1	1	1	1
SLC16A2 AND NOT-IL13RA1 AND SLAMF7	1	1	1
SLC16A2 AND NOT-IL13RA1 AND MUC13	1	1	1
SLC16A2 AND NOT-IL13RA1 AND NOT-TRPM4	1	1	1
SLC16A2 AND NOT-IL13RA1 AND ENPP3	1	1	1
SLC16A2 AND NOT-IL13RA1 AND CEACAM6	1	1	1
SLC16A2 AND NOT-IL13RA1 AND MUC1	1	1	1
SLC16A2 AND NOT-IL13RA1 AND ITGB6	1	1	1
SLC16A2 AND NOT-IL13RA1 AND ITGB3	1	1	1
SLC16A2 AND NOT-IL13RA1 AND ABCB5	1	1	1
SLC16A2 AND NOT-IL13RA1 AND CTAG2	1	1	1
SLC16A2 AND NOT-ENG AND FCRL1	1	1	1
SLC16A2 AND NOT-ENG AND SLC34A2	1	1	1
SLC16A2 AND NOT-ENG AND MSLN	1	1	1
TMEM231 AND NOT-EPCAM AND MUC13	1	1	1
TMEM231 AND NOT-EPCAM AND CD180	1	1	1
TMEM231 AND NOT-EPCAM AND ITGB6	1	1	1
TMEM231 AND NOT-EPCAM AND ABCB5	1	1	1
TMEM231 AND NOT-EPCAM AND GUCY2C	1	1	1
TMEM231 AND NOT-EPCAM AND GAGE1	1	1	1
TMEM231 AND NOT-EPCAM AND HHLA2	1	1	1
PTPRA AND NOT-IL13RA1 AND MOK	1	1	1
PTPRA AND NOT-IL13RA1 AND NOT-FOLH1	1	1	1
SLC5A4 AND NOT-CD52 AND PTK7	1	1	1
SLC5A4 AND NOT-CD52 AND SLC39A6	1	1	1
SLC5A4 AND NOT-CD52 AND CBX3	1	1	1
FADS2 AND NOT-TRPM4 AND MOK	1	1	1
FADS2 AND NOT-TRPM4 AND NOT-FOLH1	1	1	1
FADS2 AND NOT-TRPM4 AND NOT-ERBB3	1	1	1
FADS2 AND NOT-TRPM4 AND DDX3X	1	1	1
FADS2 AND NOT-TRPM4 AND CBX3	1	1	1
NOT-TRPM4 AND APCDD1 AND CD70	1	1	1

Antigen Pair Logic	F1	Precision	Recall
CDH10 AND F2R AND NOT-CD34	0.888889	1	0.8
ASTN1 AND F2R AND NOT-CD34	0.846154	1	0.733333
ASTN1 AND TNFRSF10B AND NOT-VCAM1	0.928571	1	0.866667
ASTN1 AND TNFRSF10B AND CLDN5	0.928571	1	0.866667
SCN2A AND EGFR AND NOT-CD93	0.857143	0.923077	0.8
ASTN1 AND GJD2 AND NOT-ULBP2	0.888889	1	0.8
KCNC1 AND CD1B AND ST8SIA1	0.83871	0.8125	0.866667
CDH10 AND TNFRSF10B AND NOT-CD34	0.928571	1	0.866667
COMPLEX-DLL3/CD93/KCNC1	0.814815	0.916667	0.733333
COMPLEX-DLL3/SLCO1B3/KCNC1	0.814815	0.916667	0.733333
SCN2A AND EGFR AND NOT-VAMP8	0.814815	0.916667	0.733333
COMPLEX-DLL3/ANTXR2/SCN2A	0.814815	0.916667	0.733333
COMPLEX-DLL3/ATP2B2/DISP2	0.814815	0.916667	0.733333
OPCML AND BEST3 AND NOT-RAET1E	0.8125	0.764706	0.866667
GPR158 AND TNFRSF10B AND NOT-CD34	0.8125	0.764706	0.866667
ASTN1 AND GJD2 AND NOT-RAET1E	0.888889	1	0.8
IGDCC3 AND ASTN1 AND NOT-ULBP2	0.965517	1	0.933333
CDH10 AND F2R AND NOT-RAET1E	0.888889	1	0.8
ASTN1 AND TNFRSF10B AND NOT-RAET1E	0.846154	1	0.733333
CDH10 AND TNFRSF10B AND NOT-RAET1E	0.928571	1	0.866667
ASTN1 AND TSHR AND NOT-RAET1E	0.8	0.8	0.8
OPCML AND EGFR AND NOT-SLC39A8	0.8	0.8	0.8
OPCML AND F2R AND NOT-CD34	0.8	0.8	0.8
OPCML AND F2R AND NOT-RAET1E	0.8	0.8	0.8
OPCML AND EGFR AND NOT-CD93	0.8	0.8	0.8
OPCML AND EGFR AND NOT-CD1B	0.8	0.8	0.8
ASTN1 AND TSHR AND NOT-ULBP2	0.8	0.8	0.8
ASTN1 AND TSHR AND NOT-CD34	0.8	0.8	0.8
ASTN1 AND CSPG4 AND NOT-PTPRT	0.8	0.7	0.933333
ASTN1 AND CSPG4 AND NOT-SLC30A10	0.8	0.7	0.933333
DLL3 AND NOT-SCN2A AND NOT-SLC39A8	0.8	1	0.666667
ASTN1 AND NOT-SST AND SLCO1B3	0.8	1	0.666667
BCAN AND NOT-CLDN5 AND SLC5A10	0.928571	1	0.866667
BCAN AND NOT-SMPD2 AND VCAM1	0.928571	1	0.866667
ASTN1 AND EGFR AND NOT-CD34	0.888889	1	0.8
ASTN1 AND EGFR AND NOT-IL13RA1	0.888889	1	0.8
ASTN1 AND EGFR AND NOT-CLDN1	0.888889	1	0.8
ASTN1 AND EGFR AND NOT-ULBP2	0.888889	1	0.8
ASTN1 AND EGFR AND NOT-STEAP1	0.888889	1	0.8
CDH10 AND EGFR AND NOT-STEAP1	0.857143	0.923077	0.8
COMPLEX-DLL3/ITGAV/SCN2A	0.857143	0.923077	0.8
OMG AND EGFR AND NOT-RAET1E	0.857143	0.923077	0.8
CDH10 AND EGFR AND NOT-RAET1E	0.857143	0.923077	0.8
SCN2A AND EGFR AND NOT-CD34	0.857143	0.923077	0.8
SCN2A AND EGFR AND NOT-CLDN1	0.857143	0.923077	0.8
COMPLEX-DLL3/KCNC1/VCAM1	0.857143	0.923077	0.8
CDH10 AND EGFR AND NOT-CD34	0.857143	0.923077	0.8
ASTN1 AND EGFR AND NOT-RAET1E	0.846154	1	0.733333
COMPLEX-DLL3/SCN2A/VCAM1	0.814815	0.916667	0.733333
SCN2A AND EGFR AND NOT-IL13RA1	0.814815	0.916667	0.733333
SCN2A AND EGFR AND NOT-PROM1	0.814815	0.916667	0.733333
ST8SIA1 AND DISP2 AND MUC1	0.846154	1	0.733333
SCN2A AND EGFR AND NOT-RAET1E	0.857143	0.923077	0.8
ASTN1 AND MUC1 AND CLDN5	0.866667	0.866667	0.866667
GPR158 AND EGFR AND NOT-CD34	0.8	0.8	0.8
OPCML AND EGFR AND NOT-CD34	0.8	0.8	0.8
GPR158 AND EGFR AND NOT-RAET1E	0.8	0.8	0.8
SSX1 AND NOT-AOC3 AND CLDN5	0.8	0.8	0.8
OPCML AND EGFR AND NOT-RAET1E	0.8	0.8	0.8
ASTN1 AND CSPG4 AND NOT-CLDN5	0.8	0.7	0.933333
ASTN1 AND CSPG4 AND NOT-IL13RA1	0.8	0.7	0.933333
ASTN1 AND CSPG4 AND NOT-ULBP2	0.8	0.7	0.933333
DLL3 AND NOT-SCN2A AND CLDN5	0.8	1	0.666667
ASTN1 AND CSPG4 AND NOT-RAET1E	0.8	0.7	0.933333
ASTN1 AND CA9 AND CLDN5	0.8	0.8	0.8
ASTN1 AND NOT-ABCA5 AND VCAM1	0.814815	0.916667	0.733333
ASTN1 AND CD33 AND CLDN5	0.8	0.7	0.933333
ASTN1 AND NOT-EPCAM AND IL13RA1	0.83871	0.8125	0.866667
ASTN1 AND TNC AND NOT-IL13RA1	0.827586	0.857143	0.8
MEGF11 AND CD34 AND NOT-TRPM4	0.8	1	0.666667
ASTN1 AND TNC AND RAET1E	0.827586	0.857143	0.8
ASTN1 AND MUC13 AND NOT-RAET1E	0.8	1	0.666667
SSX1 AND NOT-EPHA2 AND ITGAV	0.827586	0.857143	0.8
SCN2A AND SSX1 AND NOT-CLDN1	0.8	0.8	0.8
ASTN1 AND PTK7 AND NOT-IL13RA1	0.8	1	0.666667
ASTN1 AND PTK7 AND NOT-CLDN1	0.8	1	0.666667
BEST3 AND STEAP2 AND NOT-TRPM4	0.83871	0.8125	0.866667

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
NOT-TRPM4 AND APCDD1 AND DDX3X	1	1	1
TMEM231 AND NOT-STEAP1 AND VTCN1	1	1	1
TMEM231 AND NOT-STEAP1 AND DDX3X	1	1	1
TMEM231 AND NOT-EPCAM AND TNFRSF17	1	1	1
TMEM231 AND NOT-EPCAM AND SSTR3	1	1	1
TMEM231 AND NOT-EPCAM AND FCRL2	1	1	1
FZD3 AND NOT-ERBB4 AND NOT-BIRC5	1	1	1
FZD3 AND NOT-ERBB4 AND CD70	1	1	1
FZD3 AND NOT-ERBB4 AND NOT-MS4A1	1	1	1
FZD3 AND NOT-ERBB4 AND PROM1	1	1	1
FZD3 AND NOT-ERBB4 AND NOT-THY1	1	1	1
FZD3 AND NOT-ERBB4 AND NOT-SSTR4	1	1	1
FZD3 AND NOT-SST AND NOT-ERBB4	1	1	1
Esophageal cancer			
SLC39A4 AND NOT-ST6GALNAC6 AND CDH3	0.833333	1	0.714286
SLC39A4 AND NOT-ST6GALNAC6 AND LAMP3	0.833333	1	0.714286
SLC39A4 AND NOT-ST6GALNAC6 AND F2RL2	0.88	1	0.785714
SLC39A4 AND NOT-ST6GALNAC6 AND TACSTD2	0.814815	0.846154	0.785714
SLC39A4 AND NOT-CDHR1 AND F2RL2	0.962963	1	0.928571
SLC39A4 AND NOT-ST6GALNAC6 AND NOT-CYBRD1	0.814815	0.846154	0.785714
SLC39A4 AND NOT-CDHR1 AND CDH3	0.923077	1	0.857143
SLC39A4 AND NOT-CA12 AND GJB3	0.888889	0.923077	0.857143
SLC39A4 AND NOT-ST6GALNAC6 AND TM4SF1	0.88	1	0.785714
SLC39A4 AND NOT-TNFRSF25 AND F2RL2	0.8	0.909091	0.714286
SLC39A4 AND NOT-ABCC11 AND TACSTD2	0.923077	1	0.857143
SLC39A4 AND NOT-CA12 AND CDH3	0.857143	0.857143	0.857143
SLC39A4 AND NOT-CA12 AND CDH3	0.857143	0.857143	0.857143
SLC39A4 AND NOT-PTGDR2 AND F2RL2	0.8	0.909091	0.714286
SLC39A4 AND TACSTD2 AND NOT-LIFR	0.827586	0.8	0.857143
SLC39A4 AND NOT-C1QTNF1 AND F2RL2	0.8	0.909091	0.714286
SLC52A2 AND NOT-ST6GALNAC6 AND NOT-ACVR1B	0.833333	1	0.714286
SLC52A2 AND NOT-CD300LG AND TACSTD2	0.833333	1	0.714286
SLC52A2 AND NOT-ST6GALNAC6 AND NOT-PLXNA2	0.833333	1	0.714286
SLC52A2 AND NOT-ST6GALNAC6 AND FAM57A	0.833333	1	0.714286
SLC39A4 AND NOT-ABCC11 AND CDH3	0.88	1	0.785714
SLC39A4 AND NOT-SGCA AND F2RL2	0.8	0.909091	0.714286
SLC39A4 AND NOT-CDHR1 AND NOT-ATP10A	0.83871	0.764706	0.928571
SLC39A4 AND NOT-CA12 AND PTPRK	0.833333	1	0.714286
SLC39A4 AND NOT-MMP24 AND TACSTD2	0.846154	0.916667	0.785714
CDH17 AND TACSTD2 AND NOT-CD300LG	0.88	1	0.785714
SLC39A4 AND NOT-ST6GALNAC6 AND NOT-SLC30A10	0.846154	0.916667	0.785714
SLC39A4 AND CXCL16 AND NOT-DUOX1	0.8	0.909091	0.714286
SLC39A4 AND NOT-CD1A AND CDH3	0.923077	1	0.857143
SLC39A4 AND NOT-MMP24 AND CDH3	0.8	0.909091	0.714286
SLC39A4 AND CXCL16 AND NOT-ATP10A	0.8	0.909091	0.714286
SLC39A4 AND CXCL16 AND NOT-PTH1R	0.833333	1	0.714286
SLC39A4 AND NOT-CNGA4 AND TACSTD2	0.833333	1	0.714286
SLC39A4 AND NOT-ST6GALNAC6 AND NOT-CALHM1	0.833333	1	0.714286
SLC39A4 AND CXCL16 AND NOT-WDR19	0.8	0.909091	0.714286
SLC39A4 AND CXCL16 AND NOT-ADCY4	0.833333	1	0.714286
SLC39A4 AND NOT-ATP8B2 AND OSMR	0.833333	1	0.714286
SLC39A4 AND NOT-CNGA4 AND F2RL2	0.833333	1	0.714286
SLC39A4 AND NOT-MMP24 AND F2RL2	0.846154	0.916667	0.785714
ST14 AND APOLD1 AND NOT-CD36	0.846154	0.916667	0.785714
SLC39A4 AND NOT-ST6GALNAC6 AND NOT-KCNH4	0.814815	0.846154	0.785714
SLC39A4 AND NOT-ATP8B2 AND F2RL2	0.866667	0.8125	0.928571
SLC39A4 AND NOT-ATP8B2 AND F2RL2	0.866667	0.8125	0.928571
SLC39A4 AND CXCL16 AND NOT-ST3GAL5	0.833333	1	0.714286
SLC39A4 AND NOT-ATP8B2 AND TLR2	0.833333	1	0.714286
SLC39A4 AND NOT-KCNH4 AND ANO1	0.833333	1	0.714286
SLC39A4 AND LRRC8E AND NOT-ANO8	0.846154	0.916667	0.785714
SLC39A4 AND CXCL16 AND NOT-GLDN	0.833333	1	0.714286
SLC39A4 AND NOT-ATP8B2 AND ABCC1	0.88	1	0.785714
SLC39A4 AND NOT-CDHR1 AND NOT-ATP8B2	0.83871	0.764706	0.928571
SLC39A4 AND NOT-CD1A AND F2RL2	0.962963	1	0.928571
SLC39A4 AND NOT-GJB4 AND F2RL2	0.923077	1	0.857143
SLC39A4 AND NOT-GJB4 AND F2RL2	0.923077	1	0.857143
SLC39A4 AND NOT-CD1A AND TACSTD2	0.928571	0.928571	0.928571
SLC39A4 AND CDH3 AND NOT-ATP8B2	0.857143	0.857143	0.857143
SLC39A4 AND CDH3 AND NOT-ATP8B2	0.857143	0.857143	0.857143
SLC39A4 AND NOT-GJB4 AND TACSTD2	0.888889	0.923077	0.857143
SLC39A4 AND NOT-GJB4 AND TACSTD2	0.888889	0.923077	0.857143
SLC39A4 AND NOT-CA12 AND VANGL1	0.833333	1	0.714286
SLC39A4 AND NOT-ATP8B2 AND LDLRAD3	0.833333	1	0.714286
SLC39A4 AND NOT-ST6GALNAC6 AND CLDN1	0.88	1	0.785714
ST14 AND NOT-DUOXA1 AND CLDN1	0.962963	1	0.928571
SLC39A4 AND NOT-ST6GALNAC6 AND SLC7A5	0.88	1	0.785714

Antigen Pair Logic	F1	Precision	Recall
ST8SIA1 AND DISP2 AND MUC13	0.83871	0.8125	0.866667
SSX1 AND NOT-SCARA5 AND CLDN5	0.8	0.8	0.8
ASTN1 AND MUC13 AND NOT-IL13RA1	0.888889	1	0.8
NOT-AOC3 AND CLDN5 AND VTCN1	0.846154	1	0.733333
COMPLEX-ITGAV/OMG/MUC13	0.896552	0.928571	0.866667
NOT-AOC3 AND CLDN5 AND MUC13	0.827586	0.857143	0.8
ASTN1 AND MUC13 AND NOT-ULBP2	0.888889	1	0.8
SSX1 AND NOT-EPHA2 AND CLDN5	0.827586	0.857143	0.8
ASTN1 AND RAET1E AND ST8SIA1	0.83871	0.8125	0.866667
ASTN1 AND MUC13 AND NOT-CD34	0.888889	1	0.8
COMPLEX-ITGAV/MUC13/GPR158	0.83871	0.8125	0.866667
ASTN1 AND BEST3 AND NOT-SLC5A11	0.967742	0.9375	1
ASTN1 AND BEST3 AND NOT-PTPRT	0.967742	0.9375	1
ASTN1 AND BEST3 AND NOT-SLC30A10	0.967742	0.9375	1
ASTN1 AND BEST3 AND NOT-CNTNAP4	0.967742	0.9375	1
ASTN1 AND BEST3 AND NOT-GYPC	0.933333	0.933333	0.933333
ASTN1 AND BEST3 AND NOT-ANTXR2	0.967742	0.9375	1
ASTN1 AND BEST3 AND NOT-FMNL1	0.933333	0.933333	0.933333
COMPLEX-NKAIN4/SLCO1B3/SCN2A	0.888889	1	0.8
COMPLEX-CDH10/NKAIN4/SLC39A8	0.888889	1	0.8
OMG AND BEST3 AND NOT-SLC30A10	0.967742	0.9375	1
OPCML AND BEST3 AND NOT-CD1B	0.882353	0.789474	1
CDH10 AND BEST3 AND NOT-SLC30A10	0.9375	0.882353	1
GPR158 AND BEST3 AND NOT-SLC39A8	0.9375	0.882353	1
OPCML AND BEST3 AND NOT-SLC39A8	0.909091	0.833333	1
ASTN1 AND F2R AND NOT-CD163	0.888889	1	0.8
BEST3 AND SCN2A AND NOT-SLC5A10	0.967742	0.9375	1
OMG AND BEST3 AND NOT-SLCO1B3	0.857143	0.923077	0.8
COMPLEX-NKAIN4/SLCO1B3/KCNC1	0.846154	1	0.733333
ASTN1 AND F2R AND NOT-ANTXR2	0.846154	1	0.733333
COMPLEX-NKAIN4/CD93/KCNC1	0.846154	1	0.733333
ASTN1 AND F2R AND NOT-GYPC	0.846154	1	0.733333
COMPLEX-ANTXR2/NKAIN4/SCN2A	0.846154	1	0.733333
COMPLEX-NKAIN4/ATP2B2/DISP2	0.846154	1	0.733333
ASTN1 AND F2R AND SLC30A10	0.846154	1	0.733333
ASTN1 AND TNFRSF10B AND NOT-GYPC	0.888889	1	0.8
OMG AND TNFRSF10B AND NOT-SLC30A10	0.83871	0.8125	0.866667
OMG AND TNFRSF10B AND SLCO1B3	0.866667	0.866667	0.866667
OMG AND TNFRSF10B AND CD1B	0.83871	0.8125	0.866667
COMPLEX-SEZ6/KCNQ2/SLC30A1	0.888889	1	0.8
ASTN1 AND TNFRSF10B AND NOT-ANTXR2	0.928571	1	0.866667
COMPLEX-SEZ6/KCNQ2/TNFRSF10B	0.928571	1	0.866667
ASTN1 AND F2R AND LRRTM4	0.888889	1	0.8
CDH10 AND F2R AND SLC30A10	0.888889	1	0.8
BEST3 AND DISP2 AND NOT-SLC13A2	0.827586	0.857143	0.8
OPCML AND BEST3 AND NOT-SLCO1B3	0.827586	0.857143	0.8
ASTN1 AND F2R AND PTPRT	0.888889	1	0.8
COMPLEX-SEMA4B/SEZ6/KCNQ2	0.928571	1	0.866667
COMPLEX-SEZ6/KCNQ2/GPR37L1	0.823529	0.736842	0.933333
COMPLEX-SEZ6/BEST3/KCNQ2	0.9375	0.882353	1
OPCML AND F2R AND NOT-SLC39A8	0.888889	1	0.8
ASTN1 AND TNFRSF10B AND NOT-CD163	0.928571	1	0.866667
CDH10 AND TNFRSF10B AND SLC30A10	0.928571	1	0.866667
ASTN1 AND CXCL16 AND NOT-GYPC	0.83871	0.8125	0.866667
ASTN1 AND GJD2 AND NOT-PTPRT	0.888889	1	0.8
ASTN1 AND TNFRSF10B AND PTPRT	0.928571	1	0.866667
ASTN1 AND NOT-SLC12A5 AND SLCO1B3	0.814815	0.916667	0.733333
DTNA AND NOT-SLC5A11 AND ASTN1	0.896552	0.928571	0.866667
ASTN1 AND TNFRSF10B AND NOT-SLC31A1	0.846154	1	0.733333
ASTN1 AND TNFRSF10B AND NOT-SLC30A10	0.928571	1	0.866667
ASTN1 AND TNFRSF10B AND LRRTM4	0.928571	1	0.866667
CDH10 AND NOT-DGKE AND SLCO1B3	0.814815	0.916667	0.733333
COMPLEX-SEZ6/F2R/KCNQ2	0.928571	1	0.866667
ASTN1 AND CXCL16 AND NOT-CNTNAP4	0.875	0.823529	0.933333
ASTN1 AND GJD2 AND NOT-SLC30A10	0.888889	1	0.8
ASTN1 AND F2R AND SLCO1B3	0.888889	1	0.8
OMG AND CD1B AND PPAPDC1B	0.857143	0.923077	0.8
ASTN1 AND CXCL16 AND PTPRT	0.875	0.823529	0.933333
ASTN1 AND TNFRSF10B AND SLCO1B3	0.928571	1	0.866667
COMPLEX-SEZ6/KCNQ2/P2RX4	0.810811	0.681818	1
ASTN1 AND TSHR AND NOT-SLC30A10	0.8	0.8	0.8
OPCML AND F2R AND NOT-CD1B	0.8	0.8	0.8
BEST3 AND SLC24A2 AND NOT-SLCO1B3	0.8	0.8	0.8
ASTN1 AND TSHR AND NOT-PTPRT	0.8	0.8	0.8
COMPLEX-NKAIN4/CD93/SCN2A	0.8	1	0.666667
ASTN1 AND TSHR AND NOT-SLCO1B3	0.8	0.8	0.8
CDH10 AND NOT-ATP8A2 AND SLCO1B3	0.8	1	0.666667

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
SLC39A4 AND CLDN1 AND NOT-GPR146	0.857143	0.857143	0.857143
ST14 AND NOT-DUOX1 AND CLDN1	0.962963	1	0.928571
SLC39A4 AND CLDN1 AND NOT-FGFR1	0.888889	0.923077	0.857143
PTPRK AND NOT-WDR19 AND CLDN1	0.833333	1	0.714286
SLC39A4 AND NOT-GPR146 AND SLC7A5	0.846154	0.916667	0.785714
SLC39A4 AND NOT-CA12 AND TPBG	0.814815	0.846154	0.785714
SLC39A4 AND SLC7A5 AND NOT-ANO8	0.928571	0.928571	0.928571
SLC39A4 AND SLC7A5 AND NOT-ANO8	0.928571	0.928571	0.928571
SLC39A4 AND CLDN1 AND NOT-ADCY4	0.827586	0.8	0.857143
SLC39A4 AND CLDN1 AND NOT-ADCY4	0.827586	0.8	0.857143
SLC39A4 AND NOT-ST6GALNAC6 AND NOT-ABCA5	0.814815	0.846154	0.785714
SLC39A4 AND NOT-IL11RA AND F2RL2	0.866667	0.8125	0.928571
SLC39A4 AND NOT-IL11RA AND F2RL2	0.866667	0.8125	0.928571
EPCAM AND NOT-ABHD6 AND ST14	0.8	0.909091	0.714286
EPCAM AND NOT-ABHD6 AND GJB2	0.8	0.909091	0.714286
SLC39A4 AND CLDN1 AND NOT-CADM1	0.833333	1	0.714286
SLC39A4 AND NOT-ADCY9 AND CLDN1	0.846154	0.916667	0.785714
SLC39A4 AND CLDN1 AND NOT-TGFB3	0.814815	0.846154	0.785714
SLC39A4 AND CLDN1 AND NOT-PTH1R	0.846154	0.916667	0.785714
SLC39A4 AND NOT-CDHR1 AND NOT-IL11RA	0.866667	0.8125	0.928571
SLC39A4 AND NOT-SCTR AND CLDN1	0.833333	1	0.714286
SLC39A4 AND CXCL16 AND NOT-ATP8B2	0.833333	1	0.714286
SLC39A4 AND NOT-CNGA4 AND NOT-SLC30A10	0.8	0.909091	0.714286
SLC39A4 AND NOT-CNGA4 AND SLC6A6	0.833333	1	0.714286
SLC39A4 AND NOT-MMP24 AND SLC6A6	0.846154	0.916667	0.785714
SLC39A4 AND NOT-KCNH4 AND LRRC8E	0.857143	0.857143	0.857143
SLC39A4 AND NOT-SLC30A10 AND CDH17	0.857143	0.857143	0.857143
SLC39A4 AND NOT-CD1A AND NOT-SLC30A10	0.866667	0.8125	0.928571
SLC39A4 AND NOT-CD1A AND SLC6A6	0.928571	0.928571	0.928571
SLC39A4 AND NOT-GJB4 AND LRRC8E	0.88	1	0.785714
SLC39A4 AND CXCL16 AND NOT-SHISA9	0.833333	1	0.714286
SLC39A4 AND CXCL16 AND NOT-CHRNG	0.8	0.909091	0.714286
SLC39A4 AND NOT-KCNK4 AND SLC6A6	0.814815	0.846154	0.785714
SLC39A4 AND CXCL16 AND FUT3	0.833333	1	0.714286
SLC39A4 AND CXCL16 AND NOT-GPR22	0.8	0.909091	0.714286
SLC39A4 AND NOT-MMP24 AND SLC7A5	0.846154	0.916667	0.785714
SLC39A4 AND NOT-STAB1 AND CLDN1	0.8	0.909091	0.714286
SLC39A4 AND NOT-MMP24 AND CLDN1	0.88	1	0.785714
SLC39A4 AND CXCL16 AND NOT-IL11RA	0.8	0.909091	0.714286
SLC39A4 AND NOT-CD1A AND CLDN1	0.923077	1	0.857143
CDH17 AND CLDN1 AND NOT-CD300LG	0.88	1	0.785714
SLC39A4 AND NOT-ATP8B2 AND CLDN1	0.923077	1	0.857143
SLC39A4 AND NOT-ATP8B2 AND CLDN1	0.923077	1	0.857143
SLC39A4 AND NOT-MMP24 AND NOT-GPA33	0.8	0.909091	0.714286
SLC39A4 AND CLDN1 AND NOT-FXYD6	0.888889	0.923077	0.857143
SLC39A4 AND NOT-ATP8B2 AND SLC7A5	0.928571	0.928571	0.928571
SLC39A4 AND NOT-ATP8B2 AND SLC7A5	0.928571	0.928571	0.928571
SLC39A4 AND NOT-CNGA4 AND SLC7A5	0.8	0.909091	0.714286
SLC39A4 AND NOT-ATP8B2 AND SPON2	0.866667	0.8125	0.928571
CDH17 AND CLDN1 AND NOT-WDR19	0.962963	1	0.928571
SLC39A4 AND NOT-CNGA4 AND NOT-GPA33	0.833333	1	0.714286
SLC39A4 AND NOT-GPA33 AND CDH17	0.88	1	0.785714
SLC39A4 AND NOT-CD1A AND SLC7A5	0.962963	1	0.928571
SLC39A4 AND CXCL16 AND EPCAM	0.833333	1	0.714286
EPCAM AND NOT-SLC22A5 AND GJB2	0.88	1	0.785714
SLC39A4 AND NOT-SEZ6 AND SLC7A5	0.8	0.909091	0.714286
CDH17 AND CLDN1 AND NOT-GPR1	0.923077	1	0.857143
SLC39A4 AND NOT-KCNH4 AND CLDN1	0.923077	1	0.857143
CDH17 AND CLDN1 AND NOT-CDON	0.962963	1	0.928571
SLC39A4 AND SLC7A5 AND NOT-FXYD6	0.896552	0.866667	0.928571
EPCAM AND NOT-SLC22A5 AND SEMA4B	0.814815	0.846154	0.785714
COMPLEX-PROCR/CD82/EPCAM	0.833333	1	0.714286
EPCAM AND SLC7A5 AND NOT-FXYD6	0.8125	0.722222	0.928571
EPCAM AND SLC7A5 AND NOT-SHISA9	0.896552	0.866667	0.928571
EPCAM AND SLC7A5 AND FUT3	0.928571	0.928571	0.928571
EPCAM AND SDC1 AND NOT-FXYD6	0.88	1	0.785714
EPCAM AND SDC1 AND NOT-AOC3	0.833333	1	0.714286
EPCAM AND SLC7A5 AND SMPD2	0.833333	1	0.714286
EPCAM AND SDC1 AND NOT-SHISA9	0.88	1	0.785714
EPCAM AND SDC1 AND FUT3	0.88	1	0.785714
COMPLEX-PROCR/EPCAM/SDC1	0.923077	1	0.857143
EPCAM AND SDC1 AND SEMA4B	0.833333	1	0.714286
EPCAM AND THY1 AND SMPD2	0.8	0.909091	0.714286
EPCAM AND PROCR AND CLDN1	0.962963	1	0.928571
EPCAM AND CLDN1 AND FUT3	0.923077	1	0.857143
EPCAM AND F2R AND NOT-IL11RA	0.8125	0.722222	0.928571
EPCAM AND SLC7A5 AND NOT-FGFR1	0.896552	0.866667	0.928571

Antigen Pair Logic	F1	Precision	Recall
ASTN1 AND F2R AND NOT-SLC31A1	0.8	1	0.666667
COMPLEX-SEZ6/KCNQ2/CXCL16	0.8	0.7	0.933333
SEZ6 AND SLC30A1 AND NOT-GHR	0.8	1	0.666667
ASTN1 AND CXCL16 AND NOT-SLC5A11	0.903226	0.875	0.933333
CDH10 AND NOT-CACNA1B AND SLCO1B3	0.8	1	0.666667
CDH10 AND NOT-KIAA0319 AND SLCO1B3	0.8	1	0.666667
COMPLEX-NKAIN4/SLCO1B3/ASTN1	0.8	1	0.666667
ASTN1 AND F2R AND CNTNAP4	0.846154	1	0.733333
ASTN1 AND TSHR AND NOT-ANTXR2	0.8	0.8	0.8
ASTN1 AND CXCL16 AND LRRTM4	0.875	0.823529	0.933333
COMPLEX-NKAIN4/CD93/DISP2	0.8	1	0.666667
COMPLEX-NKAIN4/SLCO1B3/OPCML	0.8	1	0.666667
ASTN1 AND CXCL16 AND NOT-CD163	0.875	0.823529	0.933333
ASTN1 AND SCN10A AND NOT-CD163	0.827586	0.857143	0.8
IGDCC3 AND ASTN1 AND NOT-PTPRT	0.965517	1	0.933333
ASTN1 AND GJD2 AND NOT-SLC31A1	0.888889	1	0.8
CDH10 AND SLCO1B3 AND NOT-SYT4	0.846154	1	0.733333
ASTN1 AND BEST2 AND NOT-SLC30A10	0.8	0.7	0.933333
ASTN1 AND BEST2 AND NOT-PTPRT	0.8	0.7	0.933333
COMPLEX-BTN3A3/SEZ6/KCNQ2	0.965517	1	0.933333
ASTN1 AND TNFRSF10B AND NOT-CNTNAP4	0.928571	1	0.866667
ASTN1 AND TNFRSF10B AND NOT-SLC5A11	0.928571	1	0.866667
COMPLEX-SEZ6/CACNG4/KCNQ2	0.848485	0.777778	0.933333
ASTN1 AND CDH11 AND NOT-CNTNAP4	0.8	0.8	0.8
DISP2 AND NOT-PTGER4 AND SEMA4B	0.8	0.8	0.8
ASTN1 AND SLC22A11 AND NOT-SLC30A10	0.827586	0.857143	0.8
DLL3 AND NOT-TRPM4 AND SLAMF7	0.846154	1	0.733333
SSX1 AND NOT-CLDN1 AND CLDN5	0.827586	0.857143	0.8
ST8SIA1 AND NOT-EPCAM AND CLDN5	0.827586	0.857143	0.8
ST8SIA1 AND NOT-IL20RA AND CLDN5	0.83871	0.8125	0.866667
ST8SIA1 AND CLDN5 AND NOT-MST1R	0.83871	0.8125	0.866667
ST8SIA1 AND NOT-TRPM4 AND CD34	0.83871	0.8125	0.866667
ST8SIA1 AND CLDN5 AND NOT-PMEL	0.83871	0.8125	0.866667
ST8SIA1 AND CLDN5 AND NOT-ITGB6	0.83871	0.8125	0.866667
ST8SIA1 AND CLDN5 AND NOT-PROM1	0.8	0.8	0.8
ST8SIA1 AND NOT-TRPM4 AND NOT-SLAMF7	0.83871	0.8125	0.866667
Ovarian Cancer			
KCNK15 AND NOT-BMP2 AND NOT-FPR1	1	1	1
KCNK15 AND NOT-BMP2 AND NOT-KCNQ4	1	1	1
KCNK15 AND NOT-BMP2 AND NOT-FPR1	1	1	1
KCNK15 AND NOT-BMP2 AND NOT-KCNQ4	1	1	1
KCNK15 AND NOT-BMP2 AND NOT-ITM2A	1	1	1
KCNK15 AND NOT-BMP2 AND NOT-PTH1R	1	1	1
KCNK15 AND NOT-BMP2 AND NOT-TNFSF12	1	1	1
KCNK15 AND NOT-BMP2 AND ADRA1B	1	1	1
KCNK15 AND NOT-BMP2 AND NOT-PDGFRB	1	1	1
KCNK15 AND NOT-BMP2 AND NOT-ITGA5	1	1	1
KCNK15 AND NOT-BMP2 AND NOT-PDGFRB	1	1	1
KCNK15 AND NOT-BMP2 AND NOT-ITGA5	1	1	1
KCNK15 AND NOT-BMP2 AND SLC51B	1	1	1
KCNK15 AND NOT-BMP2 AND GJD4	1	1	1
KCNK15 AND NOT-BMP2 AND NOT-FGFR4	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-LRPAP1	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-IL7R	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-HMOX2	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-PNPLA2	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-ICAM3	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-ANO9	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND DPP6	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-ADRB2	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-CXCR2	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-CPT1C	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-MFSD10	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND SLC17A4	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-STOM	0.96	0.923077	1
KCNK15 AND NOT-DSC3 AND NOT-ADRB2	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-STX8	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND SCTR	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-CSF3R	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-CYP4F12	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-CCR1	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-FNDC4	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-SLC1A4	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-TGOLN2	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-TNFRSF1B	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-TMED10	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-CDHR3	0.96	0.923077	1

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
EPCAM AND SLC7A5 AND NOT-CD300LG	0.827586	0.8	0.857143
EPCAM AND SLC7A5 AND NOT-ATP10A	0.8125	0.722222	0.928571
EPCAM AND SLC7A5 AND GJB2	0.962963	1	0.928571
SLC39A4 AND NOT-IL11RA AND CLDN1	0.888889	0.923077	0.857143
SLC39A4 AND NOT-IL11RA AND CLDN1	0.888889	0.923077	0.857143
EPCAM AND SLC7A5 AND NOT-SLC15A2	0.83871	0.764706	0.928571
SLC39A4 AND NOT-ABCA5 AND CLDN1	0.846154	0.916667	0.785714
SLC39A4 AND NOT-ABCA5 AND CLDN1	0.846154	0.916667	0.785714
EPCAM AND SLC7A5 AND NOT-WDR19	0.896552	0.866667	0.928571
EPCAM AND SLC7A5 AND PPAP2C	0.923077	1	0.857143
EPCAM AND SLC7A5 AND NOT-MARVELD1	0.827586	0.8	0.857143
SLC39A4 AND SLC7A5 AND EPCAM	0.928571	0.928571	0.928571
SLC39A4 AND SLC7A5 AND EPCAM	0.928571	0.928571	0.928571
SLC39A4 AND CLDN1 AND EPCAM	0.827586	0.8	0.857143
EPCAM AND SLC7A5 AND ACVR1C	0.857143	0.857143	0.857143
EPCAM AND SLC7A5 AND ATP1B1	0.827586	0.8	0.857143
EPCAM AND SLC7A5 AND LGR4	0.8125	0.722222	0.928571
SLC39A4 AND NOT-ABCA5 AND SLC7A5	0.857143	0.857143	0.857143
SLC39A4 AND NOT-ABCA5 AND SLC7A5	0.857143	0.857143	0.857143
SLC39A4 AND NOT-IL11RA AND SPON2	0.83871	0.764706	0.928571
EPCAM AND GJB2 AND CLDN1	0.962963	1	0.928571
SLC39A4 AND NOT-IL11RA AND SLC7A5	0.928571	0.928571	0.928571
SLC39A4 AND NOT-IL11RA AND SLC7A5	0.928571	0.928571	0.928571
EPCAM AND SLC7A5 AND ST14	0.866667	0.8125	0.928571
MUC13 AND CLDN1 AND NOT-WDR19	0.928571	0.928571	0.928571
EPCAM AND SLC7A5 AND NOT-BMP2	0.8125	0.722222	0.928571
EPCAM AND SLC7A5 AND NOT-SLC13A4	0.928571	0.928571	0.928571
EPCAM AND SLC7A5 AND NOT-GRIN3B	0.814815	0.846154	0.785714
EPCAM AND SLC7A5 AND NOT-GRIK2	0.896552	0.866667	0.928571
SLC39A4 AND NOT-GPA33 AND MUC13	0.88	1	0.785714
EPCAM AND SLC7A5 AND GPR35	0.866667	0.8125	0.928571
EPCAM AND SLC7A5 AND NOT-IL11RA	0.83871	0.764706	0.928571
EPCAM AND SLC7A5 AND NOT-FOLR1	0.962963	1	0.928571
EPCAM AND SLC7A5 AND NOT-SLC34A2	0.8125	0.722222	0.928571
EPCAM AND SLC7A5 AND NOT-GPNMB	0.827586	0.8	0.857143
EPCAM AND SDC1 AND NOT-IL11RA	0.88	1	0.785714
EPCAM AND THY1 AND NOT-IL11RA	0.846154	0.916667	0.785714
EPCAM AND SDC1 AND NOT-FOLR1	0.88	1	0.785714
EPCAM AND SDC1 AND NOT-BMPR1B	0.88	1	0.785714
EPCAM AND SDC1 AND NOT-SLC34A2	0.88	1	0.785714
EPCAM AND THY1 AND NOT-FOLR1	0.866667	0.8125	0.928571
EPCAM AND NOT-FOLR1 AND CLDN1	0.962963	1	0.928571
EPCAM AND NOT-SLC34A2 AND CLDN1	0.962963	1	0.928571
EPCAM AND NOT-FOLR1 AND SPON2	0.83871	0.764706	0.928571
EPCAM AND CLDN1 AND NOT-BMPR1B	0.896552	0.866667	0.928571
Glioblastoma			
NLGN4X AND LAPTM5 AND NOT-PTPRM	0.903226	1	0.823529
GPM6A AND LAPTM5 AND P2RY1	0.914286	0.888889	0.941176
NRCAM AND NOT-PTPRM AND LAPTM5	0.9375	1	0.882353
PTPRZ1 AND SIRPG AND SYT11	0.9375	1	0.882353
GPM6B AND LAPTM5 AND RHCG	0.969697	1	0.941176
GPM6A AND LAPTM5 AND RHCG	0.969697	1	0.941176
NRCAM AND NOT-PTPRM AND BTN3A3	0.882353	0.882353	0.882353
NLGN4X AND LAPTM5 AND NOT-YIPF3	0.903226	1	0.823529
NLGN4X AND LAPTM5 AND NOT-ZDHHC5	0.903226	1	0.823529
PTPRZ1 AND LAPTM5 AND NOT-YIPF3	0.9375	1	0.882353
PTPRZ1 AND NOT-STX8 AND BST2	0.875	0.933333	0.823529
PTPRZ1 AND NOT-STX8 AND ATP7A	0.903226	1	0.823529
NRCAM AND NOT-PTPRM AND PPAPDC1B	0.903226	1	0.823529
GPM6A AND SLC2A10 AND NOT-FLVCR1	0.903226	1	0.823529
PTPRZ1 AND LAPTM5 AND NOT-SDC2	0.903226	1	0.823529
PTPRZ1 AND LAPTM5 AND NOT-ZDHHC5	0.9375	1	0.882353
GPM6B AND LAPTM5 AND NOT-APLP1	0.969697	1	0.941176
GRIA3 AND KCNK15 AND CDH11	0.882353	0.882353	0.882353
GPM6B AND F2R AND NOT-TSPAN14	0.866667	1	0.764706
GPM6B AND F2R AND CD81	0.866667	1	0.764706
GPM6B AND F2R AND OR1A1	0.866667	1	0.764706
GPM6B AND F2R AND NOT-CDHR2	0.866667	1	0.764706
GPM6B AND F2R AND NOT-TMEM127	0.866667	1	0.764706
GPM6B AND F2R AND NOT-KCNMB2	0.866667	1	0.764706
GPM6B AND F2R AND NOT-SLC44A3	0.866667	1	0.764706
COMPLEX-F2R/GPM6B/KITLG	0.866667	1	0.764706
GPM6B AND F2R AND NOT-CNNM2	0.866667	1	0.764706
GPM6B AND F2R AND C3AR1	0.866667	1	0.764706
GPM6B AND F2R AND NOT-SLC12A6	0.866667	1	0.764706
GPM6B AND F2R AND NOT-HMOX2	0.866667	1	0.764706
GPM6B AND F2R AND NOT-GDE1	0.866667	1	0.764706

Antigen Pair Logic	F1	Precision	Recall
KCNK15 AND NOT-BMP2 AND NOT-CD8B	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-PIEZO1	0.96	0.923077	1
COMPLEX-PCDHB3/KCNK15/BMP2	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-AQP4	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND SCTR	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-PTPRCAP	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-ATP2C2	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-ATP2C2	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND MARCH1	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-SLC39A4	0.96	0.923077	1
COMPLEX-PTH2R/KCNK15/BMP2	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-POPDC2	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-CAV1	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-RHOT2	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-TCIRG1	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-SLC23A1	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-LILRB3	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-CSF3R	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND TNFRSF25	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-NUCB2	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-C10orf54	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-SELPLG	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-ABCA3	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-CDHR3	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-SCN4B	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-BBS4	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-CD3D	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-TNFRSF14	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-PCDH9	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-DNAJC1	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-CX3CL1	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-SELPLG	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-RHBG	0.96	0.923077	1
KCNK15 AND NOT-SLC31A2 AND NOT-CDHR3	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-VAMP1	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-IGSF9	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-C5AR1	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-PNPLA2	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-CD2	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-DHRS3	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-ENTPD2	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-PIEZO1	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-FCGR3B	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-ICAM3	0.96	0.923077	1
KCNK15 AND NOT-SLC31A2 AND CELSR3	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-IL7R	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-COMT	0.96	0.923077	1
KCNK15 AND NOT-SLC31A2 AND TNFRSF25	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-CD8B	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-GPR20	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-GGT5	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-PCDH9	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-MFSD3	0.96	0.923077	1
KCNK15 AND NOT-SLC31A2 AND CELSR3	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-TMED10	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND MARCH1	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-SLC29A4	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-ADCY9	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-ZFYVE27	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-CSF1R	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND GAGE1	1	1	1
KCNK15 AND NOT-BMP2 AND NOT-GPC3	1	1	1
KCNK15 AND NOT-BMP2 AND HLA-DOB	1	1	1
KCNK15 AND NOT-BMP2 AND NOT-PROM1	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-ANXA1	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-PROM1	0.96	0.923077	1
COMPLEX-KCNK15/BMP2/VTCN1	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND GPNMB	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-CD52	0.96	0.923077	1
COMPLEX-KCNK15/BMP2/CLDN6	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND BMPR1B	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-TRPM4	0.96	0.923077	1
COMPLEX-ALK/KCNK15/BMP2	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-SPON2	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-CD52	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-SDC1	0.96	0.923077	1
COMPLEX-IL11RA/KCNK15/BMP2	0.96	0.923077	1

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
NLGN4X AND LAPTM5 AND NOT-SDC2	0.866667	1	0.764706
GPM6B AND F2R AND NOT-RHCG	0.866667	1	0.764706
GPM6B AND F2R AND NOT-CRIM1	0.866667	1	0.764706
COMPLEX-F2R/SLC16A9/GPM6B	0.866667	1	0.764706
GPM6B AND F2R AND NOT-ADRA2C	0.866667	1	0.764706
GPM6B AND F2R AND TMEM119	0.866667	1	0.764706
GPM6B AND F2R AND NOT-UNC5D	0.866667	1	0.764706
GPM6B AND F2R AND PTCHD1	0.866667	1	0.764706
GPM6B AND F2R AND NOT-EPHA4	0.866667	1	0.764706
GPM6B AND F2R AND NOT-TMEM65	0.866667	1	0.764706
GPM6A AND ABCA1 AND ADAM29	0.866667	1	0.764706
GPM6B AND F2R AND SIRPG	0.866667	1	0.764706
GPM6B AND F2R AND NOT-THSD7A	0.866667	1	0.764706
GPM6B AND F2R AND NOT-MFAP3L	0.866667	1	0.764706
GPM6B AND F2R AND NOT-TENM1	0.866667	1	0.764706
NRCAM AND LAPTM5 AND NOT-YIPF3	0.9375	1	0.882353
NLGN4X AND LAPTM5 AND NOT-CD40	0.903226	1	0.823529
NLGN4X AND LAPTM5 AND NOT-SLC12A6	0.903226	1	0.823529
NLGN4X AND LAPTM5 AND NOT-ITGB7	0.903226	1	0.823529
NLGN4X AND LAPTM5 AND NOT-LILRB2	0.903226	1	0.823529
NLGN4X AND LAPTM5 AND NOT-PNPLA2	0.903226	1	0.823529
GPM6A AND ADAM29 AND DISP1	0.9375	1	0.882353
GPM6B AND LAPTM5 AND KITLG	0.864865	0.8	0.941176
GPM6A AND WLS AND ADAM29	0.888889	0.842105	0.941176
NRCAM AND NOT-VAMP1 AND BTN3A3	0.875	0.933333	0.823529
NLGN4X AND LAPTM5 AND NOT-ITSN1	0.903226	1	0.823529
GPM6A AND VAMP8 AND RHCG	0.909091	0.9375	0.882353
GPM6A AND TLR2 AND ADAM29	0.909091	0.9375	0.882353
NRCAM AND NOT-STX8 AND ATP7A	0.875	0.933333	0.823529
PTPRZ1 AND NOT-UNC5D AND SYT11	0.909091	0.9375	0.882353
GPM6A AND HLA-B AND ADAM29	0.857143	0.833333	0.882353
GPM6B AND LAPTM5 AND P2RY1	0.857143	0.833333	0.882353
GPM6B AND VAMP8 AND RHCG	0.857143	0.833333	0.882353
NRCAM AND LAPTM5 AND NOT-ZDHHC5	0.9375	1	0.882353
PTPRZ1 AND LAPTM5 AND NOT-PNPLA2	0.9375	1	0.882353
NRCAM AND TLR2 AND NOT-S1PR1	0.9375	1	0.882353
GPM6A AND ATRAID AND ADAM29	0.866667	1	0.764706
GPM6A AND SLC2A10 AND NOT-ZDHHC2	0.9375	1	0.882353
GPM6A AND SLC2A10 AND ADAM29	0.9375	1	0.882353
NRCAM AND LAPTM5 AND NOT-PNPLA2	0.9375	1	0.882353
PTPRZ1 AND NOT-ATP2C1 AND GPR158	0.875	0.933333	0.823529
MLC1 AND SLC7A7 AND NOT-SLC12A6	0.848485	0.875	0.823529
MLC1 AND TLR2 AND NOT-LTB	0.848485	0.875	0.823529
GPM6A AND NOT-ATP8A2 AND RHCG	0.848485	0.875	0.823529
SYT11 AND ADAM12 AND NOT-HMOX2	0.848485	0.875	0.823529
MLC1 AND TLR2 AND NOT-GPR146	0.848485	0.875	0.823529
GPR158 AND SLC30A5 AND NOT-ATP2C1	0.848485	0.875	0.823529
GPR158 AND ABCA1 AND NOT-SYT2	0.848485	0.875	0.823529
PTPRZ1 AND NOT-TM9SF2 AND NOT-GPR158	0.875	0.933333	0.823529
GPM6B AND TCIRG1 AND SLC4A8	0.848485	0.875	0.823529
AQP4 AND RHCG AND NOT-SLC16A5	0.866667	1	0.764706
NRCAM AND LAPTM5 AND NOT-ITSN1	0.9375	1	0.882353
NRCAM AND LAPTM5 AND NOT-SLC12A6	0.9375	1	0.882353
NRCAM AND LAPTM5 AND NOT-LILRB2	0.9375	1	0.882353
GPM6A AND VAMP8 AND P2RY1	0.857143	0.833333	0.882353
MLC1 AND SLC7A7 AND NOT-ITGB7	0.848485	0.875	0.823529
MLC1 AND SLC2A10 AND NOT-ABCC10	0.866667	1	0.764706
MLC1 AND SLC2A10 AND NOT-LEPR	0.866667	1	0.764706
MLC1 AND SLC2A10 AND NOT-STS	0.866667	1	0.764706
MLC1 AND SLC2A10 AND NOT-EPHA1	0.866667	1	0.764706
PTPRZ1 AND NOT-PIK3IP1 AND ITGAM	0.875	0.933333	0.823529
FAT3 AND KCNK15 AND LAPTM5	0.842105	0.761905	0.941176
GPM6B AND SIRPG AND CD163	0.842105	0.761905	0.941176
SYT11 AND SLC7A7 AND NOT-ITGB7	0.842105	0.761905	0.941176
SYT11 AND SIRPG AND WLS	0.918919	0.85	1
PTPRZ1 AND NOT-VAMP1 AND BTN3A3	0.903226	1	0.823529
NRCAM AND CYBA AND NOT-TSPAN5	0.9375	1	0.882353
PTPRZ1 AND NOT-SLC2A12 AND FZD7	0.909091	0.9375	0.882353
PTPRZ1 AND NOT-LRP11 AND ZDHHC5	0.903226	1	0.823529
NRCAM AND RHCG AND NOT-PARM1	0.909091	0.9375	0.882353
PTPRZ1 AND SIRPG AND GPM6B	0.9375	1	0.882353
NRCAM AND CYBA AND NOT-RER1	0.9375	1	0.882353
PTPRZ1 AND NOT-SLC2A12 AND RHCG	0.9375	1	0.882353
PTPRZ1 AND SIRPG AND NOT-ADIPOR1	0.9375	1	0.882353
PTPRZ1 AND NOT-SLC2A12 AND NOT-TSPAN5	0.909091	0.9375	0.882353
GPM6A AND SLC2A10 AND NOT-PTPRM	0.9375	1	0.882353
PTPRZ1 AND KCNK15 AND FAT3	0.903226	1	0.823529

Antigen Pair Logic	F1	Precision	Recall
COMPLEX-CEACAM6/KCNK15/BMP2	0.96	0.923077	1
COMPLEX-MST1R/KCNK15/BMP2	0.96	0.923077	1
COMPLEX-KCNK15/BMP2/LGR5	0.96	0.923077	1
COMPLEX-KCNK15/BMP2/TPBG	0.96	0.923077	1
COMPLEX-IGF1R/KCNK15/BMP2	0.96	0.923077	1
COMPLEX-MET/KCNK15/BMP2	0.96	0.923077	1
COMPLEX-KCNK15/CXCR5/BMP2	0.96	0.923077	1
COMPLEX-SLC31A2/KCNK15/CLDN6	0.96	0.923077	1
COMPLEX-RNF43/KCNK15/BMP2	0.96	0.923077	1
COMPLEX-STEAP1/KCNK15/BMP2	0.96	0.923077	1
COMPLEX-ALDH1A1/KCNK15/BMP2	0.96	0.923077	1
COMPLEX-KCNK15/BMP2/MUC16	0.96	0.923077	1
COMPLEX-ERBB3/KCNK15/BMP2	0.96	0.923077	1
COMPLEX-DKK1/KCNK15/BMP2	0.96	0.923077	1
COMPLEX-KCNK15/BMP2/SSTR2	0.96	0.923077	1
COMPLEX-KCNK15/BMP2/CLDN9	0.96	0.923077	1
COMPLEX-KCNK15/BMP2/CD19	0.96	0.923077	1
COMPLEX-ERBB4/KCNK15/BMP2	0.96	0.923077	1
COMPLEX-SLC31A2/KCNK15/VTCN1	0.96	0.923077	1
COMPLEX-ABCA5/KCNK15/BMP2	0.96	0.923077	1
COMPLEX-IL2RA/KCNK15/BMP2	0.96	0.923077	1
COMPLEX-KCNK15/BMP2/FCRL5	0.96	0.923077	1
COMPLEX-DNAJB8/KCNK15/BMP2	0.96	0.923077	1
KCNK15 AND NOT-DSC3 AND GAGE1	1	1	1
KCNK15 AND NOT-DSC3 AND NOT-ENG	1	1	1
KCNK15 AND NOT-BMP2 AND NOT-IL20RA	0.96	0.923077	1
KCNK15 AND NOT-DSC3 AND NOT-ENG	1	1	1
KCNK15 AND NOT-DSC3 AND NOT-CSPG4	1	1	1
KCNK15 AND NOT-SLC31A2 AND GPNMB	0.96	0.923077	1
KCNK15 AND NOT-DSC3 AND HLA-DOB	0.956522	1	0.916667
KCNK15 AND NOT-BMP2 AND NOT-CD70	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-IL20RA	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-CD70	0.96	0.923077	1
KCNK15 AND NOT-SLC31A2 AND NOT-IL20RA	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-CLDN8	0.96	0.923077	1
KCNK15 AND NOT-AQP9 AND NOT-ENG	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-CLDN8	0.96	0.923077	1
KCNK15 AND NOT-SLC31A2 AND NOT-IL20RA	0.96	0.923077	1
KCNK15 AND NOT-DSC3 AND NOT-CLEC14A	0.96	0.923077	1
COMPLEX-SLC31A2/IL2RA/KCNK15	0.96	0.923077	1
COMPLEX-SLC31A2/DKK1/KCNK15	0.96	0.923077	1
KCNK15 AND NOT-DSC3 AND NOT-TNC	0.96	0.923077	1
COMPLEX-SLC31A2/TRPM4/KCNK15	0.96	0.923077	1
COMPLEX-SLC31A2/GAGE1/KCNK15	0.96	0.923077	1
COMPLEX-SLC31A2/STEAP1/KCNK15	0.96	0.923077	1
COMPLEX-SLC31A2/HLA-DOB/KCNK15	0.96	0.923077	1
KCNK15 AND NOT-DSC3 AND NOT-AXL	1	1	1
COMPLEX-SLC31A2/IGF1R/KCNK15	0.96	0.923077	1
KCNK15 AND NOT-DSC3 AND NOT-CD34	1	1	1
COMPLEX-SLC31A2/EGFR/KCNK15	0.96	0.923077	1
KCNK15 AND NOT-DSC3 AND NOT-CLDN5	0.96	0.923077	1
COMPLEX-SLC31A2/ALDH1A1/KCNK15	0.96	0.923077	1
COMPLEX-SLC31A2/KCNK15/ST8SIA1	0.96	0.923077	1
KCNK15 AND NOT-DSC3 AND NOT-P2RX5	0.96	0.923077	1
COMPLEX-SLC31A2/ERBB3/KCNK15	0.96	0.923077	1
KCNK15 AND NOT-DSC3 AND NOT-GPC3	1	1	1
COMPLEX-SLC31A2/ERBB4/KCNK15	0.96	0.923077	1
COMPLEX-SLC31A2/KCNK15/ULBP2	0.96	0.923077	1
COMPLEX-SLC31A2/ABCA5/KCNK15	0.96	0.923077	1
COMPLEX-SLC31A2/PCYT1A/KCNK15	0.96	0.923077	1
COMPLEX-SLC31A2/KCNK15/ULBP2	0.96	0.923077	1
COMPLEX-SLC31A2/KCNK15/MUC16	0.96	0.923077	1
COMPLEX-SLC31A2/TRPM4/KCNK15	0.96	0.923077	1
COMPLEX-SLC31A2/CEACAM6/KCNK15	0.96	0.923077	1
KCNK15 AND NOT-DSC3 AND MUC16	1	1	1
KCNK15 AND NOT-SLC31A2 AND NOT-ANXA1	0.96	0.923077	1
KCNK15 AND NOT-PTGDR AND GAGE1	0.96	0.923077	1
KCNK15 AND NOT-DSC3 AND NOT-CLDN23	0.96	0.923077	1
KCNK15 AND NOT-DSC3 AND NOT-CLDN23	0.96	0.923077	1
KCNK15 AND NOT-TNFSF13B AND HLA-DOB	0.956522	1	0.916667
KCNK15 AND NOT-DSC3 AND NOT-CD52	0.96	0.923077	1
KCNK15 AND NOT-DSC3 AND NOT-FOLR2	0.96	0.923077	1
KCNK15 AND NOT-DSC3 AND NOT-SLC39A6	0.96	0.923077	1
KCNK15 AND NOT-DSC3 AND NOT-SPON2	0.96	0.923077	1
KCNK15 AND NOT-DSC3 AND BMPR1B	0.96	0.923077	1
KCNK15 AND NOT-DSC3 AND NOT-CD52	0.96	0.923077	1
KCNK15 AND NOT-DSC3 AND NOT-ABCA5	0.96	0.923077	1

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
PTPRZ1 AND KCNK15 AND GRIA3	0.903226	1	0.823529
PTPRZ1 AND KCNK15 AND IGSF11	0.903226	1	0.823529
GPM6B AND SIRPG AND SLC3A2	0.903226	1	0.823529
GPM6B AND SIRPG AND CYBA	0.882353	0.882353	0.882353
GPM6A AND CD99 AND RHCG	0.882353	0.882353	0.882353
GPM6A AND OR1A1 AND DISP1	0.882353	0.882353	0.882353
NRCAM AND SLC22A4 AND NOT-YIPF3	0.882353	0.882353	0.882353
NRCAM AND NOT-PTPRM AND SLC7A7	0.9375	1	0.882353
GPM6A AND WLS AND OR1A1	0.888889	0.842105	0.941176
NRCAM AND CYBA AND NOT-ZDHHC5	0.9375	1	0.882353
GPM6B AND SIRPG AND WLS	0.909091	0.9375	0.882353
PTPRZ1 AND NOT-SLC2A12 AND TMEM119	0.875	0.933333	0.823529
GPM6B AND OR1A1 AND SLC3A2	0.875	0.933333	0.823529
PTPRZ1 AND NOT-STX8 AND NOT-RECK	0.875	0.933333	0.823529
GPM6A AND OR1A1 AND TLR2	0.909091	0.9375	0.882353
NRCAM AND SLC7A7 AND NOT-YIPF3	0.9375	1	0.882353
NRCAM AND SLC22A4 AND NOT-GPR146	0.882353	0.882353	0.882353
NRCAM AND CTNS AND NOT-SLC26A11	0.9375	1	0.882353
NRCAM AND CTNS AND NOT-VAMP1	0.9375	1	0.882353
GPM6A AND SLC7A7 AND NOT-SLC12A6	0.882353	0.882353	0.882353
GPM6A AND SLC7A7 AND NOT-ITGB7	0.882353	0.882353	0.882353
GPM6A AND HLA-B AND RHCG	0.909091	0.9375	0.882353
COMPLEX-SLC9A6/PTPRZ1/TLR2	0.909091	0.9375	0.882353
PTPRZ1 AND NOT-VAMP1 AND NOT-TRPM7	0.875	0.933333	0.823529
GPM6A AND HLA-B AND P2RY1	0.9375	1	0.882353
GPM6A AND SLC2A10 AND NOT-RHBG	0.9375	1	0.882353
GPM6A AND SLC2A10 AND NOT-STX8	0.903226	1	0.823529
GPM6A AND SLC2A10 AND NOT-TSPAN5	0.9375	1	0.882353
PTPRZ1 AND FZD7 AND NOT-TRPM7	0.909091	0.9375	0.882353
NRCAM AND SLC22A4 AND NOT-SLC1A1	0.882353	0.882353	0.882353
GPM6A AND OR1A1 AND SLC30A5	0.866667	1	0.764706
PTPRZ1 AND NOT-STX8 AND CD9	0.866667	1	0.764706
NRCAM AND NOT-PTPRM AND SLC2A10	0.866667	1	0.764706
PTPRZ1 AND NOT-MTUS1 AND NOT-CDHR2	0.866667	1	0.764706
NRCAM AND NOT-PTPRM AND BTN3A2	0.866667	1	0.764706
GPM6A AND SLC2A10 AND NOT-SLC9A6	0.866667	1	0.764706
GPM6A AND PLP2 AND OR1A1	0.866667	1	0.764706
PTPRZ1 AND NOT-MTUS1 AND NOT-CAV2	0.866667	1	0.764706
PTPRZ1 AND NOT-MTUS1 AND NOT-VAMP1	0.866667	1	0.764706
GPM6A AND IL10RA AND OR1A1	0.866667	1	0.764706
GPM6A AND ABCA1 AND RHCG	0.866667	1	0.764706
GPM6B AND AQP4 AND RHCG	0.875	0.933333	0.823529
PTPRZ1 AND SLC7A7 AND NOT-YIPF3	0.903226	1	0.823529
NRCAM AND BTN3A2 AND NOT-VAMP1	0.866667	1	0.764706
GPM6A AND SLC2A10 AND NOT-ABCC10	0.9375	1	0.882353
PTPRZ1 AND IL10RA AND NOT-ZDHHC5	0.903226	1	0.823529
NRCAM AND SLC7A7 AND NOT-SLC12A6	0.9375	1	0.882353
PTPRZ1 AND IL10RA AND GRIA3	0.903226	1	0.823529
NRCAM AND SLC22A4 AND NOT-AQP7	0.882353	0.882353	0.882353
COMPLEX-MTUS1/PTPRZ1/SLC33A1	0.866667	1	0.764706
PTPRZ1 AND SLC7A7 AND NOT-ZDHHC5	0.903226	1	0.823529
NRCAM AND BTN3A2 AND NOT-HEG1	0.866667	1	0.764706
PTPRZ1 AND NOT-VAMP1 AND NOT-TM9SF2	0.875	0.933333	0.823529
PTPRZ1 AND NOT-VAMP1 AND NOT-ZDHHC5	0.875	0.933333	0.823529
COMPLEX-ABCA8/PTPRZ1/STX8	0.875	0.933333	0.823529
COMPLEX-PTPRZ1/CAV2/STX8	0.875	0.933333	0.823529
COMPLEX-PTPRCAP/PTPRZ1/STX8	0.875	0.933333	0.823529
PTPRZ1 AND NOT-UNC5D AND GPM6B	0.909091	0.9375	0.882353
COMPLEX-PTPRZ1/STX8/SLC12A6	0.875	0.933333	0.823529
COMPLEX-PAG1/PTPRZ1/STX8	0.875	0.933333	0.823529
COMPLEX-TMEM150C/PTPRZ1/STX8	0.875	0.933333	0.823529
COMPLEX-TMEM65/PTPRZ1/STX8	0.875	0.933333	0.823529
GPM6A AND SLC2A10 AND NOT-STS	0.9375	1	0.882353
PTPRZ1 AND FZD7 AND NOT-LEPR	0.909091	0.9375	0.882353
PTPRZ1 AND IL10RA AND FAT3	0.903226	1	0.823529
PTPRZ1 AND FZD7 AND NOT-AQP7	0.909091	0.9375	0.882353
COMPLEX-ZDHHC5/PTPRZ1/STX8	0.875	0.933333	0.823529
GPM6B AND OR1A1 AND WLS	0.857143	0.833333	0.882353
GPM6B AND HLA-B AND RHCG	0.882353	0.882353	0.882353
GPM6A AND HLA-B AND OR1A1	0.857143	0.833333	0.882353
GPM6A AND SLC2A10 AND NOT-TMEM127	0.9375	1	0.882353
GPM6A AND SLC2A10 AND NOT-SLC41A3	0.9375	1	0.882353
NRCAM AND TLR2 AND NOT-GPR146	0.9375	1	0.882353
PTPRZ1 AND SLC7A7 AND FAT3	0.903226	1	0.823529
GPM6B AND SLC3A2 AND NOT-CDHR2	0.903226	1	0.823529
GPM6B AND TLR2 AND RHCG	0.882353	0.882353	0.882353
GPM6A AND WLS AND RHCG	0.941176	0.941176	0.941176

Antigen Pair Logic	F1	Precision	Recall
KCNK15 AND NOT-BMP2 AND PIRT	1	1	1
KCNK15 AND NOT-GHR AND NOT-FPR1	1	1	1
KCNK15 AND NOT-BMP2 AND NOT-AOC3	1	1	1
KCNK15 AND NOT-SYT4 AND NOT-RAMP3	1	1	1
KCNK15 AND NOT-GHR AND SLC51B	1	1	1
KCNK15 AND NOT-GHR AND NOT-FPR1	1	1	1
KCNK15 AND NOT-SYT4 AND NOT-RAMP3	1	1	1
KCNK15 AND NOT-BMP2 AND NOT-ESAM	1	1	1
KCNK15 AND NOT-BMP2 AND DCC	1	1	1
KCNK15 AND NOT-BMP2 AND NOT-ESAM	1	1	1
KCNK15 AND NOT-BMP2 AND NOT-EMP3	1	1	1
KCNK15 AND NOT-THBD AND NOT-LRFN5	1	1	1
KCNK15 AND NOT-THBD AND NOT-LRFN5	1	1	1
KCNK15 AND NOT-BMP2 AND NOT-CD93	0.96	0.923077	1
KCNK15 AND NOT-GHR AND NOT-KCNQ4	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-ATP4B	0.96	0.923077	1
KCNK15 AND NOT-GHR AND GJD4	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-SLC9A1	0.96	0.923077	1
KCNK15 AND NOT-GHR AND NOT-SLC31A2	0.96	0.923077	1
KCNK15 AND NOT-GHR AND NOT-IL7R	0.96	0.923077	1
KCNK15 AND NOT-GHR AND NOT-FCGR3B	0.96	0.923077	1
KCNK15 AND NOT-GHR AND NOT-PNPLA2	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND LYPD1	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-DYSF	0.96	0.923077	1
KCNK15 AND NOT-GHR AND NOT-SLC47A1	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-CD36	0.96	0.923077	1
KCNK15 AND NOT-GHR AND NOT-CD68	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-LRRC8E	0.96	0.923077	1
KCNK15 AND NOT-GHR AND NOT-ADRB2	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND DRD5	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-ABCA12	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-SLC22A18	0.96	0.923077	1
KCNK15 AND NOT-GHR AND CELSR3	0.96	0.923077	1
KCNK15 AND NOT-GHR AND NOT-PDGFRB	0.96	0.923077	1
KCNK15 AND NOT-GHR AND NOT-ENPP2	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND SLC4A5	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-CD163	0.96	0.923077	1
KCNK15 AND NOT-GHR AND SLC17A4	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-SLC9A1	0.96	0.923077	1
KCNK15 AND NOT-GHR AND NOT-PDGFRB	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-LRRC8E	0.96	0.923077	1
KCNK15 AND NOT-GHR AND CRB3	0.96	0.923077	1
KCNK15 AND NOT-GHR AND NOT-KCNQ4	0.96	0.923077	1
KCNK15 AND NOT-GHR AND NOT-ST6GALNAC6	0.96	0.923077	1
KCNK15 AND NOT-SYT4 AND NOT-ADRB2	0.96	0.923077	1
KCNK15 AND NOT-GHR AND NOT-CTSZ	0.96	0.923077	1
KCNK15 AND NOT-GHR AND NOT-SLC31A2	0.96	0.923077	1
KCNK15 AND NOT-GHR AND NOT-ITGA5	0.96	0.923077	1
KCNK15 AND NOT-GHR AND NOT-MR1	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-CD44	0.96	0.923077	1
KCNK15 AND NOT-GHR AND NOT-SLC22A4	0.96	0.923077	1
KCNK15 AND NOT-GHR AND NOT-MR1	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND CHRNG	0.96	0.923077	1
KCNK15 AND NOT-GHR AND NOT-FCGR3B	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-CD36	0.96	0.923077	1
KCNK15 AND NOT-GHR AND NOT-HEPH	0.96	0.923077	1
KCNK15 AND NOT-GHR AND NOT-PCDHGA10	0.96	0.923077	1
KCNK15 AND NOT-GHR AND NOT-ITGA5	0.96	0.923077	1
KCNK15 AND NOT-GHR AND NOT-BMP2	0.96	0.923077	1
KCNK15 AND NOT-GHR AND CEACAM1	0.96	0.923077	1
KCNK15 AND NOT-GHR AND NOT-CD68	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND FXYD7	0.96	0.923077	1
KCNK15 AND NOT-SYT4 AND NOT-IL7R	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-TTYH2	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-GPR137B	0.96	0.923077	1
KCNK15 AND NOT-GHR AND NOT-PNPLA2	0.96	0.923077	1
KCNK15 AND NOT-GHR AND NOT-TNFSF12	0.96	0.923077	1
KCNK15 AND NOT-GHR AND NOT-IL7R	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-GGTLC1	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-CD44	0.96	0.923077	1
KCNK15 AND NOT-SYT4 AND NOT-IL7R	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-MEGF11	0.96	0.923077	1
KCNK15 AND NOT-GHR AND NOT-SLC31A2	0.96	0.923077	1
KCNK15 AND NOT-GHR AND SCNN1A	0.96	0.923077	1
KCNK15 AND NOT-GHR AND NOT-ST6GALNAC6	0.96	0.923077	1
KCNK15 AND NOT-GHR AND GPR143	0.96	0.923077	1
KCNK15 AND NOT-GHR AND SUSD3	0.96	0.923077	1

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
PTPRZ1 AND ITGA2 AND NOT-ZDHHC5	0.909091	0.9375	0.882353
PTPRZ1 AND FZD7 AND NOT-JAG2	0.909091	0.9375	0.882353
GPM6A AND SLC2A10 AND NOT-EPHA1	0.9375	1	0.882353
GPM6A AND SLC2A10 AND OR1A1	0.9375	1	0.882353
PTPRZ1 AND NOT-VAMP1 AND NOT-RER1	0.875	0.933333	0.823529
NRCAM AND NOT-TMEM150C AND SLC30A5	0.866667	1	0.764706
GPM6B AND SLC7A7 AND FNDC4	0.857143	0.833333	0.882353
GPM6B AND SLC3A2 AND NOT-HMOX2	0.903226	1	0.823529
AQP4 AND RHCG AND NOT-MME	0.866667	1	0.764706
NRCAM AND NOT-PTPRM AND DIABLO	0.875	0.933333	0.823529
AQP4 AND RHCG AND NOT-SLC26A9	0.866667	1	0.764706
PTPRZ1 AND NOT-STX8 AND VCAM1	0.903226	1	0.823529
PTPRZ1 AND VCAM1 AND NOT-ITM2A	0.9375	1	0.882353
PTPRZ1 AND VCAM1 AND NOT-CYB5R1	0.9375	1	0.882353
NRCAM AND VCAM1 AND NOT-SLC12A6	0.909091	0.9375	0.882353
PTPRZ1 AND SLC7A7 AND NOT-TRPM4	0.903226	1	0.823529
GPM6B AND OR1A1 AND CBX3	0.875	0.933333	0.823529
PTPRZ1 AND VCAM1 AND NOT-TRPM7	0.9375	1	0.882353
PTPRZ1 AND NOT-GDE1 AND VCAM1	0.9375	1	0.882353
NRCAM AND VCAM1 AND NOT-YIPF3	0.909091	0.9375	0.882353
NRCAM AND VCAM1 AND NOT-GPR146	0.909091	0.9375	0.882353
PTPRZ1 AND VCAM1 AND NOT-SLC12A2	0.9375	1	0.882353
COMPLEX-GPNMB/PTPRZ1/STX8	0.875	0.933333	0.823529
PTPRZ1 AND VCAM1 AND NOT-CAV2	0.9375	1	0.882353
PTPRZ1 AND VCAM1 AND NOT-JAG2	0.9375	1	0.882353
PTPRZ1 AND VCAM1 AND NOT-EPHA1	0.9375	1	0.882353
PTPRZ1 AND VCAM1 AND NOT-PERP	0.9375	1	0.882353
PTPRZ1 AND VCAM1 AND NOT-YIPF3	0.9375	1	0.882353
PTPRZ1 AND TLR2 AND NOT-MST1R	0.903226	1	0.823529
NRCAM AND VCAM1 AND NOT-SLC12A2	0.909091	0.9375	0.882353
PTPRZ1 AND FZD7 AND NOT-TRPM4	0.909091	0.9375	0.882353
PTPRZ1 AND VCAM1 AND NOT-GPR153	0.9375	1	0.882353
AQP4 AND RHCG AND NOT-ITGB6	0.866667	1	0.764706
PTPRZ1 AND VCAM1 AND NOT-PTPRF	0.9375	1	0.882353
PTPRZ1 AND VCAM1 AND NOT-ZDHHC5	0.9375	1	0.882353
PTPRZ1 AND FZD7 AND NOT-MST1R	0.909091	0.9375	0.882353
PTPRZ1 AND VCAM1 AND NOT-AQP7	0.9375	1	0.882353
PTPRZ1 AND VCAM1 AND NOT-GJB2	0.9375	1	0.882353
PTPRZ1 AND NOT-STS AND VCAM1	0.9375	1	0.882353
PTPRZ1 AND VCAM1 AND NOT-LEPR	0.9375	1	0.882353
PTPRZ1 AND IL13RA1 AND NOT-ZDHHC5	0.903226	1	0.823529
PTPRZ1 AND VCAM1 AND NOT-CAV1	0.903226	1	0.823529
PTPRZ1 AND VCAM1 AND FAT3	0.903226	1	0.823529
AQP4 AND RHCG AND NOT-TRPM4	0.866667	1	0.764706
PTPRZ1 AND VCAM1 AND NOT-TMEM219	0.9375	1	0.882353
GPM6B AND SIRPG AND CBX3	0.875	0.933333	0.823529
PTPRZ1 AND NOT-TRPM4 AND C3AR1	0.9375	1	0.882353
PTPRZ1 AND VCAM1 AND NOT-ORAI1	0.9375	1	0.882353
PTPRZ1 AND VCAM1 AND NOT-CXADR	0.9375	1	0.882353
PTPRZ1 AND VCAM1 AND NOT-MFSD5	0.9375	1	0.882353
AQP4 AND RHCG AND NOT-SDC1	0.866667	1	0.764706
AQP4 AND RHCG AND NOT-MUC1	0.866667	1	0.764706
AQP4 AND RHCG AND NOT-CLDN23	0.866667	1	0.764706
AQP4 AND RHCG AND NOT-CLDN7	0.866667	1	0.764706
PTPRZ1 AND VCAM1 AND NOT-TSPAN14	0.9375	1	0.882353
IGSF11 AND KCNK15 AND TNC	0.83871	0.928571	0.764706
GPM6B AND TNC AND SIRPG	0.83871	0.928571	0.764706
NRCAM AND VCAM1 AND NOT-CDHR2	0.909091	0.9375	0.882353
NRCAM AND VCAM1 AND NOT-APLP2	0.875	0.933333	0.823529
COMPLEX-TSPAN5/EDNRB/PTPRZ1	0.83871	0.928571	0.764706
PTPRZ1 AND VCAM1 AND PTPRC	0.903226	1	0.823529
NRCAM AND VCAM1 AND NOT-ERMAP	0.909091	0.9375	0.882353
PTPRZ1 AND SLC2A10 AND NOT-TRPM4	0.903226	1	0.823529
PTPRZ1 AND VCAM1 AND NOT-TSPAN4	0.9375	1	0.882353
PTPRZ1 AND VCAM1 AND NOT-CDHR2	0.9375	1	0.882353
PTPRZ1 AND VCAM1 AND NOT-AQP9	0.9375	1	0.882353
PTPRZ1 AND VCAM1 AND NOT-PKN2	0.9375	1	0.882353
PTPRZ1 AND VCAM1 AND PNPLA2	0.9375	1	0.882353
AQP4 AND RHCG AND NOT-SLAMF7	0.866667	1	0.764706
AQP4 AND RHCG AND NOT-IL20RA	0.866667	1	0.764706
PTPRZ1 AND TLR2 AND VCAM1	0.903226	1	0.823529
PTPRZ1 AND NOT-SLC3A1 AND VCAM1	0.9375	1	0.882353
PTPRZ1 AND VCAM1 AND NOT-OR3A3	0.9375	1	0.882353
PTPRZ1 AND RHCG AND NOT-MST1R	0.9375	1	0.882353
PTPRZ1 AND VCAM1 AND NOT-PTPRS	0.9375	1	0.882353
NLGN1 AND SLC7A7 AND NOT-TRPM4	0.866667	1	0.764706
PTPRZ1 AND NOT-STS AND NOT-CLDN12	0.909091	0.9375	0.882353

Antigen Pair Logic	F1	Precision	Recall
KCNK15 AND NOT-BMP2 AND NOT-CD163	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-MEGF11	0.96	0.923077	1
KCNK15 AND NOT-GHR AND SUSD3	0.96	0.923077	1
KCNK15 AND NOT-BMP2 AND NOT-CD93	0.96	0.923077	1
KCNK15 AND NOT-ABCA8 AND NOT-CD44	0.956522	1	0.916667
KCNK15 AND NOT-THBD AND PTPRG	0.956522	1	0.916667
KCNK15 AND NOT-GHR AND TACR2	0.956522	1	0.916667
KCNK15 AND NOT-BMP2 AND SLC2A1	0.956522	1	0.916667
KCNK15 AND NOT-ABCA8 AND NOT-THBD	0.956522	1	0.916667
KCNK15 AND NOT-GHR AND GPR35	0.956522	1	0.916667
KCNK15 AND NOT-THBD AND NOT-ADRB2	0.956522	1	0.916667
COMPLEX-CHRNB4/KCNK15/BMP2	0.96	0.923077	1
COMPLEX-SLC31A2/KCNK15/SYT4	0.96	0.923077	1
COMPLEX-TAS2R10/KCNK15/BMP2	0.96	0.923077	1
KCNK15 AND NOT-SYT4 AND NOT-ECSCR	1	1	1
COMPLEX-SLC31A2/KCNK15/SYT4	0.96	0.923077	1
COMPLEX-CATSPER3/KCNK15/BMP2	0.96	0.923077	1
COMPLEX-KCNK15/MSMO1/BMP2	0.96	0.923077	1
COMPLEX-ATP8A1/KCNK15/BMP2	0.96	0.923077	1
COMPLEX-OXTR/KCNK15/BMP2	0.96	0.923077	1
COMPLEX-NMBR/KCNK15/BMP2	0.96	0.923077	1
COMPLEX-MTNR1B/KCNK15/BMP2	0.96	0.923077	1
KCNK15 AND NOT-GHR AND PIRT	1	1	1
KCNK15 AND NOT-GHR AND NOT-CD44	0.96	0.923077	1
KCNK15 AND NOT-GHR AND NOT-CD44	0.96	0.923077	1
KCNK15 AND NOT-GHR AND NOT-EMP3	0.96	0.923077	1
KCNK15 AND NOT-SYT4 AND NOT-SELP	1	1	1
KCNK15 AND NOT-SYT4 AND NOT-SELP	1	1	1
KCNK15 AND NOT-SYT4 AND NOT-CD93	0.96	0.923077	1
KCNK15 AND NOT-SYT4 AND NOT-CD93	0.96	0.923077	1
KCNK15 AND NOT-GHR AND GJA10	0.96	0.923077	1
KCNK15 AND NOT-SYT4 AND NOT-CD44	0.96	0.923077	1
KCNK15 AND NOT-GHR AND FXYD7	0.96	0.923077	1
KCNK15 AND NOT-SYT4 AND NOT-CD44	0.96	0.923077	1
KCNK15 AND NOT-SYT4 AND NOT-CD44	0.96	0.923077	1
KCNK15 AND SLC2A1 AND NOT-CD44	0.956522	1	0.916667
KCNK15 AND NOT-THBD AND SLC2A1	0.956522	1	0.916667
KCNK15 AND NOT-SYT4 AND NOT-CD163	0.96	0.923077	1
KCNK15 AND NOT-GABRG1 AND NOT-CD163	0.96	0.923077	1
KCNK15 AND NOT-SYT4 AND NOT-KCNK6	1	1	1
KCNK15 AND NOT-SYT4 AND NOT-CD163	0.96	0.923077	1
KCNK15 AND NOT-GABRG1 AND NOT-SELP	1	1	1
KCNK15 AND NOT-GABRG1 AND NOT-CD163	0.96	0.923077	1
KCNK15 AND NOT-GABRG1 AND NOT-SELP	1	1	1
KCNK15 AND NOT-SYT4 AND NOT-TTYH2	0.96	0.923077	1
KCNK15 AND NOT-GABRG1 AND NOT-CD93	0.96	0.923077	1
KCNK15 AND SLC2A1 AND NOT-SELP	0.956522	1	0.916667
KCNK15 AND NOT-GABRG1 AND NOT-CD44	0.96	0.923077	1
KCNK15 AND NOT-THBD AND LHFPL5	0.956522	1	0.916667
KCNK15 AND NOT-THBD AND NOT-SYT4	1	1	1
KCNK15 AND NOT-GABRG1 AND NOT-CD44	0.96	0.923077	1
KCNK15 AND NOT-THBD AND NOT-SYT4	1	1	1
KCNK15 AND SLC2A1 AND NOT-CD93	0.956522	1	0.916667
KCNK15 AND NOT-GABRG1 AND NOT-TTYH2	0.96	0.923077	1
KCNK15 AND NOT-LRRN4 AND NOT-CD163	0.96	0.923077	1
KCNK15 AND NOT-LRRN4 AND NOT-CD163	0.96	0.923077	1
KCNK15 AND NOT-LRRN4 AND NOT-SELP	1	1	1
KCNK15 AND NOT-LRRN4 AND NOT-CD93	0.956522	1	0.916667
KCNK15 AND NOT-LRRN4 AND NOT-SELP	1	1	1
KCNK15 AND NOT-LRRN4 AND NOT-CD93	0.956522	1	0.916667
KCNK15 AND NOT-LRRN4 AND NOT-CD44	0.96	0.923077	1
KCNK15 AND NOT-LRRN4 AND NOT-TTYH2	0.96	0.923077	1
KCNK15 AND NOT-CLDN15 AND NOT-KCNK6	1	1	1
KCNK15 AND NOT-CLDN15 AND NOT-CD44	0.96	0.923077	1
KCNK15 AND NOT-CLDN15 AND NOT-TTYH2	0.96	0.923077	1
KCNK15 AND NOT-CLDN15 AND NOT-CD44	0.96	0.923077	1
KCNK15 AND NOT-GHR AND NOT-PAQR8	0.923077	0.857143	1
KCNK15 AND NOT-GHR AND EFNB2	0.923077	0.857143	1
KCNK15 AND NOT-GHR AND NOT-TGFBI	0.923077	0.857143	1
KCNK15 AND NOT-GHR AND NOT-ALCAM	0.923077	0.857143	1
KCNK15 AND NOT-GHR AND NOT-MRGPRF	0.923077	0.857143	1
COMPLEX-FUT3/GHR/KCNK15	0.923077	0.857143	1
KCNK15 AND NOT-GHR AND ATP8A1	0.923077	0.857143	1
KCNK15 AND NOT-GABRG1 AND NOT-GYPC	0.923077	0.857143	1
KCNK15 AND NOT-LRRN4 AND NOT-GYPC	0.923077	0.857143	1
KCNK15 AND NOT-GHR AND NOT-MSMO1	0.923077	0.857143	1
KCNK15 AND NOT-GHR AND NOT-ZDHHC2	0.923077	0.857143	1

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
PTPRZ1 AND VCAM1 AND NOT-ERMAP	0.9375	1	0.882353
PTPRZ1 AND CD70 AND CCR1	0.903226	1	0.823529
PTPRZ1 AND VCAM1 AND NOT-SLC7A6	0.9375	1	0.882353
AQP4 AND RHCG AND NOT-MST1R	0.866667	1	0.764706
NRCAM AND VCAM1 AND NOT-SLC33A1	0.909091	0.9375	0.882353
PTPRZ1 AND VCAM1 AND CCR1	0.903226	1	0.823529
GPM6B AND OR1A1 AND IL13RA1	0.833333	0.789474	0.882353
PTPRZ1 AND VCAM1 AND NOT-KCNJ2	0.827586	1	0.705882
GPM6B AND TNC AND FNDC4	0.827586	1	0.705882
AQP4 AND RHCG AND NOT-CLDN1	0.827586	1	0.705882
COMPLEX-EMP1/NRCAM/VCAM1	0.827586	1	0.705882
PTPRZ1 AND VCAM1 AND NOT-TMEM50B	0.9375	1	0.882353
GPM6B AND SLC2A10 AND NOT-TRPM4	0.827586	1	0.705882
PTPRZ1 AND CCR1 AND NOT-TRPM4	0.903226	1	0.823529
PTPRZ1 AND NOT-MST1R AND TMEM119	0.875	0.933333	0.823529
PTPRZ1 AND VCAM1 AND NOT-SLC12A6	0.9375	1	0.882353
PTPRZ1 AND VCAM1 AND NOT-CNNM4	0.9375	1	0.882353
PTPRZ1 AND VCAM1 AND NOT-FZD4	0.9375	1	0.882353
PTPRZ1 AND RHCG AND NOT-TRPM4	0.9375	1	0.882353
PTPRZ1 AND VCAM1 AND FAM26F	0.903226	1	0.823529
PTPRZ1 AND VCAM1 AND ITGAM	0.903226	1	0.823529
PTPRZ1 AND VCAM1 AND NOT-SLC12A7	0.9375	1	0.882353
GPM6B AND SLC3A2 AND NOT-CLDN11	0.903226	1	0.823529
GPM6B AND SLC3A2 AND NOT-EDNRB	0.827586	1	0.705882
NRCAM AND VCAM1 AND NOT-FGFR2	0.909091	0.9375	0.882353
PTPRZ1 AND VCAM1 AND NOT-SLC33A1	0.9375	1	0.882353
NRCAM AND VCAM1 AND NOT-TRPM7	0.909091	0.9375	0.882353
NLGN4X AND VCAM1 AND NOT-LEPR	0.823529	0.823529	0.823529
PTPRZ1 AND VCAM1 AND NOT-TMEM127	0.9375	1	0.882353
NRCAM AND VCAM1 AND CYB5R1	0.909091	0.9375	0.882353
PTPRZ1 AND VCAM1 AND NOT-TMEM97	0.9375	1	0.882353
GPM6A AND ANXA1 AND OR1A1	0.823529	0.823529	0.823529
NRCAM AND NOT-STS AND VCAM1	0.9375	1	0.882353
NLGN4X AND LAPTM5 AND NOT-VANGL1	0.903226	1	0.823529
SYT11 AND LAPTM5 AND RHCG	1	1	1
GPM6B AND F2R AND SLC4A8	0.866667	1	0.764706
GPM6B AND F2R AND SLCO1C1	0.866667	1	0.764706
GPM6B AND F2R AND NOT-AJAP1	0.866667	1	0.764706
SYT11 AND F2R AND NOT-SLC12A5	0.866667	1	0.764706
MLC1 AND SLC2A10 AND NOT-FLVCR1	0.866667	1	0.764706
SYT11 AND F2R AND TMEM100	0.866667	1	0.764706
LYPD1 AND LAPTM5 AND NOT-ITGB7	0.903226	1	0.823529
GPR158 AND F2R AND NOT-SYT2	0.83871	0.928571	0.764706
GPR158 AND F2R AND NOT-ATP2C1	0.83871	0.928571	0.764706
GPR158 AND F2R AND NOT-HEG1	0.83871	0.928571	0.764706
GPR158 AND F2R AND NOT-TMEM127	0.83871	0.928571	0.764706
GPR158 AND F2R AND NOT-TM9SF2	0.83871	0.928571	0.764706
GPR158 AND F2R AND NOT-LRRC4	0.83871	0.928571	0.764706
GPR158 AND F2R AND OR1A1	0.83871	0.928571	0.764706
LYPD1 AND LAPTM5 AND NOT-SLC12A6	0.903226	1	0.823529
MLC1 AND SLC2A10 AND NOT-VANGL1	0.866667	1	0.764706
MLC1 AND SLC2A10 AND NOT-PIK3IP1	0.866667	1	0.764706
MLC1 AND SLC2A10 AND NOT-ZDHHC2	0.866667	1	0.764706
GPR158 AND F2R AND NOT-EPHA4	0.83871	0.928571	0.764706
GPR158 AND F2R AND NOT-ABCA3	0.83871	0.928571	0.764706
GPR158 AND F2R AND NOT-IL6ST	0.83871	0.928571	0.764706
LYPD1 AND LAPTM5 AND NOT-LILRB2	0.903226	1	0.823529
GPR158 AND OR51B6 AND NOT-AJAP1	0.842105	0.761905	0.941176
SYT11 AND CDH11 AND RHCG	0.914286	0.888889	0.941176
MLC1 AND BTN3A3 AND NOT-USP48	0.866667	1	0.764706
SYT11 AND VAMP8 AND RHCG	0.888889	0.842105	0.941176
LYPD1 AND LAPTM5 AND NOT-CD40	0.903226	1	0.823529
SYT11 AND NOT-ATP8A2 AND RHCG	0.833333	0.789474	0.882353
LYPD1 AND LAPTM5 AND NOT-SDC2	0.866667	1	0.764706
LYPD1 AND LAPTM5 AND NOT-YIPF3	0.903226	1	0.823529
GPM6A AND EMP3 AND ADAM29	0.827586	1	0.705882
GPM6A AND F2R AND ADAM29	0.827586	1	0.705882
MLC1 AND BTN3A3 AND NOT-RER1	0.903226	1	0.823529
GPR158 AND OR51B6 AND NOT-ABCG4	0.864865	0.8	0.941176
NRCAM AND ATP7A AND NOT-ATP8B1	0.909091	0.9375	0.882353
LYPD1 AND LAPTM5 AND NOT-ZDHHC5	0.903226	1	0.823529
MLC1 AND TNFSF13B AND NOT-ZDHHC2	0.9375	1	0.882353
GPR158 AND IL1RAP AND NOT-ABCG4	0.827586	1	0.705882
SYT11 AND NOT-CNTNAP2 AND RHCG	0.909091	0.9375	0.882353
MLC1 AND BTN3A3 AND NOT-TM9SF2	0.903226	1	0.823529
MLC1 AND BTN3A3 AND NOT-SNAP23	0.866667	1	0.764706
LYPD1 AND LAPTM5 AND NOT-PNPLA2	0.903226	1	0.823529

Antigen Pair Logic	F1	Precision	Recall
KCNK15 AND NOT-GHR AND NOT-ATP7A	0.923077	0.857143	1
COMPLEX-GHR/KCNK15/SLC22A16	0.923077	0.857143	1
KCNK15 AND NOT-GHR AND NOT-MSMO1	0.923077	0.857143	1
KCNK15 AND NOT-GHR AND NOT-FAT1	0.923077	0.857143	1
KCNK15 AND NOT-LRRN4 AND NOT-AOC3	0.923077	0.857143	1
KCNK15 AND NOT-GHR AND FXYD6	0.923077	0.857143	1
KCNK15 AND NOT-GHR AND HLA-DRB1	0.923077	0.857143	1
COMPLEX-SLC22A9/GHR/KCNK15	0.923077	0.857143	1
KCNK15 AND NOT-GHR AND CD163	0.923077	0.857143	1
KCNK15 AND NOT-GHR AND CHRNG	0.923077	0.857143	1
KCNK15 AND NOT-GABRG1 AND NOT-SCARA5	0.923077	0.857143	1
KCNK15 AND NOT-GHR AND NOT-MRGPRF	0.923077	0.857143	1
KCNK15 AND NOT-GHR AND CD163	0.923077	0.857143	1
KCNK15 AND NOT-SYT4 AND NOT-GYPC	0.923077	0.857143	1
KCNK15 AND NOT-LRRN4 AND NOT-LAT	0.923077	0.857143	1
KCNK15 AND NOT-GABRG1 AND NOT-GYPC	0.923077	0.857143	1
KCNK15 AND NOT-LRRN4 AND NOT-SCARA5	0.923077	0.857143	1
KCNK15 AND NOT-LRRN4 AND NOT-LAT	0.923077	0.857143	1
KCNK15 AND NOT-GHR AND NOT-ANPEP	0.923077	0.857143	1
KCNK15 AND NOT-SYT4 AND NOT-LAT	0.923077	0.857143	1
COMPLEX-GHR/SLC13A5/KCNK15	0.923077	0.857143	1
KCNK15 AND NOT-GABRG1 AND NOT-SCARA5	0.923077	0.857143	1
COMPLEX-GHR/CXCL16/KCNK15	0.923077	0.857143	1
KCNK15 AND NOT-GHR AND NOT-ALCAM	0.923077	0.857143	1
KCNK15 AND NOT-GHR AND NOT-SLC9A1	0.923077	0.857143	1
KCNK15 AND NOT-GHR AND EPHA10	0.923077	0.857143	1
KCNK15 AND NOT-SYT4 AND NOT-SCARA5	0.923077	0.857143	1
COMPLEX-GHR/KCNK15/CDH22	0.923077	0.857143	1
KCNK15 AND NOT-GHR AND NOT-MEGF11	0.923077	0.857143	1
KCNK15 AND NOT-LRRN4 AND NOT-GYPC	0.923077	0.857143	1
KCNK15 AND NOT-GHR AND NOT-MEGF11	0.923077	0.857143	1
KCNK15 AND NOT-GHR AND NOT-ATP7A	0.923077	0.857143	1
KCNK15 AND NOT-SYT4 AND NOT-SCARA5	0.923077	0.857143	1
KCNK15 AND NOT-GABRG1 AND NOT-LAT	0.923077	0.857143	1
KCNK15 AND NOT-LRRN4 AND NOT-AOC3	0.923077	0.857143	1
COMPLEX-GHR/KCNK15/SLC2A1	0.923077	0.857143	1
COMPLEX-PTPRT/GHR/KCNK15	0.923077	0.857143	1
KCNK15 AND NOT-GHR AND NOT-ATP8B2	0.923077	0.857143	1
COMPLEX-SLC17A2/GHR/KCNK15	0.923077	0.857143	1
KCNK15 AND NOT-GHR AND NOT-CSMD3	0.923077	0.857143	1
KCNK15 AND NOT-LRRN4 AND NOT-SCARA5	0.923077	0.857143	1
COMPLEX-GHR/MPL/KCNK15	0.923077	0.857143	1
COMPLEX-GHR/GRM7/KCNK15	0.923077	0.857143	1
KCNK15 AND NOT-GHR AND HLA-DOB	0.96	0.923077	1
KCNK15 AND NOT-GHR AND GAGE1	0.96	0.923077	1
KCNK15 AND NOT-GHR AND FOLH1	0.956522	1	0.916667
KCNK15 AND NOT-SYT4 AND MUC16	1	1	1
KCNK15 AND NOT-THBD AND NOT-IL20RA	1	1	1
KCNK15 AND NOT-THBD AND NOT-IL20RA	1	1	1
COMPLEX-GHR/KCNK15/VTCN1	0.923077	0.857143	1
KCNK15 AND NOT-GHR AND NOT-MOK	0.923077	0.857143	1
KCNK15 AND NOT-LRRN4 AND BMPR1B	0.923077	0.857143	1
KCNK15 AND NOT-GHR AND NOT-PCYT1A	0.923077	0.857143	1
KCNK15 AND NOT-GHR AND ALDH1A1	0.923077	0.857143	1
KCNK15 AND NOT-GHR AND GPNMB	0.923077	0.857143	1
KCNK15 AND NOT-GHR AND NOT-ANXA1	0.923077	0.857143	1
KCNK15 AND NOT-GHR AND NOT-CLDN23	0.923077	0.857143	1
KCNK15 AND NOT-GHR AND NOT-CD70	0.923077	0.857143	1
KCNK15 AND NOT-GHR AND NOT-TPBG	0.923077	0.857143	1
KCNK15 AND NOT-GHR AND NOT-ULBP2	0.923077	0.857143	1
KCNK15 AND NOT-GHR AND NOT-MET	0.923077	0.857143	1
COMPLEX-CD52/GHR/KCNK15	0.923077	0.857143	1
KCNK15 AND NOT-GHR AND NOT-SDC1	0.923077	0.857143	1
KCNK15 AND NOT-GHR AND NOT-IL11RA	0.923077	0.857143	1
KCNK15 AND NOT-TPBG AND NOT-CD163	0.923077	0.857143	1
KCNK15 AND NOT-LRRN4 AND NOT-TNC	0.923077	0.857143	1
KCNK15 AND NOT-GHR AND NOT-ST8SIA1	0.923077	0.857143	1
KCNK15 AND NOT-SYT4 AND NOT-TNC	0.923077	0.857143	1
COMPLEX-CD52/GHR/KCNK15	0.923077	0.857143	1
COMPLEX-GHR/KCNK15/CXCR5	0.923077	0.857143	1
KCNK15 AND NOT-GABRG1 AND NOT-TNC	0.923077	0.857143	1
KCNK15 AND NOT-GHR AND NOT-ULBP2	0.923077	0.857143	1
KCNK15 AND NOT-GABRG1 AND BMPR1B	0.923077	0.857143	1
COMPLEX-GHR/KCNK15/MUC16	0.923077	0.857143	1
COMPLEX-GHR/KCNK15/CLDN9	0.923077	0.857143	1
COMPLEX-ALK/GHR/KCNK15	0.923077	0.857143	1
KCNK15 AND NOT-GHR AND NOT-DKK1	0.923077	0.857143	1

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
GPM6A AND CD93 AND BST2	0.827586	1	0.705882
LYPD1 AND LAPTM5 AND NOT-ITSN1	0.903226	1	0.823529
SYT11 AND CD163 AND NOT-SLC12A6	0.842105	0.761905	0.941176
SYT11 AND NOT-GRM3 AND RHCG	0.8125	0.866667	0.764706
GPM6A AND TGFBI AND BST2	0.8125	0.866667	0.764706
GPR158 AND CD163L1 AND NOT-ABCG4	0.8125	0.866667	0.764706
MLC1 AND NOT-S1PR1 AND TLR2	0.8125	0.866667	0.764706
GPM6A AND CD163 AND ADAM29	0.827586	1	0.705882
PTPRZ1 AND NOT-ATP8B1 AND ATP7A	0.866667	1	0.764706
MLC1 AND TNFSF13B AND NOT-FLVCR1	0.848485	0.875	0.823529
GPR158 AND FAM57A AND NOT-AJAP1	0.83871	0.928571	0.764706
GPM6B AND NOT-AJAP1 AND PPAPDC1B	0.823529	0.823529	0.823529
GPR158 AND SLC11A1 AND NOT-ABCG4	0.827586	1	0.705882
SYT11 AND NOT-SYT4 AND RHCG	0.848485	0.875	0.823529
LYPD1 AND PSENEN AND CD163	0.827586	1	0.705882
PTPRZ1 AND NOT-PIK3IP1 AND NOT-FAT1	0.909091	0.9375	0.882353
GPR158 AND SLC11A1 AND NOT-AJAP1	0.827586	1	0.705882
GPR158 AND F2R AND NOT-VAMP5	0.8	0.923077	0.705882
GPM6A AND CD93 AND ADAM29	0.8	0.923077	0.705882
GPM6A AND ANTXR2 AND ADAM29	0.8	0.923077	0.705882
SYT11 AND RHCG AND BST2	0.829268	0.708333	1
NRCAM AND CD163 AND NOT-PPAPDC1B	0.9375	1	0.882353
SYT11 AND CD163 AND RHCG	0.941176	0.941176	0.941176
GPR158 AND CD58 AND NOT-VAMP1	0.827586	1	0.705882
GPR158 AND RHCG AND NOT-ABCG4	0.842105	0.761905	0.941176
GPR158 AND OR51B6 AND BST2	0.864865	0.8	0.941176
GPR158 AND FAM57A AND NOT-ABCG4	0.83871	0.928571	0.764706
FXYD6 AND LAPTM5 AND NOT-LILRB2	0.848485	0.875	0.823529
FXYD6 AND LAPTM5 AND NOT-CD40	0.848485	0.875	0.823529
CDH10 AND LAPTM5 AND NOT-ZDHHC5	0.810811	0.75	0.882353
PCDH8 AND LAPTM5 AND NOT-CD40	0.8125	0.866667	0.764706
GPR158 AND CD58 AND NOT-SYT2	0.8	0.923077	0.705882
SYT11 AND CD163 AND FNDC4	0.864865	0.8	0.941176
GPM6A AND CD163 AND NOT-PPAPDC1B	0.827586	1	0.705882
PTPRZ1 AND NOT-FLVCR1 AND BST2	0.875	0.933333	0.823529
GPM6B AND F2R AND NOT-CLDN11	0.866667	1	0.764706
PTPRZ1 AND LAPTM5 AND VCAM1	0.9375	1	0.882353
PTPRZ1 AND VCAM1 AND GPR158	0.9375	1	0.882353
NRCAM AND LAPTM5 AND VCAM1	0.9375	1	0.882353
NOT-AOC3 AND EDNRB AND ABCA1	0.848485	0.875	0.823529
PTPRZ1 AND VCAM1 AND NOT-PIK3IP1	0.9375	1	0.882353
NRCAM AND VCAM1 AND NOT-ZDHHC2	0.909091	0.9375	0.882353
NRCAM AND BTN3A3 AND VCAM1	0.857143	0.833333	0.882353
NOT-AOC3 AND EDNRB AND ABCC3	0.83871	0.928571	0.764706
CD163L1 AND NOT-AOC3 AND EDNRB	0.83871	0.928571	0.764706
MLC1 AND SLC2A10 AND NOT-TRPM4	0.866667	1	0.764706
PTPRZ1 AND VCAM1 AND NOT-SMPD2	0.9375	1	0.882353
PTPRZ1 AND VCAM1 AND NOT-ZDHHC2	0.9375	1	0.882353
SYT11 AND IL1RAP AND EDNRB	0.827586	1	0.705882
PTPRZ1 AND VCAM1 AND BST2	0.9375	1	0.882353
PTPRZ1 AND VCAM1 AND BTN3A3	0.9375	1	0.882353
OR51B6 AND EDNRB AND NOT-AOC3	0.914286	0.888889	0.941176
OR51B6 AND NOT-ATP8B1 AND EDNRB	0.833333	0.789474	0.882353
SLC11A1 AND NOT-AOC3 AND EDNRB	0.827586	1	0.705882
GPR158 AND ADAM12 AND VCAM1	0.848485	0.875	0.823529
NOT-ATP8B1 AND SLC3A2 AND NOT-CD52	0.875	0.933333	0.823529
TENM1 AND NOT-KCNK6 AND TNC	0.8125	0.866667	0.764706
GPM6A AND TNC AND ADAM29	0.8125	0.866667	0.764706
GPM6B AND TNC AND NOT-AJAP1	0.8125	0.866667	0.764706
SYT11 AND RHCG AND NOT-SST	0.848485	0.875	0.823529
SYT11 AND IL13RA1 AND NOT-SLC12A6	0.842105	0.761905	0.941176
COMPLEX-PIK3IP1/EDNRB/PTPRZ1	0.909091	0.9375	0.882353
TNC AND NOT-KCNK6 AND TSPAN5	0.875	0.933333	0.823529
SYT11 AND RHCG AND GPNMB	0.914286	0.888889	0.941176
GPR158 AND OR1A1 AND CBX3	0.909091	0.9375	0.882353
NOT-ATP8B1 AND EDNRB AND SLC3A2	0.903226	1	0.823529
FAT3 AND TNC AND KCND2	0.8125	0.866667	0.764706
SRR AND NOT-AOC3 AND EDNRB	0.827586	1	0.705882
PCDH12 AND NOT-AOC3 AND EDNRB	0.8	0.923077	0.705882
GPM6A AND BST2 AND CBX3	0.8	0.923077	0.705882
SYT11 AND ABCC3 AND EDNRB	0.8	0.923077	0.705882
NOT-ATP8B1 AND ABCA1 AND NOT-CD52	0.8	0.923077	0.705882
GPM6A AND BST2 AND PCYT1A	0.8	0.923077	0.705882
NOT-AOC3 AND EDNRB AND IL1RAP	0.8	0.923077	0.705882
SYT11 AND RHCG AND AXL	0.820513	0.727273	0.941176
TNC AND NOT-AOC3 AND GLDN	0.83871	0.928571	0.764706
GPR158 AND SLC7A7 AND NOT-TRPM4	0.820513	0.727273	0.941176

Antigen Pair Logic	F1	Precision	Recall
KCNK15 AND NOT-SYT4 AND BMPR1B	0.923077	0.857143	1
COMPLEX-GHR/KCNK15/CLDN6	0.923077	0.857143	1
KCNK15 AND NOT-GHR AND NOT-STEAP1	0.923077	0.857143	1
KCNK15 AND NOT-CLDN15 AND MUC16	0.956522	1	0.916667
KCNK15 AND NOT-GHR AND NOT-CD70	0.923077	0.857143	1
KCNK15 AND NOT-GHR AND NOT-CLDN23	0.923077	0.857143	1
KCNK15 AND NOT-GHR AND NOT-HSPA5	0.923077	0.857143	1
KCNK15 AND NOT-PROCR AND NOT-ENG	1	1	1
KCNK15 AND NOT-PROCR AND NOT-CD52	0.96	0.923077	1
KCNK15 AND NOT-SYT4 AND NOT-GPNMB	0.96	0.923077	1
KCNK15 AND NOT-PROCR AND NOT-IL20RA	0.96	0.923077	1
KCNK15 AND NOT-CLDN15 AND NOT-IL3RA	0.916667	0.916667	0.916667
KCNK15 AND NOT-SYT4 AND NOT-IL3RA	0.916667	0.916667	0.916667
KCNK15 AND NOT-GHR AND RNF43	0.916667	0.916667	0.916667
KCNK15 AND NOT-PROCR AND NOT-TRPM4	0.96	0.923077	1
KCNK15 AND NOT-PROCR AND ULBP2	0.916667	0.916667	0.916667
KCNK15 AND NOT-PROCR AND NOT-CLDN8	0.96	0.923077	1
KCNK15 AND NOT-THBD AND CLDN9	0.909091	1	0.833333
KCNK15 AND NOT-THBD AND BIRC5	0.909091	1	0.833333
KCNK15 AND NOT-THBD AND NCAM1	0.909091	1	0.833333
COMPLEX-MRGPRF/EDNRB/KCNK15	0.96	0.923077	1
KCNK15 AND NOT-PTGER4 AND NOT-ENG	0.923077	0.857143	1
KCNK15 AND NOT-GABRG1 AND NOT-GPNMB	0.96	0.923077	1
KCNK15 AND NOT-LRRN4 AND NOT-GPNMB	0.96	0.923077	1
COMPLEX-EDNRB/ALCAM/KCNK15	0.96	0.923077	1
COMPLEX-EDNRB/ATP7A/KCNK15	0.923077	0.857143	1
COMPLEX-BTN3A3/EDNRB/KCNK15	0.96	0.923077	1
KCNK15 AND NOT-CLDN15 AND NOT-TNC	0.888889	0.8	1
KCNK15 AND NOT-LRRN4 AND NOT-SPON2	0.888889	0.8	1
KCNK15 AND NOT-LRRN4 AND NOT-AXL	0.888889	0.8	1
KCNK15 AND NOT-KDR AND NOT-CD44	0.96	0.923077	1
KCNK15 AND NOT-PTGER4 AND NOT-CLDN23	0.888889	0.8	1
KCNK15 AND NOT-PTGER4 AND NOT-CD52	0.923077	0.857143	1
KCNK15 AND NOT-PIK3IP1 AND NOT-IL3RA	0.916667	0.916667	0.916667
KCNK15 AND NOT-PTGER4 AND NOT-IL20RA	0.888889	0.8	1
COMPLEX-EDNRB/KCNK15/MSMO1	0.916667	0.916667	0.916667
KCNK15 AND NOT-PTGER4 AND NOT-CLDN8	0.923077	0.857143	1
KCNK15 AND NOT-ANTXR2 AND NOT-IL20RA	1	1	1
KCNK15 AND NOT-GHR AND MS4A1	0.88	0.846154	0.916667
KCNK15 AND NOT-GHR AND SST	0.88	0.846154	0.916667
KCNK15 AND NOT-ASTN1 AND NOT-CLDN8	0.88	0.846154	0.916667
KCNK15 AND NOT-GHR AND TNC	0.88	0.846154	0.916667
KCNK15 AND NOT-FOLR2 AND NOT-NPHS2	0.96	0.923077	1
COMPLEX-ANTXR2/KCNK15/ULBP2	1	1	1
KCNK15 AND NOT-SLC39A8 AND NOT-ENG	0.909091	1	0.833333
COMPLEX-ANTXR2/KCNK15/VCAM1	1	1	1
KCNK15 AND NOT-FOLR2 AND NOT-CNTNAP4	0.96	0.923077	1
KCNK15 AND NOT-PTGER4 AND NOT-TRPM4	0.888889	0.8	1
KCNK15 AND NOT-PHLDB2 AND NOT-CD52	0.888889	0.8	1
KCNK15 AND NOT-GHR AND ROR1	0.869565	0.909091	0.833333
KCNK15 AND NOT-FOLR2 AND CD93	0.869565	0.909091	0.833333
KCNK15 AND NOT-CD163 AND NOT-DPEP1	0.869565	0.909091	0.833333
COMPLEX-EDNRB/KCNK15/CD163	0.869565	0.909091	0.833333
KCNK15 AND NOT-AMHR2 AND NOT-CLDN8	0.88	0.846154	0.916667
KCNK15 AND NOT-TTYH2 AND NOT-IL20RA	0.916667	0.916667	0.916667
KCNK15 AND NOT-NPHS2 AND NOT-GPNMB	0.96	0.923077	1
KCNK15 AND NOT-CD44 AND NOT-DPEP1	0.869565	0.909091	0.833333
KCNK15 AND NOT-MRGPRF AND NOT-GPNMB	1	1	1
KCNK15 AND NOT-CD44 AND NOT-CXCR5	0.869565	0.909091	0.833333
KCNK15 AND NOT-CD44 AND NCAM1	0.869565	0.909091	0.833333
KCNK15 AND NOT-SYT4 AND GAGE1	0.857143	0.75	1
KCNK15 AND NOT-ADAM2 AND ULBP2	0.857143	1	0.75
KCNK15 AND NOT-CD44 AND SST	0.916667	0.916667	0.916667
KCNK15 AND NOT-DPP10 AND NOT-CLDN8	0.857143	0.75	1
MUC16 AND NOT-CD74 AND NOT-THBD	0.8	1	0.666667
MUC16 AND NOT-CD74 AND NOT-SCARA5	0.8	1	0.666667
MUC16 AND NOT-CD44 AND NOT-MSMO1	0.8	1	0.666667
MUC16 AND NOT-CD44 AND NOT-LRRC8E	0.8	1	0.666667
MUC16 AND NOT-CD44 AND NOT-SYT4	0.8	1	0.666667
MUC16 AND NOT-CD44 AND NOT-CDH9	0.8	1	0.666667
NOT-CD44 AND RNF43 AND NOT-ANTXR2	0.833333	0.833333	0.833333
ERBB4 AND NOT-CD74 AND NOT-DTNA	0.8	0.769231	0.833333
NOT-ADAM2 AND LRRC8E AND NOT-PTK7	0.909091	1	0.833333
COMPLEX-CLDN23/EDNRB/KCNK15	0.96	0.923077	1
COMPLEX-EDNRB/KCNK15/VCAM1	0.916667	0.916667	0.916667
KCNK15 AND NOT-TPBG AND NOT-CLDN8	0.88	0.846154	0.916667
KCNK15 AND NOT-KDR AND NOT-CLDN8	0.88	0.846154	0.916667

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
GPR158 AND OR51B6 AND VCAM1	0.864865	0.8	0.941176
GPR158 AND PCYT1A AND NOT-ATP2C1	0.8	0.923077	0.705882
COMPLEX-EDNRB/ATP8B1/PTPRZ1	0.909091	0.9375	0.882353
GPR158 AND OR51B6 AND EDNRB	0.888889	0.842105	0.941176
FZD3 AND VCAM1 AND LAPTM5	0.8	0.923077	0.705882
GPR158 AND OR1A1 AND PCYT1A	0.8	0.923077	0.705882
COMPLEX-EDNRB/FLVCR1/PTPRZ1	0.875	0.933333	0.823529
SYT11 AND IL13RA1 AND FNDC4	0.833333	0.789474	0.882353
TNC AND NOT-KCNK6 AND PRIMA1	0.8125	0.866667	0.764706
LYPD1 AND SLC2A10 AND NOT-TRPM4	0.866667	1	0.764706
COMPLEX-PTPRZ1/BST2/CLDN12	0.909091	0.9375	0.882353
GPR158 AND PCYT1A AND NOT-TMEM127	0.8	0.923077	0.705882
SYT11 AND RHCG AND CBX3	0.83871	0.928571	0.764706
SYT11 AND PCYT1A AND NOT-SLC12A6	0.8	0.923077	0.705882
SYT11 AND WLS AND EDNRB	0.866667	1	0.764706
PAQR6 AND LAPTM5 AND VCAM1	0.8125	0.866667	0.764706
GPR158 AND RHCG AND VCAM1	0.820513	0.727273	0.941176
SLC11A1 AND NOT-IL15RA AND EDNRB	0.827586	1	0.705882
GPR158 AND PCYT1A AND NOT-ABCA3	0.8	0.923077	0.705882
SLCO1C1 AND SLC2A10 AND NOT-TRPM4	0.866667	1	0.764706
OR51B6 AND NOT-CD19 AND CD83	0.857143	0.833333	0.882353
PTPRS AND RHCG AND NOT-ERBB2	0.888889	0.842105	0.941176
PTPRZ1 AND VCAM1 AND NOT-TRPM4	0.9375	1	0.882353
NRCAM AND CD70 AND VCAM1	0.909091	0.9375	0.882353
PTPRZ1 AND VCAM1 AND CD70	0.9375	1	0.882353
GPM6B AND TNC AND NOT-CLDN11	0.8125	0.866667	0.764706
COMPLEX-EDNRB/PTPRZ1/CD70	0.909091	0.9375	0.882353
OR51B6 AND EDNRB AND NOT-RNF43	0.888889	0.842105	0.941176
BMPR1B AND EDNRB AND FAM57A	0.8	0.923077	0.705882
OR51B6 AND EDNRB AND NOT-TRPM4	0.864865	0.8	0.941176
OR51B6 AND EDNRB AND NOT-CLDN8	0.864865	0.8	0.941176
BMPR1B AND PTPRS AND CD70	0.8125	0.866667	0.764706
PTPRS AND NOT-CLDN8 AND SLC39A6	0.83871	0.928571	0.764706
GPR158 AND F2R AND NOT-AJAP1	0.83871	0.928571	0.764706
GPR158 AND F2R AND NOT-ABCG4	0.83871	0.928571	0.764706
LYPD1 AND LAPTM5 AND NOT-VANGL1	0.903226	1	0.823529
GPR158 AND F2R AND NOT-BST2	0.8	0.923077	0.705882
CDH10 AND LAPTM5 AND NOT-VANGL1	0.810811	0.75	0.882353
PCDH8 AND LAPTM5 AND NOT-VANGL1	0.8125	0.866667	0.764706
LYPD1 AND CD163 AND NOT-PPAPDC1B	0.903226	1	0.823529
LYPD1 AND BST2 AND NOT-GPR158	0.83871	0.928571	0.764706
LYPD1 AND BST2 AND NOT-PPAPDC1B	0.875	0.933333	0.823529
LYPD1 AND BST2 AND NOT-FLVCR1	0.875	0.933333	0.823529
SCN2A AND LAPTM5 AND NOT-VANGL1	0.866667	1	0.764706
MLC1 AND NOT-FLVCR1 AND BST2	0.8	0.923077	0.705882
SYT11 AND F2R AND EDNRB	0.866667	1	0.764706
BMPR1B AND NOT-VANGL1 AND LAPTM5	0.820513	0.727273	0.941176
GPR158 AND PCYT1A AND NOT-AJAP1	0.8	0.923077	0.705882
NOT-ATP8B1 AND NOT-CD52 AND LAPTM5	0.8	0.923077	0.705882
SLCO1C1 AND LAPTM5 AND VCAM1	0.823529	0.823529	0.823529
GPR158 AND BST2 AND PCYT1A	0.8	0.923077	0.705882
GPR158 AND PCYT1A AND NOT-ABCG4	0.827586	1	0.705882
LYPD1 AND BST2 AND NOT-CLDN12	0.875	0.933333	0.823529
NOT-ATP8B1 AND EDNRB AND SLC31A1	0.8	0.923077	0.705882
EDNRB AND NOT-IL15RA AND CXCL16	0.83871	0.928571	0.764706
TNC AND NOT-AOC3 AND EDNRB	0.848485	0.875	0.823529
SYT11 AND EDNRB AND CBX3	0.857143	0.833333	0.882353
BMPR1B AND VCAM1 AND LAPTM5	0.842105	0.761905	0.941176
GPR158 AND VCAM1 AND PCYT1A	0.8	0.923077	0.705882
GPR158 AND EDNRB AND PCYT1A	0.8	0.923077	0.705882
Glioma			
GPM6A AND F2R AND NOT-SLC6A5	0.985294	1	0.971014
GPM6A AND F2R AND NOT-HTR7	0.992701	1	0.985507
GPM6A AND F2R AND NOT-TNFRSF18	0.985294	1	0.971014
GPM6A AND F2R AND NOT-C14orf180	0.992701	1	0.985507
GPM6A AND F2R AND NOT-SLC6A3	0.985294	1	0.971014
GPM6A AND F2R AND NOT-OR7A17	0.992701	1	0.985507
GPM6A AND F2R AND NOT-SLC7A4	0.992701	1	0.985507
GPM6A AND F2R AND NOT-SLC30A2	0.985294	1	0.971014
GPM6A AND F2R AND NOT-HRH2	0.977778	1	0.956522
GPM6A AND F2R AND NOT-FZD10	0.977778	1	0.956522
GPM6A AND F2R AND NOT-SLC1A7	0.977778	1	0.956522
GPM6A AND F2R AND NOT-PCDHA5	0.985294	1	0.971014
GPM6A AND F2R AND NOT-MC3R	0.992701	1	0.985507
GPM6A AND F2R AND NOT-AQP8	0.992701	1	0.985507
GPM6A AND F2R AND NOT-MEP1B	0.992701	1	0.985507
GPM6A AND F2R AND NOT-LY9	0.992701	1	0.985507

Antigen Pair Logic	F1	Precision	Recall
KCNK15 AND NOT-ENG AND NOT-GPNMB	1	1	1
KCNK15 AND NOT-SPON2 AND NOT-IL20RA	0.909091	1	0.833333
KCNK15 AND NOT-CD52 AND NOT-GPNMB	0.96	0.923077	1
KCNK15 AND NOT-GPNMB AND NOT-CLDN8	0.96	0.923077	1
KCNK15 AND NOT-GPNMB AND NOT-TRPM4	0.96	0.923077	1
KCNK15 AND NOT-GPNMB AND NOT-IL20RA	0.96	0.923077	1
KCNK15 AND NOT-MET AND NOT-CLDN8	0.916667	0.916667	0.916667
KCNK15 AND NOT-ENG AND NOT-VCAM1	0.827586	0.705882	1
KCNK15 AND NOT-KDR AND NOT-CD52	0.827586	0.705882	1
KCNK15 AND NOT-MET AND NOT-CD52	0.827586	0.705882	1
KCNK15 AND NOT-ENG AND NOT-VCAM1	0.827586	0.705882	1
COMPLEX-EDNRB/IL20RA/KCNK15	0.827586	0.705882	1
KCNK15 AND NOT-ROR1 AND NOT-CLDN8	0.88	0.846154	0.916667
COMPLEX-CD52/KCNK15/SST	0.827586	0.705882	1
COMPLEX-ALDH1A1/KCNK15/VCAM1	0.818182	0.9	0.75
KCNK15 AND NOT-ENG AND NOT-MET	0.8	0.666667	1
MUC16 AND NOT-PROM1 AND CELSR3	0.8	1	0.666667
MUC16 AND NOT-MTUS1 AND NOT-MS4A1	0.8	1	0.666667
MUC16 AND NOT-CLDN23 AND CELSR3	0.8	1	0.666667
COMPLEX-CLDN23/OAS1/KCNK15	0.8	0.666667	1
MUC16 AND NOT-PROM1 AND NOT-JAM3	0.8	1	0.666667
COMPLEX-CLDN23/KCNK15/SST	0.8	0.666667	1
COMPLEX-CLDN23/ENG/KCNK15	0.8	0.666667	1
COMPLEX-CLDN23/ENG/KCNK15	0.8	0.666667	1
KCNK15 AND NOT-ENG AND NOT-ROR1	0.8	0.666667	1
MUC16 AND NOT-MTUS1 AND NOT-CD52	0.8	1	0.666667
COMPLEX-ENG/OAS1/KCNK15	0.8	0.666667	1
COMPLEX-ENG/IL20RA/KCNK15	0.8	0.666667	1
COMPLEX-ENG/STEAP1/KCNK15	0.8	0.666667	1
MUC16 AND NOT-MTUS1 AND NOT-P2RX5	0.8	1	0.666667
COMPLEX-ENG/IL20RA/KCNK15	0.8	0.666667	1
COMPLEX-ENG/KCNK15/SST	0.8	0.666667	1
COMPLEX-ENG/TNC/KCNK15	0.8	0.666667	1
KCNK15 AND NOT-KDR AND NOT-ENG	0.8	0.666667	1
COMPLEX-ENG/IL11RA/KCNK15	0.8	0.666667	1
MUC16 AND NOT-PTK7 AND CELSR3	0.8	1	0.666667
MUC16 AND NOT-IGF1R AND CELSR3	0.8	1	0.666667
MUC16 AND SUSD3 AND NOT-CD19	0.8	1	0.666667
NOT-CD44 AND LRRC8E AND NOT-ADAM2	0.909091	1	0.833333
NOT-ADAM2 AND LRRC8E AND NOT-CD74	0.909091	1	0.833333
NOT-GRID2 AND LRRC8E AND NOT-CD44	0.8	1	0.666667
NOT-CD44 AND LRRC8E AND NOT-MRGPRF	0.8	0.769231	0.833333
EPHA10 AND LRRC8E AND NOT-SELP	0.818182	0.9	0.75
NOT-EMP3 AND LRRC8E AND NOT-CD44	0.8	1	0.666667
NOT-ADAM2 AND LRRC8E AND NOT-MLANA	0.909091	1	0.833333
NOT-CD44 AND LRRC8E AND NOT-MLC1	0.8	1	0.666667
KCNK15 AND NOT-DSC3 AND NOT-ERBB2	1	1	1
KCNK15 AND NOT-TNFSF13B AND NOT-ERBB2	1	1	1
KCNK15 AND NOT-PTPRZ1 AND NOT-ERBB2	1	1	1
KCNK15 AND NOT-PTGDR AND NOT-ERBB2	0.888889	0.8	1
KCNK15 AND NOT-CNIH3 AND NOT-ERBB2	0.888889	0.8	1
KCNK15 AND NOT-SLC44A1 AND NOT-ERBB2	0.8	0.666667	1
KCNK15 AND NOT-LSAMP AND NOT-ERBB2	0.8	0.666667	1
KCNK15 AND NOT-SYT1 AND NOT-ERBB2	0.8	0.666667	1
KCNK15 AND NOT-TRPA1 AND NOT-ERBB2	0.8	0.666667	1
KCNK15 AND CELSR3 AND NOT-ERBB2	0.8	0.666667	1
KCNK15 AND NOT-CLEC4M AND NOT-ERBB2	0.8	0.666667	1
KCNK15 AND NOT-SGCG AND NOT-ERBB2	0.8	0.666667	1
KCNK15 AND NOT-CNGB3 AND NOT-ERBB2	0.8	0.666667	1
COMPLEX-STX6/ERBB2/KCNK15	0.8	0.666667	1
KCNK15 AND NOT-SCTR AND NOT-ERBB2	0.8	0.666667	1
KCNK15 AND NOT-LMBRD1 AND NOT-ERBB2	0.8	0.666667	1
KCNK15 AND NOT-BBS4 AND NOT-ERBB2	0.8	0.666667	1
NOT-SLC16A7 AND PCDH18 AND NOT-ERBB2	0.8	0.666667	1
NOT-PPAP2B AND PCDH18 AND NOT-ERBB2	0.923077	0.857143	1
KCNK15 AND NOT-CLDN15 AND NOT-ERBB2	0.8	0.666667	1
KCNK15 AND NOT-GABRG1 AND NOT-ERBB2	0.8	0.666667	1
KCNK15 AND NOT-LRRN4 AND NOT-ERBB2	0.8	0.666667	1
KCNK15 AND NOT-SYT4 AND NOT-ERBB2	0.8	0.666667	1
MUC16 AND NOT-ERBB2 AND CELSR3	0.8	1	0.666667
Prostate Cancer			
OR51E2 AND NOT-ACE AND NOT-BMP2	1	1	1
COMPLEX-NPY2R/SGCD/OR51E2	1	1	1
OR51E2 AND NOT-DAGLB AND NOT-DAGLA	1	1	1
OR51E2 AND NOT-ACE AND NOT-TLR5	1	1	1
COMPLEX-PIANP/SGCD/OR51E2	1	1	1
COMPLEX-GRPR/SGCD/OR51E2	1	1	1

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
GPM6A AND F2R AND NOT-OR10H1	0.985294	1	0.971014
GPM6A AND F2R AND NOT-GPR87	0.992701	1	0.985507
GPM6A AND F2R AND NOT-ITGA9	0.985294	1	0.971014
PTPRZ1 AND F2R AND NOT-SYT8	0.985294	1	0.971014
GPM6A AND F2R AND NOT-GPR35	0.977778	1	0.956522
GPM6A AND F2R AND NOT-OR2S2	0.977778	1	0.956522
GPM6A AND F2R AND NOT-TAAR5	0.992701	1	0.985507
GPM6A AND F2R AND NOT-GJB3	0.992701	1	0.985507
COMPLEX-SULF2/PTPRZ1/SLC1A6	0.985294	1	0.971014
GPM6A AND F2R AND NOT-ILDR1	0.992701	1	0.985507
GPM6A AND F2R AND NOT-OPN1SW	0.985294	1	0.971014
GPM6A AND F2R AND NOT-KLRD1	0.992701	1	0.985507
GPM6A AND F2R AND CD63	0.992701	1	0.985507
GPM6A AND F2R AND NOT-OR10A4	0.992701	1	0.985507
PTPRZ1 AND F2R AND NOT-WNT3	0.985294	1	0.971014
PTPRZ1 AND F2R AND NOT-GJB5	0.985294	1	0.971014
GPM6A AND F2R AND NOT-ADAM28	0.992701	1	0.985507
GPM6A AND F2R AND NOT-ICAM2	0.992701	1	0.985507
GPM6A AND F2R AND NOT-PAQR9	0.992701	1	0.985507
GPM6A AND F2R AND NOT-ENTPD2	0.992701	1	0.985507
GPM6A AND F2R AND NOT-PAQR6	0.992701	1	0.985507
GPM6A AND F2R AND NOT-C1QTNF1	0.992701	1	0.985507
GPM6A AND F2R AND NOT-ZDHHC20	0.992701	1	0.985507
GPM6A AND F2R AND NOT-KCNE2	0.992701	1	0.985507
GPM6A AND F2R AND NOT-SLC38A4	0.992701	1	0.985507
GPM6A AND F2R AND NOT-PIGR	0.992701	1	0.985507
GPM6A AND F2R AND SLC25A3	0.992701	1	0.985507
GPM6A AND F2R AND NOT-CLEC4D	0.992701	1	0.985507
GPM6A AND F2R AND NOT-MGAM	0.985294	1	0.971014
GPM6A AND F2R AND NOT-NCAM2	0.992701	1	0.985507
GPM6A AND F2R AND NOT-RXFP2	0.992701	1	0.985507
GPM6A AND F2R AND NOT-OR51M1	0.985294	1	0.971014
GPM6A AND F2R AND NOT-HCRTR1	0.985294	1	0.971014
GPM6A AND TSPAN6 AND NOT-SLC9B1	0.970588	0.985075	0.956522
GPM6A AND TSPAN6 AND NOT-FGF6	0.970588	0.985075	0.956522
GPM6A AND TSPAN6 AND NOT-ABCG5	0.970588	0.985075	0.956522
GPM6A AND TSPAN6 AND NOT-OR3A2	0.970588	0.985075	0.956522
GPM6A AND TSPAN6 AND NOT-ZP4	0.970588	0.985075	0.956522
GPM6A AND F2R AND NOT-GPR18	0.992701	1	0.985507
GPM6A AND TSPAN6 AND NOT-SLC26A8	0.970588	0.985075	0.956522
GPM6A AND TSPAN6 AND NOT-BEST2	0.970588	0.985075	0.956522
GPM6A AND TSPAN6 AND NOT-MUC17	0.970588	0.985075	0.956522
GPM6A AND TSPAN6 AND NOT-LRRN4	0.970588	0.985075	0.956522
GPM6A AND TSPAN6 AND NOT-KCNJ9	0.970588	0.985075	0.956522
GPM6A AND F2R AND NOT-CDH24	0.977778	1	0.956522
GPM6A AND F2R AND NOT-ITGAD	0.970149	1	0.942029
COMPLEX-TNFRSF12A/SULF2/PTPRZ1	0.970149	1	0.942029
COMPLEX-IFNAR2/SULF2/PTPRZ1	0.970149	1	0.942029
GPM6A AND F2R AND NOT-PTPRH	0.977778	1	0.956522
GPM6A AND PLGRKT AND NOT-FGF6	0.970149	1	0.942029
PTPRZ1 AND NOT-LRP11 AND NOT-KCNA7	0.985507	0.985507	0.985507
GPM6A AND TSPAN6 AND NOT-SLC22A11	0.977778	1	0.956522
GPM6A AND TSPAN6 AND NOT-KCNA4	0.970588	0.985075	0.956522
GPM6A AND TSPAN6 AND NOT-GJB4	0.970588	0.985075	0.956522
GPM6A AND TSPAN6 AND ATP13A5	0.970588	0.985075	0.956522
GPM6A AND PLGRKT AND NOT-MUC17	0.970149	1	0.942029
PTPRZ1 AND NOT-LRP11 AND NOT-FUT3	0.985294	1	0.971014
GPM6A AND SLC40A1 AND NOT-SLCO1A2	0.970588	0.985075	0.956522
PTPRZ1 AND F2R AND NOT-PVRL4	0.985294	1	0.971014
PTPRZ1 AND F2R AND NOT-GPR87	0.985294	1	0.971014
COMPLEX-CRB1/KIAA1324/PTPRZ1	0.970588	0.985075	0.956522
PTPRZ1 AND NOT-GJB5 AND NOT-GRM3	0.992701	1	0.985507
PTPRZ1 AND NOT-PVRL4 AND NOT-GRM3	0.992701	1	0.985507
PTPRZ1 AND NOT-GRM3 AND NOT-GPR87	0.992701	1	0.985507
PTPRZ1 AND NOT-LRP11 AND NOT-MTNR1B	0.985507	0.985507	0.985507
GPM6A AND PLGRKT AND NOT-LRRN4	0.970149	1	0.942029
GPM6A AND PLGRKT AND NOT-ZP4	0.970149	1	0.942029
GPM6A AND PLGRKT AND NOT-BEST2	0.970149	1	0.942029
PTPRZ1 AND NOT-CD3G AND NOT-LRP11	0.977778	1	0.956522
COMPLEX-TNFSF13B/SULF2/PTPRZ1	0.977778	1	0.956522
PTPRZ1 AND NOT-LRP11 AND HLA-B	0.992701	1	0.985507
COMPLEX-MSR1/SULF2/PTPRZ1	0.985294	1	0.971014
COMPLEX-MERTK/SULF2/PTPRZ1	0.977778	1	0.956522
PTPRZ1 AND NOT-EFNA5 AND NOT-PAQR6	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-EFNA5 AND IFNGR2	0.985294	1	0.971014
PTPRZ1 AND NOT-EFNA5 AND SLC39A14	0.985294	1	0.971014
PTPRZ1 AND NOT-LRP11 AND NOT-SLC6A3	0.978102	0.985294	0.971014

Antigen Pair Logic	F1	Precision	Recall
OR51E2 AND NOT-ACE AND NOT-KCNK1	1	1	1
OR51E2 AND NOT-ACE AND NOT-ADRA2C	1	1	1
OR51E2 AND NOT-SI AND NOT-MYADM	1	1	1
OR51E2 AND NOT-ABCG2 AND NOT-ACE	1	1	1
OR51E2 AND NOT-FLRT3 AND NOT-SI	1	1	1
OR51E2 AND NOT-ACE AND NOT-CD69	1	1	1
OR51E2 AND NOT-FAT4 AND NOT-SI	1	1	1
OR51E2 AND NOT-FAT4 AND NOT-ACE	1	1	1
OR51E2 AND NOT-ACE AND KCNG3	1	1	1
OR51E2 AND NOT-SLC25A4 AND PTPRCAP	1	1	1
OR51E2 AND NOT-STX2 AND NOT-SLC2A5	1	1	1
OR51E2 AND NOT-ACE AND NOT-SLC8A1	1	1	1
OR51E2 AND NOT-CX3CL1 AND NOT-SLC2A5	1	1	1
OR51E2 AND NOT-ACE AND NOT-LEPR	1	1	1
OR51E2 AND NOT-VAMP1 AND NOT-SLC2A5	1	1	1
COMPLEX-GJB6/SGCD/OR51E2	1	1	1
OR51E2 AND NOT-MCAM AND NOT-DAGLA	1	1	1
COMPLEX-GLDN/LSR/OR51E2	1	1	1
COMPLEX-SGCD/ABCC8/OR51E2	1	1	1
COMPLEX-ITPR3/SGCD/OR51E2	1	1	1
OR51E2 AND NOT-CX3CL1 AND NOT-ACE	1	1	1
OR51E2 AND NOT-KCNE4 AND NOT-ACE	1	1	1
COMPLEX-PCDHB7/SGCD/OR51E2	1	1	1
OR51E2 AND NOT-ACE AND NOT-CLCA4	1	1	1
OR51E2 AND NOT-STX4 AND NOT-DAGLA	1	1	1
OR51E2 AND NOT-ACE AND NOT-FLRT3	1	1	1
OR51E2 AND NOT-CD247 AND NOT-VIMP	1	1	1
OR51E2 AND NOT-SLC24A3 AND GABRB3	1	1	1
OR51E2 AND NOT-EFNB3 AND NPFFR2	1	1	1
OR51E2 AND NOT-STX4 AND NOT-SLC2A5	1	1	1
OR51E2 AND NOT-ACE AND NOT-ENPP2	1	1	1
COMPLEX-OR51E2/CACNA1H/CD40	1	1	1
OR51E2 AND NOT-STX7 AND NOT-DAGLA	1	1	1
COMPLEX-NTSR1/SGCD/OR51E2	1	1	1
COMPLEX-FZD2/SGCD/OR51E2	1	1	1
OR51E2 AND NOT-SEMA6D AND NOT-ACE	1	1	1
OR51E2 AND NOT-EFNB3 AND NOT-CD81	1	1	1
COMPLEX-GLDN/OR51E2/CDH1	1	1	1
COMPLEX-KCND1/SGCD/OR51E2	1	1	1
OR51E2 AND NOT-TPCN2 AND NOT-SLC2A5	1	1	1
OR51E2 AND NOT-FAT4 AND NOT-ACE	1	1	1
OR51E2 AND NOT-TM7SF3 AND NOT-ACE	1	1	1
COMPLEX-TACSTD2/SGCD/OR51E2	1	1	1
OR51E2 AND NOT-DNER AND NOT-EFNB3	1	1	1
OR51E2 AND NOT-SGCA AND NOT-DAGLA	1	1	1
COMPLEX-NRG1/SGCD/OR51E2	1	1	1
OR51E2 AND NOT-KCNJ15 AND KCNMB2	1	1	1
OR51E2 AND NOT-ADCY5 AND NOT-DAGLA	1	1	1
OR51E2 AND NOT-SLC23A1 AND NOT-DAGLA	1	1	1
OR51E2 AND NOT-SLC24A3 AND NOT-DAGLA	1	1	1
OR51E2 AND NOT-SLC23A1 AND NOT-SLC2A5	1	1	1
OR51E2 AND NOT-JPH1 AND NOT-ACE	1	1	1
OR51E2 AND NOT-TM7SF3 AND SIRPG	1	1	1
OR51E2 AND VMP1 AND NOT-MYADM	1	1	1
COMPLEX-ADRB1/GLDN/OR51E2	1	1	1
COMPLEX-SGCD/SLC8A1/OR51E2	1	1	1
OR51E2 AND NOT-ACE AND NOT-FLRT3	1	1	1
OR51E2 AND NOT-FAT4 AND NOT-SI	1	1	1
OR51E2 AND NOT-ACE AND NOT-CD69	1	1	1
OR51E2 AND NOT-ACE AND NOT-SLC16A14	1	1	1
OR51E2 AND NOT-ADCY5 AND PTPRCAP	1	1	1
OR51E2 AND NOT-SLC2A5 AND NOT-MYADM	1	1	1
OR51E2 AND NOT-SLC25A4 AND GABRB3	1	1	1
OR51E2 AND NOT-ACE AND NOT-PTPRE	1	1	1
OR51E2 AND NOT-ACE AND NOT-ADCY3	1	1	1
OR51E2 AND NOT-PDE6B AND NOT-PLP2	1	1	1
OR51E2 AND NOT-SLC2A5 AND NOT-WDR19	1	1	1
COMPLEX-OR10A5/SGCD/OR51E2	1	1	1
OR51E2 AND NOT-SDC3 AND NOT-ACE	1	1	1
OR51E2 AND NOT-ACE AND NOT-CNTN1	1	1	1
OR51E2 AND NOT-PDE6B AND KCNMB2	1	1	1
OR51E2 AND NOT-NDRG4 AND NOT-SLC2A5	1	1	1
OR51E2 AND NOT-HAS3 AND NOT-DAGLA	1	1	1
OR51E2 AND NOT-ACE AND NOT-SIRPA	1	1	1
OR51E2 AND NOT-HAS3 AND PTPRCAP	1	1	1
OR51E2 AND NOT-NRSN2 AND NOT-SLC2A5	1	1	1
OR51E2 AND NOT-DYSF AND PTPRCAP	1	1	1

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
PTPRZ1 AND NOT-LRP11 AND NOT-SLC6A5	0.978102	0.985294	0.971014
GPM6A AND TSPAN6 AND CD63	0.977778	1	0.956522
GPM6A AND HLA-B AND NOT-AQP8	0.977778	1	0.956522
PTPRZ1 AND NOT-EFNA5 AND NRP1	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-LRP11 AND NOT-OR1D2	0.985507	0.985507	0.985507
COMPLEX-ITGB8/SULF2/PTPRZ1	0.985294	1	0.971014
COMPLEX-SLCO2B1/KIAA1324/PTPRZ1	0.985507	0.985507	0.985507
GPM6A AND TSPAN6 AND NOT-GPR87	0.970588	0.985075	0.956522
GPM6A AND TSPAN6 AND NOT-C14orf180	0.970588	0.985075	0.956522
GPM6A AND TSPAN6 AND ATP12A	0.970588	0.985075	0.956522
GPM6A AND TSPAN6 AND NOT-HTR7	0.970588	0.985075	0.956522
GPM6A AND TSPAN6 AND NOT-SLC7A4	0.970588	0.985075	0.956522
COMPLEX-CNGA3/KIAA1324/PTPRZ1	0.978102	0.985294	0.971014
GPM6A AND TSPAN6 AND NOT-OR7A17	0.970588	0.985075	0.956522
GPM6A AND TSPAN6 AND NOT-TMEM219	0.970588	0.985075	0.956522
GPM6A AND SLC40A1 AND NOT-PAQR6	0.970588	0.985075	0.956522
COMPLEX-SULF2/PTPRZ1/GPR65	0.985294	1	0.971014
COMPLEX-SULF2/PTPRZ1/SLC16A1	0.970149	1	0.942029
PTPRZ1 AND NOT-EFNA5 AND NOT-PRRG1	0.978102	0.985294	0.971014
PTPRZ1 AND HLA-G AND NOT-OR7A10	0.992701	1	0.985507
PTPRZ1 AND NOT-EFNA5 AND NOT-SLC22A1	0.985294	1	0.971014
COMPLEX-GRIK3/NRCAM/PTPRZ1	0.992701	1	0.985507
PTPRZ1 AND NOT-EFNA5 AND STX6	0.985294	1	0.971014
GPM6A AND TSPAN6 AND NOT-KCNB1	0.970588	0.985075	0.956522
PTPRZ1 AND NOT-LRP11 AND TMBIM6	0.992701	1	0.985507
PTPRZ1 AND NOT-LRP11 AND TMBIM6	0.992701	1	0.985507
PTPRZ1 AND NOT-LRP11 AND NOT-SLC6A2	0.992701	1	0.985507
PTPRZ1 AND NOT-LRP11 AND NOT-CD40LG	0.978102	0.985294	0.971014
GPM6A AND TSPAN6 AND NOT-AQP8	0.977778	1	0.956522
GPM6A AND TSPAN6 AND NOT-MEP1B	0.970588	0.985075	0.956522
GPM6A AND TSPAN6 AND NOT-MC3R	0.970588	0.985075	0.956522
GPM6A AND TSPAN6 AND NOT-LY9	0.970588	0.985075	0.956522
COMPLEX-SULF2/PTPRZ1/TGFA	0.970149	1	0.942029
GPM6A AND TSPAN6 AND NOT-KLRD1	0.970588	0.985075	0.956522
GPM6A AND TSPAN6 AND NOT-PIGR	0.970588	0.985075	0.956522
GPM6A AND SLC40A1 AND NOT-RHBDL2	0.970588	0.985075	0.956522
GPM6A AND TSPAN6 AND NOT-GPR18	0.970588	0.985075	0.956522
PTPRZ1 AND NOT-LRP11 AND TSPAN6	0.977778	1	0.956522
PTPRZ1 AND NOT-LRP11 AND NOT-BPI	0.992701	1	0.985507
GPM6A AND SLC40A1 AND NOT-HTR7	0.970588	0.985075	0.956522
COMPLEX-ADORA3/KIAA1324/PTPRZ1	0.985507	0.985507	0.985507
COMPLEX-CD300A/SULF2/PTPRZ1	0.985294	1	0.971014
PTPRZ1 AND NOT-SYT2 AND NOT-LRP11	0.985507	0.985507	0.985507
GPM6A AND CD63 AND NOT-TNFSF9	0.992701	1	0.985507
PTPRZ1 AND NOT-LRP11 AND NOT-SLC5A1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-LRP11 AND NOT-ATP4A	0.978102	0.985294	0.971014
GPM6A AND PLGRKT AND NOT-HTR7	0.970149	1	0.942029
PTPRZ1 AND NOT-LRP11 AND NOT-HRH2	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-LRP11 AND NOT-MAS1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-LRP11 AND NOT-LTA	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-LRP11 AND NOT-DIO3	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-LRP11 AND NOT-MAS1	0.985507	0.985507	0.985507
COMPLEX-NRCAM/PTPRZ1/GPR65	0.977778	1	0.956522
PTPRZ1 AND NOT-LRP11 AND NOT-LTK	0.985294	1	0.971014
COMPLEX-NRCAM/PTPRZ1/SLC16A1	0.970149	1	0.942029
PTPRZ1 AND NOT-EFNA5 AND NOT-GJB1	0.978102	0.985294	0.971014
GPM6A AND TSPAN6 AND NOT-IL4R	0.970588	0.985075	0.956522
GPM6A AND TSPAN6 AND NOT-CD300LG	0.970588	0.985075	0.956522
GPM6A AND TSPAN6 AND NOT-ADTRP	0.970588	0.985075	0.956522
GPM6A AND TSPAN6 AND NOT-ZDHHC20	0.970588	0.985075	0.956522
GPM6A AND TSPAN6 AND NOT-GJB3	0.970588	0.985075	0.956522
PTPRZ1 AND NOT-LRP11 AND NOT-KCNB2	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-EFNA5 AND NOT-GPR62	0.978102	0.985294	0.971014
COMPLEX-MERTK/NRCAM/PTPRZ1	0.977778	1	0.956522
GPM6A AND TSPAN6 AND NOT-PAQR6	0.970588	0.985075	0.956522
PTPRZ1 AND NOT-LRP11 AND NOT-TRPM6	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-LRP11 AND NOT-HPN	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-LRP11 AND ITGAE	0.985294	1	0.971014
GPM6A AND TSPAN6 AND NOT-ILDR1	0.970588	0.985075	0.956522
COMPLEX-CD300A/NRCAM/PTPRZ1	0.992701	1	0.985507
GPM6A AND TSPAN6 AND NOT-SELE	0.970588	0.985075	0.956522
PTPRZ1 AND NOT-LRP11 AND NOT-HTR7	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-LRP11 AND IFNGR2	0.992701	1	0.985507
PTPRZ1 AND NOT-LRP11 AND IFNGR2	0.992701	1	0.985507
PTPRZ1 AND NOT-LRP11 AND NOT-RHAG	0.992701	1	0.985507
PTPRZ1 AND NOT-LRP11 AND NOT-CXCR6	0.977778	1	0.956522
PTPRZ1 AND HLA-B AND NOT-SLC6A14	0.992701	1	0.985507

Antigen Pair Logic	F1	Precision	Recall
OR51E2 AND NOT-UNC5B AND SLCO1A2	1	1	1
TRPM8 AND NOT-CAV1 AND APOLD1	1	1	1
TRPM8 AND NOT-SLC2A5 AND NOT-TENM2	1	1	1
OR51E2 AND NOT-SI AND SLCO1A2	1	1	1
OR51E2 AND NOT-ACE AND SLCO1A2	1	1	1
TRPM8 AND NOT-SLC2A5 AND NOT-STX4	1	1	1
TRPM8 AND NOT-SLC2A5 AND NOT-VAMP1	1	1	1
TRPM8 AND NOT-SLC2A5 AND NOT-FZD7	1	1	1
TRPM8 AND NOT-CAV1 AND CD69	1	1	1
OR51E2 AND NOT-SYT4 AND NOT-SLC2A5	1	1	1
TRPM8 AND NOT-SLC2A5 AND NOT-SLC25A4	1	1	1
OR51E2 AND NOT-KCNJ15 AND GPR19	1	1	1
OR51E2 AND NOT-CD247 AND SLCO1A2	1	1	1
OR51E2 AND NOT-IL17RA AND NOT-DAGLA	1	1	1
OR51E2 AND SLCO1A2 AND CD63	1	1	1
TRPM8 AND NOT-SLC18A2 AND NOT-CLSTN2	1	1	1
OR51E2 AND NOT-FMNL1 AND NOT-PLP2	1	1	1
OR51E2 AND NOT-SLC39A2 AND NPFFR2	1	1	1
TRPM8 AND NOT-FADS2 AND NOT-FZD1	1	1	1
TRPM8 AND NOT-ATP1A2 AND NOT-FADS2	1	1	1
OR51E2 AND NOT-SLC39A2 AND NOT-VIMP	1	1	1
TRPM8 AND NOT-GPR161 AND KCNS3	1	1	1
TRPM8 AND NOT-GPR161 AND LDLRAD3	1	1	1
OR51E2 AND NOT-PTH1R AND SLCO1A2	1	1	1
TRPM8 AND NOT-CAV1 AND NOT-CLSTN2	1	1	1
TRPM8 AND NOT-SLC2A5 AND NOT-GGT7	1	1	1
OR51E2 AND NOT-SI AND SLCO1A2	1	1	1
OR51E2 AND NOT-PTH1R AND GPR19	1	1	1
OR51E2 AND NOT-FMNL1 AND KCNMB2	1	1	1
TRPM8 AND NOT-FADS2 AND NOT-SDC3	1	1	1
OR51E2 AND NOT-FMNL1 AND NOT-MYADM	1	1	1
OR51E2 AND NOT-CD247 AND GPR19	1	1	1
TRPM8 AND NOT-SLC2A5 AND NOT-SGCE	1	1	1
OR51E2 AND NOT-DYSF AND GABRB3	1	1	1
OR51E2 AND GJA1 AND SLCO1A2	1	1	1
OR51E2 AND NOT-PDE6B AND SLCO1A2	1	1	1
TRPM8 AND NOT-CAV1 AND PANX1	1	1	1
OR51E2 AND NOT-PDE6B AND SLCO1A2	1	1	1
TRPM8 AND NOT-SLC2A5 AND NOT-MYOF	1	1	1
TRPM8 AND NOT-CAV1 AND SLC6A14	1	1	1
TRPM8 AND NOT-CAV1 AND PANX1	1	1	1
COMPLEX-PTPRF/OR51E2/DYSF	1	1	1
TRPM8 AND NOT-CAV1 AND ICAM3	1	1	1
TRPM8 AND NOT-CAV1 AND IFNGR2	1	1	1
TRPM8 AND NOT-CAV1 AND NUCB2	1	1	1
TRPM8 AND NOT-SLC2A5 AND NOT-STX2	1	1	1
TRPM8 AND NOT-SLC2A5 AND NOT-SLC16A9	1	1	1
TRPM8 AND NOT-SLC2A5 AND NOT-KCNE4	1	1	1
OR51E2 AND NOT-DYSF AND NOT-DAGLA	1	1	1
TRPM8 AND NOT-SLC2A5 AND NOT-SLC16A2	1	1	1
OR51E2 AND NOT-SLC39A2 AND NOT-PLP2	1	1	1
OR51E2 AND NOT-SYT4 AND NOT-SLC2A5	1	1	1
TRPM8 AND NOT-CAV1 AND MYADM	1	1	1
TRPM8 AND NOT-SLC2A5 AND NOT-PANX1	1	1	1
TRPM8 AND NOT-SLC2A5 AND NOT-MPZ	1	1	1
TRPM8 AND NOT-CAV1 AND CD69	1	1	1
OR51E2 AND NOT-UNC5B AND GPR19	1	1	1
COMPLEX-KCNJ6/SGCD/OR51E2	1	1	1
TRPM8 AND NOT-SLC2A5 AND NOT-PTGER2	1	1	1
TRPM8 AND NOT-SLC2A5 AND NOT-SLC24A3	1	1	1
TRPM8 AND NOT-SLC2A5 AND NOT-SGCA	1	1	1
OR51E2 AND NOT-FMNL1 AND NOT-VIMP	1	1	1
TRPM8 AND NOT-UNC5B AND NOT-PLP2	1	1	1
TRPM8 AND NOT-SLC2A5 AND NOT-ABHD6	1	1	1
TRPM8 AND NOT-SLC2A5 AND NOT-JAM3	1	1	1
TRPM8 AND NOT-SLC2A5 AND NOT-GPR161	1	1	1
TRPM8 AND NOT-GPR161 AND NOT-CLSTN2	1	1	1
TRPM8 AND NOT-PANX1 AND LDLRAD3	1	1	1
TRPM8 AND NOT-GPR161 AND SLC6A14	1	1	1
TRPM8 AND NOT-CAV1 AND NOT-UNC5B	1	1	1
TRPM8 AND NOT-UNC5B AND NOT-CD81	1	1	1
OR51E2 AND NOT-ACE AND NOT-PAQR8	1	1	1
OR51E2 AND NOT-EFNB3 AND SLCO1A2	1	1	1
COMPLEX-SYT4/OR51E2/FADS2	1	1	1
OR51E2 AND NOT-SLC2A5 AND NOT-EMP3	1	1	1
TRPM8 AND NOT-SLC2A5 AND NOT-TPCN2	1	1	1
TRPM8 AND NOT-SLC2A5 AND NOT-STX2	1	1	1

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
PTPRZ1 AND HLA-B AND GPM6A	0.992701	1	0.985507
PTPRZ1 AND HLA-B AND PCDH17	0.992701	1	0.985507
GPM6A AND PRCP AND NOT-PAQR6	0.970149	1	0.942029
COMPLEX-SULF2/PTPRZ1/LRP11	0.970588	0.985075	0.956522
PTPRZ1 AND HLA-B AND NOT-SPINT1	0.992701	1	0.985507
GPM6A AND TSPAN6 AND NOT-OR10A4	0.970588	0.985075	0.956522
PTPRZ1 AND HLA-B AND NOT-GPR87	0.992701	1	0.985507
PTPRZ1 AND NOT-LRP11 AND HLA-G	0.992701	1	0.985507
PTPRZ1 AND HLA-B AND GPM6B	0.992701	1	0.985507
PTPRZ1 AND HLA-B AND NOT-PROM2	0.992701	1	0.985507
PTPRZ1 AND HLA-B AND NOT-PERP	0.992701	1	0.985507
PTPRZ1 AND NOT-LRP11 AND HLA-G	0.992701	1	0.985507
COMPLEX-CD320/KIAA1324/PTPRZ1	0.985507	0.985507	0.985507
GPM6A AND HLA-B AND NOT-RHBDL2	0.977778	1	0.956522
PTPRZ1 AND NOT-LRP11 AND NOT-CLDN6	0.978102	0.985294	0.971014
COMPLEX-MSR1/PTPRZ1/BCAN	0.992701	1	0.985507
PTPRZ1 AND NOT-LRP11 AND CBX3	0.992701	1	0.985507
GPM6A AND TSPAN6 AND NOT-FOLR1	0.970588	0.985075	0.956522
GPM6A AND TSPAN6 AND NOT-RAET1E	0.970588	0.985075	0.956522
PTPRZ1 AND NOT-LRP11 AND NOT-GPA33	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-LRP11 AND NOT-GPA33	0.985507	0.985507	0.985507
COMPLEX-MERTK/PTPRZ1/BCAN	0.985294	1	0.971014
COMPLEX-CD300A/PTPRZ1/BCAN	0.992701	1	0.985507
PTPRZ1 AND CBX3 AND NOT-SYT8	0.977778	1	0.956522
PTPRZ1 AND CBX3 AND NOT-GJB5	0.977778	1	0.956522
PTPRZ1 AND NOT-EPCAM AND IFNGR2	0.992701	1	0.985507
PTPRZ1 AND NOT-LRP11 AND NOT-SSTR3	0.985507	0.985507	0.985507
GPM6A AND TSPAN6 AND NOT-SLAMF7	0.970588	0.985075	0.956522
PTPRZ1 AND CBX3 AND NOT-SPINT1	0.977778	1	0.956522
PTPRZ1 AND CBX3 AND NOT-PROM2	0.977778	1	0.956522
PTPRZ1 AND CBX3 AND GPM6A	0.977778	1	0.956522
COMPLEX-ABCA5/ITGB8/PTPRZ1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-EPCAM AND HLA-G	0.992701	1	0.985507
PTPRZ1 AND CBX3 AND GPM6B	0.977778	1	0.956522
PTPRZ1 AND CBX3 AND NOT-GPR87	0.977778	1	0.956522
PTPRZ1 AND CBX3 AND PCDH17	0.977778	1	0.956522
PTPRZ1 AND CBX3 AND NOT-SLC6A14	0.977778	1	0.956522
PTPRZ1 AND CBX3 AND NOT-PERP	0.977778	1	0.956522
PTPRZ1 AND HLA-B AND NOT-CLDN23	0.992701	1	0.985507
PTPRZ1 AND HLA-B AND NOT-TRPM4	0.977778	1	0.956522
GPM6A AND SLC40A1 AND NOT-FOLR1	0.970588	0.985075	0.956522
PTPRZ1 AND HLA-B AND NCAM1	0.992701	1	0.985507
COMPLEX-HRH1/PTPRZ1/BCAN	0.970149	1	0.942029
COMPLEX-PTPRZ1/BCAN/GPR65	0.992701	1	0.985507
GPM6A AND HLA-B AND NOT-RAET1E	0.963504	0.970588	0.956522
PTPRZ1 AND NOT-SYT2 AND NOT-CLDN23	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-SYT2 AND NOT-TRPM4	0.970588	0.985075	0.956522
GPM6A AND TSPAN6 AND NOT-ERBB3	0.962963	0.984848	0.942029
GPM6A AND TSPAN6 AND NOT-CLDN1	0.962963	0.984848	0.942029
PTPRZ1 AND NOT-SYT2 AND NCAM1	0.985507	0.985507	0.985507
COMPLEX-PMEPA1/PTPRZ1/BCAN	0.962406	1	0.927536
COMPLEX-ITGA7/PTPRZ1/BCAN	0.962406	1	0.927536
COMPLEX-MET/NRCAM/PTPRZ1	0.962406	1	0.927536
COMPLEX-SLCO2B1/CLDN7/PTPRZ1	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-LRP11 AND NOT-CLDN9	0.985507	0.985507	0.985507
COMPLEX-SLC6A14/PTPRZ1/BCAN	0.970149	1	0.942029
PTPRZ1 AND NOT-LRP11 AND NOT-ULBP3	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-LRP11 AND NOT-SSTR5	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-LRP11 AND NOT-DNAJB8	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-LRP11 AND NOT-SSTR5	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-LRP11 AND NOT-FCRL2	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-LRP11 AND NOT-CLDN4	0.985507	0.985507	0.985507
PTPRZ1 AND CBX3 AND NOT-F11R	0.977778	1	0.956522
PTPRZ1 AND CBX3 AND NOT-TMEM30B	0.977778	1	0.956522
PTPRZ1 AND CBX3 AND IGSF11	0.977778	1	0.956522
PTPRZ1 AND CBX3 AND NOT-MGST2	0.977778	1	0.956522
PTPRZ1 AND CBX3 AND APLP1	0.977778	1	0.956522
PTPRZ1 AND NOT-EPCAM AND NOT-AQP8	0.992701	1	0.985507
PTPRZ1 AND CBX3 AND NOT-GJB3	0.977778	1	0.956522
PTPRZ1 AND CBX3 AND ATP1B2	0.977778	1	0.956522
PTPRZ1 AND CBX3 AND NOT-PVRL4	0.977778	1	0.956522
PTPRZ1 AND HLA-B AND NOT-SDC1	0.992701	1	0.985507
PTPRZ1 AND NOT-EPCAM AND TMBIM6	0.992701	1	0.985507
PTPRZ1 AND SLC40A1 AND NOT-CLDN23	0.977778	1	0.956522
PTPRZ1 AND NOT-SYT2 AND NOT-SDC1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-RAET1E AND HLA-G	0.985294	1	0.971014
PTPRZ1 AND SLC40A1 AND NOT-TRPM4	0.962406	1	0.927536

Antigen Pair Logic	F1	Precision	Recall
OR51E2 AND NOT-KCNJ15 AND SLCO1A2	1	1	1
COMPLEX-ANO6/OXTR/OR51E2	1	1	1
TRPM8 AND NOT-CAV1 AND GOLM1	1	1	1
TRPM8 AND NOT-CAV1 AND FZD5	1	1	1
TRPM8 AND NOT-CAV1 AND SLC2A3	1	1	1
TRPM8 AND NOT-POPDC2 AND NOT-UNC5B	1	1	1
TRPM8 AND NOT-CAV1 AND SLC19A2	1	1	1
TRPM8 AND NOT-SLC2A5 AND NOT-ABCC6	1	1	1
OR51E2 AND NOT-EFNB3 AND GPR19	1	1	1
TRPM8 AND NOT-SLC2A5 AND NOT-STX7	1	1	1
OR51E2 AND NOT-DNER AND NOT-FMNL1	1	1	1
OR51E2 AND NOT-CD247 AND SLCO1A2	1	1	1
TRPM8 AND NOT-CAV1 AND NOT-CLSTN2	1	1	1
TRPM8 AND NOT-PCDH7 AND LDLRAD3	1	1	1
COMPLEX-PTCHD1/OXTR/OR51E2	1	1	1
TRPM8 AND NOT-ROR2 AND NOT-SLC2A5	1	1	1
TRPM8 AND NOT-ROR2 AND NOT-CLSTN2	1	1	1
COMPLEX-OMG/SGCD/OR51E2	1	1	1
OR51E2 AND NOT-EFNB3 AND GPR19	1	1	1
STEAP2 AND NOT-CAV1 AND SLC39A14	1	1	1
STEAP2 AND NOT-GPR161 AND SLC6A14	1	1	1
STEAP2 AND NOT-SLC16A14 AND NOT-ACE	1	1	1
STEAP2 AND NOT-PCDH7 AND SLC6A14	1	1	1
COMPLEX-SGCD/CA9/OR51E2	1	1	1
STEAP2 AND NOT-CAV1 AND CXCR4	1	1	1
STEAP2 AND NOT-CAV1 AND SLC6A14	1	1	1
STEAP2 AND NOT-CAV1 AND SLC30A5	1	1	1
STEAP2 AND NOT-CAV1 AND ICAM3	1	1	1
STEAP2 AND NOT-CAV1 AND MYADM	1	1	1
STEAP2 AND NOT-CAV1 AND VMP1	1	1	1
STEAP2 AND NOT-CAV1 AND IFNGR2	1	1	1
STEAP2 AND NOT-CAV1 AND GOLM1	1	1	1
STEAP2 AND NOT-CAV1 AND CD69	1	1	1
STEAP2 AND NOT-SLC2A5 AND NOT-SGCE	1	1	1
STEAP2 AND NOT-PCDH7 AND LDLRAD3	1	1	1
STEAP2 AND NOT-CAV1 AND SLC39A14	1	1	1
STEAP2 AND NOT-SLC2A5 AND NOT-STX7	1	1	1
STEAP2 AND NOT-ROR2 AND SLC6A14	1	1	1
STEAP2 AND NOT-PCDH7 AND LDLRAD3	1	1	1
STEAP2 AND NOT-SGCE AND LDLRAD3	1	1	1
STEAP2 AND NOT-CAV1 AND MYADM	1	1	1
STEAP2 AND NOT-CAV1 AND FZD5	1	1	1
STEAP2 AND NOT-CAV1 AND CXCR4	1	1	1
STEAP2 AND NOT-MYADM AND LDLRAD3	1	1	1
COMPLEX-KDR/NPY2R/OR51E2	1	1	1
STEAP2 AND NOT-SLC2A5 AND NOT-SGCE	1	1	1
COMPLEX-KDR/TENM2/OR51E2	1	1	1
OR51E2 AND NOT-DKK1 AND NOT-SLC2A5	1	1	1
OR51E2 AND NOT-ACE AND NOT-ITGB6	1	1	1
OR51E2 AND NOT-DKK1 AND NOT-ACE	1	1	1
STEAP2 AND NOT-CAV1 AND GOLM1	1	1	1
STEAP2 AND NOT-SLC2A5 AND NOT-VAMP1	1	1	1
STEAP2 AND NOT-SLC2A5 AND NOT-TENM2	1	1	1
STEAP2 AND NOT-ROR2 AND LDLRAD3	1	1	1
STEAP2 AND NOT-ABHD6 AND LDLRAD3	1	1	1
STEAP2 AND NOT-CAV1 AND SLC6A14	1	1	1
COMPLEX-KDR/PCDHGB6/OR51E2	1	1	1
COMPLEX-MAGEA4/SGCD/OR51E2	1	1	1
STEAP2 AND NOT-MYADM AND SLC6A14	1	1	1
STEAP2 AND NOT-SGCE AND LDLRAD3	1	1	1
STEAP2 AND NOT-GPR161 AND SLC6A14	1	1	1
STEAP2 AND NOT-ABHD6 AND LDLRAD3	1	1	1
STEAP2 AND NOT-CAV1 AND GOLM1	1	1	1
COMPLEX-KCNK3/KDR/OR51E2	1	1	1
COMPLEX-DKK1/SGCD/OR51E2	1	1	1
OR51E2 AND NOT-DKK1 AND NOT-SLC2A5	1	1	1
STEAP2 AND NOT-CAV1 AND PROS1	1	1	1
STEAP2 AND NOT-PCDH7 AND SLC6A14	1	1	1
STEAP2 AND NOT-SLC18A2 AND SLC6A14	1	1	1
STEAP2 AND NOT-GPR161 AND LDLRAD3	1	1	1
STEAP2 AND NOT-MYADM AND LDLRAD3	1	1	1
COMPLEX-KDR/IYD/OR51E2	1	1	1
STEAP2 AND NOT-SLC16A14 AND NOT-ACE	1	1	1
STEAP2 AND NOT-SLC2A5 AND NOT-STX7	1	1	1
STEAP2 AND NOT-CAV1 AND VMP1	1	1	1
OR51E2 AND NOT-ACE AND NOT-AXL	1	1	1
STEAP2 AND NOT-PANX1 AND LDLRAD3	1	1	1

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
GPM6A AND ABCA1 AND NOT-RAET1E	0.963504	0.970588	0.956522
PTPRZ1 AND SLC40A1 AND NCAM1	0.977778	1	0.956522
GPM6A AND PLGRKT AND NOT-FOLR1	0.970149	1	0.942029
PTPRZ1 AND NOT-CD3G AND NOT-CLDN23	0.977778	1	0.956522
PTPRZ1 AND NOT-KCNB2 AND NOT-TRPM4	0.970588	0.985075	0.956522
PTPRZ1 AND NOT-CD3G AND NCAM1	0.977778	1	0.956522
PTPRZ1 AND NOT-KCNB2 AND NOT-CLDN23	0.985507	0.985507	0.985507
COMPLEX-CLDN7/ADORA3/PTPRZ1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-LRP11 AND NOT-CEACAM5	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-LRP11 AND NOT-CLDN18	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-CD3G AND NOT-TRPM4	0.962406	1	0.927536
PTPRZ1 AND NOT-KCNB2 AND NCAM1	0.985507	0.985507	0.985507
COMPLEX-TNFSF13B/CLDN7/PTPRZ1	0.978102	0.985294	0.971014
PTPRZ1 AND CBX3 AND NOT-WNT3	0.977778	1	0.956522
PTPRZ1 AND CBX3 AND NOT-BTC	0.977778	1	0.956522
PTPRZ1 AND NOT-EPCAM AND NOT-ESYT3	0.985507	0.985507	0.985507
PTPRZ1 AND CBX3 AND NOT-RECK	0.977778	1	0.956522
PTPRZ1 AND NOT-EPCAM AND SLC39A14	0.992701	1	0.985507
PTPRZ1 AND CBX3 AND NOT-SLC15A1	0.977778	1	0.956522
PTPRZ1 AND CBX3 AND NOT-ACVR2A	0.977778	1	0.956522
PTPRZ1 AND CBX3 AND SLC6A1	0.977778	1	0.956522
GPM6A AND SLC40A1 AND NOT-CLDN1	0.970588	0.985075	0.956522
GPM6A AND SLC40A1 AND NOT-ERBB3	0.962963	0.984848	0.942029
PTPRZ1 AND NOT-LRP11 AND NOT-RAET1E	0.985294	1	0.971014
COMPLEX-PTPRZ1/RYR1/BCAN	0.970588	0.985075	0.956522
COMPLEX-PTPRZ1/BCAN/SLC22A4	0.962406	1	0.927536
GPM6A AND SLC40A1 AND NOT-RAET1E	0.970588	0.985075	0.956522
GPM6A AND SLC40A1 AND NOT-SLAMF7	0.962963	0.984848	0.942029
COMPLEX-ITGB8/PTPRZ1/BCAN	0.977778	1	0.956522
COMPLEX-PTPRZ1/BCAN/NRP2	0.962406	1	0.927536
GPM6A AND PLGRKT AND NOT-RAET1E	0.970149	1	0.942029
PTPRZ1 AND JAM2 AND NOT-GPA33	0.985294	1	0.971014
COMPLEX-TNFSF13B/PTPRZ1/BCAN	0.992701	1	0.985507
PTPRZ1 AND JAM2 AND NCAM1	0.985294	1	0.971014
PTPRZ1 AND JAM2 AND NOT-SDC1	0.985294	1	0.971014
PTPRZ1 AND SLC40A1 AND NOT-SDC1	0.977778	1	0.956522
COMPLEX-PTPRZ1/BCAN/SIGLEC10	0.992701	1	0.985507
PTPRZ1 AND JAM2 AND NOT-MUC1	0.985294	1	0.971014
PTPRZ1 AND NOT-CD3G AND NOT-MUC1	0.977778	1	0.956522
PTPRZ1 AND NOT-SYT2 AND NOT-MUC1	0.985507	0.985507	0.985507
PTPRZ1 AND TSPAN6 AND NOT-MUC1	0.962406	1	0.927536
PTPRZ1 AND NOT-KCNB2 AND NOT-MUC1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-HAS1 AND NOT-MUC1	0.955224	0.984615	0.927536
PTPRZ1 AND NOT-CD5 AND NOT-MUC1	0.985294	1	0.971014
PTPRZ1 AND NOT-S1PR5 AND NOT-MUC1	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-ACVR1B AND NOT-MUC1	0.978102	0.985294	0.971014
PTPRZ1 AND BFAR AND NOT-MUC1	0.992701	1	0.985507
PTPRZ1 AND XPR1 AND NOT-MUC1	0.970149	1	0.942029
COMPLEX-FAT2/MUC1/PTPRZ1	0.985507	0.985507	0.985507
PTPRZ1 AND HLA-G AND NOT-ERBB2	0.985294	1	0.971014
COMPLEX-C1orf210/MUC1/PTPRZ1	0.992701	1	0.985507
PTPRZ1 AND NOT-FLT3 AND NOT-MUC1	0.985294	1	0.971014
PTPRZ1 AND NOT-SYT1 AND NOT-MUC1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-TNFSF9 AND NOT-MUC1	0.985294	1	0.971014
PTPRZ1 AND NOT-RHBDL2 AND NOT-MUC1	0.970149	1	0.942029
PTPRZ1 AND NOT-ESYT3 AND NOT-EPHA2	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-ESYT3 AND NOT-MUC1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-ESYT3 AND NOT-MUC1	0.985507	0.985507	0.985507
COMPLEX-FUT1/MUC1/PTPRZ1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-SLC18A2 AND NOT-MUC1	0.985294	1	0.971014
PTPRZ1 AND NOT-PRPH2 AND NOT-MUC1	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-OR7A10 AND NOT-MUC1	0.955224	0.984615	0.927536
PTPRZ1 AND NOT-SIGLEC6 AND NOT-MUC1	0.992701	1	0.985507
COMPLEX-DSC3/MUC1/PTPRZ1	0.985294	1	0.971014
PTPRZ1 AND NOT-SYT8 AND NOT-EPHA2	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-KCNG3 AND NOT-MUC1	0.977778	1	0.956522
PTPRZ1 AND NOT-CD40LG AND NOT-MUC1	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-ESYT3 AND NOT-ERBB2	0.978102	0.985294	0.971014
COMPLEX-MUC1/PTPRZ1/GPR68	0.985507	0.985507	0.985507
PTPRZ1 AND TMBIM6 AND NOT-ERBB2	0.978102	0.985294	0.971014
COMPLEX-MUC1/PTGER3/PTPRZ1	0.985294	1	0.971014
PTPRZ1 AND NOT-UPK1A AND NOT-MUC1	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-CXCR3 AND NOT-MUC1	0.970149	1	0.942029
PTPRZ1 AND NOT-LY6K AND NOT-MUC1	0.970588	0.985075	0.956522
PTPRZ1 AND NOT-BPI AND NOT-MUC1	0.992701	1	0.985507
PTPRZ1 AND TMBIM6 AND NOT-MUC1	0.985507	0.985507	0.985507
COMPLEX-MUC1/SLC46A2/PTPRZ1	0.978102	0.985294	0.971014

Antigen Pair Logic	F1	Precision	Recall
COMPLEX-CD180/SGCD/OR51E2	1	1	1
STEAP2 AND NOT-SLC2A5 AND NOT-VAMP1	1	1	1
STEAP2 AND NOT-SLC18A2 AND SLC6A14	1	1	1
STEAP2 AND NOT-ROR2 AND LDLRAD3	1	1	1
STEAP2 AND NOT-CAV1 AND IFNGR2	1	1	1
STEAP2 AND NOT-MYADM AND SLC6A14	1	1	1
COMPLEX-CLEC14A/FNDC4/OR51E2	1	1	1
OR51E2 AND NOT-ACE AND NOT-ITGB6	1	1	1
OR51E2 AND NOT-SLC2A5 AND TNFRSF13C	1	1	1
OR51E2 AND NOT-SLC2A5 AND BIRC5	1	1	1
STEAP2 AND NOT-GGT7 AND SLC6A14	1	1	1
STEAP2 AND NOT-CAV1 AND PANX1	1	1	1
STEAP2 AND NOT-SLC16A2 AND SLC6A14	1	1	1
STEAP2 AND NOT-CAV1 AND NOT-ICAM4	1	1	1
STEAP2 AND LDLRAD3 AND NOT-ITM2C	1	1	1
STEAP2 AND NOT-SLC2A5 AND NOT-SLC16A2	1	1	1
COMPLEX-STEAP2/FURIN/FNDC4	1	1	1
STEAP2 AND NOT-PLP2 AND LDLRAD3	1	1	1
STEAP2 AND NOT-FADS2 AND GOLM1	1	1	1
STEAP2 AND NOT-FZD1 AND LILRB2	1	1	1
STEAP2 AND NOT-SLC2A5 AND NOT-CNTN1	1	1	1
STEAP2 AND NOT-GPR161 AND GOLM1	1	1	1
STEAP2 AND NOT-MYADM AND GOLM1	1	1	1
STEAP2 AND NPFFR2 AND NOT-UNC5B	1	1	1
OR51E2 AND NOT-ACE AND NOT-CLDN11	1	1	1
STEAP2 AND NOT-ROR2 AND NOT-UNC5B	1	1	1
STEAP2 AND NOT-SLC18A2 AND NOT-UNC5B	1	1	1
STEAP2 AND NOT-MAL AND NOT-SLC52A3	1	1	1
STEAP2 AND NOT-STX2 AND LDLRAD3	1	1	1
STEAP2 AND NOT-SLC25A4 AND LDLRAD3	1	1	1
OR51E2 AND NOT-ACE AND NOT-SLC34A2	1	1	1
OR51E2 AND NOT-ACE AND CD70	1	1	1
STEAP2 AND NOT-ABCC6 AND LDLRAD3	1	1	1
STEAP2 AND NOT-SLC18A2 AND GOLM1	1	1	1
STEAP2 AND NOT-NPY1R AND LDLRAD3	1	1	1
OR51E2 AND NOT-ACE AND NOT-MST1R	1	1	1
TRPM8 AND NOT-CAV1 AND VCAM1	1	1	1
STEAP2 AND NOT-CAV1 AND STEAP4	1	1	1
STEAP2 AND SLCO1A2 AND LDLRAD3	1	1	1
TRPM8 AND NOT-TNC AND NOT-UNC5B	1	1	1
STEAP2 AND SLCO1A2 AND LDLRAD3	1	1	1
TRPM8 AND NOT-CAV1 AND CLDN8	1	1	1
STEAP2 AND NOT-CAV1 AND PTGER4	1	1	1
STEAP2 AND SLCO1A2 AND NOT-SLC2A5	1	1	1
STEAP2 AND NOT-CAV1 AND CD163	1	1	1
STEAP2 AND NOT-SYT4 AND NOT-SLC2A5	1	1	1
STEAP2 AND NOT-SYT4 AND NOT-SLC2A5	1	1	1
STEAP2 AND NOT-ROR2 AND F2R	1	1	1
COMPLEX-SLC39A6/SLC22A18/OR51E2	1	1	1
STEAP2 AND NOT-PCDH7 AND PTGER4	1	1	1
STEAP2 AND NOT-GPR161 AND PTGER4	1	1	1
STEAP2 AND NOT-ATP1A2 AND F2R	1	1	1
TRPM8 AND NOT-PCDH7 AND MUC1	1	1	1
TRPM8 AND NOT-SGCE AND CLDN8	1	1	1
TRPM8 AND NOT-FZD1 AND ABCA5	1	1	1
STEAP2 AND SLCO1A2 AND NOT-UNC5B	1	1	1
STEAP2 AND NOT-SLC2A5 AND NOT-DYSF	1	1	1
TRPM8 AND NOT-FZD1 AND RNF43	1	1	1
STEAP2 AND NOT-DNER AND NOT-SLC39A2	1	1	1
TRPM8 AND NOT-SLC52A3 AND CLDN8	1	1	1
TRPM8 AND NOT-ATP1A2 AND VCAM1	1	1	1
TRPM8 AND NOT-FADS2 AND NOT-LGR5	1	1	1
STEAP2 AND NOT-SLC2A5 AND NOT-PAQR8	1	1	1
TRPM8 AND NOT-CSPG4 AND NOT-UNC5B	1	1	1
STEAP2 AND SLCO1A2 AND GOLM1	1	1	1
STEAP2 AND NOT-FZD1 AND THBD	1	1	1
TRPM8 AND NOT-SLC2A5 AND NOT-EGFR	1	1	1
TRPM8 AND NOT-LSAMP AND CLDN8	1	1	1
STEAP2 AND NOT-SLC2A5 AND NOT-SYT11	1	1	1
STEAP2 AND NOT-SLC2A5 AND NOT-DYSF	1	1	1
STEAP2 AND NOT-PCDH7 AND F2R	1	1	1
STEAP2 AND NOT-DYSF AND LDLRAD3	1	1	1
STEAP2 AND NPFFR2 AND NOT-SLC39A2	1	1	1
STEAP2 AND NOT-SLC2A5 AND NOT-GYPC	1	1	1
STEAP2 AND NOT-PLP2 AND NOT-SLC39A2	1	1	1
TRPM8 AND NOT-SLC2A5 AND NOT-DKK1	1	1	1
STEAP2 AND SLCO1A2 AND NOT-CLSTN2	1	1	1

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
COMPLEX-MUC1/PTPRZ1/SLC28A3	0.985507	0.985507	0.985507
COMPLEX-EDAR/MUC1/PTPRZ1	0.962963	0.984848	0.942029
PTPRZ1 AND HLA-C AND NOT-MUC1	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-SYT5 AND NOT-MUC1	0.985507	0.985507	0.985507
PTPRZ1 AND IFNGR2 AND NOT-ERBB2	0.985294	1	0.971014
PTPRZ1 AND NOT-CD8B AND NOT-MUC1	0.992701	1	0.985507
PTPRZ1 AND NOT-OR1D2 AND NOT-MUC1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-F10 AND NOT-MUC1	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-GPR143 AND NOT-MUC1	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-SYT8 AND ERBB2	0.985507	0.985507	0.985507
COMPLEX-MUC1/PTPRZ1/SCNN1A	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-GJB5 AND ERBB2	0.985507	0.985507	0.985507
PTPRZ1 AND TNFRSF21 AND NOT-MUC1	0.977778	1	0.956522
COMPLEX-MUC1/ATP2A3/PTPRZ1	0.970588	0.985075	0.956522
COMPLEX-MUC1/PTPRZ1/PVRL4	0.992701	1	0.985507
COMPLEX-GJB5/MUC1/PTPRZ1	0.985507	0.985507	0.985507
COMPLEX-MUC1/PTPRZ1/SYT8	0.985507	0.985507	0.985507
COMPLEX-MUC1/PTPRZ1/PVRL4	0.992701	1	0.985507
COMPLEX-GJB5/MUC1/PTPRZ1	0.985507	0.985507	0.985507
COMPLEX-EPHA2/HLA-G/PTPRZ1	0.992701	1	0.985507
COMPLEX-MUC1/PTPRZ1/SYT8	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-DIO3 AND NOT-MUC1	0.978102	0.985294	0.971014
COMPLEX-COL17A1/MUC1/PTPRZ1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-OR1J2 AND NOT-MUC1	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-ADRA2B AND NOT-MUC1	0.985507	0.985507	0.985507
COMPLEX-MUC1/PTPRZ1/ITM2C	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-MRGPRX1 AND NOT-ERBB2	0.978102	0.985294	0.971014
COMPLEX-MUC1/PON2/PTPRZ1	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-RHCG AND NOT-MUC1	0.978102	0.985294	0.971014
COMPLEX-MUC15/MUC1/PTPRZ1	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-ISLR2 AND NOT-ERBB2	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-CLCNKB AND NOT-MUC1	0.970588	0.985075	0.956522
PTPRZ1 AND NOT-ADAM11 AND NOT-MUC1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-GPR87 AND ERBB2	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-LY6G6D AND NOT-MUC1	0.978102	0.985294	0.971014
COMPLEX-MUC1/PTPRZ1/SLC1A3	0.970588	0.985075	0.956522
PTPRZ1 AND NOT-SLC6A4 AND NOT-MUC1	0.985294	1	0.971014
COMPLEX-MUC1/PTPRZ1/CD1E	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-P2RY2 AND NOT-MUC1	0.962963	0.984848	0.942029
PTPRZ1 AND NOT-STAB2 AND NOT-MUC1	0.970588	0.985075	0.956522
PTPRZ1 AND SRR AND NOT-MUC1	0.977778	1	0.956522
PTPRZ1 AND NOT-KLB AND NOT-MUC1	0.970588	0.985075	0.956522
COMPLEX-CFTR/MUC1/PTPRZ1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-SCN2B AND NOT-MUC1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-ISLR2 AND NOT-MUC1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-MRGPRX1 AND NOT-MUC1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-OR4D1 AND NOT-MUC1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-MRGPRX1 AND NOT-MUC1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-SLC6A5 AND NOT-MUC1	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-SLC6A3 AND NOT-MUC1	0.978102	0.985294	0.971014
PTPRZ1 AND VAMP3 AND NOT-MUC1	0.985507	0.985507	0.985507
COMPLEX-MARVELD2/MUC1/PTPRZ1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-BDKRB1 AND NOT-MUC1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-SLC1A7 AND NOT-MUC1	0.970588	0.985075	0.956522
PTPRZ1 AND NOT-TRPV6 AND NOT-MUC1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-CRHR1 AND NOT-MUC1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-OR4D1 AND NOT-ERBB2	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-LTA AND NOT-ERBB2	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-IL1RL2 AND NOT-MUC1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-OR1F1 AND NOT-ERBB2	0.985294	1	0.971014
GPM6A AND F2R AND NOT-FOLR1	0.992701	1	0.985507
GPM6A AND F2R AND NOT-RAET1E	0.985294	1	0.971014
GPM6A AND F2R AND NOT-MUC16	0.970149	1	0.942029
PTPRZ1 AND CBX3 AND ASTN1	0.977778	1	0.956522
PTPRZ1 AND NOT-SLC6A15 AND NOT-CLDN23	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-SLC6A15 AND NOT-TRPM4	0.970588	0.985075	0.956522
COMPLEX-PTPRZ1/BCAN/SLC1A6	0.970149	1	0.942029
PTPRZ1 AND NOT-SLC6A15 AND NCAM1	0.978102	0.985294	0.971014
PTPRZ1 AND F2R AND NOT-SDC1	0.985294	1	0.971014
PTPRZ1 AND F2R AND NOT-TRPM4	0.985294	1	0.971014
PTPRZ1 AND F2R AND NCAM1	0.985294	1	0.971014
PTPRZ1 AND NOT-EPCAM AND NOT-EPHA10	0.985507	0.985507	0.985507
PTPRZ1 AND CBX3 AND GPR158	0.977778	1	0.956522
PTPRZ1 AND CBX3 AND NOT-ATP8B1	0.977778	1	0.956522
PTPRZ1 AND NOT-SLC6A15 AND NOT-SDC1	0.978102	0.985294	0.971014
PTPRZ1 AND F2R AND NOT-CLDN23	0.985294	1	0.971014
PTPRZ1 AND NOT-ABCG4 AND NOT-TRPM4	0.962963	0.984848	0.942029

Antigen Pair Logic	F1	Precision	Recall
STEAP2 AND NOT-PLP2 AND NOT-SLC39A2	1	1	1
TRPM8 AND NOT-ATP1A2 AND MUC1	1	1	1
TRPM8 AND NOT-SLC18A2 AND CLDN8	1	1	1
STEAP2 AND NOT-MYADM AND PTGER4	1	1	1
TRPM8 AND NOT-SLC2A5 AND BIRC5	1	1	1
TRPM8 AND NOT-ROR2 AND MUC1	1	1	1
TRPM8 AND NOT-ATP1A2 AND MUC1	1	1	1
TRPM8 AND NOT-SLC2A5 AND P2RX5	1	1	1
TRPM8 AND NOT-SLC2A5 AND NOT-IL13RA2	1	1	1
TRPM8 AND NOT-SLC2A5 AND NOT-IL11RA	1	1	1
TRPM8 AND NOT-PANX1 AND CLDN8	1	1	1
TRPM8 AND NOT-PCDH7 AND MUC1	1	1	1
STEAP2 AND NOT-SLC2A5 AND NOT-SLC4A8	1	1	1
TRPM8 AND NOT-FADS2 AND CLDN8	1	1	1
STEAP2 AND NOT-POPDC2 AND NOT-SLC39A2	1	1	1
TRPM8 AND NOT-TNC AND SLC6A14	1	1	1
TRPM8 AND NOT-POPDC2 AND VCAM1	1	1	1
STEAP2 AND NOT-POPDC2 AND NOT-SLC39A2	1	1	1
TRPM8 AND NOT-TNC AND NOT-CLSTN2	1	1	1
TRPM8 AND NOT-TNC AND NOT-CLSTN2	1	1	1
TRPM8 AND NOT-PCDH7 AND CLDN8	1	1	1
STEAP2 AND NOT-POPDC2 AND PTGER4	1	1	1
STEAP2 AND SLCO1A2 AND KCNS3	1	1	1
TRPM8 AND NOT-GPR161 AND VCAM1	1	1	1
TRPM8 AND NOT-LSAMP AND BIRC5	1	1	1
TRPM8 AND NOT-PCDH7 AND VCAM1	1	1	1
TRPM8 AND NOT-GPR161 AND CLDN8	1	1	1
STEAP2 AND SLCO1A2 AND SLC6A14	1	1	1
STEAP2 AND SLCO1A2 AND GOLM1	1	1	1
TRPM8 AND NOT-SLC2A5 AND NOT-LGR5	1	1	1
STEAP2 AND SLCO1A2 AND NOT-CLSTN2	1	1	1
STEAP2 AND NOT-GPR161 AND F2R	1	1	1
STEAP2 AND NOT-ATP1A2 AND PTGER4	1	1	1
STEAP2 AND NOT-SLC2A5 AND SLC2A2	1	1	1
STEAP2 AND NOT-SLC18A2 AND F2R	1	1	1
STEAP2 AND NOT-SYT4 AND NOT-LSAMP	1	1	1
TRPM8 AND NOT-ROR2 AND VCAM1	1	1	1
TRPM8 AND NOT-POPDC2 AND CLDN8	1	1	1
TRPM8 AND NOT-FADS2 AND NOT-LGR5	1	1	1
STEAP2 AND NOT-SLC2A5 AND NOT-EMP3	1	1	1
TRPM8 AND NOT-SLC2A5 AND NOT-CSPG4	1	1	1
TRPM8 AND NOT-ROR2 AND CLDN8	1	1	1
TRPM8 AND NOT-LSAMP AND BIRC5	1	1	1
TRPM8 AND NOT-SLC2A5 AND NOT-ITGB6	1	1	1
TRPM8 AND NOT-SLC2A5 AND NOT-LGR5	1	1	1
STEAP2 AND NOT-ROR2 AND PTGER4	1	1	1
STEAP2 AND NOT-DYSF AND LDLRAD3	1	1	1
TRPM8 AND NOT-SLC2A5 AND NOT-IL11RA	1	1	1
STEAP2 AND SLCO1A2 AND NOT-UNC5B	1	1	1
TRPM8 AND NOT-ABHD6 AND CLDN8	1	1	1
STEAP2 AND NOT-POPDC2 AND CD163	1	1	1
STEAP2 AND NOT-SLC2A5 AND NOT-PAQR8	1	1	1
STEAP2 AND NOT-POPDC2 AND STEAP4	1	1	1
TRPM8 AND NOT-FZD1 AND CLDN8	1	1	1
TRPM8 AND NOT-GPR161 AND MUC1	1	1	1
STEAP2 AND NOT-SLC2A5 AND NOT-GYPC	1	1	1
TRPM8 AND NOT-SLC2A5 AND NOT-CSPG4	1	1	1
STEAP2 AND NOT-SLC2A5 AND NOT-TTYH2	1	1	1
STEAP2 AND NOT-CAV1 AND ABCA5	1	1	1
STEAP2 AND NOT-CAV1 AND CLDN8	1	1	1
STEAP2 AND NOT-CAV1 AND VCAM1	1	1	1
COMPLEX-KDR/CLDN5/OR51E2	1	1	1
STEAP2 AND NOT-UNC5B AND NOT-TNC	1	1	1
STEAP2 AND NOT-SLC18A2 AND CLDN8	1	1	1
STEAP2 AND NOT-FADS2 AND CLDN8	1	1	1
STEAP2 AND NOT-ROR2 AND VCAM1	1	1	1
STEAP2 AND NOT-FZD1 AND ABCA5	1	1	1
STEAP2 AND NOT-POPDC2 AND VCAM1	1	1	1
STEAP2 AND BIRC5 AND NOT-LSAMP	1	1	1
STEAP2 AND BIRC5 AND NOT-KCNJ15	1	1	1
STEAP2 AND NOT-MYADM AND MUC1	1	1	1
STEAP2 AND NOT-POPDC2 AND VCAM1	1	1	1
STEAP2 AND NOT-ROR2 AND CLDN8	1	1	1
STEAP2 AND NOT-POPDC2 AND ABCA5	1	1	1
STEAP2 AND NOT-ATP1A2 AND VCAM1	1	1	1
STEAP2 AND NOT-GPR161 AND MUC1	1	1	1
STEAP2 AND NOT-GPR161 AND MUC1	1	1	1

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
COMPLEX-TNFRSF12A/PTPRZ1/BCAN	0.962406	1	0.927536
PTPRZ1 AND CBX3 AND NOT-SLC13A2	0.977778	1	0.956522
PTPRZ1 AND CBX3 AND NOT-KCNK6	0.977778	1	0.956522
PTPRZ1 AND CBX3 AND SLC1A2	0.977778	1	0.956522
PTPRZ1 AND F2R AND NOT-SSTR3	0.985294	1	0.971014
PTPRZ1 AND NOT-NKAIN2 AND NOT-TRPM4	0.970588	0.985075	0.956522
PTPRZ1 AND NOT-NKAIN2 AND NOT-CLDN23	0.985507	0.985507	0.985507
PTPRZ1 AND F2R AND NOT-SSTR5	0.985294	1	0.971014
PTPRZ1 AND F2R AND NOT-GPA33	0.985294	1	0.971014
PTPRZ1 AND NOT-NKAIN2 AND NCAM1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-ABCG4 AND NOT-CLDN23	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-ABCG4 AND NCAM1	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-EPCAM AND NOT-LRRC52	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-EPCAM AND NOT-DIO1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-EPCAM AND NOT-SLC17A1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-EPCAM AND NOT-SLC17A3	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-EPCAM AND NOT-ABCG5	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-EPCAM AND NOT-GRIN2B	0.992701	1	0.985507
PTPRZ1 AND CBX3 AND NOT-SCARA5	0.977778	1	0.956522
PTPRZ1 AND NOT-EPCAM AND NOT-ADAM30	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-HCN1 AND NOT-TRPM4	0.970588	0.985075	0.956522
PTPRZ1 AND NOT-HCN1 AND NOT-CLDN23	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-NKAIN2 AND NOT-SDC1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-HCN1 AND NCAM1	0.978102	0.985294	0.971014
PTPRZ1 AND F2R AND NOT-SSTR4	0.985294	1	0.971014
PTPRZ1 AND NOT-ATP8A2 AND NOT-TRPM4	0.970588	0.985075	0.956522
PTPRZ1 AND NOT-SCN8A AND NOT-TRPM4	0.962963	0.984848	0.942029
PTPRZ1 AND NOT-ATP8A2 AND NOT-CLDN23	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-SCN8A AND NOT-CLDN23	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-SCN8A AND NCAM1	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-ABCG4 AND NOT-SDC1	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-ATP8A2 AND NCAM1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-KIAA0319 AND NOT-SDC1	0.985507	0.985507	0.985507
PTPRZ1 AND CBX3 AND NOT-SLC43A1	0.977778	1	0.956522
PTPRZ1 AND CBX3 AND NOT-GUCY2D	0.977778	1	0.956522
PTPRZ1 AND CBX3 AND NOT-THBD	0.977778	1	0.956522
PTPRZ1 AND NOT-HCN1 AND NOT-SDC1	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-ATP8A2 AND NOT-SDC1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-SCN8A AND NOT-SDC1	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-CDH8 AND NOT-CLDN23	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-CDH8 AND NOT-TRPM4	0.970588	0.985075	0.956522
COMPLEX-PTPRZ1/BCAN/TNFRSF10B	0.962406	1	0.927536
PTPRZ1 AND NOT-CACNA1B AND NOT-CLDN23	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-CACNA1B AND NOT-TRPM4	0.970588	0.985075	0.956522
PTPRZ1 AND NOT-CDH8 AND NCAM1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-CACNA1B AND NCAM1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-KIAA0319 AND NOT-CLDN23	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-KIAA0319 AND NOT-TRPM4	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-KIAA0319 AND NCAM1	0.985507	0.985507	0.985507
PTPRZ1 AND CBX3 AND NOT-MRGPRF	0.977778	1	0.956522
PTPRZ1 AND CBX3 AND SLC22A14	0.977778	1	0.956522
PTPRZ1 AND NOT-HTR5A AND NCAM1	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-HTR5A AND NOT-TRPM4	0.970588	0.985075	0.956522
PTPRZ1 AND CBX3 AND GABRB1	0.977778	1	0.956522
PTPRZ1 AND NOT-HTR5A AND NOT-CLDN23	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-SLC32A1 AND NOT-CLDN23	0.970588	0.985075	0.956522
PTPRZ1 AND NOT-GRM3 AND NOT-CLDN23	0.992701	1	0.985507
PTPRZ1 AND NOT-GRM3 AND NOT-TRPM4	0.977778	1	0.956522
PTPRZ1 AND NOT-SLC32A1 AND NOT-TRPM4	0.955224	0.984615	0.927536
PTPRZ1 AND NOT-DGKE AND NCAM1	0.985507	0.985507	0.985507
PTPRZ1 AND CBX3 AND CRB1	0.962406	1	0.927536
PTPRZ1 AND NOT-CDH8 AND NOT-SDC1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-GABRA2 AND NOT-CLDN23	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-GRM3 AND NCAM1	0.992701	1	0.985507
PTPRZ1 AND NOT-DGKE AND NOT-TRPM4	0.970588	0.985075	0.956522
PTPRZ1 AND NOT-SLC32A1 AND NCAM1	0.970588	0.985075	0.956522
PTPRZ1 AND NOT-CACNA1B AND NOT-SDC1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-GABRA2 AND NOT-TRPM4	0.970588	0.985075	0.956522
PTPRZ1 AND NOT-GABRA2 AND NCAM1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-CLDN6 AND ASTN1	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-GRM1 AND NOT-CLDN23	0.970588	0.985075	0.956522
PTPRZ1 AND NOT-GRM1 AND NOT-TRPM4	0.955224	0.984615	0.927536
PTPRZ1 AND NOT-SLC26A8 AND NOT-CLDN23	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-SLC26A8 AND NOT-TRPM4	0.970588	0.985075	0.956522
PTPRZ1 AND NOT-SLC26A8 AND NCAM1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-GRM1 AND NCAM1	0.970588	0.985075	0.956522
PTPRZ1 AND CBX3 AND SLC8A3	0.977778	1	0.956522

Antigen Pair Logic	F1	Precision	Recall
STEAP2 AND BIRC5 AND NOT-SLC2A5	1	1	1
STEAP2 AND BIRC5 AND NOT-EFNB1	1	1	1
STEAP2 AND NOT-ROR2 AND MUC1	1	1	1
STEAP2 AND NOT-PCDH7 AND CLDN8	1	1	1
STEAP2 AND NOT-MYADM AND MUC1	1	1	1
STEAP2 AND NOT-PCDH7 AND CLDN8	1	1	1
STEAP2 AND NOT-PANX1 AND CLDN8	1	1	1
STEAP2 AND NOT-ATP1A2 AND MUC1	1	1	1
STEAP2 AND NOT-ROR2 AND CLDN8	1	1	1
STEAP2 AND NOT-ROR2 AND MUC1	1	1	1
STEAP2 AND NOT-FZD1 AND RNF43	1	1	1
STEAP2 AND NOT-GPR161 AND VCAM1	1	1	1
STEAP2 AND NOT-ATP1A2 AND VCAM1	1	1	1
STEAP2 AND NOT-ABHD6 AND CLDN8	1	1	1
STEAP2 AND NOT-ABHD6 AND CLDN8	1	1	1
STEAP2 AND NOT-PCDH7 AND MUC1	1	1	1
STEAP2 AND NOT-POPDC2 AND CLDN8	1	1	1
STEAP2 AND NOT-MYADM AND VCAM1	1	1	1
STEAP2 AND NOT-TNC AND SLC6A14	1	1	1
STEAP2 AND NOT-MYADM AND CLDN8	1	1	1
STEAP2 AND NOT-MYADM AND VCAM1	1	1	1
STEAP2 AND NOT-PCDH7 AND MUC1	1	1	1
STEAP2 AND BIRC5 AND NOT-SLC2A5	1	1	1
STEAP2 AND NOT-FADS2 AND CLDN8	1	1	1
STEAP2 AND NOT-SLC52A3 AND CLDN8	1	1	1
STEAP2 AND NOT-TNC AND SLC6A14	1	1	1
STEAP2 AND NOT-PCDH7 AND VCAM1	1	1	1
STEAP2 AND NOT-SLC18A2 AND CLDN8	1	1	1
STEAP2 AND NOT-PCDH7 AND VCAM1	1	1	1
STEAP2 AND NOT-SLC52A3 AND CLDN8	1	1	1
STEAP2 AND NOT-ATP1A2 AND MUC1	1	1	1
STEAP2 AND NOT-ROR2 AND VCAM1	1	1	1
STEAP2 AND NOT-GPR161 AND CLDN8	1	1	1
STEAP2 AND NOT-POPDC2 AND CLDN8	1	1	1
STEAP2 AND NOT-FZD1 AND CLDN8	1	1	1
STEAP2 AND NOT-MYADM AND CLDN8	1	1	1
STEAP2 AND NOT-SGCE AND CLDN8	1	1	1
STEAP2 AND NOT-SGCE AND CLDN8	1	1	1
STEAP2 AND NOT-SGCD AND EPHA3	1	1	1
STEAP2 AND NOT-MYADM AND RNF43	1	1	1
STEAP2 AND NOT-GPR161 AND RNF43	1	1	1
STEAP2 AND BIRC5 AND NOT-DUOXA1	1	1	1
STEAP2 AND BIRC5 AND NOT-PTPRU	1	1	1
STEAP2 AND BIRC5 AND NOT-SGCD	1	1	1
STEAP2 AND NOT-MYADM AND ABCA5	1	1	1
STEAP2 AND NOT-MYADM AND TRPM4	1	1	1
STEAP2 AND BIRC5 AND NOT-CD40	1	1	1
STEAP2 AND CLDN8 AND NOT-SLC3A2	1	1	1
STEAP2 AND NOT-ROR2 AND RNF43	1	1	1
STEAP2 AND NOT-FADS2 AND TRPM4	1	1	1
STEAP2 AND NOT-SGCD AND VCAM1	1	1	1
STEAP2 AND NOT-CAV1 AND TRPM4	1	1	1
STEAP2 AND NOT-PCDH7 AND RNF43	1	1	1
STEAP2 AND NOT-ROR2 AND TRPM4	1	1	1
STEAP2 AND NOT-GPR161 AND ABCA5	1	1	1
STEAP2 AND NOT-SLC52A3 AND TRPM4	1	1	1
STEAP2 AND NOT-SGCD AND MUC1	1	1	1
STEAP2 AND NOT-SGCE AND TRPM4	1	1	1
STEAP2 AND NOT-FADS2 AND NOT-NCAM1	1	1	1
STEAP2 AND NOT-ROR2 AND ABCA5	1	1	1
STEAP2 AND NOT-GPR161 AND TRPM4	1	1	1
STEAP2 AND NOT-PCDH7 AND ABCA5	1	1	1
STEAP2 AND NOT-PCDH7 AND TRPM4	1	1	1
STEAP2 AND NOT-SLC16A2 AND MUC1	1	1	1
STEAP2 AND NOT-SGCE AND MUC1	1	1	1
STEAP2 AND NOT-STX4 AND CLDN8	1	1	1
STEAP2 AND NOT-SLC2A5 AND NOT-L1CAM	1	1	1
COMPLEX-TMC1/STEAP2/EPCAM	1	1	1
STEAP2 AND CLDN8 AND NOT-SLC20A2	1	1	1
STEAP2 AND NOT-SLC24A3 AND VCAM1	1	1	1
STEAP2 AND NOT-PLP2 AND MUC1	1	1	1
STEAP2 AND NOT-STX2 AND VCAM1	1	1	1
STEAP2 AND NOT-MCAM AND VCAM1	1	1	1
STEAP2 AND NOT-ADCY5 AND CLDN8	1	1	1
STEAP2 AND NOT-SLC25A4 AND VCAM1	1	1	1
STEAP2 AND NOT-ADCY5 AND MUC1	1	1	1
STEAP2 AND NOT-UNC5B AND NOT-CSPG4	1	1	1

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
GPM6A AND ITGAV AND PTGER4	0.956522	0.956522	0.956522
PTPRZ1 AND NOT-HTR5A AND NOT-SDC1	0.978102	0.985294	0.971014
PTPRZ1 AND CBX3 AND NKAIN4	0.970149	1	0.942029
COMPLEX-PTPRZ1/BCAN/CD58	0.985294	1	0.971014
GPM6A AND ITGAV AND BEST2	0.956522	0.956522	0.956522
GPM6A AND ITGAV AND STAB1	0.94964	0.942857	0.956522
PTPRZ1 AND CBX3 AND NOT-CLDN23	0.977778	1	0.956522
PTPRZ1 AND CBX3 AND NOT-TRPM4	0.962406	1	0.927536
PTPRZ1 AND CBX3 AND NCAM1	0.977778	1	0.956522
PTPRZ1 AND CBX3 AND NOT-SDC1	0.977778	1	0.956522
PTPRZ1 AND NOT-EPCAM AND NOT-GPA33	0.985507	0.985507	0.985507
COMPLEX-TRPM4/RNF43/PTPRZ1	0.970588	0.985075	0.956522
COMPLEX-CLDN7/PTPRZ1/CD37	0.985507	0.985507	0.985507
PTPRZ1 AND CBX3 AND NOT-FOLR1	0.977778	1	0.956522
PTPRZ1 AND NOT-CLDN6 AND NCAM1	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-CLDN6 AND NOT-TRPM4	0.962963	0.984848	0.942029
PTPRZ1 AND NOT-CLDN6 AND NOT-CLDN23	0.978102	0.985294	0.971014
COMPLEX-EGFR/PTPRZ1/CLDN8	0.985507	0.985507	0.985507
GPM6A AND ITGAV AND MUC16	0.956522	0.956522	0.956522
PTPRZ1 AND NOT-CLDN6 AND NOT-SDC1	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-GPA33 AND NOT-CLDN1	0.978102	0.985294	0.971014
COMPLEX-MET/PTPRZ1/CLDN8	0.978102	0.985294	0.971014
COMPLEX-TRPM4/PTPRZ1/CLDN8	0.985507	0.985507	0.985507
GPM6A AND ITGAV AND FOLR1	0.94964	0.942857	0.956522
PTPRZ1 AND NOT-CLDN8 AND NOT-GPA33	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-RAET1E AND NOT-GPA33	0.985294	1	0.971014
PTPRZ1 AND NOT-CEACAM6 AND NOT-GPA33	0.985294	1	0.971014
PTPRZ1 AND NOT-GPA33 AND NOT-CLDN23	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-GPA33 AND NOT-TRPM4	0.970588	0.985075	0.956522
PTPRZ1 AND NOT-GPA33 AND NOT-CLDN23	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-GPA33 AND NOT-TRPM4	0.970588	0.985075	0.956522
PTPRZ1 AND NOT-GPA33 AND NCAM1	0.985507	0.985507	0.985507
COMPLEX-CSPG4/TRPM4/PTPRZ1	0.962406	1	0.927536
COMPLEX-FOLR1/PTPRZ1/CLDN8	0.978102	0.985294	0.971014
COMPLEX-GPNMB/PTPRZ1/BCAN	0.985294	1	0.971014
GPM6A AND ITGAV AND RAET1E	0.94964	0.942857	0.956522
PTPRZ1 AND ITGAV AND NOT-SDC1	0.985294	1	0.971014
COMPLEX-IL20RA/PTPRZ1/CLDN12	0.962963	0.984848	0.942029
COMPLEX-ITGAV/MET/PTPRZ1	0.985294	1	0.971014
PTPRZ1 AND NOT-CEACAM6 AND NOT-SSTR5	0.985294	1	0.971014
PTPRZ1 AND NOT-CEACAM6 AND NOT-SSTR3	0.985294	1	0.971014
COMPLEX-PTPRZ1/TPBG/CLDN8	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-GPA33 AND NOT-SDC1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-GPA33 AND NOT-SDC1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-MST1R AND NOT-GPA33	0.985507	0.985507	0.985507
COMPLEX-CBX3/PTPRZ1/TPBG	0.977778	1	0.956522
PTPRZ1 AND NOT-ITGB6 AND NOT-GPA33	0.985294	1	0.971014
PTPRZ1 AND NOT-CLDN4 AND NOT-GPA33	0.985294	1	0.971014
COMPLEX-ITGAV/PTPRZ1/BMPR1B	0.992701	1	0.985507
GPM6A AND ITGAV AND CLDN1	0.94964	0.942857	0.956522
GPM6A AND ITGAV AND SLAMF7	0.94964	0.942857	0.956522
PTPRZ1 AND NOT-TYR AND NOT-GPA33	0.977778	1	0.956522
PTPRZ1 AND ITGAV AND NOT-TRPM4	0.985294	1	0.971014
PTPRZ1 AND ITGAV AND NOT-CLDN23	0.985294	1	0.971014
PTPRZ1 AND ITGAV AND NCAM1	0.985294	1	0.971014
PTPRZ1 AND NOT-CLDN1 AND NOT-SSTR3	0.978102	0.985294	0.971014
COMPLEX-IL20RA/TRPM4/PTPRZ1	0.970588	0.985075	0.956522
PTPRZ1 AND NOT-GPA33 AND NOT-VTCN1	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-SLAMF7 AND NOT-GPA33	0.977778	1	0.956522
COMPLEX-PTPRZ1/BCAN/BMPR1B	0.938462	1	0.884058
COMPLEX-TRPM4/PTPRZ1/BCAN	0.938462	1	0.884058
PTPRZ1 AND NOT-CLDN4 AND SDC1	0.962963	0.984848	0.942029
COMPLEX-CLDN11/PTPRZ1/CD37	0.970588	0.985075	0.956522
COMPLEX-CLDN7/TRPM4/PTPRZ1	0.970588	0.985075	0.956522
COMPLEX-HSPA5/TRPM4/PTPRZ1	0.977778	1	0.956522
PTPRZ1 AND NOT-GPA33 AND NOT-STEAP2	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-GPA33 AND CD276	0.992701	1	0.985507
PTPRZ1 AND NOT-CXCR5 AND NOT-CLDN23	0.962406	1	0.927536
PTPRZ1 AND NOT-CXCR5 AND NCAM1	0.962406	1	0.927536
PTPRZ1 AND NCAM1 AND NOT-SSTR3	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-CLDN23 AND NOT-SSTR3	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-CLDN1 AND NOT-SSTR5	0.978102	0.985294	0.971014
COMPLEX-ROR1/PTPRZ1/PMEL	0.978102	0.985294	0.971014
PTPRZ1 AND NCAM1 AND NOT-SSTR3	0.985507	0.985507	0.985507
COMPLEX-CEACAM6/TRPM4/PTPRZ1	0.992701	1	0.985507
PTPRZ1 AND NOT-CLDN23 AND NOT-CLDN18	0.978102	0.985294	0.971014
COMPLEX-ABCB5/PTPRZ1/BCAN	0.992701	1	0.985507

Antigen Pair Logic	F1	Precision	Recall
STEAP2 AND NOT-ABCC6 AND VCAM1	1	1	1
COMPLEX-PRRT2/STEAP2/VCAM1	1	1	1
STEAP2 AND NOT-SLC2A5 AND P2RX5	1	1	1
STEAP2 AND CLDN8 AND NOT-DNER	1	1	1
FOLH1 AND HPN AND NOT-ABHD6	1	1	1
FOLH1 AND HPN AND NOT-ABHD6	1	1	1
FOLH1 AND FZD4 AND NOT-NPY1R	1	1	1
FOLH1 AND HPN AND NOT-NPY1R	1	1	1
FOLH1 AND NOT-ABCC6 AND FZD4	1	1	1
FOLH1 AND FZD4 AND NOT-NPY1R	1	1	1
FOLH1 AND NOT-ABCC6 AND HPN	1	1	1
FOLH1 AND HPN AND NOT-NPY1R	1	1	1
FOLH1 AND NOT-ABCC6 AND FZD4	1	1	1
FOLH1 AND HPN AND NOT-SLC38A4	1	1	1
FOLH1 AND HPN AND NOT-SLC38A4	1	1	1
FOLH1 AND BCAM AND NOT-LSAMP	1	1	1
FOLH1 AND NOT-ABCC6 AND FZD5	1	1	1
FOLH1 AND ADRB2 AND NOT-SLC52A3	1	1	1
FOLH1 AND ADRB2 AND NOT-CAV1	1	1	1
FOLH1 AND GOLM1 AND NOT-VIMP	1	1	1
FOLH1 AND NOT-DNER AND LDLRAD3	1	1	1
FOLH1 AND ADRB2 AND NOT-CAV1	1	1	1
FOLH1 AND ADRB2 AND NOT-ABHD6	1	1	1
FOLH1 AND HPN AND NOT-ABCG2	1	1	1
FOLH1 AND ADRB2 AND NOT-LSAMP	1	1	1
FOLH1 AND NOT-DNER AND LDLRAD3	1	1	1
FOLH1 AND NOT-ABHD6 AND FZD5	1	1	1
FOLH1 AND ADRB2 AND NOT-SLC52A3	1	1	1
FOLH1 AND NOT-ABCC6 AND ICAM3	1	1	1
FOLH1 AND NOT-ABHD6 AND IFNGR2	1	1	1
FOLH1 AND HPN AND NOT-ABCG2	1	1	1
FOLH1 AND ADRB2 AND NOT-ABHD6	1	1	1
FOLH1 AND NOT-ABCC6 AND FZD5	1	1	1
FOLH1 AND GOLM1 AND NOT-VIMP	1	1	1
FOLH1 AND NOT-ABCC6 AND SLC6A14	1	1	1
FOLH1 AND NOT-ABHD6 AND FZD5	1	1	1
FOLH1 AND NOT-ABCC6 AND ICAM3	1	1	1
FOLH1 AND GOLM1 AND NOT-ABHD6	1	1	1
ANO7 AND NOT-LSAMP AND FOLH1	1	1	1
FOLH1 AND CD320 AND NOT-LSAMP	1	1	1
FOLH1 AND ITGA5 AND NOT-CAV1	1	1	1
FOLH1 AND POPDC2 AND NOT-CAV1	1	1	1
FOLH1 AND BCAM AND NOT-SGCD	1	1	1
FOLH1 AND IL1RAPL1 AND NOT-PLP1	1	1	1
FOLH1 AND HPN AND NOT-SLC47A1	1	1	1
FOLH1 AND HPN AND NOT-SLC47A1	1	1	1
FOLH1 AND ADRB2 AND NOT-NPY1R	1	1	1
FOLH1 AND ADRB2 AND NOT-ROR2	1	1	1
FOLH1 AND GOLM1 AND NOT-ABCC6	1	1	1
FOLH1 AND TMPRSS2 AND NOT-SLC2A5	1	1	1
FOLH1 AND ADRB2 AND NOT-ROR2	1	1	1
FOLH1 AND ADRB2 AND NOT-SLC18A2	1	1	1
COMPLEX-FOLH1/DAGLA/OR51E2	1	1	1
FOLH1 AND GOLM1 AND NOT-ABCC6	1	1	1
FOLH1 AND ADRB2 AND NOT-NPY1R	1	1	1
FOLH1 AND FZD4 AND NOT-SGCE	1	1	1
FOLH1 AND AGTRAP AND NOT-SLC2A5	1	1	1
FOLH1 AND SLC39A10 AND NOT-LSAMP	1	1	1
FOLH1 AND ADRB2 AND NOT-ABCC6	1	1	1
FOLH1 AND NOT-ABCC6 AND SLC19A2	1	1	1
FOLH1 AND NOT-ABHD6 AND PANX1	1	1	1
FOLH1 AND TMPRSS2 AND NOT-SLC2A5	1	1	1
FOLH1 AND BCAM AND NOT-SLC2A5	1	1	1
FOLH1 AND GOLM1 AND NOT-SEMA6D	1	1	1
FOLH1 AND NOT-ABCC6 AND SLC30A5	1	1	1
FOLH1 AND NOT-ABCC6 AND IFNGR2	1	1	1
FOLH1 AND ADRB2 AND NOT-SLC18A2	1	1	1
FOLH1 AND ADRB2 AND NOT-SGCE	1	1	1
FOLH1 AND ADRB2 AND NOT-ATP1A2	1	1	1
FOLH1 AND ADRB2 AND NOT-SLC16A2	1	1	1
FOLH1 AND HPN AND NOT-TGFBR3	1	1	1
FOLH1 AND ADRB2 AND NOT-PCDH7	1	1	1
FOLH1 AND ADRB2 AND NOT-PCDH7	1	1	1
FOLH1 AND GOLM1 AND NOT-ABCC6	1	1	1
FOLH1 AND ADRB2 AND NOT-ABCC6	1	1	1
FOLH1 AND GOLM1 AND NOT-SEMA6D	1	1	1
FOLH1 AND FZD4 AND NOT-DNER	1	1	1

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
PTPRZ1 AND NOT-CEACAM5 AND NOT-GPA33	0.985507	0.985507	0.985507
PTPRZ1 AND NCAM1 AND NOT-CLDN18	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-GPA33 AND SLC39A6	0.970149	1	0.942029
PTPRZ1 AND NOT-GPA33 AND NOT-TNFRSF10A	0.978102	0.985294	0.971014
PTPRZ1 AND SLC39A6 AND NCAM1	0.970149	1	0.942029
COMPLEX-GPNMB/CLDN7/PTPRZ1	0.985507	0.985507	0.985507
COMPLEX-EDNRB/TRPM4/PTPRZ1	0.938462	1	0.884058
PTPRZ1 AND NOT-TYR AND NOT-SSTR5	0.985294	1	0.971014
COMPLEX-PTPRZ1/BCAN/SSTR3	0.977778	1	0.956522
PTPRZ1 AND NOT-TYR AND NOT-SSTR3	0.985294	1	0.971014
PTPRZ1 AND NOT-TYR AND NOT-FOLR1	0.985294	1	0.971014
PTPRZ1 AND NOT-HHLA2 AND NOT-SDC1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-STEAP2 AND NOT-SSTR5	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-STEAP2 AND NOT-SSTR3	0.985507	0.985507	0.985507
COMPLEX-IL13RA1/MET/PTPRZ1	0.977778	1	0.956522
COMPLEX-GPNMB/PTPRZ1/CLDN8	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-CXCR5 AND NOT-SDC1	0.962406	1	0.927536
PTPRZ1 AND NOT-CEACAM6 AND NOT-SSTR4	0.985294	1	0.971014
GPM6A AND CBX3 AND NOT-FOLR1	0.962406	1	0.927536
PTPRZ1 AND NOT-CLDN23 AND NOT-DNAJB8	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-SSTR3 AND NOT-SDC1	0.985507	0.985507	0.985507
PTPRZ1 AND NCAM1 AND NOT-SSTR5	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-CLDN23 AND NOT-SSTR5	0.985507	0.985507	0.985507
COMPLEX-FOLR1/PTPRZ1/PMEL	0.970588	0.985075	0.956522
PTPRZ1 AND NCAM1 AND NOT-ULBP3	0.985507	0.985507	0.985507
PTPRZ1 AND NCAM1 AND NOT-DNAJB8	0.985507	0.985507	0.985507
COMPLEX-TRPM4/SLAMF7/PTPRZ1	0.985294	1	0.971014
COMPLEX-TRPM4/PTPRZ1/PMEL	0.978102	0.985294	0.971014
GPM6A AND F2R AND NOT-ZP4	0.992701	1	0.985507
GPM6A AND F2R AND NOT-LRRN4	0.992701	1	0.985507
GPM6A AND F2R AND NOT-OR3A2	0.992701	1	0.985507
GPM6A AND F2R AND NOT-TRPC3	0.985294	1	0.971014
GPM6A AND F2R AND NOT-BEST2	0.992701	1	0.985507
GPM6A AND F2R AND NOT-FGF6	0.992701	1	0.985507
GPM6A AND F2R AND NOT-ABCG5	0.992701	1	0.985507
GPM6A AND F2R AND NOT-GPR78	0.985294	1	0.971014
GPM6A AND F2R AND NOT-MUC17	0.992701	1	0.985507
GPM6A AND F2R AND NOT-GJB4	0.992701	1	0.985507
GPM6A AND F2R AND NOT-SLC22A11	0.992701	1	0.985507
GPM6A AND F2R AND NOT-SLC26A8	0.992701	1	0.985507
GPM6A AND F2R AND NOT-CHRND	0.992701	1	0.985507
GPM6A AND F2R AND NOT-OR8B2	0.992701	1	0.985507
GPM6A AND F2R AND NOT-SLC9B1	0.992701	1	0.985507
GPM6A AND F2R AND NOT-GABRG3	0.992701	1	0.985507
GPM6A AND F2R AND NOT-HTR6	0.985294	1	0.971014
GPM6A AND F2R AND NOT-ATP8B2	0.992701	1	0.985507
GPM6A AND F2R AND NOT-STAB1	0.992701	1	0.985507
GPM6A AND F2R AND NOT-PTGER4	0.992701	1	0.985507
GPM6A AND F2R AND NOT-KCNU1	0.992701	1	0.985507
GPM6A AND F2R AND NOT-KCNA4	0.992701	1	0.985507
GPM6A AND F2R AND NOT-ADAM21	0.977778	1	0.956522
PTPRZ1 AND NOT-SLC6A15 AND ASTN1	0.978102	0.985294	0.971014
GPM6A AND F2R AND NOT-KCNJ9	0.992701	1	0.985507
PTPRZ1 AND NOT-SLC6A15 AND GPR158	0.978102	0.985294	0.971014
PTPRZ1 AND F2R AND GPR158	0.985294	1	0.971014
PTPRZ1 AND F2R AND NOT-SLC43A1	0.985294	1	0.971014
PTPRZ1 AND F2R AND NOT-KCNK6	0.985294	1	0.971014
PTPRZ1 AND F2R AND NOT-SCARA5	0.985294	1	0.971014
PTPRZ1 AND F2R AND ASTN1	0.985294	1	0.971014
PTPRZ1 AND NOT-SLC6A15 AND NOT-KCNK6	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-SLC6A15 AND NOT-ATP8B1	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-SLC6A15 AND SLC1A2	0.978102	0.985294	0.971014
PTPRZ1 AND F2R AND NOT-SLC34A1	0.985294	1	0.971014
PTPRZ1 AND F2R AND NOT-SLC13A2	0.985294	1	0.971014
PTPRZ1 AND F2R AND NOT-ATP8B1	0.985294	1	0.971014
PTPRZ1 AND F2R AND NOT-THBD	0.985294	1	0.971014
PTPRZ1 AND F2R AND NOT-GUCY2D	0.985294	1	0.971014
PTPRZ1 AND NOT-NKAIN2 AND ASTN1	0.985507	0.985507	0.985507
PTPRZ1 AND F2R AND SLC1A2	0.985294	1	0.971014
PTPRZ1 AND NOT-ABCG4 AND ASTN1	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-HCN1 AND ASTN1	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-SLC6A15 AND NOT-SCARA5	0.978102	0.985294	0.971014
PTPRZ1 AND F2R AND NOT-MRGPRF	0.985294	1	0.971014
PTPRZ1 AND F2R AND CD74	0.985294	1	0.971014
PTPRZ1 AND F2R AND NOT-BEST2	0.985294	1	0.971014
PTPRZ1 AND F2R AND NOT-LRFN2	0.985294	1	0.971014
PTPRZ1 AND NOT-NKAIN2 AND GPR158	0.985507	0.985507	0.985507

Antigen Pair Logic	F1	Precision	Recall
FOLH1 AND GOLM1 AND NOT-ITGA6	1	1	1
FOLH1 AND SLC39A10 AND NOT-LSAMP	1	1	1
FOLH1 AND AGTRAP AND NOT-LSAMP	1	1	1
FOLH1 AND HPN AND NOT-TGFBR3	1	1	1
FOLH1 AND ADRB2 AND NOT-SGCE	1	1	1
FOLH1 AND NOT-ABHD6 AND VMP1	1	1	1
FOLH1 AND FZD4 AND NOT-SGCE	1	1	1
FOLH1 AND ADRB2 AND NOT-GPR161	1	1	1
FOLH1 AND ADRB2 AND NOT-PANX1	1	1	1
FOLH1 AND FZD4 AND NOT-DNER	1	1	1
FOLH1 AND NOT-ABHD6 AND PANX1	1	1	1
FOLH1 AND ADRB2 AND NOT-PANX1	1	1	1
FOLH1 AND NOT-ABHD6 AND VMP1	1	1	1
FOLH1 AND NOT-ABHD6 AND DHRS3	1	1	1
FOLH1 AND AGTRAP AND NOT-LSAMP	1	1	1
FOLH1 AND NOT-ABCC6 AND IFNGR2	1	1	1
FOLH1 AND NOT-ABHD6 AND DHRS3	1	1	1
FOLH1 AND AGTRAP AND NOT-SLC2A5	1	1	1
FOLH1 AND ADRB2 AND NOT-SLC2A5	1	1	1
FOLH1 AND ADRB2 AND NOT-POPDC2	1	1	1
FOLH1 AND STYK1 AND NOT-SLC52A3	0.888889	0.8	1
COMPLEX-FOLH1/OR51E2/CD69	0.888889	0.8	1
FOLH1 AND NOT-ABHD6 AND CD69	0.888889	0.8	1
FOLH1 AND NOT-ABHD6 AND ATP11B	0.888889	0.8	1
COMPLEX-CLCA4/FOLH1/OR51E2	0.888889	0.8	1
TRPM8 AND NOT-POPDC2 AND NOT-SLC39A2	1	1	1
TRPM8 AND NOT-SLC2A5 AND NOT-DYSF	1	1	1
TRPM8 AND NOT-CAV1 AND STEAP4	1	1	1
TRPM8 AND NOT-SLC2A5 AND SLCO1A2	1	1	1
OR51E2 AND NOT-FMNL1 AND SLCO1A2	1	1	1
TRPM8 AND NOT-CAV1 AND PTGER4	1	1	1
TRPM8 AND NOT-SLC39A2 AND NOT-PLP2	1	1	1
OR51E2 AND NOT-SLC39A2 AND SLCO1A2	1	1	1
TRPM8 AND NOT-SLC39A2 AND NOT-MYADM	1	1	1
TRPM8 AND NOT-SLC2A5 AND NOT-DYSF	1	1	1
OR51E2 AND NOT-FMNL1 AND GPR19	1	1	1
OR51E2 AND NOT-FMNL1 AND GPR19	1	1	1
OR51E2 AND NOT-SLC39A2 AND GPR19	1	1	1
TRPM8 AND NOT-SLC2A5 AND NOT-SYT4	1	1	1
TRPM8 AND NOT-SLC39A2 AND NOT-CD81	1	1	1
OR51E2 AND NOT-FMNL1 AND SLCO1A2	1	1	1
TRPM8 AND NOT-CAV1 AND CD163	1	1	1
COMPLEX-BFAR/TRPM8/SLC16A5	1	1	1
TRPM8 AND NOT-SLC2A5 AND NOT-GYPC	1	1	1
TRPM8 AND NOT-SLC2A5 AND NOT-EMP3	1	1	1
OR51E2 AND NOT-SLC39A2 AND GPR19	1	1	1
TRPM8 AND NOT-SLC2A5 AND NOT-SLC16A5	1	1	1
TRPM8 AND NOT-SLC2A5 AND NOT-SYT4	1	1	1
TRPM8 AND NOT-SLC39A2 AND NOT-MYADM	1	1	1
OR51E2 AND NOT-SLC39A2 AND SLCO1A2	1	1	1
TRPM8 AND NOT-SLC2A5 AND NOT-GYPC	1	1	1
TRPM8 AND NOT-SLC2A5 AND SLCO1A2	1	1	1
TRPM8 AND NOT-SLC39A2 AND NOT-ITM2C	1	1	1
TRPM8 AND NOT-SLC39A2 AND NPFFR2	1	1	1
TRPM8 AND NOT-SLC2A5 AND NOT-IL17RA	1	1	1
TRPM8 AND NOT-LSAMP AND NOT-SLC39A2	1	1	1
TRPM8 AND NOT-ATP1A2 AND PTGER4	1	1	1
COMPLEX-PAM/TRPM8/SLC16A5	1	1	1
TRPM8 AND NOT-HAS3 AND NOT-SLC39A2	1	1	1
TRPM8 AND NOT-LSAMP AND F2R	1	1	1
TRPM8 AND NOT-SLC2A5 AND NOT-EPHA2	1	1	1
TRPM8 AND NOT-LSAMP AND NOT-SLC4A8	1	1	1
TRPM8 AND NOT-LSAMP AND ATP8B1	1	1	1
TRPM8 AND NOT-DYSF AND LDLRAD3	1	1	1
TRPM8 AND NOT-UNC5B AND SLCO1A2	1	1	1
OR51E2 AND NOT-CHRNA2 AND NOT-SLC39A2	1	1	1
TRPM8 AND NOT-POPDC2 AND NOT-FMNL1	1	1	1
OR51E2 AND NOT-CHRNA2 AND NOT-SLC39A2	1	1	1
TRPM8 AND NOT-PCDH7 AND PTGER4	1	1	1
TRPM8 AND NOT-LSAMP AND NOT-SYT4	1	1	1
TRPM8 AND NOT-PRIMA1 AND NOT-SLC39A2	1	1	1
TRPM8 AND NOT-CLSTN2 AND SLCO1A2	1	1	1
TRPM8 AND NOT-POPDC2 AND NOT-FMNL1	1	1	1
TRPM8 AND NOT-SLC2A5 AND NOT-SLC4A8	1	1	1
OR51E2 AND NOT-CHRNA2 AND NOT-FMNL1	1	1	1
TRPM8 AND NOT-SLC18A2 AND F2R	1	1	1
TRPM8 AND NOT-SLC2A5 AND NOT-SLC39A2	1	1	1

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
PTPRZ1 AND F2R AND NOT-ZP4	0.985294	1	0.971014
PTPRZ1 AND F2R AND NOT-LRRN4	0.985294	1	0.971014
PTPRZ1 AND F2R AND NOT-OR3A2	0.985294	1	0.971014
PTPRZ1 AND F2R AND NOT-GRIN2B	0.985294	1	0.971014
PTPRZ1 AND F2R AND NOT-GPR135	0.985294	1	0.971014
PTPRZ1 AND F2R AND NOT-SLC17A3	0.985294	1	0.971014
PTPRZ1 AND F2R AND ATP13A5	0.985294	1	0.971014
PTPRZ1 AND F2R AND NOT-RXFP3	0.985294	1	0.971014
PTPRZ1 AND F2R AND CRB1	0.985294	1	0.971014
PTPRZ1 AND NOT-SCN8A AND ASTN1	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-ATP8A2 AND ASTN1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-ABCG4 AND GPR158	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-SLC6A15 AND NOT-THBD	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-SLC6A15 AND NOT-SLC43A1	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-HCN1 AND GPR158	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-NKAIN2 AND NOT-ATP8B1	0.985507	0.985507	0.985507
PTPRZ1 AND F2R AND NOT-EPHA10	0.985294	1	0.971014
PTPRZ1 AND NOT-NKAIN2 AND NOT-KCNK6	0.985507	0.985507	0.985507
PTPRZ1 AND F2R AND NOT-CACNG7	0.985294	1	0.971014
PTPRZ1 AND F2R AND NOT-ABCG5	0.985294	1	0.971014
PTPRZ1 AND NOT-ATP8A2 AND GPR158	0.985507	0.985507	0.985507
PTPRZ1 AND F2R AND NOT-NMBR	0.985294	1	0.971014
PTPRZ1 AND NOT-SCN8A AND GPR158	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-NKAIN2 AND SLC1A2	0.985507	0.985507	0.985507
PTPRZ1 AND F2R AND NOT-SLC22A12	0.985294	1	0.971014
PTPRZ1 AND NOT-ABCG4 AND NOT-KCNK6	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-ABCG4 AND SLC1A2	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-ABCG4 AND NOT-ATP8B1	0.970588	0.985075	0.956522
PTPRZ1 AND NOT-CDH8 AND ASTN1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-CACNA1B AND ASTN1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-KIAA0319 AND ASTN1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-KIAA0319 AND NOT-KCNK6	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-SLC16A5 AND NOT-SLCO1A2	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-SLC16A5 AND ZP4	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-HTR5A AND ASTN1	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-SLC16A5 AND TNFRSF10B	0.985294	1	0.971014
PTPRZ1 AND NOT-HCN1 AND NOT-KCNK6	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-HCN1 AND NOT-ATP8B1	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-HCN1 AND SLC1A2	0.978102	0.985294	0.971014
PTPRZ1 AND F2R AND NOT-TRPC7	0.985294	1	0.971014
PTPRZ1 AND NOT-SCN8A AND NOT-ATP8B1	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-SLC6A15 AND NOT-MUC1	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-HCN1 AND NOT-MUC1	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-SCN8A AND NOT-MUC1	0.978102	0.985294	0.971014
COMPLEX-MUC1/PTPRZ1/SLC16A5	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-HTR5A AND NOT-MUC1	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-GRM3 AND NOT-MUC1	0.992701	1	0.985507
PTPRZ1 AND NOT-GABRA2 AND NOT-MUC1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-SLC26A8 AND NOT-MUC1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-KCNK9 AND NOT-MUC1	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-MTNR1B AND NOT-MUC1	0.977778	1	0.956522
PTPRZ1 AND NOT-EPHA10 AND NOT-ERBB2	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-KCNA7 AND NOT-MUC1	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-PVRL1 AND NOT-MUC1	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-EPHA10 AND NOT-MUC1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-EPHA10 AND NOT-MUC1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-GPR83 AND NOT-MUC1	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-HCN4 AND NOT-MUC1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-NPHS1 AND NOT-MUC1	0.962963	0.984848	0.942029
COMPLEX-FUT3/MUC1/PTPRZ1	0.985294	1	0.971014
PTPRZ1 AND NOT-SLC22A9 AND NOT-MUC1	0.977778	1	0.956522
COMPLEX-MUC1/PTPRZ1/CD1A	0.992701	1	0.985507
PTPRZ1 AND NOT-FGF6 AND NOT-MUC1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-GUCY2F AND NOT-MUC1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-HTR1E AND NOT-MUC1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-CACNA1S AND NOT-MUC1	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-TSPAN16 AND NOT-MUC1	0.985294	1	0.971014
COMPLEX-TMPRSS11E/MUC1/PTPRZ1	0.985507	0.985507	0.985507
COMPLEX-MUC1/PTPRZ1/SMPD2	0.970588	0.985075	0.956522
PTPRZ1 AND NOT-SLC13A2 AND NOT-ERBB2	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-SLC17A3 AND NOT-ERBB2	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-GPR156 AND NOT-MUC1	0.962963	0.984848	0.942029
PTPRZ1 AND NOT-CLSTN3 AND NOT-MUC1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-MUC17 AND NOT-MUC1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-RGSL1 AND NOT-MUC1	0.962963	0.984848	0.942029
PTPRZ1 AND NOT-KCNA10 AND NOT-MUC1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-MRGPRX2 AND NOT-ERBB2	0.978102	0.985294	0.971014

Antigen Pair Logic	F1	Precision	Recall
OR51E2 AND NOT-FMNL1 AND MEGF11	1	1	1
TRPM8 AND GOLM1 AND SLCO1A2	1	1	1
TRPM8 AND NOT-ROR2 AND F2R	1	1	1
OR51E2 AND NOT-FMNL1 AND MEGF11	1	1	1
TRPM8 AND NOT-SLC2A5 AND NOT-STAB1	1	1	1
TRPM8 AND NOT-SLC2A5 AND SLC2A2	1	1	1
OR51E2 AND NOT-CHRNA2 AND NOT-FMNL1	1	1	1
TRPM8 AND NOT-SLC39A2 AND NOT-ITM2C	1	1	1
TRPM8 AND NOT-SLC2A5 AND NOT-SLC5A2	1	1	1
TRPM8 AND NOT-SLC2A5 AND NOT-STAB1	1	1	1
TRPM8 AND NOT-SLC2A5 AND NOT-TTYH2	1	1	1
TRPM8 AND NOT-SLC2A5 AND NOT-SLC4A8	1	1	1
COMPLEX-RABAC1/TRPM8/SLC16A5	1	1	1
TRPM8 AND NOT-SLC2A5 AND NOT-LAT	1	1	1
TRPM8 AND NOT-SLC2A5 AND NOT-SLC39A2	1	1	1
TRPM8 AND NOT-SLC2A5 AND NOT-TTYH2	1	1	1
TRPM8 AND NOT-ROR2 AND PTGER4	1	1	1
COMPLEX-GRIN2C/SLC22A18/OR51E2	1	1	1
TRPM8 AND NOT-UNC5B AND GPR19	1	1	1
TRPM8 AND NOT-SLC39A2 AND NOT-DNER	1	1	1
TRPM8 AND NOT-POPDC2 AND PTGER4	1	1	1
COMPLEX-TRPM8/NRSN2/SLC16A5	1	1	1
TRPM8 AND NOT-FZD1 AND THBD	1	1	1
TRPM8 AND NOT-SLC2A5 AND NOT-PAQR8	1	1	1
TRPM8 AND NOT-SLC2A5 AND NOT-BST2	1	1	1
TRPM8 AND NOT-POPDC2 AND CD163	1	1	1
TRPM8 AND NOT-POPDC2 AND STEAP4	1	1	1
TRPM8 AND NOT-PRIMA1 AND NOT-SLC39A2	1	1	1
OR51E2 AND NOT-SLC39A2 AND MEGF11	1	1	1
TRPM8 AND NOT-SLC39A2 AND NOT-VIMP	1	1	1
TRPM8 AND NOT-SLC2A5 AND NOT-FXYD6	1	1	1
TRPM8 AND GOLM1 AND SLCO1A2	1	1	1
OR51E2 AND NOT-GUCY2D AND NOT-SLC39A2	1	1	1
TRPM8 AND NOT-SLC2A5 AND NOT-SYT11	1	1	1
TRPM8 AND GOLM1 AND SLCO1A2	1	1	1
TRPM8 AND NOT-SLC2A5 AND NOT-PAQR8	1	1	1
TRPM8 AND NOT-GPR161 AND PTGER4	1	1	1
TRPM8 AND NOT-LSAMP AND NOT-SLC4A8	1	1	1
OR51E2 AND NOT-GUCY2D AND NOT-SLC39A2	1	1	1
COMPLEX-CDH17/SLC22A18/OR51E2	1	1	1
OR51E2 AND NOT-GUCY2D AND NOT-FMNL1	1	1	1
TRPM8 AND NOT-ATP1A2 AND F2R	1	1	1
OR51E2 AND NOT-SLC39A2 AND NOT-EBP	1	1	1
TRPM8 AND NOT-UNC5B AND SLCO1A2	1	1	1
TRPM8 AND NOT-SLC2A5 AND NOT-GRIN2C	1	1	1
TRPM8 AND NOT-SLC39A2 AND KCNMB2	1	1	1
COMPLEX-ABHD3/SLC13A1/OR51E2	1	1	1
TRPM8 AND NOT-SLC2A5 AND NOT-PHLDB2	1	1	1
FOLH1 AND EPHA3 AND NOT-SGCD	1	1	1
FOLH1 AND CLDN8 AND NOT-DNER	1	1	1
FOLH1 AND NOT-ABCC6 AND CLDN8	1	1	1
FOLH1 AND CLDN8 AND NOT-DNER	1	1	1
FOLH1 AND NOT-ABCC6 AND CLDN8	1	1	1
FOLH1 AND EPHA3 AND NOT-LSAMP	1	1	1
FOLH1 AND EPHA3 AND NOT-FZD1	1	1	1
FOLH1 AND EPHA3 AND NOT-SEMA6D	1	1	1
FOLH1 AND EPHA3 AND NOT-SLC2A5	1	1	1
FOLH1 AND NOT-ABCC6 AND MUC1	1	1	1
FOLH1 AND NOT-ABCC6 AND MUC1	1	1	1
FOLH1 AND EPHA3 AND NOT-TENM2	1	1	1
FOLH1 AND EPHA3 AND NOT-SLC52A3	1	1	1
FOLH1 AND EPHA3 AND NOT-TENM2	1	1	1
FOLH1 AND EPHA3 AND NOT-SEMA6D	1	1	1
FOLH1 AND ADRB2 AND NOT-ST8SIA1	1	1	1
FOLH1 AND EPHA3 AND NOT-SLC52A3	1	1	1
FOLH1 AND EPHA3 AND NOT-CAV1	0.888889	0.8	1
COMPLEX-FOLH1/AXL/OR51E2	0.888889	0.8	1
STEAP2 AND NOT-FZD1 AND FOLH1	0.888889	0.8	1
COMPLEX-FOLH1/AXL/OR51E2	0.888889	0.8	1
FOLH1 AND EPHA3 AND NOT-ATP1A2	0.888889	0.8	1
FOLH1 AND EPHA3 AND NOT-GPR161	0.888889	0.8	1
FOLH1 AND EPHA3 AND NOT-ITGA7	0.888889	0.8	1
FOLH1 AND TNC AND NOT-SGCE	0.888889	0.8	1
FOLH1 AND EPHA3 AND NOT-NDRG4	0.888889	0.8	1
FOLH1 AND EPHA3 AND NOT-CAV1	0.888889	0.8	1
FOLH1 AND EPHA3 AND NOT-FADS2	0.888889	0.8	1
FOLH1 AND TNC AND NOT-CAV1	0.888889	0.8	1

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
PTPRZ1 AND NOT-MRAP AND NOT-MUC1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-GRIN2B AND NOT-ERBB2	0.985294	1	0.971014
PTPRZ1 AND NOT-SLCO1B1 AND NOT-MUC1	0.985294	1	0.971014
PTPRZ1 AND NOT-ABCG8 AND NOT-MUC1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-GRIN2B AND NOT-MUC1	0.992701	1	0.985507
PTPRZ1 AND NOT-MRGPRX2 AND NOT-MUC1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-GRIN2B AND NOT-MUC1	0.992701	1	0.985507
PTPRZ1 AND NOT-MRGPRX2 AND NOT-MUC1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-GPR22 AND NOT-MUC1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-KCNJ9 AND NOT-MUC1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-CCKBR AND NOT-MUC1	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-SLC10A2 AND NOT-MUC1	0.978102	0.985294	0.971014
COMPLEX-MUC1/CD207/PTPRZ1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-AMHR2 AND NOT-MUC1	0.970588	0.985075	0.956522
PTPRZ1 AND NOT-CASR AND NOT-ERBB2	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-ABCG5 AND NOT-ERBB2	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-LRFN2 AND NOT-ERBB2	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-SLC22A16 AND NOT-MUC1	0.978102	0.985294	0.971014
PTPRZ1 AND CD74 AND NOT-ERBB2	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-GPR135 AND NOT-ERBB2	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-PCDHAC2 AND NOT-MUC1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-CNTNAP2 AND NOT-MUC1	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-SLC22A8 AND NOT-MUC1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-HTR6 AND NOT-MUC1	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-GRM7 AND NOT-MUC1	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-GALR2 AND NOT-MUC1	0.978102	0.985294	0.971014
PTPRZ1 AND CD74 AND NOT-MUC1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-LRFN2 AND NOT-MUC1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-CASR AND NOT-MUC1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-ABCG5 AND NOT-MUC1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-DSC1 AND NOT-MUC1	0.970588	0.985075	0.956522
COMPLEX-EPHA2/PTPRZ1/SLC13A2	0.985507	0.985507	0.985507
COMPLEX-MUC1/PTPRZ1/SLC13A2	0.985507	0.985507	0.985507
PTPRZ1 AND CD74 AND NOT-MUC1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-LRFN2 AND NOT-MUC1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-CASR AND NOT-MUC1	0.985507	0.985507	0.985507
COMPLEX-MUC1/PTPRZ1/SLC13A2	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-GPR135 AND NOT-MUC1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-GPR135 AND NOT-MUC1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-ABCG5 AND NOT-MUC1	0.985507	0.985507	0.985507
COMPLEX-MUC1/PTPRZ1/SELP	0.970588	0.985075	0.956522
PTPRZ1 AND NOT-SLC10A1 AND NOT-MUC1	0.977778	1	0.956522
PTPRZ1 AND NOT-MPL AND NOT-MUC1	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-LRRN4 AND NOT-ERBB2	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-DIO1 AND NOT-ERBB2	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-NMBR AND NOT-ERBB2	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-RXFP3 AND NOT-ERBB2	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-SLC22A12 AND NOT-ERBB2	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-GUCY2D AND NOT-ERBB2	0.978102	0.985294	0.971014
COMPLEX-MUC1/PTPRZ1/GGTLC1	0.992701	1	0.985507
PTPRZ1 AND NOT-SLC17A1 AND NOT-ERBB2	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-SLC34A1 AND NOT-ERBB2	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-ADCY10 AND NOT-MUC1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-BEST2 AND NOT-ERBB2	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-SLC22A2 AND NOT-ERBB2	0.978102	0.985294	0.971014
PTPRZ1 AND ATP13A5 AND NOT-ERBB2	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-LAT AND NOT-MUC1	0.962963	0.984848	0.942029
PTPRZ1 AND NOT-WNT7A AND NOT-MUC1	0.977778	1	0.956522
PTPRZ1 AND NOT-EPHA8 AND NOT-MUC1	0.985507	0.985507	0.985507
COMPLEX-ABCA12/MUC1/PTPRZ1	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-UPK3A AND NOT-MUC1	0.985507	0.985507	0.985507
PTPRZ1 AND CBX3 AND NOT-MUC1	0.977778	1	0.956522
PTPRZ1 AND NOT-CLDN6 AND NOT-MUC1	0.978102	0.985294	0.971014
COMPLEX-MUC1/PTPRZ1/CLDN8	0.978102	0.985294	0.971014
COMPLEX-CLDN4/MUC1/PTPRZ1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-GPA33 AND NOT-ERBB2	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-GPA33 AND NOT-MUC1	0.985507	0.985507	0.985507
COMPLEX-MST1R/MUC1/PTPRZ1	0.985507	0.985507	0.985507
COMPLEX-MUC1/PTPRZ1/TYR	0.985294	1	0.971014
COMPLEX-MUC1/PTPRZ1/PMEL	0.970588	0.985075	0.956522
COMPLEX-ITGB6/MUC1/PTPRZ1	0.985507	0.985507	0.985507
COMPLEX-CEACAM5/MUC1/PTPRZ1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-SSTR3 AND NOT-ERBB2	0.978102	0.985294	0.971014
COMPLEX-MUC1/CEACAM6/PTPRZ1	0.985294	1	0.971014
PTPRZ1 AND NOT-CXCR5 AND NOT-MUC1	0.962406	1	0.927536
PTPRZ1 AND NOT-SSTR3 AND NOT-MUC1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-SSTR3 AND NOT-MUC1	0.985507	0.985507	0.985507

Antigen Pair Logic	F1	Precision	Recall
FOLH1 AND ROR1 AND NOT-CAV1	0.888889	0.8	1
FOLH1 AND EPHA3 AND NOT-SLC18A2	0.888889	0.8	1
FOLH1 AND EPHA3 AND NOT-SLC18A2	0.888889	0.8	1
FOLH1 AND NOT-ABHD6 AND ROR1	0.888889	0.8	1
STEAP2 AND NOT-FZD1 AND FOLH1	0.888889	0.8	1
FOLH1 AND NOT-ABCC6 AND ROR1	0.888889	0.8	1
FOLH1 AND EPHA3 AND NOT-FGFR1	0.888889	0.8	1
STEAP2 AND NOT-FADS2 AND FOLH1	0.888889	0.8	1
STEAP2 AND NOT-SLC52A3 AND FOLH1	0.888889	0.8	1
STEAP2 AND NOT-FADS2 AND FOLH1	0.888889	0.8	1
STEAP2 AND NOT-SLC52A3 AND FOLH1	0.888889	0.8	1
FOLH1 AND EPHA3 AND NOT-PTPRZ1	0.888889	0.8	1
FOLH1 AND EPHA3 AND NOT-PTPRZ1	0.888889	0.8	1
STEAP2 AND NOT-UNC5B AND FOLH1	1	1	1
STEAP2 AND NOT-UNC5B AND FOLH1	1	1	1
STEAP2 AND NOT-UNC5B AND FOLH1	1	1	1
FOLH1 AND EPHA3 AND NOT-ACE	1	1	1
FOLH1 AND ADRB2 AND NOT-TNC	1	1	1
FOLH1 AND CLDN8 AND NOT-TGFBR3	1	1	1
FOLH1 AND EPHA3 AND NOT-KCNN3	1	1	1
FOLH1 AND CLDN8 AND NOT-TGFBR3	1	1	1
FOLH1 AND EPHA3 AND NOT-SDC3	0.888889	0.8	1
COMPLEX-FOLH1/MST1R/OR51E2	0.888889	0.8	1
FOLH1 AND EPHA3 AND NOT-ACE	1	1	1
FOLH1 AND NOT-ABCC6 AND ABCA5	1	1	1
FOLH1 AND EPHA3 AND NOT-SDC3	0.888889	0.8	1
FOLH1 AND CLDN8 AND NOT-KCNJ15	0.857143	1	0.75
FOLH1 AND EPHA3 AND NOT-PTCHD1	0.857143	1	0.75
FOLH1 AND FZD4 AND NOT-ERBB4	0.857143	1	0.75
FOLH1 AND NOT-GPRC5B AND CLDN8	0.857143	1	0.75
FOLH1 AND EPCAM AND NOT-SLC7A9	0.857143	1	0.75
FOLH1 AND EPCAM AND NOT-MGAM	0.857143	1	0.75
FOLH1 AND CLDN8 AND NOT-HS3ST3B1	0.857143	1	0.75
FOLH1 AND NOT-MET AND ATP11B	0.888889	0.8	1
FOLH1 AND EPCAM AND NOT-SLC51B	0.857143	1	0.75
STEAP2 AND CACNA1D AND FOLH1	0.857143	1	0.75
FOLH1 AND EPCAM AND NOT-DSG2	0.857143	1	0.75
FOLH1 AND EPHA3 AND NOT-NAALADL1	0.857143	1	0.75
FOLH1 AND EPCAM AND SLC19A2	0.857143	1	0.75
FOLH1 AND EPCAM AND JAG2	0.857143	1	0.75
FOLH1 AND NOT-MET AND VMP1	0.888889	0.8	1
FOLH1 AND EPCAM AND NOT-CDHR5	0.857143	1	0.75
FOLH1 AND EPCAM AND SLC14A1	0.857143	1	0.75
FOLH1 AND EPCAM AND NOT-CD164	0.857143	1	0.75
FOLH1 AND EPHA3 AND NOT-PCDH9	0.857143	1	0.75
FOLH1 AND EPHA3 AND NOT-LGR6	0.857143	1	0.75
FOLH1 AND EPHA3 AND NOT-PTPRF	0.857143	1	0.75
FOLH1 AND EPCAM AND SLC24A3	0.857143	1	0.75
FOLH1 AND EPHA3 AND NOT-IL6R	0.857143	1	0.75
FOLH1 AND EPHA3 AND NOT-BMPR2	0.857143	1	0.75
FOLH1 AND ROR1 AND NOT-SLC52A3	1	1	1
FOLH1 AND TNC AND NOT-PRNP	0.857143	1	0.75
FOLH1 AND ADRB2 AND NOT-PCYT1A	0.857143	1	0.75
FOLH1 AND GOLM1 AND NOT-PCYT1A	0.857143	1	0.75
FOLH1 AND CACNA1D AND NOT-CLDN11	0.857143	1	0.75
FOLH1 AND NOT-ABCG2 AND EPCAM	0.857143	1	0.75
FOLH1 AND EPCAM AND NOT-VIMP	0.857143	1	0.75
FOLH1 AND EPCAM AND NOT-SI	0.857143	1	0.75
FOLH1 AND EPHA3 AND NOT-ANTXR1	0.857143	1	0.75
FOLH1 AND EPCAM AND NOT-SUSD2	0.857143	1	0.75
FOLH1 AND EPCAM AND NOT-ABCB1	0.857143	1	0.75
FOLH1 AND NOT-PAQR5 AND BIRC5	0.857143	1	0.75
FOLH1 AND EPCAM AND NOT-TLR3	0.857143	1	0.75
FOLH1 AND BMPR1B AND NOT-LSAMP	0.857143	1	0.75
FOLH1 AND EPCAM AND NOT-TRHDE	0.857143	1	0.75
FOLH1 AND FLOT2 AND NOT-SLC2A5	0.888889	0.8	1
FOLH1 AND EPCAM AND NOT-SLC1A1	0.857143	1	0.75
FOLH1 AND CACNA1D AND NOT-L1CAM	0.857143	1	0.75
FOLH1 AND EPCAM AND MCAM	0.857143	1	0.75
FOLH1 AND EPCAM AND NOT-TRPM6	0.857143	1	0.75
FOLH1 AND CDH10 AND NOT-LSAMP	1	1	1
FOLH1 AND CDH10 AND NOT-LSAMP	1	1	1
FOLH1 AND HPN AND NOT-CD36	1	1	1
FOLH1 AND NOT-CD36 AND FZD4	1	1	1
FOLH1 AND NOT-ABCC6 AND SLC43A1	1	1	1
FOLH1 AND NOT-ABHD6 AND SLC43A1	1	1	1
FOLH1 AND NOT-CD36 AND ICAM3	1	1	1

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
PTPRZ1 AND NOT-CLDN18 AND NOT-MUC1	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-SSTR5 AND NOT-ERBB2	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-IL3RA AND NOT-MUC1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-ULBP3 AND NOT-MUC1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-SSTR5 AND NOT-MUC1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-DNAJB8 AND NOT-MUC1	0.985507	0.985507	0.985507
PTPRZ1 AND SLC39A6 AND NOT-MUC1	0.970149	1	0.942029
PTPRZ1 AND NOT-MAGEA1 AND NOT-MUC1	0.955224	0.984615	0.927536
PTPRZ1 AND NOT-CLDN9 AND NOT-MUC1	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-HHLA2 AND NOT-MUC1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-GPA33 AND NOT-EPHA2	0.970588	0.985075	0.956522
COMPLEX-MUC1/SEMA5B/PTPRZ1	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-CD79B AND NOT-MUC1	0.970588	0.985075	0.956522
PTPRZ1 AND NOT-GUCY2C AND NOT-MUC1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-TNFRSF8 AND NOT-MUC1	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-STEAP2 AND NOT-MUC1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-FCRL5 AND NOT-MUC1	0.962963	0.984848	0.942029
PTPRZ1 AND CD276 AND NOT-MUC1	0.985294	1	0.971014
COMPLEX-MUC1/SLAMF7/PTPRZ1	0.985294	1	0.971014
GPM6A AND ERBB2 AND NOT-RAET1E	0.924138	0.881579	0.971014
PTPRZ1 AND NOT-FCRL2 AND NOT-MUC1	0.977778	1	0.956522
PTPRZ1 AND NOT-CLDN23 AND ERBB2	0.985507	0.985507	0.985507
PTPRZ1 AND NCAM1 AND ERBB2	0.985507	0.985507	0.985507
COMPLEX-MUC1/PTPRZ1/CLDN1	0.978102	0.985294	0.971014
COMPLEX-HLA-DOB/MUC1/PTPRZ1	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-CD22 AND NOT-MUC1	0.985294	1	0.971014
PTPRZ1 AND NOT-ERBB2 AND NOT-SSTR4	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-CD160 AND NOT-MUC1	0.970588	0.985075	0.956522
PTPRZ1 AND NOT-GAGE1 AND NOT-MUC1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-MUC1 AND NOT-SSTR4	0.985507	0.985507	0.985507
COMPLEX-CLDN23/MUC1/PTPRZ1	0.985507	0.985507	0.985507
COMPLEX-MUC1/NCAM1/PTPRZ1	0.985507	0.985507	0.985507
COMPLEX-CLDN23/MUC1/PTPRZ1	0.985507	0.985507	0.985507
COMPLEX-ERBB2/PTPRZ1/TPBG	0.970588	0.985075	0.956522
PTPRZ1 AND NOT-TNFSF11 AND NOT-MUC1	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-MUC1 AND NOT-SSTR4	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-MUC1 AND NOT-CD19	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-SSTR3 AND NOT-EPHA2	0.970588	0.985075	0.956522
PTPRZ1 AND NOT-P2RX5 AND NOT-MUC1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-CD79A AND NOT-MUC1	0.947368	0.984375	0.913043
PTPRZ1 AND NOT-CLDN2 AND NOT-MUC1	0.978102	0.985294	0.971014
COMPLEX-MUC1/PTPRZ1/TNFRSF10A	0.962963	0.984848	0.942029
PTPRZ1 AND NOT-PCYT1A AND NOT-MUC1	0.985507	0.985507	0.985507
COMPLEX-MUC1/ENPP3/PTPRZ1	0.985507	0.985507	0.985507
COMPLEX-MUC1/PTPRZ1/LGR5	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-MUC1 AND NOT-TNFRSF13C	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-SSTR5 AND NOT-EPHA2	0.970588	0.985075	0.956522
PTPRZ1 AND NOT-MS4A1 AND NOT-MUC1	0.977778	1	0.956522
COMPLEX-MSLN/MUC1/PTPRZ1	0.970588	0.985075	0.956522
PTPRZ1 AND NOT-MAGEA11 AND NOT-MUC1	0.962963	0.984848	0.942029
PTPRZ1 AND DDX3X AND NOT-MUC1	0.946565	1	0.898551
COMPLEX-MUC1/PTPRZ1/TPBG	0.978102	0.985294	0.971014
COMPLEX-MUC1/PTPRZ1/TPBG	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-ALDH1A1 AND NOT-MUC1	0.955224	0.984615	0.927536
COMPLEX-EPHA2/ABCB5/PTPRZ1	0.970588	0.985075	0.956522
PTPRZ1 AND NOT-MUC1 AND SPON2	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-MUC1 AND NOT-FOLR1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-MUC1 AND NOT-FOLR1	0.985507	0.985507	0.985507
PTPRZ1 AND NOT-MUC1 AND EPHA3	0.985507	0.985507	0.985507
COMPLEX-EPHA2/PTPRZ1/BMPR1B	0.985507	0.985507	0.985507
COMPLEX-EPHA2/PTPRZ1/ULBP2	0.970588	0.985075	0.956522
PTPRZ1 AND NOT-MUC1 AND NOT-MUC16	0.962963	0.984848	0.942029
PTPRZ1 AND TNC AND NOT-MUC1	0.970149	1	0.942029
PTPRZ1 AND NOT-SSTR2 AND NOT-MUC1	0.962963	0.984848	0.942029
COMPLEX-EPHA2/PTPRZ1/CD33	0.978102	0.985294	0.971014
COMPLEX-ERBB2/ITGB3/PTPRZ1	0.978102	0.985294	0.971014
COMPLEX-EPHA2/PTPRZ1/CLDN12	0.978102	0.985294	0.971014
COMPLEX-ERBB2/MET/PTPRZ1	0.978102	0.985294	0.971014
COMPLEX-ERBB2/PTPRZ1/CD37	0.978102	0.985294	0.971014
COMPLEX-ERBB2/PTPRZ1/BMPR1B	0.978102	0.985294	0.971014
COMPLEX-GPNMB/ERBB2/PTPRZ1	0.978102	0.985294	0.971014
COMPLEX-ERBB2/PTPRZ1/CLDN12	0.978102	0.985294	0.971014
COMPLEX-ERBB2/ABCB5/PTPRZ1	0.978102	0.985294	0.971014
COMPLEX-EPHA2/PTPRZ1/TPBG	0.962963	0.984848	0.942029
COMPLEX-MUC1/PTPRZ1/CD34	0.978102	0.985294	0.971014
COMPLEX-EPHA2/IL2RA/PTPRZ1	0.978102	0.985294	0.971014
PTPRZ1 AND NOT-MUC1 AND NOT-FOLH1	0.970588	0.985075	0.956522

Antigen Pair Logic	F1	Precision	Recall
FOLH1 AND SLC43A1 AND NOT-FCGRT	1	1	1
FOLH1 AND HPN AND NOT-DYSF	1	1	1
FOLH1 AND ANTXR2 AND NOT-SLC2A5	1	1	1
FOLH1 AND NOT-GRIN2C AND LDLRAD3	1	1	1
FOLH1 AND ADRB2 AND NOT-DYSF	1	1	1
FOLH1 AND NOT-GRIN2C AND LDLRAD3	1	1	1
FOLH1 AND ANTXR2 AND NOT-SLC2A5	1	1	1
FOLH1 AND HPN AND NOT-DYSF	1	1	1
FOLH1 AND ADRB2 AND NOT-DYSF	1	1	1
FOLH1 AND ATP7A AND NOT-TLR3	0.888889	0.8	1
FOLH1 AND NOT-CD36 AND ATP11B	0.888889	0.8	1
FOLH1 AND NOT-GRIN2C AND NOX4	0.888889	0.8	1
FOLH1 AND SLC43A1 AND NOT-NPY1R	1	1	1
FOLH1 AND ANTXR2 AND NOT-SLC52A3	0.888889	0.8	1
FOLH1 AND ANTXR2 AND NOT-SLC52A3	0.888889	0.8	1
COMPLEX-EMP3/FOLH1/OR51E2	0.888889	0.8	1
FOLH1 AND PTGIS AND NOT-CAV1	0.888889	0.8	1
FOLH1 AND NOT-CD36 AND SLC19A2	1	1	1
COMPLEX-FOLH1/OR51E2/PAQR8	0.888889	0.8	1
FOLH1 AND CDH10 AND NOT-ATP1A2	0.888889	0.8	1
FOLH1 AND NOT-ABHD6 AND F2R	0.888889	0.8	1
FOLH1 AND ATP7A AND NOT-TLR3	0.888889	0.8	1
FOLH1 AND CDH10 AND NOT-ATP1A2	0.888889	0.8	1
FOLH1 AND NOT-ABHD6 AND GGTLC1	0.888889	0.8	1
FOLH1 AND GGTLC1 AND NOT-SGCE	1	1	1
FOLH1 AND NOT-ABHD6 AND F2R	0.888889	0.8	1
FOLH1 AND NOT-CD36 AND CD69	0.888889	0.8	1
FOLH1 AND CDH10 AND NOT-NDRG4	0.888889	0.8	1
COMPLEX-CDH10/FOLH1/OR51E2	0.888889	0.8	1
FOLH1 AND CDH10 AND NOT-NDRG4	0.888889	0.8	1
FOLH1 AND TMEFF2 AND NOT-ATP1A2	0.888889	0.8	1
FOLH1 AND NOT-ABCG2 AND SLC43A1	1	1	1
FOLH1 AND NOT-ABCC6 AND TGFBI	0.888889	0.8	1
FOLH1 AND TMEFF2 AND NOT-ATP1A2	0.888889	0.8	1
FOLH1 AND NOT-CD36 AND FZD5	1	1	1
FOLH1 AND NOT-ABHD6 AND ATP7A	0.888889	0.8	1
FOLH1 AND PTGIS AND NOT-CAV1	0.888889	0.8	1
FOLH1 AND ADRB2 AND NOT-CD36	1	1	1
FOLH1 AND NOT-ABCC6 AND GGTLC1	0.888889	0.8	1
FOLH1 AND PTGIS AND NOT-SLC2A5	1	1	1
FOLH1 AND SLC43A1 AND NOT-SLC38A4	1	1	1
FOLH1 AND IL1RAPL1 AND NOT-TTYH2	0.888889	0.8	1
FOLH1 AND PTGIS AND NOT-SLC2A5	1	1	1
FOLH1 AND ADRB2 AND NOT-SLC22A18	1	1	1
FOLH1 AND NOT-CD36 AND EMB	0.888889	0.8	1
FOLH1 AND ITGA5 AND NOT-EMP3	0.888889	0.8	1
FOLH1 AND ADRB2 AND NOT-SLC22A18	1	1	1
FOLH1 AND ATP7A AND NOT-SLC2A5	1	1	1
FOLH1 AND ATP7A AND NOT-SLC2A5	1	1	1
FOLH1 AND FLVCR1 AND NOT-SLC2A5	0.888889	0.8	1
FOLH1 AND FLVCR1 AND NOT-SLC2A5	0.888889	0.8	1
TRPM8 AND NOT-FADS2 AND FOLH1	0.888889	0.8	1
FOLH1 AND NOT-SLC4A8 AND IL1RAPL1	1	1	1
TRPM8 AND NOT-SLC52A3 AND FOLH1	0.888889	0.8	1
FOLH1 AND PTGIS AND NOT-SEMA6D	1	1	1
FOLH1 AND ADRB2 AND NOT-SYT4	1	1	1
FOLH1 AND ATP7A AND NOT-CAV1	0.888889	0.8	1
FOLH1 AND BCAM AND NOT-GRIN2C	0.888889	0.8	1
FOLH1 AND NOT-GRIN2C AND ANO8	0.888889	0.8	1
FOLH1 AND ADRB2 AND NOT-SYT4	1	1	1
FOLH1 AND NOT-ABCC6 AND F2R	1	1	1
FOLH1 AND HPN AND NOT-SLC22A18	1	1	1
FOLH1 AND PTGIS AND NOT-SEMA6D	1	1	1
FOLH1 AND NOT-GRIN2C AND CCR2	0.888889	0.8	1
FOLH1 AND ADRB2 AND NOT-GYPC	1	1	1
FOLH1 AND NOT-ABCC6 AND ATP7A	0.888889	0.8	1
FOLH1 AND ATP7A AND NOT-SLC52A3	1	1	1
FOLH1 AND BCAM AND NOT-SLC39A2	1	1	1
TRPM8 AND NOT-UNC5B AND FOLH1	1	1	1
FOLH1 AND ANTXR2 AND NOT-FADS2	0.888889	0.8	1
FOLH1 AND SEMA4B AND NOT-SLC52A3	0.888889	0.8	1
FOLH1 AND NOT-ABCC6 AND F2R	1	1	1
FOLH1 AND NOT-SLC4A8 AND NOX4	0.888889	0.8	1
FOLH1 AND NOT-SLC4A8 AND CCR2	0.888889	0.8	1
FOLH1 AND PTGIS AND NOT-SLC52A3	1	1	1
FOLH1 AND HPN AND NOT-SLC22A18	1	1	1
FOLH1 AND NOT-GRIN2C AND GPR176	0.888889	0.8	1

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
PTPRZ1 AND NOT-B4GALNT1 AND NOT-MUC1	0.970588	0.985075	0.956522
COMPLEX-DKK1/MUC1/PTPRZ1	0.985507	0.985507	0.985507
COMPLEX-MUC1/PTPRZ1/BMPR1B	0.985507	0.985507	0.985507
COMPLEX-MET/MUC1/PTPRZ1	0.985507	0.985507	0.985507
COMPLEX-MUC1/PTPRZ1/VCAM1	0.985507	0.985507	0.985507
COMPLEX-ITGB3/MUC1/PTPRZ1	0.985507	0.985507	0.985507
COMPLEX-GPNMB/MUC1/PTPRZ1	0.985507	0.985507	0.985507
SYT11 AND NOT-SLC6A15 AND EPHA8	0.948148	0.969697	0.927536
COMPLEX-F2R/SLCO1C1/GPR158	0.957746	0.931507	0.985507
SYT11 AND NOT-SLC6A15 AND GJD2	0.948148	0.969697	0.927536
SYT11 AND NOT-SLC6A15 AND CCR9	0.948148	0.969697	0.927536
SYT11 AND NOT-GRM1 AND GJD2	0.94964	0.942857	0.956522
SYT11 AND NOT-SLC6A15 AND MRAP	0.932331	0.96875	0.898551
COMPLEX-ABHD3/SYT11/SLC6A15	0.948148	0.969697	0.927536
SYT11 AND NOT-CNTNAP2 AND GYPC	0.971014	0.971014	0.971014
SYT11 AND NOT-GRM1 AND CATSPER3	0.94964	0.942857	0.956522
SYT11 AND NOT-GRM1 AND PIK3IP1	0.94964	0.942857	0.956522
SYT11 AND CD93 AND ATP13A5	0.957143	0.943662	0.971014
SYT11 AND NOT-CNTNAP2 AND TGFBI	0.971014	0.971014	0.971014
SYT11 AND NOT-CNTNAP2 AND SLC39A8	0.978102	0.985294	0.971014
SYT11 AND NOT-GRM1 AND NOT-SLC31A1	0.926471	0.940299	0.913043
SYT11 AND NOT-CNTNAP2 AND ATP13A5	0.971014	0.971014	0.971014
COMPLEX-SYT11/GRM1/SLC39A8	0.94964	0.942857	0.956522
SYT11 AND NOT-SLC6A15 AND NOT-SLC31A1	0.916031	0.967742	0.869565
SYT11 AND HTR1F AND NOT-SLC9B1	0.964539	0.944444	0.985507
COMPLEX-SLC31A1/SYT11/GJD2	0.933333	0.954545	0.913043
SYT11 AND ATP13A5 AND NOT-MRAP	0.978417	0.971429	0.985507
SYT11 AND NOT-ZACN AND ATP13A5	0.971429	0.957746	0.985507
COMPLEX-SYT11/IL17RA/GABRA5	0.94964	0.942857	0.956522
GPR158 AND CD93 AND ATP13A5	0.892086	0.885714	0.898551
COMPLEX-SYT11/IFI6/CD58	0.956522	0.956522	0.956522
COMPLEX-SYT11/IL17RA/SLC39A8	0.955882	0.970149	0.942029
COMPLEX-BTN3A1/SLCO1C1/GPR158	0.878378	0.822785	0.942029
COMPLEX-SYT11/IL17RA/BST2	0.957143	0.943662	0.971014
COMPLEX-SLCO1C1/GPR158/CD163	0.879433	0.861111	0.898551
COMPLEX-CD93/SLCO1C1/GPR158	0.876712	0.831169	0.927536
GPR19 AND CD93 AND NOT-IL17RA	0.887097	1	0.797101
GPR19 AND CD93 AND ATP13A5	0.887097	1	0.797101
GPR19 AND SLC31A1 AND NOT-SLC6A15	0.904762	1	0.826087
ASTN1 AND NOT-MRAP AND SLC31A1	0.904762	1	0.826087
COMPLEX-CLDN15/SLCO1C1/GPR158	0.92517	0.871795	0.985507
SYT11 AND SLC31A1 AND NOT-ZACN	0.938462	1	0.884058
COMPLEX-SLCO1C1/GPR158/PPAPDC1B	0.888889	0.853333	0.927536
SYT11 AND SLC31A1 AND NOT-MRAP	0.938462	1	0.884058
GPR19 AND SLC31A1 AND NOT-MRAP	0.921875	1	0.855072
GPR19 AND HTR1F AND NOT-ROS1	0.857143	0.84507	0.869565
GPR19 AND SLC31A1 AND NOT-GRM1	0.921875	1	0.855072
GPR158 AND NOT-OPRK1 AND ABHD3	0.855172	0.815789	0.898551
CDH10 AND SLC31A1 AND NOT-MRAP	0.878049	1	0.782609
ASTN1 AND NOT-SLC6A15 AND ABHD3	0.852459	0.981132	0.753623
GPR158 AND SLC31A1 AND NOT-MRAP	0.930233	1	0.869565
GPR158 AND SLC31A1 AND NOT-ZACN	0.930233	1	0.869565
COMPLEX-BTN3A3/SLCO1C1/GPR158	0.851613	0.767442	0.956522
GPR19 AND HTR1F AND NOT-SLC9B1	0.851064	0.833333	0.869565
GPR19 AND SLC31A1 AND NOT-CLDN17	0.913386	1	0.84058
GPR158 AND SLC31A1 AND NOT-GRM1	0.930233	1	0.869565
GPR19 AND SLC31A1 AND NOT-GJD2	0.921875	1	0.855072
LYPD1 AND NOT-CNTNAP2 AND NOT-ATP13A5	0.875	0.949153	0.811594
SCN1A AND NOT-SLC6A15 AND SLC31A1	0.852459	0.981132	0.753623
GPR19 AND SLC31A1 AND NOT-ZACN	0.921875	1	0.855072
GPR19 AND GYPC AND NOT-CNTNAP2	0.850746	0.876923	0.826087
CDH10 AND SLC31A1 AND NOT-ZACN	0.878049	1	0.782609
LYPD1 AND NOT-ZACN AND ATP13A5	0.876923	0.934426	0.826087
LYPD1 AND NOT-HTR1F AND ROS1	0.875	0.949153	0.811594
COMPLEX-IFNAR2/SLCO1C1/GPR158	0.87218	0.90625	0.84058
FXYD6 AND SLC31A1 AND NOT-MRAP	0.842975	0.980769	0.73913
FXYD6 AND SLC31A1 AND NOT-ZACN	0.842975	0.980769	0.73913
COMPLEX-GPR22/SLCO1C1/GPR158	0.842767	0.744444	0.971014
LYPD1 AND NOT-SLC9B1 AND PCDHGC4	0.876923	0.934426	0.826087
GPR158 AND SLC31A1 AND NOT-SLC22A5	0.930233	1	0.869565
KCNJ10 AND NOT-GRM1 AND SLC31A1	0.841379	0.802632	0.884058
COMPLEX-SEMA4B/SLCO1C1/GPR158	0.860927	0.792683	0.942029
LYPD1 AND NOT-GRM1 AND GJD2	0.876923	0.934426	0.826087
CDH10 AND SLC31A1 AND NOT-CLDN17	0.868852	1	0.768116
CDH10 AND CD93 AND ATP13A5	0.840336	1	0.724638
LYPD1 AND XCR1 AND NOT-MRAP	0.897638	0.982759	0.826087
GPR19 AND SLC31A1 AND NOT-CATSPER3	0.921875	1	0.855072

Antigen Pair Logic	F1	Precision	Recall
FOLH1 AND NOT-GRIN2C AND CCR2	0.888889	0.8	1
FOLH1 AND NOT-GRIN2C AND ANO8	0.888889	0.8	1
COMPLEX-PIK3IP1/FOLH1/OR51E2	0.888889	0.8	1
TRPM8 AND NOT-UNC5B AND FOLH1	1	1	1
FOLH1 AND ATP7A AND NOT-CAV1	0.888889	0.8	1
FOLH1 AND SEMA4B AND NOT-SLC52A3	0.888889	0.8	1
FOLH1 AND NOT-SLC4A8 AND CCR2	0.888889	0.8	1
FOLH1 AND ADRB2 AND NOT-SLC16A5	1	1	1
FOLH1 AND ATP8B2 AND NOT-SLC2A5	0.888889	0.8	1
FOLH1 AND CDH10 AND NOT-PTPRZ1	0.888889	0.8	1
FOLH1 AND NOT-ABCC6 AND ATP7A	0.888889	0.8	1
FOLH1 AND ATP7A AND NOT-SLC52A3	1	1	1
FOLH1 AND NOT-SEMA6D AND FLVCR1	0.888889	0.8	1
FOLH1 AND ATP8B2 AND NOT-SLC2A5	0.888889	0.8	1
FOLH1 AND NOT-SEMA6D AND FLVCR1	0.888889	0.8	1
FOLH1 AND SLC43A1 AND NOT-SLC22A18	1	1	1
FOLH1 AND NOT-GRIN2C AND SELP	0.888889	0.8	1
TRPM8 AND NOT-SLC39A2 AND FOLH1	1	1	1
FOLH1 AND NOT-NRG3 AND ATP8A2	0.857143	1	0.75
FOLH1 AND PTGIS AND NOT-SLC9A1	0.857143	1	0.75
FOLH1 AND STEAP4 AND NOT-GYPC	0.857143	1	0.75
FOLH1 AND STEAP4 AND NOT-EPHA2	0.857143	1	0.75
FOLH1 AND ATP7A AND NOT-SLC9A1	0.857143	1	0.75
FOLH1 AND STEAP4 AND NOT-SLC22A18	0.857143	1	0.75
FOLH1 AND NOT-CD36 AND ATP7A	0.888889	0.8	1
FOLH1 AND SLC43A1 AND NOT-DYSF	1	1	1
FOLH1 AND STEAP4 AND NOT-SYT4	0.857143	1	0.75
FOLH1 AND STEAP4 AND NOT-DYSF	0.857143	1	0.75
FOLH1 AND STEAP4 AND NOT-EMP3	0.857143	1	0.75
FOLH1 AND NOT-GRIN2C AND PCDH8	0.857143	1	0.75
FOLH1 AND STEAP4 AND NOT-SEMA4D	0.857143	1	0.75
FOLH1 AND NOT-SLC9A1 AND SEMA4B	0.857143	1	0.75
FOLH1 AND GHR AND NOT-DYSF	0.857143	1	0.75
FOLH1 AND CDH10 AND NOT-NRG3	0.857143	1	0.75
FOLH1 AND NOT-GRIN2C AND GRIA4	0.888889	0.8	1
FOLH1 AND STEAP4 AND NOT-TNFRSF12A	0.857143	1	0.75
FOLH1 AND NOT-SLC4A8 AND CHRNB4	0.888889	0.8	1
FOLH1 AND STEAP4 AND NOT-STAB1	0.857143	1	0.75
FOLH1 AND NOT-GRIN2C AND CDH10	1	1	1
FOLH1 AND STEAP4 AND NOT-GRIN2C	0.857143	1	0.75
FOLH1 AND STEAP4 AND NOT-BST2	0.857143	1	0.75
FOLH1 AND NOT-SLC4A8 AND CDH10	1	1	1
FOLH1 AND ANTXR2 AND NOT-SLC39A2	0.857143	1	0.75
FOLH1 AND STEAP4 AND NOT-SLC4A8	0.857143	1	0.75
FOLH1 AND STEAP4 AND NOT-PAQR8	0.857143	1	0.75
FOLH1 AND STEAP4 AND NOT-GABRA1	0.857143	1	0.75
FOLH1 AND NOT-SLC4A8 AND CHRNB4	0.888889	0.8	1
FOLH1 AND NOT-GRIN2C AND MDGA2	0.857143	1	0.75
FOLH1 AND NOT-GRIN2C AND MDGA2	0.857143	1	0.75
FOLH1 AND NOT-CD36 AND ATP8B2	0.857143	1	0.75
FOLH1 AND STEAP4 AND NOT-CDH10	0.857143	1	0.75
FOLH1 AND STEAP4 AND NOT-NKAIN2	0.857143	1	0.75
FOLH1 AND SLC43A1 AND NOT-SLC30A10	1	1	1
FOLH1 AND SLC43A1 AND NOT-BST2	1	1	1
FOLH1 AND SLC43A1 AND NOT-SCARA5	1	1	1
FOLH1 AND NOT-GABRA2 AND ATP8A2	0.857143	1	0.75
FOLH1 AND ATP8A2 AND NOT-SYT4	0.857143	1	0.75
FOLH1 AND NOT-SLC4A8 AND ATP8A2	0.857143	1	0.75
FOLH1 AND NOT-PAQR8 AND ATP8A2	0.888889	0.8	1
FOLH1 AND NOT-GABRA2 AND ATP8A2	0.857143	1	0.75
FOLH1 AND NOT-CD36 AND F2R	1	1	1
FOLH1 AND NOT-SLC4A8 AND ATP8A2	0.857143	1	0.75
FOLH1 AND NOT-SLC4A8 AND CACNA1E	0.888889	0.8	1
FOLH1 AND NOT-SLC4A8 AND PCDH8	0.857143	1	0.75
FOLH1 AND ATP7A AND NOT-SLC16A5	1	1	1
FOLH1 AND NOT-GRIN2C AND TMPRSS11E	0.888889	0.8	1
FOLH1 AND NOT-SLC4A8 AND TAS2R10	0.888889	0.8	1
FOLH1 AND NOT-GRIN2C AND CACNA1E	0.888889	0.8	1
FOLH1 AND GHR AND NOT-SLC22A18	0.857143	1	0.75
FOLH1 AND NOT-GRIN2C AND PPAPDC1B	0.888889	0.8	1
FOLH1 AND PTGIS AND NOT-SLC4A8	1	1	1
FOLH1 AND ATP8B2 AND NOT-SLC39A2	0.857143	1	0.75
FOLH1 AND NOT-SLC4A8 AND TMEFF2	0.888889	0.8	1
FOLH1 AND NOT-GRIN2C AND PPAPDC1B	0.888889	0.8	1
FOLH1 AND NOT-SLC4A8 AND USH2A	0.888889	0.8	1
FOLH1 AND PTGIS AND NOT-SYT4	0.888889	0.8	1
FOLH1 AND SELP AND NOT-FMNL1	0.857143	1	0.75

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
GPR19 AND SLC31A1 AND NOT-GABRA5	0.913386	1	0.84058
GPR158 AND SLC31A1 AND GJD2	0.930233	1	0.869565
FXYD6 AND SLC31A1 AND NOT-SLC22A5	0.842975	0.980769	0.73913
SCN1A AND NOT-OPRK1 AND ABHD3	0.839695	0.887097	0.797101
SCN2A AND SLC31A1 AND NOT-MRAP	0.880597	0.907692	0.855072
OMG AND ABHD3 AND NOT-OPRK1	0.835821	0.861538	0.811594
CDH10 AND SLC31A1 AND NOT-GRM1	0.840336	1	0.724638
FXYD6 AND SLC31A1 AND GJD2	0.842975	0.980769	0.73913
OMG AND SLC31A1 AND NOT-MRAP	0.857143	0.890625	0.826087
GPR158 AND TGFBI AND NOT-CNTNAP2	0.836879	0.819444	0.855072
COMPLEX-SLCO1C1/GPR158/CD58	0.845638	0.7875	0.913043
COMPLEX-CDH10/ABHD3/SLC6A15	0.832	0.928571	0.753623
GPR19 AND SLC9B1 AND NOT-PCDHGC4	0.830986	0.808219	0.855072
KCNJ10 AND SLC31A1 AND NOT-ZACN	0.834532	0.828571	0.84058
CDH10 AND SLC31A1 AND NOT-GJD2	0.878049	1	0.782609
OMG AND SLC31A1 AND NOT-CLDN17	0.857143	0.890625	0.826087
KCNJ10 AND SLC31A1 AND NOT-MRAP	0.84058	0.84058	0.84058
CDH10 AND SLC31A1 AND NOT-SLC22A5	0.878049	1	0.782609
FXYD6 AND SLC31A1 AND CLDN17	0.833333	0.980392	0.724638
KCNJ10 AND SLC31A1 AND NOT-CLDN17	0.84058	0.84058	0.84058
GPR19 AND NOT-SLC6A15 AND GJD2	0.827068	0.859375	0.797101
GPR19 AND NOT-SLC6A15 AND MRAP	0.827068	0.859375	0.797101
LYPD1 AND ABHD3 AND NOT-OPRK1	0.878049	1	0.782609
COMPLEX-CDH11/SLCO1C1/GPR158	0.826667	0.765432	0.898551
CDH10 AND NOT-OPRK1 AND ABHD3	0.826446	0.961538	0.724638
NKAIN4 AND SLC31A1 AND NOT-CLDN17	0.826446	0.961538	0.724638
ASTN1 AND NOT-OPRK1 AND ABHD3	0.826446	0.961538	0.724638
SCN1A AND NOT-GRM1 AND SLC31A1	0.854839	0.963636	0.768116
SYT11 AND NOT-SLC6A15 AND CD37	0.948148	0.969697	0.927536
SYT11 AND NOT-SLC6A15 AND CD19	0.940299	0.969231	0.913043
SYT11 AND NOT-GRM1 AND CD19	0.942029	0.942029	0.942029
SYT11 AND NOT-CNTNAP2 AND CD33	0.971014	0.971014	0.971014
SYT11 AND NOT-GRM1 AND LGR5	0.942029	0.942029	0.942029
COMPLEX-ITGAV/SLCO1C1/GPR158	0.930556	0.893333	0.971014
SYT11 AND TGFBI AND NOT-ULBP2	0.94964	0.942857	0.956522
SYT11 AND ABCB5 AND NOT-MRAP	0.971014	0.971014	0.971014
COMPLEX-CBX3/SLCO1C1/GPR158	0.911765	0.925373	0.898551
COMPLEX-GPNMB/SYT11/CNTNAP2	0.971014	0.971014	0.971014
SYT11 AND PROCR AND NOT-CD19	0.956522	0.956522	0.956522
SYT11 AND ATP13A5 AND LGR5	0.942029	0.942029	0.942029
SYT11 AND ATP13A5 AND NOT-ULBP2	0.957143	0.943662	0.971014
SYT11 AND PLVAP AND NOT-CD19	0.910448	0.938462	0.884058
SYT11 AND NOT-MRAP AND FOLR1	0.978417	0.971429	0.985507
SYT11 AND HTR1F AND NOT-CD19	0.964539	0.944444	0.985507
SYT11 AND CLDN16 AND NOT-CD19	0.964539	0.944444	0.985507
SYT11 AND NOT-MRAP AND CD19	0.955882	0.970149	0.942029
SYT11 AND NOT-GUCY2D AND CD19	0.942029	0.942029	0.942029
COMPLEX-SEMA4D/SYT11/STEAP2	0.964539	0.944444	0.985507
COMPLEX-SLC31A1/SYT11/P2RX5	0.957143	0.943662	0.971014
COMPLEX-SYT11/STEAP2/SLC39A8	0.964539	0.944444	0.985507
COMPLEX-SYT11/IL17RA/SSTR1	0.942029	0.942029	0.942029
SYT11 AND NOT-ZACN AND NOT-ULBP2	0.892308	0.95082	0.84058
COMPLEX-GPNMB/SYT11/ZACN	0.971429	0.957746	0.985507
SYT11 AND NOT-GABRA5 AND ULBP2	0.875	0.949153	0.811594
SYT11 AND ATP13A5 AND NOT-SSTR1	0.918519	0.939394	0.898551
BCAN AND NOT-CLDN17 AND SLC31A1	0.868852	1	0.768116
BCAN AND NOT-MRAP AND SLC31A1	0.868852	1	0.768116
BCAN AND NOT-CATSPER3 AND SLC31A1	0.868852	1	0.768116
BCAN AND NOT-ZACN AND SLC31A1	0.868852	1	0.768116
BCAN AND NOT-GJD2 AND SLC31A1	0.868852	1	0.768116
SYT11 AND NOT-GABRA5 AND CD19	0.866142	0.948276	0.797101
SYT11 AND CD58 AND NOT-ULBP2	0.925373	0.953846	0.898551
GPR19 AND SLC31A1 AND NOT-CD19	0.921875	1	0.855072
COMPLEX-SYT11/SLC22A5/SSTR1	0.918519	0.939394	0.898551
BCAN AND NOT-GABRA5 AND SLC31A1	0.859504	1	0.753623
COMPLEX-HSPA5/SLCO1C1/GPR158	0.928571	0.915493	0.942029
LYPD1 AND NOT-KCNA7 AND CD19	0.873016	0.964912	0.797101
COMPLEX-TNC/SLCO1C1/GPR158	0.882759	0.842105	0.927536
SYT11 AND IL17RA AND NOT-ULBP2	0.907692	0.967213	0.855072
GPR19 AND NOT-P2RX5 AND SLC31A1	0.887097	1	0.797101
LYPD1 AND NOT-HTR1F AND CD19	0.875	0.949153	0.811594
COMPLEX-SLC39A6/SLCO1C1/GPR158	0.839161	0.810811	0.869565
LYPD1 AND ABCB5 AND NOT-MRAP	0.897638	0.982759	0.826087
LYPD1 AND NOT-GRM1 AND CD19	0.868217	0.933333	0.811594
GPR158 AND SLC31A1 AND NOT-P2RX5	0.930233	1	0.869565
CDH10 AND SLC31A1 AND NOT-P2RX5	0.868852	1	0.768116
LYPD1 AND NOT-GUCY2D AND CD19	0.868217	0.933333	0.811594

Antigen Pair Logic	F1	Precision	Recall
FOLH1 AND NOT-SLC4A8 AND TMEFF2	0.888889	0.8	1
FOLH1 AND STEAP4 AND NOT-NFASC	0.857143	1	0.75
FOLH1 AND NOT-NRG3 AND CHRNB4	0.857143	1	0.75
FOLH1 AND ANTXR2 AND NOT-GRIN2C	0.888889	0.8	1
FOLH1 AND PTGIS AND NOT-GRIN2C	1	1	1
FOLH1 AND STEAP4 AND NOT-TTYH2	0.857143	1	0.75
FOLH1 AND PTGIS AND NOT-NRG3	0.857143	1	0.75
FOLH1 AND NOT-NRG3 AND GABRA3	0.857143	1	0.75
FOLH1 AND NOT-SLC4A8 AND NFASC	0.888889	0.8	1
FOLH1 AND PTGIS AND NOT-SLC39A2	0.857143	1	0.75
FOLH1 AND PTGIS AND NOT-GRIN2C	1	1	1
FOLH1 AND NOT-GRIN2C AND TMEFF2	0.888889	0.8	1
FOLH1 AND STEAP4 AND NOT-CNTNAP2	0.857143	1	0.75
FOLH1 AND PTGIS AND NOT-SYT4	0.888889	0.8	1
FOLH1 AND PTGIS AND NOT-UMOD	0.857143	1	0.75
FOLH1 AND NOT-NRG3 AND NPHS1	0.857143	1	0.75
FOLH1 AND NOT-SLC4A8 AND NPHS1	0.888889	0.8	1
TRPM8 AND NOT-GRIN2C AND FOLH1	0.888889	0.8	1
FOLH1 AND NOT-GRIN2C AND PCDH15	0.888889	0.8	1
FOLH1 AND ANTXR2 AND NOT-GRIN2C	0.888889	0.8	1
FOLH1 AND NOT-GRIN2C AND USH2A	0.888889	0.8	1
FOLH1 AND NOT-GRIN2C AND TMEFF2	0.888889	0.8	1
FOLH1 AND NOT-GRIN2C AND PCDH15	0.888889	0.8	1
FOLH1 AND STEAP4 AND NOT-BTN3A1	0.857143	1	0.75
FOLH1 AND NOT-SLC4A8 AND GABRA3	0.888889	0.8	1
FOLH1 AND NOT-GRIN2C AND SCN2A	0.888889	0.8	1
FOLH1 AND STEAP4 AND NOT-SCN8A	0.857143	1	0.75
TRPM8 AND NOT-GRIN2C AND FOLH1	0.888889	0.8	1
FOLH1 AND NOT-SLC4A8 AND GABRA3	0.888889	0.8	1
FOLH1 AND PTGIS AND NOT-SLC4A8	1	1	1
FOLH1 AND ATP7A AND NOT-SLC22A18	0.888889	0.8	1
FOLH1 AND NOT-GRIN2C AND NOT-SLC22A18	0.888889	0.8	1
FOLH1 AND GGTLC1 AND NOT-DYSF	1	1	1
FOLH1 AND SLC43A1 AND NOT-STAB1	1	1	1
FOLH1 AND ATP7A AND NOT-SLC22A18	0.888889	0.8	1
FOLH1 AND NOT-GRIN2C AND NMBR	0.888889	0.8	1
FOLH1 AND SELP AND NOT-SLC4A8	1	1	1
FOLH1 AND NOT-GRIN2C AND EPHA10	0.888889	0.8	1
FOLH1 AND EPHA3 AND NOT-SYT4	0.888889	0.8	1
FOLH1 AND EPHA3 AND NOT-SYT4	0.888889	0.8	1
FOLH1 AND ROR1 AND NOT-GYPC	0.888889	0.8	1
FOLH1 AND ROR1 AND NOT-EMP3	0.888889	0.8	1
STEAP2 AND NOT-SLC39A2 AND FOLH1	1	1	1
FOLH1 AND EPHA3 AND NOT-GRIN2C	1	1	1
STEAP2 AND NOT-SLC39A2 AND FOLH1	1	1	1
FOLH1 AND EPHA3 AND NOT-GRIN2C	1	1	1
FOLH1 AND STEAP4 AND NOT-CSPG4	0.857143	1	0.75
FOLH1 AND EPCAM AND NOT-PAQR8	0.857143	1	0.75
FOLH1 AND NOT-EDNRB AND ATP7A	0.857143	1	0.75
FOLH1 AND NOT-CD36 AND MUC1	0.888889	0.8	1
FOLH1 AND NOT-EDNRB AND PTGIS	0.857143	1	0.75
FOLH1 AND EPHA3 AND NOT-SLC9A1	0.857143	1	0.75
FOLH1 AND EPCAM AND TMEFF2	0.857143	1	0.75
FOLH1 AND EPCAM AND NOT-CD58	0.857143	1	0.75
FOLH1 AND EPHA3 AND NOT-SLC39A2	0.857143	1	0.75
FOLH1 AND PCDH8 AND NOT-NCAM1	0.857143	1	0.75
FOLH1 AND CDH10 AND NOT-NCAM1	1	1	1
FOLH1 AND EPCAM AND LRRC8E	0.857143	1	0.75
FOLH1 AND CDH10 AND NOT-NCAM1	1	1	1
FOLH1 AND EPCAM AND CDH10	0.857143	1	0.75
COMPLEX-FOLH1/STEAP2/SLC16A5	0.857143	1	0.75
FOLH1 AND NOT-EDNRB AND CD163	0.857143	1	0.75
FOLH1 AND STEAP4 AND NOT-AXL	0.857143	1	0.75
FOLH1 AND NOT-GRIN2C AND BIRC5	0.888889	0.8	1
FOLH1 AND CLDN8 AND NOT-CLDN10	1	1	1
FOLH1 AND EPHA3 AND NOT-CALN1	0.857143	1	0.75
FOLH1 AND EPHA3 AND NOT-UMOD	0.857143	1	0.75
FOLH1 AND EPHA3 AND NOT-NRG3	0.857143	1	0.75
FOLH1 AND NOT-GRIN2C AND BIRC5	0.888889	0.8	1
FOLH1 AND CLDN8 AND NOT-SLC13A2	0.857143	1	0.75
FOLH1 AND NOT-DPEP1 AND ATP8B1	0.857143	1	0.75
FOLH1 AND EPHA3 AND NOT-SLC22A18	0.888889	0.8	1
FOLH1 AND EPHA3 AND NOT-SLC22A18	0.888889	0.8	1
FOLH1 AND NOT-CD36 AND IL20RA	0.857143	1	0.75
FOLH1 AND EPCAM AND NOT-SLC30A10	0.857143	1	0.75
FOLH1 AND NOT-MET AND ATP7A	0.888889	0.8	1
FOLH1 AND TMEFF2 AND NOT-NCAM1	0.888889	0.8	1

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
LYPD1 AND NOT-MRAP AND CD19	0.888889	0.982456	0.811594
COMPLEX-OMG/GPR158/BCAN	0.832117	0.838235	0.826087
NCAM1 AND NOT-SLC6A15 AND SLC31A1	0.830769	0.885246	0.782609
KCNJ10 AND SLC31A1 AND NOT-P2RX5	0.87218	0.90625	0.84058
FXYD6 AND SLC31A1 AND CD19	0.833333	0.980392	0.724638
GPR19 AND HTR1F AND NOT-CD19	0.827586	0.789474	0.869565
CDH10 AND SLC31A1 AND NOT-CD19	0.878049	1	0.782609
GPR19 AND NOT-GRM1 AND STEAP2	0.827068	0.859375	0.797101
LYPD1 AND PLVAP AND NOT-CD19	0.845528	0.962963	0.753623
LYPD1 AND TGFBI AND NOT-ULBP2	0.876923	0.934426	0.826087
GPR158 AND NOT-CNTNAP2 AND GPNMB	0.863309	0.857143	0.869565
COMPLEX-SLCO1C1/GPR158/CD276	0.887417	0.817073	0.971014
LYPD1 AND NOT-CATSPER3 AND ULBP2	0.890625	0.966102	0.826087
GPR19 AND PROCR AND NOT-CD19	0.822695	0.805556	0.84058
GPR19 AND ATP1B4 AND NOT-CD19	0.821918	0.779221	0.869565
NKAIN4 AND SLC31A1 AND NOT-CD19	0.836066	0.962264	0.73913
GPR19 AND LGR5 AND NOT-GRM1	0.820896	0.846154	0.797101
OMG AND SLC31A1 AND NOT-P2RX5	0.850746	0.876923	0.826087
CDH10 AND TGFBI AND NOT-ULBP2	0.848	0.946429	0.768116
GPR19 AND PLVAP AND NOT-CD19	0.819444	0.786667	0.855072
GPR19 AND CLDN16 AND NOT-CD19	0.821918	0.779221	0.869565
SYT11 AND SLC39A8 AND NOT-SSTR1	0.896	1	0.811594
GPR158 AND GPNMB AND NOT-MRAP	0.857143	0.947368	0.782609
GPR19 AND ABCB5 AND NOT-MRAP	0.848921	0.842857	0.855072
GPR19 AND RXFP3 AND NOT-CD19	0.827586	0.789474	0.869565
GPR19 AND NOT-STEAP2 AND GABRA5	0.813793	0.776316	0.855072
COMPLEX-LYPD1/ULBP2/CD58	0.876923	0.934426	0.826087
COMPLEX-SLCO1C1/GPR158/ST8SIA1	0.867647	0.880597	0.855072
ASTN1 AND SLC31A1 AND NOT-CD19	0.896	1	0.811594
COMPLEX-IL13RA1/SLCO1C1/GPR158	0.861314	0.867647	0.855072
GPR158 AND GPNMB AND NOT-ZACN	0.80597	0.830769	0.782609
COMPLEX-ASTN1/TGFBI/ULBP2	0.859375	0.932203	0.797101
SYT11 AND NOT-SSTR1 AND MRAP	0.900763	0.951613	0.855072
CDH10 AND CD58 AND NOT-ULBP2	0.830508	1	0.710145
NCAM1 AND ABHD3 AND NOT-OPRK1	0.878049	1	0.782609
GPR19 AND GPNMB AND NOT-CNTNAP2	0.803419	0.979167	0.681159
NKAIN4 AND SLC31A1 AND NOT-P2RX5	0.803279	0.924528	0.710145
MLC1 AND CD58 AND NOT-ULBP2	0.80315	0.87931	0.73913
NCAM1 AND SLC31A1 AND NOT-MRAP	0.868852	1	0.768116
GPR19 AND NOT-CNTNAP2 AND STEAP2	0.825397	0.912281	0.753623
GPR19 AND GPNMB AND NOT-ZACN	0.803419	0.979167	0.681159
NOT-ATP8B1 AND SLC31A1 AND NOT-CD19	0.823529	0.835821	0.811594
GPR19 AND LGR5 AND ATP13A5	0.8	0.763158	0.84058
GPR19 AND GPNMB AND NOT-MRAP	0.810345	1	0.681159
COMPLEX-SLCO1C1/GPR158/CLDN12	0.902778	0.866667	0.942029
LYPD1 AND NOT-ULBP2 AND IL17RA	0.859375	0.932203	0.797101
CDH10 AND NOT-CNTNAP2 AND GPNMB	0.803419	0.979167	0.681159
COMPLEX-SYT11/SSTR1/ULBP2	0.956522	0.956522	0.956522
SYT11 AND NOT-STEAP2 AND ULBP2	0.964539	0.944444	0.985507
SYT11 AND LGR5 AND CD19	0.934307	0.941176	0.927536
BCAN AND NOT-CD19 AND SLC31A1	0.868852	1	0.768116
GPR19 AND LGR5 AND NOT-CD19	0.805556	0.773333	0.84058
NOT-EPCAM AND SLC31A1 AND NOT-P2RX5	0.892086	0.885714	0.898551
ITGAV AND NOT-GPNMB AND NOT-MRAP	0.852713	0.916667	0.797101
ITGAV AND NOT-PROCR AND NOT-CD19	0.848485	0.888889	0.811594
ITGAV AND NOT-GPNMB AND CNTNAP2	0.833333	0.873016	0.797101
NOT-EPCAM AND SLC31A1 AND NOT-CD19	0.882759	0.842105	0.927536
COMPLEX-GPR19/SSTR1/ULBP2	0.834532	0.828571	0.84058
NOT-CEACAM6 AND SLC39A8 AND STEAP2	0.874074	0.893939	0.855072
ITGAV AND NOT-MRAP AND SSTR1	0.955882	0.970149	0.942029
NOT-CEACAM6 AND SLC39A8 AND SSTR1	0.863014	0.818182	0.913043
ITGAV AND NOT-MRAP AND CD19	0.876923	0.934426	0.826087
NOT-RNF43 AND SLC31A1 AND NOT-CD19	0.867133	0.837838	0.898551
EDNRB AND NOT-MRAP AND NOT-FOLR1	0.8	0.892857	0.724638
ITGAV AND RXFP3 AND CD19	0.859259	0.878788	0.84058
NOT-RNF43 AND SLC31A1 AND NOT-P2RX5	0.875912	0.882353	0.869565
ITGAV AND NOT-MRAP AND FOLR1	0.918519	0.939394	0.898551
ITGAV AND NOT-MRAP AND NOT-ABCB5	0.910448	0.938462	0.884058
NOT-CEACAM6 AND SLC31A1 AND NOT-P2RX5	0.93617	0.916667	0.956522
ITGAV AND CD19 AND SLC6A15	0.852941	0.865672	0.84058
ITGAV AND CD19 AND ATP1B4	0.84058	0.84058	0.84058
ITGAV AND CD19 AND NOT-KCNA7	0.863636	0.904762	0.826087
ITGAV AND PCDHGC4 AND CD19	0.846715	0.852941	0.84058
ITGAV AND CD19 AND NOT-XCR1	0.876923	0.934426	0.826087
ITGAV AND CD19 AND GABRA5	0.846715	0.852941	0.84058
ITGAV AND CD19 AND HTR1F	0.846715	0.852941	0.84058
ITGAV AND CD19 AND SLC2A2	0.852941	0.865672	0.84058

Antigen Pair Logic	F1	Precision	Recall
FOLH1 AND IL20RA AND NOT-SLC9A1	0.857143	1	0.75
FOLH1 AND NOT-GRIN2C AND ROR1	1	1	1
FOLH1 AND TNC AND NOT-SYT4	0.888889	0.8	1
FOLH1 AND CLDN8 AND NOT-SLC9A1	0.857143	1	0.75
FOLH1 AND TMEFF2 AND NOT-NCAM1	0.888889	0.8	1
FOLH1 AND ROR1 AND NOT-PAQR8	0.888889	0.8	1
FOLH1 AND STEAP4 AND NOT-NCAM1	0.857143	1	0.75
FOLH1 AND PTGIS AND NOT-ITGB3	0.857143	1	0.75
FOLH1 AND ANTXR2 AND EPCAM	0.857143	1	0.75
COMPLEX-FOLH1/STEAP2/SLC43A1	1	1	1
FOLH1 AND STEAP4 AND NOT-ITGB6	0.857143	1	0.75
FOLH1 AND ROR1 AND NOT-SLC22A18	0.888889	0.8	1
FOLH1 AND TNC AND NOT-EMP3	0.888889	0.8	1
FOLH1 AND EPCAM AND NOT-SLC22A18	0.857143	1	0.75
FOLH1 AND NOT-SLC4A8 AND SLC39A6	0.888889	0.8	1
FOLH1 AND EPHA3 AND NOT-CLSTN3	0.857143	1	0.75
FOLH1 AND EPHA3 AND NOT-ABCA12	0.888889	0.8	1
FOLH1 AND EPHA3 AND NOT-GRIN2A	0.857143	1	0.75
STEAP2 AND NOT-GRIN2C AND FOLH1	0.888889	0.8	1
FOLH1 AND NOT-DPEP1 AND BTN3A1	0.857143	1	0.75
FOLH1 AND STEAP4 AND NOT-LGR5	0.857143	1	0.75
FOLH1 AND PTGIS AND NOT-B4GALNT1	0.857143	1	0.75
FOLH1 AND NOT-GRIN2C AND CLDN8	1	1	1
FOLH1 AND NOT-DPEP1 AND LRRC8E	0.857143	1	0.75
FOLH1 AND NOT-GRIN2C AND TPBG	0.857143	1	0.75
FOLH1 AND EPHA3 AND NOT-SLC4A8	1	1	1
FOLH1 AND CDH10 AND NOT-LGR5	1	1	1
FOLH1 AND ATP7A AND NOT-ITGB6	0.888889	0.8	1
FOLH1 AND NOT-GRIN2C AND MOK	0.888889	0.8	1
FOLH1 AND STEAP4 AND NOT-CD180	0.857143	1	0.75
FOLH1 AND ATP7A AND NOT-ITGB6	0.888889	0.8	1
FOLH1 AND STEAP4 AND NOT-MUC1	0.857143	1	0.75
FOLH1 AND STEAP4 AND NOT-MUC16	0.857143	1	0.75
FOLH1 AND NOT-GRIN2C AND TNC	0.888889	0.8	1
FOLH1 AND PTGIS AND NOT-DPEP1	0.857143	1	0.75
STEAP2 AND NOT-GRIN2C AND FOLH1	0.888889	0.8	1
STEAP2 AND NOT-GRIN2C AND FOLH1	0.888889	0.8	1
FOLH1 AND PTGIS AND NOT-PMEL	0.857143	1	0.75
FOLH1 AND STEAP4 AND NOT-L1CAM	0.857143	1	0.75
FOLH1 AND STEAP4 AND NOT-CLDN11	0.857143	1	0.75
FOLH1 AND STEAP4 AND NOT-SLAMF7	0.857143	1	0.75
FOLH1 AND EPHA3 AND NOT-SLC4A8	1	1	1
FOLH1 AND NOT-DPEP1 AND FLVCR1	0.857143	1	0.75
FOLH1 AND STEAP4 AND NOT-EPHA3	0.857143	1	0.75
FOLH1 AND CDH10 AND NOT-LGR5	1	1	1
FOLH1 AND SELP AND NOT-PMEL	0.857143	1	0.75
FOLH1 AND STEAP4 AND NOT-CD33	0.857143	1	0.75
FOLH1 AND SELP AND NOT-ITGB3	0.857143	1	0.75
FOLH1 AND ROR1 AND NOT-DYSF	0.888889	0.8	1
FOLH1 AND NOT-CD36 AND NOT-LGR5	0.888889	0.8	1
FOLH1 AND NOT-GRIN2C AND IL2RA	0.888889	0.8	1
FOLH1 AND CLDN8 AND NOT-ATP6V0A4	1	1	1
FOLH1 AND CLDN8 AND NOT-KCNC2	0.857143	1	0.75
FOLH1 AND NOT-MET AND NOT-SLC4A8	0.888889	0.8	1
FOLH1 AND CLDN8 AND NOT-SLC39A8	0.857143	1	0.75
FOLH1 AND ROR1 AND NOT-STAB1	0.888889	0.8	1
FOLH1 AND GGTLC1 AND NOT-ITGB6	1	1	1
FOLH1 AND EPCAM AND PPAPDC1B	0.857143	1	0.75
FOLH1 AND EPHA3 AND NOT-TNFRSF10B	0.8	0.666667	1
FOLH1 AND NOT-DPEP1 AND CLDN8	0.857143	1	0.75
FOLH1 AND EPHA3 AND NOT-B4GALNT1	0.857143	1	0.75
FOLH1 AND NOT-EDNRB AND CLDN8	0.857143	1	0.75
FOLH1 AND EPCAM AND IL20RA	0.857143	1	0.75
FOLH1 AND EPHA3 AND EPCAM	0.857143	1	0.75
FOLH1 AND EPHA3 AND NOT-EDNRB	0.857143	1	0.75
FOLH1 AND EPHA3 AND NOT-ITGB3	0.857143	1	0.75
FOLH1 AND NOT-EDNRB AND IL20RA	0.857143	1	0.75
FOLH1 AND CLDN8 AND NOT-PROM1	0.857143	1	0.75
FOLH1 AND EPHA3 AND NOT-LGR5	0.888889	0.8	1
FOLH1 AND EPHA3 AND NOT-LGR5	0.888889	0.8	1
FOLH1 AND EPHA3 AND NOT-NCAM1	0.857143	1	0.75
FOLH1 AND EPHA3 AND NOT-NCAM1	0.857143	1	0.75
FOLH1 AND EPCAM AND CLDN8	0.857143	1	0.75
FOLH1 AND EPCAM AND NOT-GUCY2C	0.857143	1	0.75
FOLH1 AND EPHA3 AND NOT-IL3RA	0.857143	1	0.75
FOLH1 AND CLDN8 AND NOT-TNFRSF17	0.888889	0.8	1
FOLH1 AND EPHA3 AND NOT-CD33	0.888889	0.8	1

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
ITGAV AND CD19 AND ROS1	0.842105	0.875	0.811594
NOT-CEACAM6 AND CNTNAP2 AND GPNMB	0.827586	0.789474	0.869565
ITGAV AND CD19 AND CLDN16	0.84058	0.84058	0.84058
ITGAV AND CD19 AND GRM1	0.834532	0.828571	0.84058
ITGAV AND CD19 AND SLC12A3	0.846715	0.852941	0.84058
ITGAV AND CD19 AND ABCG5	0.859259	0.878788	0.84058
ITGAV AND CD19 AND SLC22A2	0.850746	0.876923	0.826087
ITGAV AND CD19 AND GUCY2D	0.814815	0.833333	0.797101
ITGAV AND CD19 AND NOT-SLC10A2	0.819672	0.943396	0.724638
COMPLEX-ITGAV/TGFBI/ULBP2	0.842857	0.830986	0.855072
ITGAV AND ULBP2 AND NOT-ZACN	0.909091	0.878378	0.942029
NOT-CLDN4 AND SLC31A1 AND NOT-P2RX5	0.859155	0.835616	0.884058
NOT-CLDN4 AND SLC31A1 AND NOT-CD19	0.857143	0.807692	0.913043
ITGAV AND SSTR1 AND NOT-SLC22A5	0.916667	0.88	0.956522
ITGAV AND ULBP2 AND NOT-CATSPER3	0.942029	0.942029	0.942029
ITGAV AND SSTR1 AND NOT-CATSPER3	0.956522	0.956522	0.956522
ITGAV AND CD19 AND LGR5	0.833333	0.873016	0.797101
ITGAV AND ULBP2 AND SSTR1	0.902778	0.866667	0.942029
Leiomyosarcoma			
BVES AND NOT-ATP7B AND KIAA0754	0.857143	0.8	0.923077
DDR2 AND NOT-TM7SF2 AND NOT-TNFRSF13B	0.869565	1	0.769231
DDR2 AND NOT-TM7SF2 AND NOT-COL17A1	0.869565	1	0.769231
NOT-MAL AND BVES AND KIAA0754	0.96	1	0.923077
DDR2 AND NOT-TM7SF2 AND NOT-KIAA1919	0.869565	1	0.769231
DDR2 AND NOT-TM7SF2 AND NOT-TNFSF14	0.869565	1	0.769231
DDR2 AND NOT-TM7SF2 AND NOT-FPR2	0.869565	1	0.769231
DDR2 AND NOT-TM7SF2 AND NOT-CFTR	0.869565	1	0.769231
DDR2 AND NOT-SLC5A1 AND TAS2R4	0.869565	1	0.769231
DDR2 AND NOT-TM7SF2 AND NOT-GJB6	0.818182	1	0.692308
DDR2 AND NOT-SLC5A1 AND GPR84	0.818182	1	0.692308
BVES AND NOT-CHL1 AND KIAA0754	0.814815	0.785714	0.846154
NOT-TSPAN8 AND BVES AND KIAA0754	0.814815	0.785714	0.846154
NOT-F11R AND ATP2A2 AND NOT-NLGN3	0.923077	0.923077	0.923077
BVES AND NOT-SLC5A1 AND KIAA0754	0.857143	0.8	0.923077
DDR2 AND NOT-F11R AND NOT-ROM1	0.833333	0.909091	0.769231
DDR2 AND NOT-TM7SF2 AND CD244	0.818182	1	0.692308
DDR2 AND NOT-TM7SF2 AND KCNJ1	0.869565	1	0.769231
DDR2 AND NOT-TM7SF2 AND CHRM3	0.869565	1	0.769231
NOT-F11R AND ATP2A2 AND NOT-GJB1	0.923077	0.923077	0.923077
DDR2 AND NOT-TM7SF2 AND NOT-SLC16A4	0.869565	1	0.769231
DDR2 AND NOT-GDPD2 AND KCNG3	0.869565	1	0.769231
DDR2 AND NOT-F11R AND NOT-SYP	0.833333	0.909091	0.769231
DDR2 AND NOT-F11R AND NOT-TM7SF2	0.869565	1	0.769231
DDR2 AND NOT-F11R AND TMEM63B	0.8	0.833333	0.769231
DDR2 AND NOT-F11R AND TMIGD2	0.8	0.833333	0.769231
DDR2 AND NOT-F11R AND ANO7	0.8	0.833333	0.769231
BVES AND NOT-MFAP3L AND KIAA0754	0.8	0.833333	0.769231
DDR2 AND NOT-F11R AND NOT-EMCN	0.8	0.833333	0.769231
DDR2 AND NOT-F11R AND SCNN1A	0.8	0.833333	0.769231
VMP1 AND NOT-F11R AND SLC5A7	0.833333	0.909091	0.769231
DDR2 AND NOT-TM7SF2 AND NOT-SPN	0.869565	1	0.769231
DDR2 AND NOT-TM7SF2 AND CCR6	0.818182	1	0.692308
DDR2 AND NOT-TM7SF2 AND CXCR1	0.869565	1	0.769231
DDR2 AND NOT-TM7SF2 AND PIEZO1	0.869565	1	0.769231
DDR2 AND NOT-TM7SF2 AND GPR52	0.869565	1	0.769231
DDR2 AND NOT-TM7SF2 AND NOT-GPR160	0.869565	1	0.769231
DDR2 AND NOT-TM7SF2 AND ATP12A	0.869565	1	0.769231
DDR2 AND NOT-GDPD2 AND CCR6	0.818182	1	0.692308
DDR2 AND NOT-F11R AND CLSTN1	0.869565	1	0.769231
DDR2 AND NOT-TM7SF2 AND NOT-HAVCR2	0.818182	1	0.692308
DDR2 AND NOT-F11R AND NOT-GJB1	0.833333	0.909091	0.769231
PCDH18 AND NOT-SLC5A1 AND TAS2R4	0.818182	1	0.692308
DDR2 AND NOT-TM7SF2 AND NOT-MGAM	0.869565	1	0.769231
DDR2 AND NOT-TM7SF2 AND NOT-ICAM4	0.869565	1	0.769231
DDR2 AND NOT-TM7SF2 AND ADRB3	0.818182	1	0.692308
DDR2 AND NOT-F11R AND NOT-AQP7	0.869565	1	0.769231
DDR2 AND NOT-F11R AND NOT-NLGN3	0.833333	0.909091	0.769231
VMP1 AND NOT-F11R AND SYT9	0.869565	1	0.769231
VMP1 AND NOT-F11R AND KCNJ16	0.833333	0.909091	0.769231
SLC4A7 AND BVES AND NOT-RHBDL2	0.869565	1	0.769231
VMP1 AND NOT-F11R AND NOT-EMCN	0.833333	0.909091	0.769231
DDR2 AND NOT-TM7SF2 AND C2orf83	0.869565	1	0.769231
DDR2 AND NOT-TM7SF2 AND NOT-EVI2B	0.869565	1	0.769231
DDR2 AND NOT-F11R AND NOT-SLC24A4	0.869565	1	0.769231
VMP1 AND NOT-F11R AND SLC5A12	0.869565	1	0.769231
VMP1 AND NOT-F11R AND IL23R	0.869565	1	0.769231
BVES AND NOT-SLC27A5 AND KIAA0754	0.8	0.705882	0.923077

Antigen Pair Logic	F1	Precision	Recall
FOLH1 AND ROR1 AND NOT-LGR5	1	1	1
FOLH1 AND EPCAM AND NOT-SST	0.857143	1	0.75
FOLH1 AND EPCAM AND NOT-CEACAM5	0.857143	1	0.75
FOLH1 AND EPCAM AND CLDN11	0.857143	1	0.75
FOLH1 AND NOT-MET AND CLDN8	0.8	0.666667	1
COMPLEX-FOLH1/STEAP2/PSCA	0.8	0.666667	1
FOLH1 AND EPHA3 AND NOT-MUC1	0.8	0.666667	1
FOLH1 AND EPHA3 AND NOT-TNC	0.8	0.666667	1
FOLH1 AND EPHA3 AND NOT-KDR	0.8	0.666667	1
FOLH1 AND EPHA3 AND NOT-ST8SIA1	0.8	0.666667	1
FOLH1 AND EPHA3 AND NOT-AXL	0.8	0.666667	1
FOLH1 AND EPHA3 AND NOT-TNFRSF17	0.8	0.666667	1
FOLH1 AND EPHA3 AND NOT-L1CAM	0.8	0.666667	1
FOLH1 AND EPHA3 AND NOT-L1CAM	0.8	0.666667	1
FOLH1 AND EPHA3 AND NOT-CEACAM6	0.8	0.666667	1
FOLH1 AND EPHA3 AND NOT-DKK1	0.8	0.666667	1
FOLH1 AND EPHA3 AND NOT-MUC1	0.8	0.666667	1
FOLH1 AND EPHA3 AND NOT-CSPG4	0.8	0.666667	1
COMPLEX-FOLH1/STEAP2/ENPP3	0.8	0.666667	1
FOLH1 AND EPHA3 AND NOT-AXL	0.8	0.666667	1
FOLH1 AND EPHA3 AND NOT-CSPG4	0.8	0.666667	1
STEAP2 AND FOLH1 AND NOT-DPEP1	0.857143	1	0.75
FOLH1 AND CLDN8 AND NOT-CD33	0.888889	0.8	1
FOLH1 AND CLDN8 AND NOT-CD33	0.888889	0.8	1
FOLH1 AND CLDN8 AND NOT-CEACAM6	0.857143	1	0.75
FOLH1 AND NOT-ALDH1A1 AND MUC1	0.857143	1	0.75
STEAP2 AND NOT-KDR AND FOLH1	0.8	0.666667	1
FOLH1 AND NOT-MET AND IL20RA	0.857143	1	0.75
FOLH1 AND NOT-PROM1 AND MUC1	0.857143	1	0.75
FOLH1 AND ROR1 AND NOT-CD33	1	1	1
FOLH1 AND ROR1 AND NOT-SLAMF7	0.888889	0.8	1
FOLH1 AND CLDN8 AND NOT-FCRL5	0.8	0.666667	1
FOLH1 AND CLDN8 AND NOT-CD22	0.857143	1	0.75
FOLH1 AND ROR1 AND NOT-ITGB6	0.857143	1	0.75
FOLH1 AND EPCAM AND NOT-ITGB6	0.857143	1	0.75
FOLH1 AND NOT-ALDH1A1 AND CLDN8	0.857143	1	0.75
FOLH1 AND ROR1 AND NOT-NCAM1	0.857143	1	0.75
FOLH1 AND ROR1 AND NOT-FOLR2	0.888889	0.8	1
FOLH1 AND ROR1 AND NOT-SST	0.888889	0.8	1
FOLH1 AND ROR1 AND NOT-CSPG4	0.888889	0.8	1
FOLH1 AND ROR1 AND NOT-GUCY2C	0.888889	0.8	1
FOLH1 AND NOT-MET AND NOT-LGR5	0.888889	0.8	1
FOLH1 AND IL20RA AND NOT-CD33	0.888889	0.8	1
FOLH1 AND CLDN8 AND NOT-SLAMF7	0.888889	0.8	1
FOLH1 AND NOT-DPEP1 AND ERBB2	0.857143	1	0.75
FOLH1 AND CLDN8 AND NOT-SLAMF7	0.888889	0.8	1
FOLH1 AND IL20RA AND NOT-TNFRSF17	0.8	0.666667	1
FOLH1 AND CLDN8 AND NOT-CD180	0.888889	0.8	1
FOLH1 AND NOT-CEACAM6 AND MUC1	0.857143	1	0.75
FOLH1 AND IL20RA AND NOT-CD33	0.888889	0.8	1
FOLH1 AND CLDN8 AND NOT-CD180	0.888889	0.8	1
FOLH1 AND NOT-LGR5 AND FAP	0.857143	1	0.75
FOLH1 AND EPCAM AND IL13RA2	0.857143	1	0.75
FOLH1 AND ROR1 AND NOT-AXL	0.888889	0.8	1
STEAP2 AND CLDN8 AND NOT-FOLH1	0.857143	1	0.75
FOLH1 AND ROR1 AND NOT-CEACAM5	0.888889	0.8	1
FOLH1 AND ROR1 AND NOT-CD160	0.888889	0.8	1
STEAP2 AND CLDN8 AND NOT-FOLH1	0.857143	1	0.75
TRPM8 AND NOT-SLC39A2 AND NOT-TNC	1	1	1
TRPM8 AND NOT-SLC39A2 AND NOT-TNC	1	1	1
TRPM8 AND NOT-CSPG4 AND NOT-SLC39A2	1	1	1
TRPM8 AND BIRC5 AND NOT-FMNL1	1	1	1
TRPM8 AND BIRC5 AND NOT-FMNL1	1	1	1
COMPLEX-IGF1R/TRPM8/SLC16A5	1	1	1
STEAP2 AND EPHA10 AND NOT-SLC39A2	1	1	1
TRPM8 AND NOT-IL17RA AND TRPM4	1	1	1
STEAP2 AND SLC43A1 AND NOT-IL17RA	1	1	1
STEAP2 AND SLCO1A2 AND SLC43A1	1	1	1
TRPM8 AND NOT-GYPC AND VCAM1	1	1	1
TRPM8 AND NOT-GYPC AND MUC1	1	1	1
TRPM8 AND NOT-SYT4 AND CLDN8	1	1	1
STEAP2 AND NOT-GYPC AND PTGER4	1	1	1
STEAP2 AND GPR19 AND NOT-SLC39A2	1	1	1
TRPM8 AND NOT-GYPC AND MUC1	1	1	1
STEAP2 AND NOT-GYPC AND F2R	1	1	1
STEAP2 AND GPR19 AND NOT-SLC39A2	1	1	1
TRPM8 AND NOT-IL17RA AND CLDN8	1	1	1

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
VMP1 AND NOT-F11R AND EQTN	0.869565	1	0.769231
VMP1 AND NOT-F11R AND KCNH7	0.869565	1	0.769231
DDR2 AND NOT-F11R AND NOT-TNFSF9	0.833333	0.909091	0.769231
VMP1 AND NOT-F11R AND GABRR3	0.869565	1	0.769231
VMP1 AND NOT-F11R AND RRH	0.869565	1	0.769231
VMP1 AND NOT-F11R AND AKAP6	0.869565	1	0.769231
NOT-F11R AND ATP2A2 AND NOT-SYP	0.96	1	0.923077
DDR2 AND NOT-TM7SF2 AND NOT-OR51E1	0.869565	1	0.769231
DDR2 AND NOT-TM7SF2 AND TAS2R4	0.869565	1	0.769231
DDR2 AND NOT-TM7SF2 AND NOT-IGSF6	0.869565	1	0.769231
DDR2 AND NOT-TM7SF2 AND CLEC4A	0.869565	1	0.769231
DDR2 AND NOT-TM7SF2 AND NOT-SEMA4G	0.869565	1	0.769231
DDR2 AND NOT-F11R AND NOT-ASGR1	0.833333	0.909091	0.769231
VMP1 AND BVES AND NOT-RHBDL2	0.88	0.916667	0.846154
DDR2 AND NOT-F11R AND NOT-BEST1	0.833333	0.909091	0.769231
DDR2 AND NOT-F11R AND P2RY1	0.8	0.833333	0.769231
BVES AND NOT-GDPD2 AND KIAA0754	0.888889	0.857143	0.923077
VMP1 AND NOT-F11R AND TMIGD2	0.833333	0.909091	0.769231
VMP1 AND NOT-F11R AND PCDHGB6	0.869565	1	0.769231
VMP1 AND NOT-F11R AND C2orf83	0.833333	0.909091	0.769231
VMP1 AND NOT-F11R AND IL20RB	0.869565	1	0.769231
BVES AND HLA-B AND NOT-RHBDL2	0.869565	1	0.769231
VMP1 AND NOT-F11R AND GPR3	0.869565	1	0.769231
DDR2 AND NOT-GDPD2 AND GJB6	0.869565	1	0.769231
DDR2 AND NOT-GDPD2 AND TAS2R4	0.869565	1	0.769231
DDR2 AND NOT-GDPD2 AND GPR82	0.869565	1	0.769231
DDR2 AND NOT-GDPD2 AND OR2B2	0.869565	1	0.769231
NOT-F11R AND BRCA1 AND NOT-ICAM4	0.818182	1	0.692308
NOT-CRB3 AND BVES AND KIAA0754	0.846154	0.846154	0.846154
BVES AND KIAA0754 AND NOT-DSC3	0.88	0.916667	0.846154
DDR2 AND NOT-F11R AND NOT-MTUS1	0.833333	0.909091	0.769231
BVES AND KIAA0754 AND NOT-SLC7A10	0.8125	0.684211	1
VMP1 AND NOT-F11R AND TRDN	0.869565	1	0.769231
VMP1 AND NOT-F11R AND TAS2R5	0.869565	1	0.769231
VMP1 AND NOT-F11R AND OR2K2	0.869565	1	0.769231
VMP1 AND NOT-F11R AND SLC5A1	0.869565	1	0.769231
VMP1 AND NOT-F11R AND GJB5	0.869565	1	0.769231
VMP1 AND NOT-F11R AND F2RL2	0.833333	0.909091	0.769231
VMP1 AND NOT-F11R AND SLC6A2	0.833333	0.909091	0.769231
DDR2 AND NOT-TM7SF2 AND NOT-MLC1	0.869565	1	0.769231
DDR2 AND NOT-TM7SF2 AND NOT-SLC5A11	0.869565	1	0.769231
DDR2 AND NOT-TM7SF2 AND GJD2	0.869565	1	0.769231
DDR2 AND NOT-F11R AND NOT-SLC6A11	0.818182	1	0.692308
DDR2 AND NOT-TM7SF2 AND NOT-KIAA0319	0.869565	1	0.769231
DDR2 AND NOT-TM7SF2 AND NOT-LRRN4	0.869565	1	0.769231
DDR2 AND NOT-TM7SF2 AND CACNG1	0.869565	1	0.769231
ATP8B2 AND NOT-PIGR AND EMP2	0.869565	1	0.769231
ATP8B2 AND NOT-PIGR AND CDH5	0.869565	1	0.769231
ATP8B2 AND NOT-PIGR AND PMP22	0.833333	0.909091	0.769231
ATP8B2 AND NOT-PIGR AND LRP11	0.869565	1	0.769231
DDR2 AND NOT-TM7SF2 AND ATP13A5	0.869565	1	0.769231
DDR2 AND NOT-F11R AND NOT-MLC1	0.833333	0.909091	0.769231
DDR2 AND NOT-F11R AND NOT-GRID2	0.833333	0.909091	0.769231
DDR2 AND NOT-F11R AND NOT-SLC5A11	0.833333	0.909091	0.769231
DDR2 AND NOT-F11R AND NOT-KIAA0319	0.833333	0.909091	0.769231
DDR2 AND NOT-F11R AND ABCA12	0.8	0.833333	0.769231
DDR2 AND NOT-F11R AND CYP4A11	0.8	0.833333	0.769231
VMP1 AND NOT-F11R AND NOT-SELP	0.833333	0.909091	0.769231
NOT-F11R AND VANGL1 AND NOT-GDPD2	1	1	1
NOT-F11R AND VANGL1 AND NOT-ICAM4	1	1	1
DDR2 AND NOT-F11R AND NOT-GABRB1	0.833333	0.909091	0.769231
NOT-F11R AND VANGL1 AND NOT-ICAM4	1	1	1
NOT-F11R AND VANGL1 AND NOT-TNFRSF13B	1	1	1
NOT-F11R AND ATP2A2 AND NOT-SLC5A11	0.923077	0.923077	0.923077
DDR2 AND NOT-TM7SF2 AND SLC22A18	0.869565	1	0.769231
DDR2 AND NOT-F11R AND NOT-SLCO1A2	0.869565	1	0.769231
DDR2 AND NOT-F11R AND NOT-MAG	0.833333	0.909091	0.769231
DDR2 AND NOT-F11R AND NOT-KCNK10	0.833333	0.909091	0.769231
NOT-F11R AND VANGL1 AND NOT-ABCB1	0.962963	0.928571	1
TGFBI AND NOT-F11R AND SLC5A7	0.818182	1	0.692308
VMP1 AND NOT-F11R AND SLC39A12	0.833333	0.909091	0.769231
DDR2 AND NOT-F11R AND NOT-GABRA4	0.833333	0.909091	0.769231
VMP1 AND NOT-F11R AND PCDHA9	0.833333	0.909091	0.769231
NOT-F11R AND VANGL1 AND NOT-DIO3	1	1	1
NOT-F11R AND VANGL1 AND NOT-ATP7B	1	1	1
NOT-F11R AND VANGL1 AND NOT-PCDHA10	1	1	1
NOT-F11R AND VANGL1 AND NOT-PRR7	0.923077	0.923077	0.923077

Antigen Pair Logic	F1	Precision	Recall
TRPM8 AND NOT-TNC AND F2R	1	1	1
TRPM8 AND NOT-DYSF AND CLDN8	1	1	1
TRPM8 AND NOT-SLC39A2 AND BIRC5	1	1	1
TRPM8 AND NOT-SLC39A2 AND WT1	1	1	1
TRPM8 AND NOT-GUCY2D AND TRPM4	1	1	1
TRPM8 AND NOT-DYSF AND TRPM4	1	1	1
TRPM8 AND NOT-SLC5A2 AND TRPM4	1	1	1
STEAP2 AND EPHA10 AND NOT-FMNL1	1	1	1
TRPM8 AND NOT-GYPC AND TRPM4	1	1	1
STEAP2 AND SLC43A1 AND NOT-GUCY2D	1	1	1
STEAP2 AND SLC43A1 AND NOT-DYSF	1	1	1
STEAP2 AND NOT-GYPC AND SLC43A1	1	1	1
STEAP2 AND SLC43A1 AND NOT-EBP	1	1	1
STEAP2 AND NOT-SLC4A8 AND NOT-ANPEP	1	1	1
TRPM8 AND NOT-SLC39A2 AND ABCB5	1	1	1
TRPM8 AND NOT-SLC39A2 AND BIRC5	1	1	1
STEAP2 AND NOT-SYT4 AND SLC43A1	1	1	1
STEAP2 AND SLC43A1 AND NOT-SLC5A2	1	1	1
TRPM8 AND NOT-TNC AND SLC43A1	1	1	1
TRPM8 AND NOT-SYT4 AND TRPM4	1	1	1
STEAP2 AND SLC43A1 AND NOT-CHRNA2	1	1	1
STEAP2 AND GPR19 AND SLC43A1	1	1	1
TRPM8 AND NOT-CHRNA2 AND TRPM4	1	1	1
STEAP2 AND NOT-SLC39A2 AND NOT-SLC31A1	1	1	1
STEAP2 AND NOT-SYT4 AND NOT-FMNL1	1	1	1
STEAP2 AND GPR19 AND NOT-FMNL1	1	1	1
STEAP2 AND GPR19 AND F2R	1	1	1
STEAP2 AND NOT-DYSF AND NOT-SLC39A2	1	1	1
COMPLEX-STEAP2/SCN2A/SLC16A5	1	1	1
TRPM8 AND NOT-DYSF AND VCAM1	1	1	1
STEAP2 AND NOT-SLC39A2 AND RGSL1	1	1	1
STEAP2 AND NOT-SLC39A2 AND NOT-EBP	1	1	1
TRPM8 AND NOT-SYT4 AND ABCA5	1	1	1
TRPM8 AND NOT-EPHA2 AND CLDN8	1	1	1
STEAP2 AND NOT-DYSF AND PTGER4	1	1	1
STEAP2 AND NOT-SLC39A2 AND TMPRSS11D	1	1	1
STEAP2 AND NOT-GYPC AND STEAP4	1	1	1
STEAP2 AND NOT-GYPC AND CD163	1	1	1
TRPM8 AND NOT-TNC AND PTGER4	1	1	1
STEAP2 AND GPR19 AND NOT-FMNL1	1	1	1
STEAP2 AND NOT-SLC39A2 AND UMODL1	1	1	1
STEAP2 AND NOT-SYT4 AND NOT-FMNL1	1	1	1
STEAP2 AND NOT-SLC39A2 AND NOT-EPHA2	1	1	1
COMPLEX-STEAP2/TMPRSS11E/SLC16A5	1	1	1
TRPM8 AND NOT-SLC5A2 AND CLDN8	1	1	1
STEAP2 AND NOT-SLC39A2 AND PCDHA9	1	1	1
STEAP2 AND SLC2A2 AND NOT-SLC39A2	1	1	1
COMPLEX-GABRB2/STEAP2/SLC16A5	1	1	1
TRPM8 AND NOT-SLC39A2 AND P2RX5	1	1	1
TRPM8 AND NOT-GUCY2D AND CLDN8	1	1	1
TRPM8 AND NOT-DYSF AND ABCA5	1	1	1
TRPM8 AND NOT-CSPG4 AND PTGER4	1	1	1
STEAP2 AND NOT-SLC39A2 AND NOT-IL17RA	1	1	1
STEAP2 AND GPR19 AND PTGER4	1	1	1
STEAP2 AND NOT-SLC39A2 AND NOT-CHRNA2	1	1	1
STEAP2 AND NOT-SLC39A2 AND CACNG7	1	1	1
STEAP2 AND NOT-SLC39A2 AND NOT-IL17RA	1	1	1
STEAP2 AND NOT-SLC39A2 AND CACNG7	1	1	1
STEAP2 AND NOT-SLC39A2 AND NOT-CHRNA2	1	1	1
TRPM8 AND NOT-DYSF AND MUC1	1	1	1
TRPM8 AND NOT-SLC39A2 AND FAP	1	1	1
TRPM8 AND NOT-CSPG4 AND F2R	1	1	1
STEAP2 AND SLC2A2 AND NOT-SLC39A2	1	1	1
STEAP2 AND NOT-SLC39A2 AND NOT-GUCY2D	1	1	1
STEAP2 AND NOT-SLC39A2 AND TMPRSS11D	1	1	1
STEAP2 AND NOT-SLC39A2 AND NOT-GABRA1	1	1	1
STEAP2 AND NOT-SLC39A2 AND NOT-GUCY2D	1	1	1
STEAP2 AND NOT-SLC39A2 AND PCDHA9	1	1	1
STEAP2 AND NOT-SLC39A2 AND MEGF11	1	1	1
TRPM8 AND NOT-FMNL1 AND P2RX5	1	1	1
STEAP2 AND NOT-SLC39A2 AND NPHS2	1	1	1
STEAP2 AND NOT-SLC39A2 AND NPHS1	1	1	1
TRPM8 AND NOT-SLC39A2 AND NOT-HSPA5	1	1	1
STEAP2 AND NOT-SLC39A2 AND NMBR	1	1	1
STEAP2 AND SLC2A2 AND NOT-FMNL1	1	1	1
TRPM8 AND NOT-CHRNA2 AND CLDN8	1	1	1
STEAP2 AND NOT-SLC39A2 AND MEGF11	1	1	1

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
DDR2 AND NOT-GDPD2 AND SV2C	0.869565	1	0.769231
ATP8B2 AND NOT-EREG AND NOT-ICAM4	0.8	0.833333	0.769231
DDR2 AND NOT-TM7SF2 AND NOT-SLC17A3	0.818182	1	0.692308
NOT-F11R AND VANGL1 AND NOT-ACE2	0.88	0.916667	0.846154
DDR2 AND NOT-F11R AND NOT-ADCY8	0.833333	0.909091	0.769231
NOT-F11R AND VANGL1 AND NOT-CRHR1	0.962963	0.928571	1
VMP1 AND NOT-F11R AND OPRK1	0.869565	1	0.769231
DDR2 AND NOT-F11R AND NOT-CHRNG	0.833333	0.909091	0.769231
VMP1 AND NOT-F11R AND ADCY8	0.869565	1	0.769231
VMP1 AND NOT-F11R AND AJAP1	0.869565	1	0.769231
DDR2 AND NOT-F11R AND NOT-HCN2	0.833333	0.909091	0.769231
NOT-F11R AND VANGL1 AND NOT-SLC8A2	1	1	1
VMP1 AND NOT-F11R AND SLC13A5	0.833333	0.909091	0.769231
ATP8B2 AND NOT-GDPD2 AND NOT-GPR160	0.833333	0.909091	0.769231
VMP1 AND NOT-F11R AND GRM1	0.869565	1	0.769231
ATP8B2 AND NOT-TNFRSF13B AND NOT-EREG	0.833333	0.909091	0.769231
VMP1 AND NOT-F11R AND LRRTM4	0.869565	1	0.769231
NOT-F11R AND ATP2A2 AND NOT-KIAA0319	0.923077	0.923077	0.923077
VMP1 AND BVES AND LRRC52	0.88	0.916667	0.846154
DDR2 AND NOT-EPCAM AND TAS2R4	0.833333	0.909091	0.769231
DDR2 AND NOT-TM7SF2 AND NOT-FCRL2	0.869565	1	0.769231
DDR2 AND NOT-EPCAM AND C2orf83	0.8	0.833333	0.769231
DDR2 AND NOT-EPCAM AND SLC5A7	0.8	0.833333	0.769231
DDR2 AND NOT-F11R AND SSTR5	0.8	0.833333	0.769231
DDR2 AND NOT-EPCAM AND IL23R	0.8	0.833333	0.769231
SLC4A7 AND NOT-ITGB6 AND CDH5	0.8	0.833333	0.769231
DDR2 AND NOT-F11R AND NOT-ERBB4	0.833333	0.909091	0.769231
DDR2 AND NOT-F11R AND NOT-CD22	0.833333	0.909091	0.769231
BIRC5 AND NOT-F11R AND CDH5	0.818182	1	0.692308
SLC4A7 AND NOT-EPCAM AND ZNRF3	0.833333	0.909091	0.769231
NOT-F11R AND ATP2A2 AND NOT-CD22	0.96	1	0.923077
BIRC5 AND NOT-SMAGP AND PMP22	0.818182	1	0.692308
VMP1 AND NOT-F11R AND FCRL1	0.833333	0.909091	0.769231
VMP1 AND NOT-F11R AND LGR5	0.869565	1	0.769231
VMP1 AND NOT-F11R AND RAET1E	0.833333	0.909091	0.769231
BVES AND NOT-ERBB4 AND KIAA0754	0.866667	0.764706	1
BIRC5 AND NOT-F11R AND EMCN	0.818182	1	0.692308
NOT-CEACAM6 AND BVES AND MYOF	0.827586	0.75	0.923077
VMP1 AND NOT-F11R AND BMPR1B	0.833333	0.909091	0.769231
VMP1 AND NOT-F11R AND NOT-MSLN	0.833333	0.909091	0.769231
VMP1 AND NOT-F11R AND NOT-CD79B	0.833333	0.909091	0.769231
BIRC5 AND NOT-F11R AND NOTCH4	0.818182	1	0.692308
NOT-ERBB3 AND GPR17 AND BVES	0.818182	1	0.692308
NOT-F11R AND SLC10A3 AND NOT-CLDN6	0.916667	1	0.846154
NOT-EPCAM AND OXGR1 AND NOT-IL27RA	0.857143	0.8	0.923077
BIRC5 AND NOT-GDPD2 AND AQP4	0.818182	1	0.692308
BIRC5 AND NOT-F11R AND LIFR	0.818182	1	0.692308
BIRC5 AND NOT-GDPD2 AND CADM2	0.818182	1	0.692308
BIRC5 AND NOT-F11R AND PMP22	0.818182	1	0.692308
BIRC5 AND NOT-F11R AND DSC2	0.818182	1	0.692308
BIRC5 AND NOT-F11R AND SLC2A13	0.818182	1	0.692308
BIRC5 AND NOT-F11R AND PODXL	0.818182	1	0.692308
VMP1 AND NOT-F11R AND SSTR5	0.869565	1	0.769231
VMP1 AND NOT-F11R AND NOT-CD22	0.833333	0.909091	0.769231
SLC4A7 AND NOT-EPCAM AND CDH5	0.833333	0.909091	0.769231
SLC4A7 AND BVES AND NOT-CLDN2	0.869565	1	0.769231
DDR2 AND NOT-GDPD2 AND FCRL2	0.869565	1	0.769231
VMP1 AND NOT-F11R AND NOT-CLDN6	0.833333	0.909091	0.769231
NOT-EPCAM AND BRCA1 AND FZD7	0.818182	1	0.692308
NOT-F11R AND ATP2A2 AND NOT-ERBB4	0.96	1	0.923077
NOT-F11R AND STX2 AND NOT-CD22	0.869565	1	0.769231
NOT-EPCAM AND OXGR1 AND NOT-MAL	0.923077	0.923077	0.923077
NOT-EPCAM AND ATP2A2 AND NOT-CCR6	0.923077	0.923077	0.923077
VMP1 AND NOT-F11R AND NOT-TNFRSF17	0.833333	0.909091	0.769231
VMP1 AND NOT-F11R AND NOT-FCRL2	0.833333	0.909091	0.769231
NOT-ERBB3 AND BVES AND NOT-PAQR9	0.846154	0.846154	0.846154
NOT-F11R AND SLC10A3 AND NOT-FCRL2	0.916667	1	0.846154
BIRC5 AND NOT-GDPD2 AND CDH5	0.818182	1	0.692308
BIRC5 AND NOT-MGST2 AND PMP22	0.818182	1	0.692308
BIRC5 AND NOT-SEMA4G AND PMP22	0.818182	1	0.692308
NOT-EPCAM AND BRCA1 AND OXGR1	0.818182	1	0.692308
SLC4A7 AND NOT-CCR6 AND NOT-EPCAM	0.869565	1	0.769231
BIRC5 AND NOT-F11R AND RTN4	0.818182	1	0.692308
BIRC5 AND NOT-F11R AND SLC38A2	0.818182	1	0.692308
BIRC5 AND NOT-F11R AND BCAP31	0.818182	1	0.692308
BIRC5 AND NOT-F11R AND NOT-SLAMF1	0.818182	1	0.692308
BIRC5 AND NOT-F11R AND CLSTN1	0.818182	1	0.692308

Antigen Pair Logic	F1	Precision	Recall
STEAP2 AND NOT-SLC39A2 AND NOT-GABRA1	1	1	1
STEAP2 AND SLC2A2 AND NOT-FMNL1	1	1	1
TRPM8 AND NOT-DYSF AND MUC1	1	1	1
TRPM8 AND NOT-DKK1 AND F2R	1	1	1
STEAP2 AND NOT-TNC AND NOT-SLC39A2	1	1	1
STEAP2 AND SLCO1A2 AND CLDN8	1	1	1
STEAP2 AND BIRC5 AND NOT-FMNL1	1	1	1
STEAP2 AND SLCO1A2 AND CLDN8	1	1	1
STEAP2 AND BIRC5 AND NOT-FMNL1	1	1	1
STEAP2 AND NOT-TNC AND NOT-SLC39A2	1	1	1
STEAP2 AND SLCO1A2 AND TRPM4	1	1	1
STEAP2 AND NOT-SYT4 AND CLDN8	1	1	1
STEAP2 AND CLDN8 AND NOT-IL17RA	1	1	1
TRPM8 AND NOT-DKK1 AND CLDN8	1	1	1
STEAP2 AND NOT-GYPC AND MUC1	1	1	1
STEAP2 AND BIRC5 AND NOT-SLC39A2	1	1	1
STEAP2 AND NOT-SLC39A2 AND NOT-CSPG4	1	1	1
STEAP2 AND NOT-GYPC AND MUC1	1	1	1
TRPM8 AND NOT-TNC AND CLDN8	1	1	1
STEAP2 AND NOT-SLC39A2 AND NOT-CSPG4	1	1	1
STEAP2 AND CLDN8 AND NOT-DYSF	1	1	1
STEAP2 AND NOT-GYPC AND VCAM1	1	1	1
STEAP2 AND CLDN8 AND NOT-DYSF	1	1	1
COMPLEX-IGF1R/EPCAM/TRPM8	1	1	1
STEAP2 AND CLDN8 AND NOT-IL17RA	1	1	1
STEAP2 AND NOT-SYT4 AND CLDN8	1	1	1
STEAP2 AND GPR19 AND CLDN8	1	1	1
STEAP2 AND NOT-TNC AND F2R	1	1	1
STEAP2 AND NOT-GYPC AND VCAM1	1	1	1
COMPLEX-STEAP2/IGF1R/SLC16A5	1	1	1
TRPM8 AND NOT-TNC AND TRPM4	1	1	1
STEAP2 AND CLDN8 AND NOT-CD36	1	1	1
STEAP2 AND NOT-SLC39A2 AND WT1	1	1	1
STEAP2 AND NOT-SLC39A2 AND ABCB5	1	1	1
STEAP2 AND SLCO1A2 AND RNF43	1	1	1
STEAP2 AND NOT-SYT4 AND TRPM4	1	1	1
STEAP2 AND EPHA10 AND MUC1	1	1	1
STEAP2 AND NOT-EMP3 AND VCAM1	1	1	1
STEAP2 AND EPHA10 AND VCAM1	1	1	1
STEAP2 AND SLCO1A2 AND ABCA5	1	1	1
STEAP2 AND CLDN8 AND NOT-PTGER4	1	1	1
STEAP2 AND NOT-EMP3 AND CLDN8	1	1	1
STEAP2 AND SLC43A1 AND NOT-TNC	1	1	1
STEAP2 AND CLDN8 AND NOT-EPHA2	1	1	1
STEAP2 AND CLDN8 AND MEGF11	1	1	1
STEAP2 AND GPR19 AND VCAM1	1	1	1
STEAP2 AND GPR19 AND ABCA5	1	1	1
STEAP2 AND NOT-GYPC AND CLDN8	1	1	1
TRPM8 AND NOT-TNC AND MUC1	1	1	1
TRPM8 AND NOT-CSPG4 AND VCAM1	1	1	1
STEAP2 AND CLDN8 AND NOT-CHRNA2	1	1	1
TRPM8 AND NOT-TNC AND ABCA5	1	1	1
STEAP2 AND CLDN8 AND NOT-SLC5A2	1	1	1
TRPM8 AND NOT-TNC AND VCAM1	1	1	1
STEAP2 AND CLDN8 AND NOT-GUCY2D	1	1	1
STEAP2 AND NOT-SYT4 AND ABCA5	1	1	1
STEAP2 AND NOT-SLC39A2 AND NOT-DKK1	1	1	1
STEAP2 AND NOT-DYSF AND VCAM1	1	1	1
STEAP2 AND CLDN8 AND NOT-EPHA2	1	1	1
STEAP2 AND NOT-SLC39A2 AND NOT-DKK1	1	1	1
STEAP2 AND NOT-DKK1 AND F2R	1	1	1
STEAP2 AND NOT-TNC AND PTGER4	1	1	1
STEAP2 AND NOT-GYPC AND CLDN8	1	1	1
STEAP2 AND GPR19 AND MUC1	1	1	1
TRPM8 AND NOT-CSPG4 AND CLDN8	1	1	1
STEAP2 AND NOT-SLC39A2 AND FAP	1	1	1
TRPM8 AND NOT-TNC AND MUC1	1	1	1
STEAP2 AND CLDN8 AND MEGF11	1	1	1
STEAP2 AND GPR19 AND MUC1	1	1	1
STEAP2 AND NOT-CSPG4 AND PTGER4	1	1	1
TRPM8 AND NOT-CSPG4 AND ABCA5	1	1	1
TRPM8 AND NOT-DKK1 AND ABCA5	1	1	1
TRPM8 AND NOT-CSPG4 AND MUC1	1	1	1
STEAP2 AND CLDN8 AND NOT-CHRNA2	1	1	1
STEAP2 AND NOT-DYSF AND ABCA5	1	1	1
STEAP2 AND NOT-CSPG4 AND F2R	1	1	1
STEAP2 AND NOT-GYPC AND ABCA5	1	1	1

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
BIRC5 AND NOT-F11R AND VAPA	0.818182	1	0.692308
VMP1 AND NOT-F11R AND NOT-FCRL5	0.833333	0.909091	0.769231
VMP1 AND NOT-F11R AND NOT-ERBB4	0.833333	0.909091	0.769231
NOT-F11R AND ATP2A2 AND NOT-CLDN6	0.96	1	0.923077
CD276 AND NOT-F11R AND NOT-ICAM4	0.818182	1	0.692308
DDR2 AND NOT-GDPD2 AND CR2	0.818182	1	0.692308
BIRC5 AND NOT-F11R AND AQP4	0.818182	1	0.692308
NOT-F11R AND CBX3 AND NOT-KCNA3	0.96	1	0.923077
BIRC5 AND NOT-MARVELD2 AND PMP22	0.818182	1	0.692308
NOT-F11R AND SLC10A3 AND NOT-CD22	0.916667	1	0.846154
BVES AND NOT-ITGB6 AND KIAA0754	0.857143	0.8	0.923077
VMP1 AND NOT-EPCAM AND ZNRF3	0.814815	0.785714	0.846154
NOT-F11R AND VANGL1 AND NOT-CLDN6	1	1	1
NOT-EPCAM AND VANGL1 AND NOT-ICAM4	0.962963	0.928571	1
NOT-EPCAM AND VANGL1 AND NOT-TM4SF1	0.857143	0.8	0.923077
NOT-RNF43 AND VANGL1 AND NOT-TSPAN3	0.888889	0.857143	0.923077
NOT-RNF43 AND VANGL1 AND GPR160	0.8	0.833333	0.769231
NOT-EPCAM AND VANGL1 AND NOT-CRHR1	0.896552	0.8125	1
NOT-RNF43 AND VANGL1 AND NOT-S1PR4	0.923077	0.923077	0.923077
NOT-RNF43 AND VANGL1 AND NPY5R	0.88	0.916667	0.846154
NOT-EPCAM AND VANGL1 AND NOT-AVPR2	0.928571	0.866667	1
NOT-EPCAM AND VANGL1 AND NOT-DIO3	0.896552	0.8125	1
NOT-EPCAM AND VANGL1 AND NOT-PCDHA10	0.962963	0.928571	1
NOT-EPCAM AND VANGL1 AND NOT-CCR6	0.928571	0.866667	1
NOT-ERBB3 AND VANGL1 AND NOT-S1PR4	0.88	0.916667	0.846154
BIRC5 AND NOT-F11R AND FXYD6	0.818182	1	0.692308
NOT-EPCAM AND VANGL1 AND NOT-GPR182	0.928571	0.866667	1
NOT-EPCAM AND VANGL1 AND IGSF6	0.814815	0.785714	0.846154
NOT-EPCAM AND VANGL1 AND NOT-NPY5R	0.962963	0.928571	1
NOT-SCNN1A AND VANGL1 AND NOT-CLDN6	0.923077	0.923077	0.923077
ATP8B2 AND NOT-EREG AND NOT-CD22	0.869565	1	0.769231
NOT-F11R AND VANGL1 AND NOT-CD22	1	1	1
BIRC5 AND NOT-GDPD2 AND CLDN20	0.818182	1	0.692308
NOT-F11R AND VANGL1 AND NOT-FCRL2	1	1	1
ATP8B2 AND NOT-F11R AND NOT-CD22	0.869565	1	0.769231
NOT-ERBB3 AND VANGL1 AND NOT-ICAM4	0.88	0.916667	0.846154
NOT-F11R AND VANGL1 AND NOT-CD79B	0.962963	0.928571	1
NOT-F11R AND VANGL1 AND NOT-MSLN	0.962963	0.928571	1
ATP8B2 AND NOT-F11R AND NOT-CLDN6	0.8	0.833333	0.769231
NOT-F11R AND VANGL1 AND NOT-CLDN2	0.962963	0.928571	1
NOT-F11R AND VANGL1 AND NOT-ERBB4	1	1	1
ATP8B2 AND NOT-F11R AND NOT-FCRL2	0.869565	1	0.769231
BIRC5 AND NOT-F11R AND LRRTM4	0.818182	1	0.692308
NOT-F11R AND CBX3 AND NOT-SORL1	0.923077	0.923077	0.923077
TGFBI AND NOT-F11R AND NOT-TNFRSF17	0.818182	1	0.692308
NOT-CEACAM6 AND VANGL1 AND NOT-TSPAN3	0.827586	0.75	0.923077
BIRC5 AND NOT-F11R AND PAQR7	0.818182	1	0.692308
ATP8B2 AND NOT-FCRL2 AND NOT-TM7SF2	0.869565	1	0.769231
BIRC5 AND NOT-F11R AND SLC13A1	0.818182	1	0.692308
NOT-F11R AND VANGL1 AND NOT-LGR5	0.869565	1	0.769231
ATP8B2 AND NOT-FCRL2 AND NOT-PERP	0.869565	1	0.769231
ATP8B2 AND NOT-FCRL2 AND NOT-TACSTD2	0.869565	1	0.769231
ATP8B2 AND NOT-FCRL2 AND NOT-MGST2	0.869565	1	0.769231
NOT-ERBB3 AND CEACAM7 AND PMP22	0.818182	1	0.692308
NOT-F11R AND VANGL1 AND NOT-CLDN18	0.962963	0.928571	1
NOT-EPCAM AND BRCA1 AND SLC13A1	0.818182	1	0.692308
NOT-F11R AND VANGL1 AND NOT-KDR	0.888889	0.857143	0.923077
NOT-CEACAM6 AND VANGL1 AND NOT-ABCB8	0.827586	0.75	0.923077
NOT-SCNN1B AND VANGL1 AND NOT-CLDN6	0.923077	0.923077	0.923077
BIRC5 AND NOT-F11R AND SCN2A	0.818182	1	0.692308
BIRC5 AND NOT-F11R AND OPRK1	0.818182	1	0.692308
BIRC5 AND NOT-F11R AND KCND2	0.818182	1	0.692308
BIRC5 AND NOT-F11R AND NOT-GABRB1	0.818182	1	0.692308
BIRC5 AND NOT-F11R AND NOT-KCNK12	0.818182	1	0.692308
BIRC5 AND NOT-F11R AND ATP13A5	0.818182	1	0.692308
NOT-CRB3 AND VANGL1 AND NOT-CLDN6	0.923077	0.923077	0.923077
ATP8B2 AND NOT-COL17A1 AND NOT-CD22	0.869565	1	0.769231
NOT-F11R AND VANGL1 AND NOT-SSTR5	0.962963	0.928571	1
ATP8B2 AND NOT-KIAA1324 AND NOT-CLDN1	0.8	0.833333	0.769231
NOT-ERBB3 AND VANGL1 AND NOT-CRHR1	0.846154	0.846154	0.846154
BIRC5 AND NOT-GDPD2 AND PHLDB2	0.818182	1	0.692308
BIRC5 AND NOT-F11R AND CYP4A11	0.818182	1	0.692308
BIRC5 AND NOT-F11R AND SLC39A12	0.818182	1	0.692308
BIRC5 AND NOT-F11R AND HCN2	0.818182	1	0.692308
BIRC5 AND NOT-F11R AND ASTN1	0.818182	1	0.692308
BIRC5 AND NOT-F11R AND GUCY2F	0.818182	1	0.692308
BIRC5 AND NOT-F11R AND GABRA4	0.818182	1	0.692308

Antigen Pair Logic	F1	Precision	Recall
STEAP2 AND NOT-SLC39A2 AND FAP	1	1	1
STEAP2 AND NOT-DYSF AND MUC1	1	1	1
STEAP2 AND NOT-DYSF AND VCAM1	1	1	1
STEAP2 AND CLDN8 AND NOT-SLC5A2	1	1	1
TRPM8 AND NOT-CSPG4 AND MUC1	1	1	1
STEAP2 AND NOT-SLC4A8 AND WT1	0.888889	0.8	1
STEAP2 AND NOT-PAQR7 AND MUC1	1	1	1
STEAP2 AND NOT-SLC4A8 AND ABCB5	0.888889	0.8	1
STEAP2 AND NOT-SLC4A8 AND P2RX5	0.888889	0.8	1
STEAP2 AND NOT-NCAM1 AND SLC17A1	0.888889	0.8	1
STEAP2 AND NOT-CD36 AND VCAM1	1	1	1
TRPM8 AND NOT-CSPG4 AND RNF43	0.888889	0.8	1
STEAP2 AND NOT-SLC4A8 AND NOT-CSPG4	0.888889	0.8	1
TRPM8 AND NOT-TNC AND GPNMB	0.888889	0.8	1
STEAP2 AND NOT-EMP3 AND MUC1	0.888889	0.8	1
STEAP2 AND NOT-DYSF AND RNF43	0.888889	0.8	1
TRPM8 AND NOT-NCAM1 AND NOT-IGF1R	0.888889	0.8	1
TRPM8 AND NOT-LGR5 AND IL2RA	0.888889	0.8	1
STEAP2 AND BIRC5 AND NOT-GRIN2C	0.888889	0.8	1
STEAP2 AND NOT-CD36 AND MUC1	1	1	1
STEAP2 AND NOT-SLC22A18 AND CLDN8	1	1	1
STEAP2 AND SLC43A1 AND FAP	1	1	1
STEAP2 AND NOT-NCAM1 AND SLC13A2	0.888889	0.8	1
STEAP2 AND NOT-SLC4A8 AND NOT-CSPG4	0.888889	0.8	1
STEAP2 AND NOT-GYPC AND GPNMB	0.888889	0.8	1
STEAP2 AND SLC2A2 AND NOT-LGR5	0.888889	0.8	1
STEAP2 AND NOT-NCAM1 AND SELP	0.888889	0.8	1
STEAP2 AND BIRC5 AND CLDN8	1	1	1
STEAP2 AND CLDN8 AND NOT-DKK1	1	1	1
STEAP2 AND CLDN8 AND NOT-DKK1	1	1	1
STEAP2 AND BIRC5 AND CLDN8	1	1	1
STEAP2 AND CLDN8 AND NOT-TNC	1	1	1
STEAP2 AND NOT-CSPG4 AND VCAM1	1	1	1
STEAP2 AND NOT-DKK1 AND ABCA5	1	1	1
STEAP2 AND NOT-CSPG4 AND VCAM1	1	1	1
STEAP2 AND NOT-CSPG4 AND MUC1	1	1	1
STEAP2 AND NOT-TNC AND MUC1	1	1	1
STEAP2 AND NOT-TNC AND MUC1	1	1	1
STEAP2 AND NOT-CSPG4 AND MUC1	1	1	1
STEAP2 AND NOT-CSPG4 AND ABCA5	1	1	1
STEAP2 AND NOT-TNC AND VCAM1	1	1	1
STEAP2 AND NOT-TNC AND ABCA5	1	1	1
STEAP2 AND P2RX5 AND NOT-NCAM1	0.888889	0.8	1
STEAP2 AND CLDN8 AND WT1	1	1	1
STEAP2 AND NOT-NCAM1 AND SSTR1	0.888889	0.8	1
STEAP2 AND NOT-NCAM1 AND CD160	0.888889	0.8	1
STEAP2 AND BIRC5 AND NOT-LGR5	0.888889	0.8	1
COMPLEX-EDNRB/STEAP2/P2RX5	0.888889	0.8	1
STEAP2 AND NOT-NCAM1 AND NOT-IGF1R	0.888889	0.8	1
STEAP2 AND BIRC5 AND NOT-LGR5	0.888889	0.8	1
STEAP2 AND NOT-CSPG4 AND GPNMB	0.888889	0.8	1
COMPLEX-EDNRB/STEAP2/SSTR1	0.888889	0.8	1
STEAP2 AND BIRC5 AND NOT-NCAM1	0.888889	0.8	1
STEAP2 AND NOT-TNC AND GPNMB	0.888889	0.8	1
STEAP2 AND NOT-CSPG4 AND RNF43	0.888889	0.8	1
STEAP2 AND NOT-TNC AND RNF43	0.888889	0.8	1
STEAP2 AND NOT-NCAM1 AND MOK	0.888889	0.8	1
STEAP2 AND IL2RA AND NOT-LGR5	0.888889	0.8	1
STEAP2 AND NOT-DKK1 AND RNF43	0.888889	0.8	1
STEAP2 AND NOT-CSPG4 AND CLDN8	1	1	1
STEAP2 AND CLDN8 AND NOT-KDR	1	1	1
STEAP2 AND CLDN8 AND ABCB5	1	1	1
STEAP2 AND NOT-CSPG4 AND CLDN8	1	1	1
STEAP2 AND BIRC5 AND TRPM4	1	1	1
STEAP2 AND NOT-DKK1 AND GPNMB	0.888889	0.8	1
STEAP2 AND CLDN8 AND FAP	1	1	1
STEAP2 AND NOT-TNC AND IL13RA1	0.888889	0.8	1
STEAP2 AND NOT-DKK1 AND IL13RA1	0.888889	0.8	1
STEAP2 AND P2RX5 AND VCAM1	1	1	1
STEAP2 AND NOT-DKK1 AND VCAM1	1	1	1
STEAP2 AND CLDN8 AND FAP	1	1	1
STEAP2 AND P2RX5 AND MUC1	1	1	1
STEAP2 AND P2RX5 AND MUC1	1	1	1
STEAP2 AND P2RX5 AND GPNMB	0.888889	0.8	1
STEAP2 AND NOT-DKK1 AND MUC1	0.888889	0.8	1
STEAP2 AND NOT-DKK1 AND MUC1	0.888889	0.8	1
STEAP2 AND NOT-DKK1 AND VCAM1	1	1	1

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
TGFBI AND NOT-F11R AND KDR	0.818182	1	0.692308
BIRC5 AND NOT-F11R AND SLC17A3	0.818182	1	0.692308
BIRC5 AND NOT-F11R AND GJD2	0.818182	1	0.692308
BIRC5 AND NOT-F11R AND SLCO1A2	0.818182	1	0.692308
BIRC5 AND NOT-F11R AND NOT-MTNR1B	0.818182	1	0.692308
BIRC5 AND NOT-F11R AND SLC6A11	0.818182	1	0.692308
ATP8B2 AND NOT-GDPD2 AND NOT-TNFRSF17	0.833333	0.909091	0.769231
BIRC5 AND NOT-F11R AND KDR	0.818182	1	0.692308
BIRC5 AND NOT-ITGB6 AND CDH5	0.818182	1	0.692308
BIRC5 AND NOT-F11R AND NOT-CD79B	0.818182	1	0.692308
BIRC5 AND NOT-EPCAM AND TMPRSS5	0.818182	1	0.692308
BIRC5 AND NOT-EPCAM AND SLC5A7	0.818182	1	0.692308
BIRC5 AND NOT-EPCAM AND CDH5	0.818182	1	0.692308
BIRC5 AND NOT-EPCAM AND EQTN	0.818182	1	0.692308
BIRC5 AND NOT-EPCAM AND EMP1	0.818182	1	0.692308
BIRC5 AND NOT-EPCAM AND FZD7	0.818182	1	0.692308
BIRC5 AND NOT-EPCAM AND PCDHAC1	0.818182	1	0.692308
BIRC5 AND NOT-F11R AND NOT-MS4A1	0.818182	1	0.692308
BIRC5 AND NOT-EPCAM AND ZNRF3	0.818182	1	0.692308
BIRC5 AND NOT-ITGB6 AND PMP22	0.818182	1	0.692308
BIRC5 AND NOT-ITGB6 AND EMP2	0.818182	1	0.692308
BIRC5 AND NOT-F11R AND NOT-CD22	0.818182	1	0.692308
BIRC5 AND NOT-F11R AND NOT-FCRL5	0.818182	1	0.692308
BIRC5 AND NOT-F11R AND NOT-FCRL1	0.818182	1	0.692308
BIRC5 AND NOT-EPCAM AND NOT-CCR6	0.818182	1	0.692308
BIRC5 AND NOT-ITGB6 AND CD151	0.818182	1	0.692308
BIRC5 AND NOT-EREG AND NOT-CLDN6	0.818182	1	0.692308
NOT-EPCAM AND BRCA1 AND KDR	0.818182	1	0.692308
BIRC5 AND NOT-MGST2 AND IL13RA1	0.818182	1	0.692308
BIRC5 AND NOT-F11R AND MSLN	0.818182	1	0.692308
BIRC5 AND NOT-GDPD2 AND NOT-P2RX5	0.818182	1	0.692308
BIRC5 AND NOT-F11R AND NOT-FCRL2	0.818182	1	0.692308
BIRC5 AND NOT-MGST2 AND NOT-FCRL2	0.818182	1	0.692308
NOT-F11R AND CBX3 AND KDR	0.96	1	0.923077
NOT-EPCAM AND CBX3 AND NOT-CD40LG	0.962963	0.928571	1
BIRC5 AND NOT-F11R AND SSTR5	0.818182	1	0.692308
BIRC5 AND NOT-COL17A1 AND NOT-P2RX5	0.818182	1	0.692308
NOT-F11R AND CBX3 AND NOT-CD79B	0.96	1	0.923077
BIRC5 AND NOT-GDPD2 AND KDR	0.818182	1	0.692308
BIRC5 AND NOT-MST1R AND CDH5	0.818182	1	0.692308
CD276 AND NOT-GDPD2 AND CR2	0.818182	1	0.692308
CBX3 AND NOT-ITGB6 AND CDH5	0.923077	0.923077	0.923077
NOT-F11R AND CBX3 AND NOT-CD22	0.96	1	0.923077
NOT-EPCAM AND CBX3 AND ZNRF3	1	1	1
BIRC5 AND NOT-F11R AND NOT-CLDN6	0.818182	1	0.692308
BIRC5 AND NOT-GDPD2 AND NOT-CD79A	0.818182	1	0.692308
BIRC5 AND NOT-EREG AND NOT-CD22	0.818182	1	0.692308
BIRC5 AND NOT-F11R AND NOT-TNFRSF17	0.818182	1	0.692308
NOT-ERBB3 AND ITGB5 AND MSLN	0.814815	0.785714	0.846154
NOT-EPCAM AND CBX3 AND CDH5	1	1	1
NOT-F11R AND CBX3 AND NOT-FCRL2	0.96	1	0.923077
NOT-EPCAM AND CBX3 AND NOT-CCR6	1	1	1
COMPLEX-FGFR1/CLDN1/CD79B	0.818182	1	0.692308
BIRC5 AND NOT-MST1R AND GPR84	0.818182	1	0.692308
BIRC5 AND NOT-F11R AND LGR5	0.818182	1	0.692308
SLC4A7 AND NOT-CD79A AND NOT-MST1R	0.833333	0.909091	0.769231
NOT-F11R AND VANGL1 AND NOT-PAQR7	0.827586	0.75	0.923077
NOT-F11R AND VANGL1 AND NOT-SLC6A11	0.916667	1	0.846154
NOT-F11R AND VANGL1 AND NOT-SLC5A11	0.962963	0.928571	1
NOT-F11R AND VANGL1 AND NOT-MTNR1B	1	1	1
NOT-F11R AND VANGL1 AND NOT-ATP13A5	0.96	1	0.923077
NOT-F11R AND VANGL1 AND NOT-SLC1A6	1	1	1
NOT-F11R AND VANGL1 AND NOT-SELP	0.896552	0.8125	1
NOT-F11R AND VANGL1 AND NOT-GRIN1	0.962963	0.928571	1
NOT-F11R AND VANGL1 AND NOT-GUCY2F	0.962963	0.928571	1
NOT-F11R AND VANGL1 AND NOT-HRH3	0.962963	0.928571	1
NOT-F11R AND VANGL1 AND NOT-WNT7A	0.869565	1	0.769231
NOT-F11R AND VANGL1 AND NOT-ABCG8	0.962963	0.928571	1
NOT-F11R AND VANGL1 AND NOT-KCNQ2	1	1	1
NOT-F11R AND VANGL1 AND NOT-GPR6	0.916667	1	0.846154
NOT-F11R AND VANGL1 AND NOT-KCNA10	0.962963	0.928571	1
NOT-F11R AND VANGL1 AND NOT-SEZ6	0.962963	0.928571	1
NOT-F11R AND VANGL1 AND NOT-DGKE	0.896552	0.8125	1
NOT-F11R AND VANGL1 AND NOT-KIAA0319	0.962963	0.928571	1
NOT-F11R AND VANGL1 AND NOT-OR8D1	0.962963	0.928571	1
TGFBI AND NOT-F11R AND SLCO1A2	0.818182	1	0.692308
NOT-F11R AND VANGL1 AND NOT-FGF6	0.962963	0.928571	1

Antigen Pair Logic	F1	Precision	Recall
COMPLEX-CBX3/STEAP2/IGF1R	1	1	1
STEAP2 AND CLDN8 AND NOT-CLDN5	1	1	1
STEAP2 AND P2RX5 AND VCAM1	1	1	1
STEAP2 AND NOT-DKK1 AND IL13RA1	0.888889	0.8	1
COMPLEX-FAP/STEAP2/CLDN5	0.888889	0.8	1
STEAP2 AND NOT-TNC AND IL13RA1	0.888889	0.8	1
STEAP2 AND NOT-ERBB4 AND CLDN8	0.857143	1	0.75
STEAP2 AND CLDN8 AND NOT-DDX3X	0.857143	1	0.75
STEAP2 AND CLDN8 AND CLDN6	1	1	1
STEAP2 AND EPCAM AND FAP	0.857143	1	0.75
STEAP2 AND NOT-EDNRB AND CLDN8	0.857143	1	0.75
STEAP2 AND CLDN8 AND NOT-SLC7A5	0.857143	1	0.75
STEAP2 AND CLDN8 AND NOT-ANXA1	0.857143	1	0.75
STEAP2 AND NOT-TPBG AND CLDN8	0.857143	1	0.75
STEAP2 AND CLDN8 AND NOT-IL13RA1	0.857143	1	0.75
STEAP2 AND NOT-B4GALNT1 AND SSTR1	0.857143	1	0.75
STEAP2 AND NOT-B4GALNT1 AND CLDN23	0.857143	1	0.75
STEAP2 AND NOT-B4GALNT1 AND NOT-IGF1R	0.857143	1	0.75
STEAP2 AND NOT-NCAM1 AND NOT-HSPA5	0.888889	0.8	1
STEAP2 AND NOT-B4GALNT1 AND MOK	0.857143	1	0.75
STEAP2 AND EPCAM AND LGR5	0.857143	1	0.75
STEAP2 AND P2RX5 AND NOT-DPEP1	0.857143	1	0.75
STEAP2 AND NOT-EDNRB AND EPHA3	0.857143	1	0.75
STEAP2 AND NOT-FLOT2 AND VCAM1	0.857143	1	0.75
COMPLEX-STEAP2/EPCAM/VCAM1	0.857143	1	0.75
STEAP2 AND NOT-NCAM1 AND VCAM1	1	1	1
STEAP2 AND NOT-PCYT1A AND VCAM1	0.857143	1	0.75
STEAP2 AND NOT-CLDN12 AND VCAM1	0.857143	1	0.75
STEAP2 AND NOT-CD34 AND VCAM1	0.857143	1	0.75
STEAP2 AND ABCB5 AND MUC1	1	1	1
COMPLEX-STEAP2/EPCAM/SSTR1	0.857143	1	0.75
COMPLEX-STEAP2/IL11RA/EPCAM	0.857143	1	0.75
STEAP2 AND NOT-B4GALNT1 AND TNFRSF10A	0.857143	1	0.75
COMPLEX-CD160/STEAP2/EPCAM	0.857143	1	0.75
COMPLEX-STEAP2/HLA-DOB/EPCAM	0.857143	1	0.75
STEAP2 AND NOT-SLC7A5 AND MUC1	0.857143	1	0.75
STEAP2 AND WT1 AND NOT-LGR5	0.888889	0.8	1
STEAP2 AND EPCAM AND NOT-HSPA5	0.857143	1	0.75
STEAP2 AND ABCB5 AND NOT-LGR5	0.888889	0.8	1
STEAP2 AND NOT-B4GALNT1 AND CD160	0.857143	1	0.75
STEAP2 AND EPCAM AND NOT-IGF1R	0.857143	1	0.75
STEAP2 AND CLDN8 AND NOT-DPEP1	0.857143	1	0.75
STEAP2 AND NOT-ANXA1 AND VCAM1	0.857143	1	0.75
STEAP2 AND NOT-NCAM1 AND TNFRSF10A	0.888889	0.8	1
STEAP2 AND CLDN8 AND NOT-CLDN12	0.857143	1	0.75
STEAP2 AND NOT-PCYT1A AND MUC1	0.857143	1	0.75
STEAP2 AND NOT-B4GALNT1 AND MUC1	0.857143	1	0.75
TRPM8 AND NOT-SLC39A2 AND SLCO1A2	1	1	1
TRPM8 AND NOT-SLC39A2 AND GPR19	1	1	1
TRPM8 AND NOT-IL17RA AND SLC43A1	1	1	1
TRPM8 AND NOT-SLC39A2 AND EPHA10	1	1	1
TRPM8 AND SLCO1A2 AND NOT-FMNL1	1	1	1
TRPM8 AND NOT-SLC39A2 AND NOT-IL17RA	1	1	1
TRPM8 AND NOT-SLC39A2 AND NOT-GUCY2D	1	1	1
TRPM8 AND NOT-SLC39A2 AND NOT-CHRNA2	1	1	1
TRPM8 AND NOT-SLC39A2 AND NOT-EPHA2	1	1	1
TRPM8 AND NOT-SLC39A2 AND NOT-CHRNA2	1	1	1
TRPM8 AND NOT-SLC39A2 AND MEGF11	1	1	1
TRPM8 AND NOT-GYPC AND PTGER4	1	1	1
TRPM8 AND SLCO1A2 AND NOT-FMNL1	1	1	1
TRPM8 AND NOT-GYPC AND F2R	1	1	1
TRPM8 AND NOT-SLC39A2 AND NOT-GUCY2D	1	1	1
TRPM8 AND NOT-FMNL1 AND GPR19	1	1	1
TRPM8 AND NOT-SLC39A2 AND NOT-EBP	1	1	1
TRPM8 AND NOT-SLC39A2 AND NOT-EPHA2	1	1	1
TRPM8 AND NOT-SLC39A2 AND NOT-PTGER4	1	1	1
TRPM8 AND NOT-GYPC AND SLC43A1	1	1	1
TRPM8 AND NOT-GUCY2D AND SLC43A1	1	1	1
COMPLEX-SLC5A2/TRPM8/SLC16A5	1	1	1
TRPM8 AND NOT-FMNL1 AND EPHA10	1	1	1
TRPM8 AND NOT-SYT4 AND SLC43A1	1	1	1
TRPM8 AND NOT-SLC22A18 AND NOT-SLC39A2	1	1	1
TRPM8 AND NOT-CHRNA2 AND SLC43A1	1	1	1
TRPM8 AND NOT-SLC39A2 AND NOT-ANPEP	1	1	1
TRPM8 AND NOT-SLC39A2 AND NOT-CD36	1	1	1
TRPM8 AND NOT-DYSF AND SLC43A1	1	1	1
TRPM8 AND NOT-SLC4A8 AND NOT-ANPEP	1	1	1

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
NOT-F11R AND VANGL1 AND NOT-SV2C	1	1	1
NOT-F11R AND VANGL1 AND NOT-IGDCC3	0.869565	1	0.769231
NOT-F11R AND VANGL1 AND NOT-ADCY8	0.962963	0.928571	1
NOT-F11R AND VANGL1 AND NOT-MC2R	1	1	1
NOT-F11R AND VANGL1 AND NOT-HTR1E	1	1	1
NOT-F11R AND VANGL1 AND NOT-SLC22A6	0.962963	0.928571	1
NOT-F11R AND VANGL1 AND NOT-CACNG6	0.96	1	0.923077
NOT-F11R AND VANGL1 AND NOT-LRRN4	0.962963	0.928571	1
NOT-F11R AND VANGL1 AND NOT-HCN2	0.88	0.916667	0.846154
NOT-F11R AND VANGL1 AND NOT-MLC1	0.962963	0.928571	1
NOT-F11R AND VANGL1 AND NOT-SLC22A11	1	1	1
NOT-F11R AND VANGL1 AND NOT-LYPD1	0.923077	0.923077	0.923077
NOT-F11R AND VANGL1 AND NOT-GPR37L1	0.962963	0.928571	1
NOT-ADRA2A AND VANGL1 AND MLANA	0.869565	1	0.769231
NOT-F11R AND VANGL1 AND NOT-CRB2	0.96	1	0.923077
NOT-F11R AND VANGL1 AND NOT-RXFP3	0.962963	0.928571	1
ATP8B2 AND NOT-F11R AND NOT-DGKE	0.8	0.833333	0.769231
NOT-F11R AND VANGL1 AND NOT-SLC1A2	0.833333	0.909091	0.769231
NOT-F11R AND VANGL1 AND NOT-FXYD7	0.962963	0.928571	1
NOT-F11R AND VANGL1 AND NOT-CACNA1S	0.88	0.916667	0.846154
NOT-F11R AND VANGL1 AND NOT-CACNG7	0.96	1	0.923077
NOT-F11R AND VANGL1 AND NOT-CLCN1	0.962963	0.928571	1
NOT-F11R AND VANGL1 AND NOT-MAG	0.962963	0.928571	1
NOT-F11R AND VANGL1 AND NOT-ZP4	0.923077	0.923077	0.923077
NOT-F11R AND VANGL1 AND NOT-GABRA4	1	1	1
NOT-F11R AND VANGL1 AND NOT-CLCN1	0.962963	0.928571	1
EMP3 AND NOT-MLANA AND VANGL1	0.869565	1	0.769231
ATP8B2 AND NOT-MLC1 AND CYP4A11	0.833333	0.909091	0.769231
NOT-SMPD2 AND VANGL1 AND LRRN4	0.869565	1	0.769231
ATP8B2 AND NOT-MLC1 AND SV2C	0.833333	0.909091	0.769231
ATP8B2 AND NOT-GUCY2F AND VANGL1	0.869565	1	0.769231
ATP8B2 AND NOT-SLC39A2 AND VANGL1	0.869565	1	0.769231
GYPC AND VANGL1 AND NOT-GRIN1	0.833333	0.909091	0.769231
NOT-ATP8A1 AND LRRN4 AND NOT-SV2C	0.8	0.833333	0.769231
NOT-SORL1 AND VANGL1 AND NOT-SLC39A2	0.869565	1	0.769231
ATP8B2 AND NOT-GPR12 AND VANGL1	0.833333	0.909091	0.769231
GYPC AND VANGL1 AND NOT-KCNA10	0.88	0.916667	0.846154
PHLDB2 AND NOT-SLC39A2 AND VANGL1	0.928571	0.866667	1
GYPC AND VANGL1 AND NOT-CLCN1	0.88	0.916667	0.846154
GYPC AND VANGL1 AND NOT-GPR12	0.88	0.916667	0.846154
GYPC AND VANGL1 AND NOT-SLC13A5	0.88	0.916667	0.846154
PHLDB2 AND NOT-MLANA AND VANGL1	0.818182	1	0.692308
GYPC AND VANGL1 AND NOT-SLC39A2	0.88	0.916667	0.846154
EMP3 AND NOT-GPR12 AND VANGL1	0.88	0.916667	0.846154
NOT-EPCAM AND VANGL1 AND CLDN20	0.962963	0.928571	1
NOT-EPCAM AND VANGL1 AND NOT-KCNA10	0.896552	0.8125	1
NOT-EPCAM AND VANGL1 AND NOT-GRIN1	0.896552	0.8125	1
NOT-EPCAM AND VANGL1 AND NOT-CLCN1	0.896552	0.8125	1
NOT-EPCAM AND VANGL1 AND NOT-HRH3	0.896552	0.8125	1
NOT-EPCAM AND VANGL1 AND NOT-GUCY2F	0.896552	0.8125	1
NOT-ERBB3 AND VANGL1 AND NOT-HRH3	0.846154	0.846154	0.846154
NOT-ERBB3 AND VANGL1 AND NOT-GRIN1	0.846154	0.846154	0.846154
NOT-ERBB3 AND VANGL1 AND NOT-KCNA10	0.846154	0.846154	0.846154
NOT-ERBB3 AND VANGL1 AND NOT-GUCY2F	0.846154	0.846154	0.846154
NOT-CEACAM6 AND VANGL1 AND NOT-LRRN4	0.888889	0.857143	0.923077
BIRC5 AND NOT-MLC1 AND CYP4A11	0.818182	1	0.692308
NOT-ERBB3 AND VANGL1 AND NOT-CLCN1	0.846154	0.846154	0.846154
NOT-ERBB3 AND VANGL1 AND NOT-SLC13A5	0.846154	0.846154	0.846154
ATP8B2 AND NOT-FCRL2 AND NOT-SLC22A18	0.869565	1	0.769231
ATP8B2 AND NOT-CD79B AND NOT-SLC13A1	0.833333	0.909091	0.769231
NOT-CEACAM6 AND VANGL1 AND NOT-GRIN1	0.827586	0.75	0.923077
NOT-ERBB3 AND VANGL1 AND NOT-SLC22A11	0.846154	0.846154	0.846154
BIRC5 AND NOT-MLC1 AND NOT-SLC22A18	0.818182	1	0.692308
NOT-CEACAM6 AND VANGL1 AND SLC13A5	0.827586	0.75	0.923077
NOT-CEACAM6 AND VANGL1 AND GPR12	0.814815	0.785714	0.846154
NOT-ERBB3 AND VANGL1 AND SLC39A2	0.869565	1	0.769231
NOT-CEACAM6 AND VANGL1 AND GUCY2F	0.827586	0.75	0.923077
NOT-CEACAM6 AND VANGL1 AND NOT-SLC22A11	0.888889	0.857143	0.923077
ATP8B2 AND NOT-CD79A AND NOT-KIAA0319	0.833333	0.909091	0.769231
NOT-CLDN4 AND VANGL1 AND NOT-GRIN1	0.888889	0.857143	0.923077
ATP8B2 AND NOT-KIAA0319 AND NOT-P2RX5	0.833333	0.909091	0.769231
BIRC5 AND NOT-MLC1 AND SLC13A1	0.818182	1	0.692308
NOT-CLDN4 AND VANGL1 AND NOT-HRH3	0.888889	0.857143	0.923077
NOT-CLDN4 AND VANGL1 AND NOT-GUCY2F	0.888889	0.857143	0.923077
NOT-CLDN4 AND VANGL1 AND NOT-KCNA10	0.888889	0.857143	0.923077
NOT-CLDN4 AND VANGL1 AND NOT-CLCN1	0.888889	0.857143	0.923077
NOT-SMPD2 AND VANGL1 AND CLDN18	0.818182	1	0.692308

Antigen Pair Logic	F1	Precision	Recall
TRPM8 AND NOT-SLC5A2 AND SLC43A1	1	1	1
TRPM8 AND NOT-SLC39A2 AND SLCO6A1	1	1	1
TRPM8 AND NOT-SLC39A2 AND NOT-SEMA4D	1	1	1
TRPM8 AND NOT-SLC39A2 AND NOT-STAB1	1	1	1
TRPM8 AND NOT-SLC39A2 AND NOT-TNFRSF12A	1	1	1
TRPM8 AND NOT-SLC39A2 AND NOT-STAB1	1	1	1
TRPM8 AND NOT-SLC39A2 AND PCDHA9	1	1	1
TRPM8 AND NOT-IL17RA AND F2R	1	1	1
TRPM8 AND NOT-SLC39A2 AND TMPRSS11D	1	1	1
TRPM8 AND NOT-SLC39A2 AND NPHS1	1	1	1
TRPM8 AND NOT-SLC39A2 AND NOT-GABRA1	1	1	1
TRPM8 AND NOT-SLC39A2 AND NMBR	1	1	1
TRPM8 AND NOT-SYT4 AND NOT-FMNL1	1	1	1
TRPM8 AND NOT-FMNL1 AND SLC2A2	1	1	1
TRPM8 AND NOT-SLC39A2 AND NOT-TNFRSF12A	1	1	1
TRPM8 AND NOT-SLC39A2 AND NOT-GABRA1	1	1	1
TRPM8 AND NOT-SLC39A2 AND TMEM235	1	1	1
TRPM8 AND NOT-SLC39A2 AND RGSL1	1	1	1
TRPM8 AND NOT-SLC39A2 AND NPHS2	1	1	1
TRPM8 AND NOT-SLC39A2 AND ADAM2	1	1	1
TRPM8 AND NOT-SLC39A2 AND NOT-SLC5A2	1	1	1
TRPM8 AND NOT-DYSF AND PTGER4	1	1	1
TRPM8 AND NOT-SLC39A2 AND CACNG7	1	1	1
TRPM8 AND NOT-SLC39A2 AND SLC2A2	1	1	1
TRPM8 AND NOT-SLC39A2 AND UMODL1	1	1	1
TRPM8 AND NOT-SLC39A2 AND NOT-SLC5A2	1	1	1
TRPM8 AND NOT-EPHA2 AND F2R	1	1	1
TRPM8 AND NOT-SLC39A2 AND NOT-SHISA9	1	1	1
TRPM8 AND NOT-GYPC AND NOT-SLC39A2	0.888889	0.8	1
TRPM8 AND NOT-SLC4A8 AND NMBR	0.888889	0.8	1
TRPM8 AND NOT-SYT4 AND FLVCR1	0.888889	0.8	1
TRPM8 AND NOT-SLC4A8 AND USH2A	0.888889	0.8	1
TRPM8 AND NOT-GYPC AND CD93	0.888889	0.8	1
TRPM8 AND NOT-SLC4A8 AND SLC2A2	0.888889	0.8	1
TRPM8 AND NOT-SLC4A8 AND EPHA10	0.888889	0.8	1
TRPM8 AND NOT-EPHA2 AND SLC43A1	1	1	1
TRPM8 AND NOT-SLC4A8 AND NOT-GUCY2D	0.888889	0.8	1
TRPM8 AND NOT-GRIN2C AND GPR19	0.888889	0.8	1
TRPM8 AND NOT-SLC22A18 AND NOT-GRIN2C	0.888889	0.8	1
TRPM8 AND NOT-GRIN2C AND SLCO1A2	0.888889	0.8	1
TRPM8 AND NOT-SLC4A8 AND AGTR2	0.888889	0.8	1
TRPM8 AND NOT-SLC4A8 AND SCN2A	0.888889	0.8	1
TRPM8 AND NOT-SLC4A8 AND MEGF11	0.888889	0.8	1
TRPM8 AND NOT-SLC4A8 AND SLCO1A2	0.888889	0.8	1
TRPM8 AND NOT-GRIN2C AND ATP8A2	0.888889	0.8	1
TRPM8 AND NOT-SLC4A8 AND GRIA4	0.888889	0.8	1
TRPM8 AND NOT-SLC4A8 AND CACNA1E	0.888889	0.8	1
TRPM8 AND NOT-SLC4A8 AND SLCO1A2	0.888889	0.8	1
TRPM8 AND NOT-SLC4A8 AND SLCO6A1	0.888889	0.8	1
TRPM8 AND NOT-SLC39A2 AND NOT-CDH10	1	1	1
TRPM8 AND NOT-SLC4A8 AND NOT-CHRNA2	0.888889	0.8	1
TRPM8 AND NOT-SYT4 AND NOT-ANPEP	0.888889	0.8	1
TRPM8 AND NOT-SLC39A2 AND NOT-VAMP8	1	1	1
TRPM8 AND NOT-SLC4A8 AND NOT-CHRNA2	0.888889	0.8	1
TRPM8 AND NOT-GRIN2C AND SLC2A2	0.888889	0.8	1
TRPM8 AND NOT-GABRA1 AND SLC43A1	1	1	1
TRPM8 AND NOT-GRIN2C AND MEGF11	0.888889	0.8	1
TRPM8 AND NOT-GYPC AND CDH10	0.888889	0.8	1
TRPM8 AND NOT-SLC4A8 AND ABCA12	0.888889	0.8	1
TRPM8 AND NOT-SYT4 AND NOT-SLC39A2	0.888889	0.8	1
TRPM8 AND NOT-DYSF AND CD93	0.888889	0.8	1
TRPM8 AND NOT-GRIN2C AND SLC43A1	1	1	1
TRPM8 AND NOT-GRIN2C AND GRIA4	0.888889	0.8	1
TRPM8 AND NOT-SLC4A8 AND NOT-EBP	0.888889	0.8	1
TRPM8 AND NOT-STAB1 AND SLC43A1	1	1	1
TRPM8 AND NOT-IL17RA AND NOT-SLC4A8	0.888889	0.8	1
TRPM8 AND NOT-SLC4A8 AND ADAM2	0.888889	0.8	1
TRPM8 AND NOT-SLC4A8 AND GPR19	0.888889	0.8	1
TRPM8 AND NOT-SLC4A8 AND NOT-GUCY2D	0.888889	0.8	1
Sarcoma			
TGFBI AND NOT-SPINT1 AND PRLR	0.903226	1	0.823529
ADAM12 AND NOT-ACPP AND CD74	0.944444	0.894737	1
TGFBI AND NOT-SPINT1 AND KCNH7	0.909091	0.9375	0.882353
TGFBI AND NOT-SPINT1 AND KIR2DL3	0.909091	0.9375	0.882353
ADAM12 AND NOT-SPINT1 AND GABRB1	0.941176	0.941176	0.941176
ADAM12 AND NOT-ABCB1 AND OR4N4	0.894737	0.809524	1
TGFBI AND NOT-SPINT1 AND TAS2R9	0.903226	1	0.823529

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
NOT-CLDN4 AND VANGL1 AND NOT-SLC22A11	0.888889	0.857143	0.923077
NOT-CLDN4 AND VANGL1 AND MLANA	0.916667	1	0.846154
NOT-CLDN4 AND VANGL1 AND SLC13A5	0.923077	0.923077	0.923077
FAP AND NOT-GRIN1 AND SV2C	0.818182	1	0.692308
GYPC AND VANGL1 AND NOT-CLDN6	0.88	0.916667	0.846154
ATP8B2 AND NOT-CD22 AND SLC13A1	0.869565	1	0.769231
BIRC5 AND NOT-KIAA0319 AND CYP4A11	0.818182	1	0.692308
NOT-RNF43 AND VANGL1 AND NOT-CLDN6	0.923077	0.923077	0.923077
NOT-EPCAM AND VANGL1 AND NOT-CLDN6	0.962963	0.928571	1
NOT-ERBB3 AND VANGL1 AND NOT-CLDN6	0.88	0.916667	0.846154
NOT-EPCAM AND VANGL1 AND NOT-MSLN	0.896552	0.8125	1
NOT-EPCAM AND VANGL1 AND NOT-KDR	0.857143	0.8	0.923077
BIRC5 AND NOT-EPCAM AND CLDN20	0.818182	1	0.692308
NOT-CLDN4 AND VANGL1 AND NOT-CLDN6	0.923077	0.923077	0.923077
BIRC5 AND NOT-EPCAM AND SLC13A1	0.818182	1	0.692308
NOT-CEACAM6 AND VANGL1 AND NOT-CLDN6	0.857143	0.8	0.923077
NOT-CEACAM6 AND VANGL1 AND NOT-SLC34A2	0.857143	0.8	0.923077
NOT-ERBB3 AND VANGL1 AND NOT-CLDN2	0.846154	0.846154	0.846154
NOT-CEACAM6 AND VANGL1 AND NOT-CLDN2	0.888889	0.857143	0.923077
ATP8B2 AND NOT-ITGB6 AND KDR	0.833333	0.909091	0.769231
NOT-ERBB3 AND VANGL1 AND SLC34A2	0.818182	1	0.692308
NOT-CEACAM6 AND VANGL1 AND NOT-MSLN	0.827586	0.75	0.923077
ATP8B2 AND NOT-ERBB4 AND NOT-P2RX5	0.869565	1	0.769231
ATP8B2 AND NOT-ERBB4 AND KDR	0.869565	1	0.769231
ATP8B2 AND NOT-EPCAM AND KDR	0.833333	0.909091	0.769231
ATP8B2 AND NOT-CD79A AND NOT-MST1R	0.833333	0.909091	0.769231
BIRC5 AND NOT-SLC22A18 AND NOT-FCRL2	0.818182	1	0.692308
BIRC5 AND NOT-MST1R AND NOT-MLC1	0.818182	1	0.692308
NOT-CLDN4 AND VANGL1 AND NOT-CLDN2	0.888889	0.857143	0.923077
BIRC5 AND NOT-MST1R AND SLC6A11	0.818182	1	0.692308
FAP AND NOT-CLDN6 AND SLC13A1	0.818182	1	0.692308
BIRC5 AND NOT-MST1R AND GRID2	0.818182	1	0.692308
NOT-ITGB6 AND SLC6A18 AND KDR	0.8	0.833333	0.769231
NOT-ATP8A1 AND SSTR5 AND NOT-ERBB4	0.8	0.833333	0.769231
FAP AND NOT-CD22 AND SLC13A1	0.818182	1	0.692308
BIRC5 AND NOT-EPCAM AND KDR	0.818182	1	0.692308
BIRC5 AND NOT-ITGB6 AND KDR	0.818182	1	0.692308
BIRC5 AND NOT-CD79B AND NOT-MST1R	0.818182	1	0.692308
NOT-EPCAM AND CBX3 AND KDR	0.8125	0.684211	1
BIRC5 AND NOT-MST1R AND KDR	0.818182	1	0.692308
CBX3 AND NOT-ITGB6 AND KDR	0.827586	0.75	0.923077
BIRC5 AND NOT-MST1R AND NOT-MS4A1	0.818182	1	0.692308
BIRC5 AND NOT-MST1R AND NOT-FCRL1	0.818182	1	0.692308
BIRC5 AND NOT-MST1R AND SSTR5	0.818182	1	0.692308
FAP AND NOT-MSLN AND NOT-MST1R	0.818182	1	0.692308
CBX3 AND NOT-MST1R AND NOT-MS4A1	0.916667	1	0.846154
CBX3 AND NOT-MST1R AND NOT-CD79A	0.96	1	0.923077
CBX3 AND NOT-MST1R AND NOT-CD79B	0.96	1	0.923077
FAP AND NOT-MST1R AND FCRL1	0.818182	1	0.692308
CBX3 AND NOT-MST1R AND NOT-FCRL1	0.916667	1	0.846154
FAP AND NOT-MST1R AND MS4A1	0.818182	1	0.692308
AML			
NOT-PTPRK AND NOT-GPR171 AND HCST	0.963107	0.939394	0.988048
NOT-PTPRK AND NOT-GPR171 AND HCST	0.963107	0.939394	0.988048
NOT-PODXL AND NOT-GPR171 AND HCST	0.963107	0.939394	0.988048
NOT-PODXL AND NOT-GPR171 AND HCST	0.963107	0.939394	0.988048
NOT-PTPRK AND NOT-GPR171 AND PTGER2	0.962963	0.942748	0.984064
NOT-SGCE AND HCST AND NOT-GPR171	0.962818	0.946154	0.98008
COMPLEX-FLT3/HEG1/RHAG	0.96124	0.935849	0.988048
COMPLEX-FLT3/SIT1/RHAG	0.96124	0.935849	0.988048
COMPLEX-TIGIT/FLT3/RHAG	0.961089	0.939163	0.984064
COMPLEX-FLT3/PODXL/RHAG	0.963107	0.939394	0.988048
NOT-PTPRK AND CD46 AND NOT-P2RY10	0.960474	0.952941	0.968127
NOT-PTPRK AND CD46 AND NOT-P2RY10	0.960474	0.952941	0.968127
NOT-PTPRK AND CSF3R AND NOT-BTN3A2	0.960474	0.952941	0.968127
NOT-PTPRK AND CSF3R AND NOT-BTN3A2	0.960474	0.952941	0.968127
COMPLEX-ABCC5/FLT3/RHAG	0.961089	0.939163	0.984064
NOT-WLS AND HCST AND NOT-GPR171	0.96	0.963855	0.956175
COMPLEX-FLT3/RHAG/BMPR2	0.961089	0.939163	0.984064
COMPLEX-FLT3/LAMP3/RHAG	0.96124	0.935849	0.988048
COMPLEX-FLT3/SLC9A9/RHAG	0.96124	0.935849	0.988048
NOT-PTPRK AND CSF3R AND NOT-P2RY10	0.965116	0.939623	0.992032
NOT-PTPRK AND CSF3R AND NOT-P2RY10	0.965116	0.939623	0.992032
COMPLEX-FLT3/RHAG/TSPAN6	0.959381	0.932331	0.988048
COMPLEX-FLT3/TMEM150C/RHAG	0.959381	0.932331	0.988048
COMPLEX-AGTR1/FLT3/RHAG	0.959381	0.932331	0.988048
COMPLEX-FLT3/AMIGO2/RHAG	0.959381	0.932331	0.988048

Antigen Pair Logic	F1	Precision	Recall
ADAM12 AND NOT-ST14 AND SV2C	0.85	0.73913	1
ADAM12 AND NOT-ST14 AND PCDHGC4	0.85	0.73913	1
ADAM12 AND NOT-ST14 AND GABRB1	0.864865	0.8	0.941176
ADAM12 AND NOT-ABCB1 AND GJA10	0.829268	0.708333	1
ADAM12 AND NOT-ST14 AND GABRG1	0.864865	0.8	0.941176
ADAM12 AND NOT-ST14 AND BEST3	0.85	0.73913	1
ADAM12 AND NOT-SPINT1 AND CRB1	0.944444	0.894737	1
ADAM12 AND NOT-SPINT1 AND SV2C	0.944444	0.894737	1
ADAM12 AND NOT-ST14 AND OR4N4	0.894737	0.809524	1
ADAM12 AND NOT-ST14 AND MLANA	0.85	0.73913	1
TGFBI AND NOT-SPINT1 AND PRRG2	0.875	0.933333	0.823529
ADAM12 AND NOT-HPN AND MLANA	0.882353	0.882353	0.882353
ADAM12 AND NOT-SPINT1 AND ACSL6	0.842105	0.761905	0.941176
ADAM12 AND NOT-ILDR1 AND CALN1	0.894737	0.809524	1
ADAM12 AND NOT-ST14 AND LCT	0.85	0.73913	1
ADAM12 AND NOT-ACPP AND SLC39A12	0.918919	0.85	1
PLXND1 AND NOT-ABCB1 AND GJA10	0.903226	1	0.823529
TGFBI AND NOT-SPINT1 AND KCNK7	0.909091	0.9375	0.882353
ATP8B2 AND NOT-ABCB1 AND NOT-CLCA4	0.8	0.923077	0.705882
ADAM12 AND NOT-ST14 AND GRIN2A	0.809524	0.68	1
ADAM12 AND NOT-ILDR1 AND GUCY2F	0.888889	0.842105	0.941176
TGFBI AND NOT-SPINT1 AND OR7A17	0.909091	0.9375	0.882353
ADAM12 AND NOT-ST14 AND ASTN1	0.829268	0.708333	1
TGFBI AND NOT-SPINT1 AND TAS2R14	0.909091	0.9375	0.882353
ADAM12 AND GP5 AND NOT-OR8D1	0.829268	0.708333	1
ADAM12 AND GP5 AND NOT-LRFN2	0.871795	0.772727	1
ADAM12 AND GP5 AND NOT-HTR6	0.842105	0.761905	0.941176
ADAM12 AND GP5 AND NOT-AGTR2	0.85	0.73913	1
OSMR AND NOT-ACPP AND NOT-AMHR2	0.903226	1	0.823529
DDR2 AND NOT-WNT4 AND LCT	0.8	0.923077	0.705882
ADAM12 AND NOT-ACPP AND PCDHGC4	0.918919	0.85	1
ADAM12 AND NOT-ILDR1 AND PCDHAC2	0.918919	0.85	1
ADAM12 AND NOT-ABCB1 AND FSHR	0.85	0.73913	1
ADAM12 AND NOT-ST14 AND CALN1	0.829268	0.708333	1
ADAM12 AND NOT-ILDR1 AND HCN4	0.944444	0.894737	1
ATP8B2 AND NOT-ABCB1 AND NOT-SPINT1	0.8	0.923077	0.705882
ATP8B2 AND NOT-ABCB1 AND GLRA1	0.8	0.923077	0.705882
ATP8B2 AND NOT-ABCB1 AND NOT-ST14	0.8	0.923077	0.705882
ATP8B2 AND NOT-ABCB1 AND CORIN	0.8	0.923077	0.705882
ATP8B2 AND NOT-ABCB1 AND NOT-FGFR3	0.8	0.923077	0.705882
ATP8B2 AND NOT-ABCB1 AND GP5	0.8	0.923077	0.705882
ADAM12 AND NOT-ST14 AND KCNV1	0.810811	0.75	0.882353
ATP8B2 AND NOT-ABCB1 AND OR10H2	0.8	0.923077	0.705882
TGFBI AND NOT-SPINT1 AND APP	0.9375	1	0.882353
ADAM12 AND NOT-ST14 AND GJA10	0.809524	0.68	1
ADAM12 AND NOT-SPINT1 AND CSMD2	0.944444	0.894737	1
ADAM12 AND NOT-SPINT1 AND GUCY2F	0.918919	0.85	1
ADAM12 AND NOT-ST14 AND GABRA2	0.894737	0.809524	1
ADAM12 AND NOT-ST14 AND CRB2	0.85	0.73913	1
TGFBI AND NOT-SPINT1 AND CDH5	0.9375	1	0.882353
ADAM12 AND NOT-ST14 AND NETO1	0.894737	0.809524	1
TGFBI AND NOT-SPINT1 AND GJD4	0.909091	0.9375	0.882353
ADAM12 AND NOT-SPINT1 AND ADCY10	0.944444	0.894737	1
ADAM12 AND NOT-SPINT1 AND NOT-ATP8B1	0.810811	0.75	0.882353
OSMR AND NOT-ACPP AND NOT-ADAM30	0.866667	1	0.764706
SLC2A10 AND NOT-CXADR AND GRIK4	0.827586	1	0.705882
TGFBI AND NOT-SPINT1 AND AQP4	0.827586	1	0.705882
TGFBI AND NOT-SPINT1 AND ERVFRD-1	0.909091	0.9375	0.882353
ADAM12 AND NOT-ST14 AND GABRA5	0.871795	0.772727	1
ADAM12 AND NOT-SPINT1 AND MPL	0.871795	0.772727	1
ADAM12 AND NOT-SPINT1 AND PCDHAC2	0.944444	0.894737	1
ADAM12 AND GP5 AND NOT-OPALIN	0.829268	0.708333	1
OSMR AND NOT-ACPP AND NOT-SLC22A6	0.903226	1	0.823529
OSMR AND NOT-CXADR AND GRIK4	0.8125	0.866667	0.764706
OSMR AND NOT-ACPP AND NOT-GPR135	0.903226	1	0.823529
OSMR AND NOT-ACPP AND NOT-GUCY2F	0.903226	1	0.823529
TGFBI AND NOT-SPINT1 AND ATP12A	0.909091	0.9375	0.882353
ADAM12 AND NOT-ST14 AND GUCY2F	0.829268	0.708333	1
ADAM12 AND NOT-ACPP AND ASTN1	0.894737	0.809524	1
ADAM12 AND NOT-ST14 AND CNTNAP4	0.829268	0.708333	1
ADAM12 AND NOT-ILDR1 AND LYPD1	0.85	0.73913	1
ADAM12 AND NOT-ACPP AND SV2C	0.918919	0.85	1
OSMR AND NOT-STEAP4 AND NOT-ACPP	0.827586	1	0.705882
ADAM12 AND NOT-CXADR AND GRIK4	0.944444	0.894737	1
TGFBI AND NOT-ACPP AND PRLR	0.8	0.923077	0.705882
ADAM12 AND GP5 AND NOT-SLC39A2	0.85	0.73913	1
ADAM12 AND NOT-SPINT1 AND NOT-ATP8A2	0.871795	0.772727	1

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
COMPLEX-FLT3/KLRF1/RHAG	0.959381	0.932331	0.988048
COMPLEX-FLT3/EMCN/RHAG	0.959381	0.932331	0.988048
COMPLEX-FLT3/CRIM1/RHAG	0.959381	0.932331	0.988048
COMPLEX-FLT3/PROS1/RHAG	0.959381	0.932331	0.988048
COMPLEX-FLT3/KLRD1/RHAG	0.959381	0.932331	0.988048
COMPLEX-FLT3/RHAG/ST3GAL5	0.959381	0.932331	0.988048
COMPLEX-FLT3/RHAG/FAT4	0.959381	0.932331	0.988048
NOT-PTPRK AND NOT-GPR171 AND CSF3R	0.959223	0.935606	0.984064
NOT-PTPRK AND NOT-GPR171 AND CSF3R	0.959223	0.935606	0.984064
COMPLEX-FLT3/RHAG/PPAP2B	0.959064	0.938931	0.98008
COMPLEX-FLT3/RHAG/APOLD1	0.959064	0.938931	0.98008
COMPLEX-FLT3/RHAG/TRPC1	0.959064	0.938931	0.98008
COMPLEX-FLT3/RHAG/CAV1	0.959381	0.932331	0.988048
COMPLEX-FLT3/ITGB5/RHAG	0.959381	0.932331	0.988048
COMPLEX-DAG1/FLT3/RHAG	0.959381	0.932331	0.988048
COMPLEX-FLT3/RHAG/TEK	0.959381	0.932331	0.988048
COMPLEX-FLT3/ENPP2/RHAG	0.963107	0.939394	0.988048
COMPLEX-FLT3/PTPRM/RHAG	0.96124	0.935849	0.988048
COMPLEX-FLT3/RHAG/HEPH	0.963107	0.939394	0.988048
COMPLEX-FLT3/RHAG/ACVR2A	0.958904	0.942308	0.976096
COMPLEX-FLT3/ABHD6/RHAG	0.963107	0.939394	0.988048
COMPLEX-FLT3/KCNMA1/RHAG	0.963107	0.939394	0.988048
COMPLEX-FLT3/FPR3/RHAG	0.959381	0.932331	0.988048
NOT-PODXL AND NOT-GPR171 AND CSF3R	0.959223	0.935606	0.984064
NOT-PODXL AND NOT-GPR171 AND CSF3R	0.959223	0.935606	0.984064
NOT-PTPRK AND NOT-KLRB1 AND HCST	0.95825	0.956349	0.960159
COMPLEX-FLT3/RHAG/JAM3	0.959381	0.932331	0.988048
NOT-PTPRK AND NOT-GPR171 AND ESYT2	0.958084	0.96	0.956175
COMPLEX-FLT3/GOLM1/RHAG	0.961089	0.939163	0.984064
COMPLEX-FLT3/ST6GALNAC6/RHAG	0.959381	0.932331	0.988048
COMPLEX-TSPAN9/FLT3/RHAG	0.959381	0.932331	0.988048
COMPLEX-FLT3/SLC17A5/RHAG	0.959381	0.932331	0.988048
COMPLEX-FLT3/CHIC1/RHAG	0.959381	0.932331	0.988048
COMPLEX-FLT3/SLC25A4/RHAG	0.957529	0.928839	0.988048
COMPLEX-ADRA2A/FLT3/RHAG	0.957529	0.928839	0.988048
COMPLEX-FLT3/MR1/RHAG	0.957529	0.928839	0.988048
NOT-PTPRK AND P2RX1 AND NOT-P2RY10	0.957529	0.928839	0.988048
NOT-PTPRK AND P2RX1 AND NOT-P2RY10	0.957529	0.928839	0.988048
COMPLEX-FLT3/RHAG/ABCG1	0.957529	0.928839	0.988048
NOT-WLS AND HCST AND NOT-CD3G	0.965795	0.97561	0.956175
NOT-PODXL AND NOT-LTB AND HCST	0.957404	0.975207	0.940239
NOT-PODXL AND NOT-LTB AND HCST	0.957404	0.975207	0.940239
COMPLEX-FLT3/IFNAR1/RHAG	0.957364	0.932075	0.984064
COMPLEX-FLT3/SLC2A8/RHAG	0.957364	0.932075	0.984064
COMPLEX-SLC16A10/FLT3/RHAG	0.957364	0.932075	0.984064
COMPLEX-FLT3/RHAG/DST	0.957364	0.932075	0.984064
COMPLEX-FLT3/RHAG/CD40	0.957364	0.932075	0.984064
COMPLEX-FLT3/RHAG/KL	0.957364	0.932075	0.984064
NOT-PTPRK AND P2RX1 AND NOT-CLEC4A	0.958904	0.942308	0.976096
COMPLEX-FLT3/RHAG/STX7	0.957198	0.935361	0.98008
COMPLEX-FLT3/CLEC4A/RHAG	0.957198	0.935361	0.98008
COMPLEX-FLT3/SIRPG/RHAG	0.957198	0.935361	0.98008
COMPLEX-FLT3/GPR171/RHAG	0.957198	0.935361	0.98008
COMPLEX-FLT3/HMOX2/RHAG	0.957198	0.935361	0.98008
COMPLEX-FLT3/RHAG/CD27	0.957031	0.938697	0.976096
NOT-PERP AND HCST AND NOT-GPR171	0.956863	0.942085	0.972112
COMPLEX-FLT3/IL7R/RHAG	0.959223	0.935606	0.984064
NOT-PTPRK AND CXCR4 AND NOT-P2RY10	0.959693	0.925926	0.996016
NOT-PTPRK AND CXCR4 AND NOT-P2RY10	0.959693	0.925926	0.996016
NOT-PODXL AND HCST AND NOT-KLRB1	0.956349	0.952569	0.960159
NOT-WLS AND CSF3R AND NOT-GPR171	0.96	0.963855	0.956175
NOT-WLS AND CSF3R AND NOT-KLRB1	0.955823	0.963563	0.948207
COMPLEX-FLT3/SLC22A17/RHAG	0.955684	0.925373	0.988048
COMPLEX-FLT3/PLXNA2/RHAG	0.955684	0.925373	0.988048
COMPLEX-GPR176/FLT3/RHAG	0.955684	0.925373	0.988048
COMPLEX-ABCA3/FLT3/RHAG	0.955684	0.925373	0.988048
COMPLEX-FLT3/JAM2/RHAG	0.955684	0.925373	0.988048
COMPLEX-FLT3/RHAG/ZNRF3	0.955684	0.925373	0.988048
COMPLEX-FLT3/RHAG/FZD7	0.955684	0.925373	0.988048
COMPLEX-FLT3/NPC1/RHAG	0.955684	0.925373	0.988048
NOT-PODXL AND HCST AND NOT-CD28	0.957031	0.938697	0.976096
COMPLEX-FLT3/SLC39A1/RHAG	0.955513	0.928571	0.984064
COMPLEX-FLT3/LGR6/RHAG	0.955513	0.928571	0.984064
NOT-PTPRK AND LTBR AND NOT-CLEC4A	0.955466	0.971193	0.940239
NOT-PTPRK AND NOT-GPR171 AND P2RX4	0.968254	0.964427	0.972112
NOT-PTPRK AND P2RX1 AND NOT-BTN3A1	0.96699	0.943182	0.992032
NOT-PTPRK AND NOT-IL2RB AND P2RX4	0.963563	0.979424	0.948207

Antigen Pair Logic	F1	Precision	Recall
DDR2 AND NOT-WNT4 AND OR4N4	0.8	0.923077	0.705882
ADAM12 AND GP5 AND NOT-GABRB1	0.809524	0.68	1
ADAM12 AND NOT-ST14 AND SCN2A	0.894737	0.809524	1
ADAM12 AND OR4N4 AND NOT-CLEC4M	0.871795	0.772727	1
TSPAN4 AND NOT-GDPD2 AND VANGL1	0.9375	1	0.882353
ADAM12 AND NOT-ST14 AND MPL	0.809524	0.68	1
ADAM12 AND NOT-ST14 AND ADCY10	0.85	0.73913	1
ADAM12 AND NOT-ST14 AND OR5P3	0.85	0.73913	1
ADAM12 AND NOT-ACPP AND ACSL6	0.820513	0.727273	0.941176
OSMR AND NOT-SPINT1 AND NOT-GUCY2F	0.875	0.933333	0.823529
TGFBI AND NOT-CXADR AND NOT-ABCB1	0.9375	1	0.882353
ADAM12 AND GP5 AND NOT-ACSL6	0.829268	0.708333	1
TGFBI AND NOT-SPINT1 AND ADCY10	0.909091	0.9375	0.882353
TGFBI AND NOT-SPINT1 AND CRB1	0.875	0.933333	0.823529
TGFBI AND NOT-SPINT1 AND CASR	0.909091	0.9375	0.882353
TGFBI AND NOT-SPINT1 AND GRIK4	0.909091	0.9375	0.882353
TGFBI AND NOT-SPINT1 AND PCDHAC2	0.909091	0.9375	0.882353
TGFBI AND NOT-SPINT1 AND GUCY2F	0.909091	0.9375	0.882353
TGFBI AND NOT-SPINT1 AND NRG3	0.875	0.933333	0.823529
TGFBI AND NOT-SPINT1 AND GHSR	0.909091	0.9375	0.882353
TGFBI AND NOT-SPINT1 AND SLC5A8	0.909091	0.9375	0.882353
TGFBI AND NOT-SPINT1 AND SV2C	0.875	0.933333	0.823529
ATP8B2 AND NOT-ABCB1 AND GJA10	0.8	0.923077	0.705882
TGFBI AND NOT-SPINT1 AND MPL	0.909091	0.9375	0.882353
TGFBI AND NOT-SPINT1 AND CSMD2	0.9375	1	0.882353
TGFBI AND NOT-ACPP AND GPR135	0.8	0.923077	0.705882
TGFBI AND NOT-SPINT1 AND CCKAR	0.909091	0.9375	0.882353
TGFBI AND NOT-SPINT1 AND GABRB1	0.875	0.933333	0.823529
TGFBI AND NOT-SPINT1 AND ACSL6	0.875	0.933333	0.823529
TGFBI AND NOT-ACPP AND GHSR	0.8	0.923077	0.705882
TGFBI AND NOT-ACPP AND EPHA10	0.8	0.923077	0.705882
ADAM12 AND NOT-MMP24 AND ASTN1	0.842105	0.761905	0.941176
ADAM12 AND NOT-PCDHAC2 AND AGTR2	0.848485	0.875	0.823529
TGFBI AND NOT-ACPP AND CALN1	0.8	0.923077	0.705882
TGFBI AND NOT-ACPP AND ADCY10	0.8	0.923077	0.705882
TGFBI AND NOT-ACPP AND ASTN1	0.8	0.923077	0.705882
TGFBI AND NOT-ACPP AND SHISA9	0.8	0.923077	0.705882
TGFBI AND NOT-ACPP AND ADAM30	0.8	0.923077	0.705882
TGFBI AND NOT-ACPP AND SV2C	0.8	0.923077	0.705882
OSMR AND NOT-LRRN4 AND GRIK4	0.83871	0.928571	0.764706
TGFBI AND NOT-SPINT1 AND ATP8B1	0.9375	1	0.882353
ADAM12 AND PCDHGC4 AND NOT-HTR6	0.864865	0.8	0.941176
ADAM12 AND NOT-MMP24 AND CRB2	0.894737	0.809524	1
TGFBI AND NOT-ACPP AND OR1C1	0.8	0.923077	0.705882
ADAM12 AND PCDHGC4 AND NOT-SLC39A2	0.918919	0.85	1
ADAM12 AND SV2C AND NOT-HTR6	0.864865	0.8	0.941176
ADAM12 AND NOT-PCDHAC2 AND TSPAN16	0.857143	0.833333	0.882353
TGFBI AND NOT-ACPP AND PIRT	0.8	0.923077	0.705882
OSMR AND NOT-AGTR2 AND GRIK4	0.83871	0.928571	0.764706
ADAM12 AND OR4N4 AND NOT-HTR6	0.914286	0.888889	0.941176
ADAM12 AND NOT-MMP24 AND CALN1	0.871795	0.772727	1
ADAM12 AND SV2C AND NOT-AGTR2	0.871795	0.772727	1
TGFBI AND NOT-ACPP AND GPR156	0.8	0.923077	0.705882
ADAM12 AND GRIK4 AND NOT-HTR6	0.888889	0.842105	0.941176
ADAM12 AND PCDHGC4 AND NOT-AGTR2	0.871795	0.772727	1
ADAM12 AND CALN1 AND NOT-SLC22A6	1	1	1
ADAM12 AND OR4N4 AND NOT-SLC39A2	0.918919	0.85	1
ADAM12 AND PCDHGC4 AND NOT-NPBWR1	0.944444	0.894737	1
ADAM12 AND SV2C AND NOT-NPBWR1	0.944444	0.894737	1
ADAM12 AND GRIK4 AND NOT-AGER	0.871795	0.772727	1
ADAM12 AND GJA10 AND NOT-HTR6	0.820513	0.727273	0.941176
ADAM12 AND LRRC52 AND NOT-FXYD7	0.894737	0.809524	1
ADAM12 AND CRB2 AND NOT-HTR6	0.864865	0.8	0.941176
ADAM12 AND NOT-SLC22A6 AND JPH3	1	1	1
ADAM12 AND GJA10 AND NOT-SLC39A2	0.871795	0.772727	1
ADAM12 AND CALN1 AND NOT-HTR6	0.842105	0.761905	0.941176
ADAM12 AND NOT-SLC22A6 AND PLVAP	1	1	1
FAP AND NOT-SLMAP AND CASR	0.866667	1	0.764706
FAP AND NOT-SLMAP AND NOT-GABRB1	0.866667	1	0.764706
FAP AND NOT-ABCB1 AND GJA10	0.866667	1	0.764706
ADAM12 AND NOT-EPCAM AND GRIK4	0.918919	0.85	1
FAP AND NOT-SLMAP AND GJA10	0.866667	1	0.764706
FAP AND NOT-SLMAP AND ADCY10	0.866667	1	0.764706
FAP AND NOT-SLMAP AND PCDHGC4	0.866667	1	0.764706
FAP AND NOT-SLMAP AND PCDHAC2	0.866667	1	0.764706
FAP AND NOT-SLMAP AND SV2C	0.866667	1	0.764706
FAP AND NOT-ABCB1 AND OR4N4	0.866667	1	0.764706

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
NOT-PHLDB2 AND HCST AND TAAR5	0.964	0.967871	0.960159
NOT-PHLDB2 AND NOT-KIAA1324 AND LTBR	0.967742	0.979592	0.956175
NOT-PHLDB2 AND NOT-KIAA1324 AND LTBR	0.967742	0.979592	0.956175
NOT-PTPRK AND LTBR AND NOT-BTN3A1	0.963855	0.97166	0.956175
NOT-WLS AND SLC22A16 AND P2RX1	0.961303	0.983333	0.940239
NOT-PHLDB2 AND HCST AND NOT-CD27	0.961089	0.939163	0.984064
NOT-PHLDB2 AND CXCR4 AND NOT-P2RY10	0.961089	0.939163	0.984064
COMPLEX-S1PR1/FLT3/RHAG	0.961089	0.939163	0.984064
NOT-PHLDB2 AND HCST AND NOT-SIRPG	0.961089	0.939163	0.984064
NOT-PTPRK AND NOT-GPR171 AND SLC39A8	0.96063	0.949416	0.972112
NOT-PTPRK AND NOT-PHLDB2 AND TAAR5	0.960317	0.956522	0.964143
NOT-PTPRK AND NOT-PHLDB2 AND TAAR5	0.960317	0.956522	0.964143
NOT-PHLDB2 AND CXCR4 AND TAAR5	0.960317	0.956522	0.964143
NOT-PTPRK AND P2RX4 AND NOT-KLRD1	0.959677	0.971429	0.948207
NOT-WLS AND SLC22A16 AND HCST	0.959514	0.975309	0.944223
COMPLEX-FLT3/ATP8B1/RHAG	0.959381	0.932331	0.988048
NOT-PTPRK AND NOT-GPR171 AND CD44	0.959381	0.932331	0.988048
NOT-PTPRK AND NOT-GPR171 AND CD44	0.959381	0.932331	0.988048
COMPLEX-FLT3/GHR/RHAG	0.959381	0.932331	0.988048
NOT-WLS AND CSF3R AND NOT-PHLDB2	0.961924	0.967742	0.956175
NOT-PTPRK AND CD44 AND NOT-P2RY10	0.970646	0.953846	0.988048
NOT-ANTXR1 AND EMP3 AND NOT-P2RY10	0.960938	0.942529	0.98008
NOT-WLS AND LTBR AND NOT-PHLDB2	0.960825	0.995726	0.928287
COMPLEX-BTN3A1/FLT3/RHAG	0.96124	0.935849	0.988048
NOT-PODXL AND P2RX1 AND NOT-BTN3A1	0.957692	0.925651	0.992032
NOT-WLS AND SLC22A16 AND ATP6AP2	0.957576	0.971311	0.944223
NOT-WLS AND SLC22A16 AND ATP13A1	0.957576	0.971311	0.944223
NOT-WLS AND SLC22A16 AND LAMP1	0.957576	0.971311	0.944223
NOT-WLS AND SLC22A16 AND TGOLN2	0.957404	0.975207	0.940239
NOT-PTPRK AND P2RX4 AND NOT-CD247	0.957404	0.975207	0.940239
COMPLEX-FLT3/IFNAR2/RHAG	0.957364	0.932075	0.984064
NOT-WLS AND SLC22A16 AND VAMP2	0.95723	0.979167	0.936255
COMPLEX-FLT3/RHAG/PHLDB2	0.957198	0.935361	0.98008
COMPLEX-EFNB2/FLT3/RHAG	0.957031	0.938697	0.976096
NOT-PHLDB2 AND HCST AND NOT-GPR171	0.957031	0.938697	0.976096
NOT-PTPRK AND NOT-SIRPG AND P2RX4	0.957031	0.938697	0.976096
NOT-PHLDB2 AND HCST AND NOT-CD3G	0.957031	0.938697	0.976096
NOT-PTPRK AND P2RX4 AND NOT-CD8B	0.959184	0.983264	0.936255
NOT-PTPRK AND NOT-CD3G AND P2RX4	0.96063	0.949416	0.972112
NOT-PHLDB2 AND HCST AND LTBR	0.958743	0.945736	0.972112
NOT-PHLDB2 AND HCST AND LTBR	0.958743	0.945736	0.972112
COMPLEX-BTN3A3/FLT3/RHAG	0.959223	0.935606	0.984064
NOT-PHLDB2 AND HCST AND ATP2A2	0.962376	0.956693	0.968127
NOT-PHLDB2 AND HCST AND PIEZO1	0.956349	0.952569	0.960159
NOT-PHLDB2 AND CLEC12A AND NOT-KIAA1324	0.956	0.959839	0.952191
NOT-PTPRK AND EMP3 AND NOT-P2RY10	0.955854	0.922222	0.992032
NOT-PHLDB2 AND HCST AND NOT-KIAA1324	0.984127	0.980237	0.988048
NOT-PHLDB2 AND HCST AND NOT-KIAA1324	0.984127	0.980237	0.988048
NOT-PHLDB2 AND HCST AND NOT-PLXDC1	0.959381	0.932331	0.988048
COMPLEX-SLC31A1/FLT3/RHAG	0.955513	0.928571	0.984064
NOT-WLS AND SLC22A16 AND CSF3R	0.955466	0.971193	0.940239
NOT-PHLDB2 AND HCST AND TSPAN3	0.955285	0.975104	0.936255
NOT-PHLDB2 AND HCST AND TSPAN3	0.955285	0.975104	0.936255
NOT-PHLDB2 AND CXCR4 AND ESYT2	0.955166	0.935115	0.976096
NOT-PHLDB2 AND HCST AND NOT-CD2	0.955166	0.935115	0.976096
NOT-PTPRK AND P2RX4 AND NOT-CD8A	0.955102	0.979079	0.932271
NOT-PTPRK AND NOT-CD2 AND P2RX4	0.954918	0.983122	0.928287
NOT-MTUS1 AND CD46 AND NOT-BTN3A1	0.954813	0.94186	0.968127
NOT-SGCE AND EMP3 AND NOT-P2RY10	0.954635	0.945313	0.964143
FLT3 AND EMP3 AND NOT-LAMP3	0.954357	0.995671	0.916335
NOT-WLS AND CSF3R AND NOT-BTN3A3	0.954455	0.948819	0.960159
NOT-PHLDB2 AND HCST AND NOT-TIGIT	0.961089	0.939163	0.984064
NOT-PTPRK AND CD44 AND NOT-KLRB1	0.954092	0.956	0.952191
NOT-PTPRK AND CD44 AND NOT-KLRB1	0.954092	0.956	0.952191
NOT-PHLDB2 AND NOT-KIAA1324 AND PTGER2	0.966601	0.953488	0.98008
COMPLEX-FLT3/RHAG/SLC8A3	0.953846	0.921933	0.988048
COMPLEX-SEMA4D/FLT3/RHAG	0.953846	0.921933	0.988048
COMPLEX-FLT3/RHAG/KCNK6	0.953846	0.921933	0.988048
NOT-PHLDB2 AND HCST AND NOT-BTN3A2	0.956522	0.94902	0.964143
NOT-PTPRK AND P2RX4 AND NOT-P2RY10	0.953722	0.963415	0.944223
NOT-WLS AND SLC22A16 AND ITGB2	0.953722	0.963415	0.944223
NOT-PHLDB2 AND HCST AND KIT	0.953722	0.963415	0.944223
NOT-WLS AND SLC22A16 AND PLP2	0.953722	0.963415	0.944223
NOT-WLS AND SLC22A16 AND CXCR4	0.953722	0.963415	0.944223
NOT-WLS AND SLC22A16 AND CD53	0.953722	0.963415	0.944223
NOT-WLS AND SLC22A16 AND EVI2B	0.953722	0.963415	0.944223
COMPLEX-EMP3/FLT3/RHAG	0.953668	0.925094	0.984064

Antigen Pair Logic	F1	Precision	Recall
NOT-EPCAM AND VANGL1 AND NOT-ABCB1	0.882353	0.882353	0.882353
FAP AND NOT-SLMAP AND OR8D1	0.866667	1	0.764706
FAP AND NOT-SLMAP AND FXYD7	0.866667	1	0.764706
FAP AND NOT-SLMAP AND CRB1	0.866667	1	0.764706
FAP AND NOT-SLMAP AND ROS1	0.866667	1	0.764706
FAP AND NOT-SLMAP AND GHSR	0.866667	1	0.764706
FAP AND NOT-SLMAP AND LCT	0.866667	1	0.764706
FAP AND NOT-SLMAP AND NOT-DIO1	0.827586	1	0.705882
NOT-EPCAM AND SLC2A10 AND GRIK4	0.827586	1	0.705882
FAP AND NOT-SLMAP AND ACSL6	0.827586	1	0.705882
NOT-EPCAM AND VANGL1 AND NOT-LTA	0.882353	0.882353	0.882353
NOT-EPCAM AND SLC31A1 AND NOT-ABCB1	0.833333	0.789474	0.882353
FAP AND NOT-ATP13A5 AND NOT-CXADR	0.827586	1	0.705882
FAP AND NOT-SLMAP AND CCKAR	0.866667	1	0.764706
ADAM12 AND NOT-EPCAM AND ACSL6	0.820513	0.727273	0.941176
NOT-EPCAM AND VANGL1 AND NOT-CORIN	0.833333	0.789474	0.882353
FAP AND NOT-SLMAP AND FSHR	0.866667	1	0.764706
FAP AND NOT-SLMAP AND CRB2	0.866667	1	0.764706
FAP AND NOT-SLMAP AND NOT-OR4N4	0.866667	1	0.764706
ADAM12 AND NOT-CLDN8 AND PCDHGC4	0.871795	0.772727	1
NOT-EPCAM AND FAM57A AND NOT-GUCY2F	0.823529	0.823529	0.823529
TGFBI AND NOT-EPCAM AND NOT-ABCB1	0.9375	1	0.882353
NOT-EPCAM AND VANGL1 AND NOT-TSPAN32	0.909091	0.9375	0.882353
ADAM12 AND NOT-CLDN8 AND LCT	0.871795	0.772727	1
NOT-EPCAM AND SLC2A10 AND CALN1	0.827586	1	0.705882
NOT-EPCAM AND VANGL1 AND CD4	0.875	0.933333	0.823529
NOT-EPCAM AND SLC2A10 AND CRB1	0.827586	1	0.705882
NOT-EPCAM AND SLC2A10 AND GJA10	0.827586	1	0.705882
NOT-EPCAM AND FAM57A AND GJA10	0.823529	0.823529	0.823529
FAP AND NOT-ATP13A5 AND NOT-IL22RA1	0.827586	1	0.705882
DDR2 AND NOT-CLDN8 AND GJA10	0.8	0.923077	0.705882
DDR2 AND NOT-CLDN8 AND LCT	0.8	0.923077	0.705882
ATP8B2 AND NOT-ABCB1 AND NOT-EPCAM	0.8	0.923077	0.705882
ATP8B2 AND NOT-ABCB1 AND NOT-CLDN8	0.8	0.923077	0.705882
FAP AND NOT-AGTR2 AND GP5	0.827586	1	0.705882
FAP AND NOT-LTA AND LRRC52	0.866667	1	0.764706
NOT-EPCAM AND SLC2A10 AND KCNK10	0.827586	1	0.705882
FAP AND NOT-GUCY2F AND GPR160	0.827586	1	0.705882
FAP AND NOT-OR8D1 AND EDA	0.866667	1	0.764706
ADAM12 AND NOT-CLDN8 AND OR4N4	0.894737	0.809524	1
ADAM12 AND NOT-CLDN8 AND GJA10	0.829268	0.708333	1
FAP AND NOT-LTA AND EPHA10	0.866667	1	0.764706
FAP AND NOT-LTA AND CALN1	0.866667	1	0.764706
FAP AND NOT-LTA AND PCDHAC2	0.866667	1	0.764706
ADAM12 AND NOT-SLC34A2 AND PCDHGC4	0.842105	0.761905	0.941176
NOT-EPCAM AND SLC2A10 AND GUCY2F	0.827586	1	0.705882
NOT-EPCAM AND SLC2A10 AND GABRA5	0.827586	1	0.705882
OSMR AND NOT-EPCAM AND GRIK4	0.8125	0.866667	0.764706
DDR2 AND NOT-CLDN8 AND OR4N4	0.8	0.923077	0.705882
ADAM12 AND NOT-CLDN7 AND GABRB1	0.864865	0.8	0.941176
FAP AND NOT-OR8D1 AND ICOS	0.866667	1	0.764706
NOT-EPCAM AND SLC2A10 AND MPL	0.827586	1	0.705882
DDR2 AND NOT-CLDN8 AND GUCY2F	0.8	0.923077	0.705882
FAP AND NOT-OR8D1 AND KIR2DL3	0.866667	1	0.764706
FAP AND NOT-OR8D1 AND CADM2	0.866667	1	0.764706
FAP AND NOT-XCR1 AND NOT-ST14	0.827586	1	0.705882
FAP AND NOT-NPC1L1 AND LRRC52	0.866667	1	0.764706
FAP AND NOT-OR8D1 AND GPR63	0.866667	1	0.764706
FAP AND NOT-AGTR2 AND TAS2R9	0.827586	1	0.705882
OSMR AND NOT-CLDN7 AND NOT-GUCY2F	0.875	0.933333	0.823529
OSMR AND NOT-EPCAM AND ACSL6	0.8125	0.866667	0.764706
TGFBI AND NOT-SPINT1 AND CD34	0.9375	1	0.882353
FAP AND NOT-NPC1L1 AND SLC34A1	0.866667	1	0.764706
FAP AND NOT-NPC1L1 AND JPH3	0.866667	1	0.764706
NOT-EPCAM AND SLC2A10 AND OR4N4	0.827586	1	0.705882
ADAM12 AND NOT-ABCB1 AND CORIN	0.85	0.73913	1
ADAM12 AND NOT-ABCB1 AND GP5	0.871795	0.772727	1
ADAM12 AND NOT-ST14 AND CORIN	0.85	0.73913	1
ADAM12 AND NOT-ABCB1 AND AQP4	0.894737	0.809524	1
ADAM12 AND NOT-HPN AND CORIN	0.944444	0.894737	1
ADAM12 AND NOT-ST14 AND TAS2R40	0.894737	0.809524	1
ADAM12 AND NOT-ST14 AND AQP4	0.894737	0.809524	1
TSPAN4 AND NOT-GDPD2 AND GPR173	0.820513	0.727273	0.941176
ADAM12 AND GP5 AND NOT-OR7A17	0.918919	0.85	1
PLXND1 AND NOT-ABCB1 AND GP5	0.866667	1	0.764706
ADAM12 AND NOT-ST14 AND GPR63	0.85	0.73913	1
ADAM12 AND NOT-ABCB1 AND ICOS	0.85	0.73913	1

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
COMPLEX-NKAIN2/FLT3/RHAG	0.953668	0.925094	0.984064
NOT-PHLDB2 AND HCST AND NOT-KLRB1	0.953908	0.959677	0.948207
NOT-PTPRK AND CSF3R AND NOT-BTN3A1	0.972763	0.95057	0.996016
NOT-WLS AND CSF3R AND NOT-BTN3A1	0.956522	0.94902	0.964143
NOT-PHLDB2 AND NOT-KIAA1324 AND ESYT2	0.965657	0.979508	0.952191
NOT-PTPRK AND P2RX4 AND NOT-TIGIT	0.96146	0.979339	0.944223
NOT-PTPRK AND NOT-CD2 AND CD44	0.952941	0.938224	0.968127
NOT-PTPRK AND NOT-CD2 AND CD44	0.952941	0.938224	0.968127
COMPLEX-FLT3/FXYD6/RHAG	0.952941	0.938224	0.968127
P2RY8 AND NOT-PHLDB2 AND TAAR5	0.953722	0.963415	0.944223
P2RY8 AND NOT-PHLDB2 AND TAAR5	0.953722	0.963415	0.944223
NOT-PTPRK AND NOT-TIGIT AND CD44	0.957364	0.932075	0.984064
NOT-PTPRK AND NOT-TIGIT AND CD44	0.957364	0.932075	0.984064
FLT3 AND EMP3 AND NOT-ABHD6	0.952381	0.991379	0.916335
FLT3 AND NOT-TSPAN6 AND EMP3	0.952381	0.991379	0.916335
FLT3 AND NOT-PODXL AND EMP3	0.952381	0.991379	0.916335
FLT3 AND EMP3 AND NOT-FAT4	0.952381	0.991379	0.916335
FLT3 AND EMP3 AND NOT-HEPH	0.952381	0.991379	0.916335
NOT-EMP2 AND LTBR AND NOT-BTN3A1	0.952381	0.948617	0.956175
COMPLEX-EDNRB/FLT3/RHAG	0.963107	0.939394	0.988048
COMPLEX-FLT3/HLA-DOB/RHAG	0.959381	0.932331	0.988048
COMPLEX-ALDH1A1/FLT3/RHAG	0.958904	0.942308	0.976096
COMPLEX-FLT3/AXL/RHAG	0.957198	0.935361	0.98008
COMPLEX-FLT3/ABCA5/RHAG	0.957198	0.935361	0.98008
COMPLEX-FLT3/PCYT1A/RHAG	0.96124	0.935849	0.988048
COMPLEX-FLT3/RHAG/CLDN12	0.955513	0.928571	0.984064
COMPLEX-FLT3/OAS1/RHAG	0.95534	0.931818	0.98008
COMPLEX-EDNRB/FLT3/ST14	0.954545	0.991416	0.920319
COMPLEX-EDNRB/FLT3/PLXNA2	0.954545	0.991416	0.920319
COMPLEX-EDNRB/FLT3/TMPRSS2	0.954545	0.991416	0.920319
COMPLEX-SPON2/FLT3/RHAG	0.953846	0.921933	0.988048
COMPLEX-FLT3/RNF43/RHAG	0.953846	0.921933	0.988048
NOT-PTPRK AND CD37 AND NOT-P2RY10	0.953125	0.934866	0.972112
NOT-PTPRK AND CD37 AND NOT-P2RY10	0.953125	0.934866	0.972112
NOT-WLS AND CSF3R AND NOT-MS4A1	0.952756	0.941634	0.964143
COMPLEX-FLT3/RHAG/TNFRSF10A	0.952015	0.918519	0.988048
COMPLEX-FLT3/IL11RA/RHAG	0.952015	0.918519	0.988048
COMPLEX-FLT3/RHAG/VTCN1	0.952015	0.918519	0.988048
COMPLEX-FLT3/HSPA5/RHAG	0.952015	0.918519	0.988048
NOT-PTPRK AND CSF3R AND NOT-SLAMF7	0.950382	0.912088	0.992032
NOT-SGCE AND HCST AND NOT-CD160	0.95	0.918216	0.984064
COMPLEX-FLT3/RHAG/CD79B	0.949219	0.931034	0.968127
NOT-WLS AND CSF3R AND NOT-HLA-DOB	0.94902	0.934363	0.964143
NOT-WLS AND P2RX1 AND NOT-FCRL2	0.948819	0.937743	0.960159
NOT-WLS AND P2RX1 AND NOT-CD22	0.948819	0.937743	0.960159
NOT-WLS AND P2RX1 AND NOT-FCRL5	0.948819	0.937743	0.960159
NOT-WLS AND P2RX1 AND NOT-HLA-DOB	0.948819	0.937743	0.960159
NOT-PTPRK AND P2RX1 AND NOT-SLAMF7	0.948375	0.911765	0.988048
NOT-PODXL AND CSF3R AND NOT-CD160	0.948177	0.914815	0.984064
NOT-WLS AND P2RX1 AND NOT-CR2	0.946955	0.934109	0.960159
COMPLEX-FLT3/TMPRSS2/CLDN5	0.946721	0.974684	0.920319
NOT-PODXL AND HCST AND NOT-CD160	0.946565	0.908425	0.988048
NOT-PTPRK AND NOT-CD160 AND HCST	0.94636	0.911439	0.984064
COMPLEX-FLT3/RHAG/CD37	0.945525	0.923954	0.968127
NOT-DAG1 AND DDX3X AND NOT-P2RY10	0.954813	0.94186	0.968127
NOT-WLS AND CSF3R AND NOT-CR2	0.945313	0.927203	0.964143
NOT-WLS AND P2RX1 AND NOT-MS4A1	0.945098	0.930502	0.960159
NOT-WLS AND P2RX1 AND NOT-FCRL1	0.945098	0.930502	0.960159
NOT-PTPRK AND P2RX1 AND NOT-HLA-DOB	0.944972	0.902174	0.992032
NOT-PTPRK AND CSF3R AND NOT-MS4A1	0.944551	0.908088	0.984064
NOT-PTPRK AND CD70 AND NOT-CD247	0.944551	0.908088	0.984064
NOT-PTPRK AND CD70 AND NOT-CD247	0.944551	0.908088	0.984064
NOT-PODXL AND CSF3R AND NOT-MS4A1	0.944551	0.908088	0.984064
FLT3 AND CD37 AND NOT-PTPRK	0.943633	0.991228	0.900398
FLT3 AND CD37 AND NOT-PODXL	0.943633	0.991228	0.900398
NOT-WLS AND CSF3R AND NOT-FCRL2	0.94347	0.923664	0.964143
NOT-WLS AND CSF3R AND NOT-FCRL5	0.94347	0.923664	0.964143
NOT-PODXL AND P2RX1 AND NOT-HLA-DOB	0.943182	0.898917	0.992032
COMPLEX-FLT3/FOLR2/RHAG	0.943026	0.930233	0.956175
NOT-PTPRK AND P2RX1 AND NOT-FCRL5	0.942966	0.901818	0.988048
NOT-PODXL AND P2RX1 AND NOT-FCRL5	0.942966	0.901818	0.988048
COMPLEX-GPNMB/FLT3/RHAG	0.942801	0.933594	0.952191
COMPLEX-FLT3/ITGAV/RHAG	0.942574	0.937008	0.948207
NOT-PTPRK AND P2RX1 AND NOT-MS4A1	0.942529	0.907749	0.98008
NOT-PTPRK AND CD37 AND NOT-TIGIT	0.942085	0.913858	0.972112
NOT-PTPRK AND CD37 AND NOT-TIGIT	0.942085	0.913858	0.972112
NOT-WLS AND CSF3R AND NOT-FCRL1	0.941634	0.920152	0.964143

Antigen Pair Logic	F1	Precision	Recall
OSMR AND NOT-SLC19A3 AND NOT-ST14	0.848485	0.875	0.823529
ADAM12 AND NOT-ILDR1 AND KIR2DL3	0.944444	0.894737	1
ADAM12 AND NOT-ST14 AND TAS2R14	0.871795	0.772727	1
ADAM12 AND NOT-ST14 AND KCNH7	0.829268	0.708333	1
ADAM12 AND NOT-SPINT1 AND PRLR	0.971429	0.944444	1
ADAM12 AND NOT-SPINT1 AND TAS2R14	0.971429	0.944444	1
OSMR AND NOT-ABCB1 AND NOT-CXADR	0.875	0.933333	0.823529
OSMR AND NOT-SLC19A3 AND NOT-ACPP	0.875	0.933333	0.823529
ADAM12 AND NOT-UPK1B AND CORIN	0.848485	0.875	0.823529
ADAM12 AND GP5 AND NOT-ACPP	0.85	0.73913	1
ADAM12 AND NOT-ST14 AND OR10A4	0.829268	0.708333	1
DDR2 AND NOT-ABCB1 AND CORIN	0.827586	1	0.705882
ADAM12 AND NOT-ABCB1 AND GLRA1	0.810811	0.75	0.882353
ADAM12 AND NOT-SPINT1 AND KCNH7	0.918919	0.85	1
ADAM12 AND GP5 AND NOT-WNT1	0.85	0.73913	1
ADAM12 AND NOT-SPINT1 AND KIR2DL3	0.971429	0.944444	1
ADAM12 AND NOT-ST14 AND FLT3LG	0.85	0.73913	1
DDR2 AND NOT-WNT4 AND CORIN	0.8	0.923077	0.705882
ADAM12 AND NOT-ILDR1 AND CADM2	0.944444	0.894737	1
ADAM12 AND NOT-ST14 AND TAAR1	0.85	0.73913	1
OSMR AND NOT-ABCB1 AND NOT-SPINT1	0.875	0.933333	0.823529
ADAM12 AND NOT-ST14 AND GPR17	0.871795	0.772727	1
ADAM12 AND NOT-ST14 AND CADM2	0.894737	0.809524	1
ADAM12 AND NOT-ACPP AND OR5P2	0.918919	0.85	1
ADAM12 AND NOT-SPINT1 AND NOT-PAM	0.842105	0.761905	0.941176
ADAM12 AND NOT-SIGLEC6 AND NOT-GPR160	0.85	0.73913	1
PLXND1 AND NOT-ABCB1 AND CORIN	0.903226	1	0.823529
OSMR AND NOT-ABCB1 AND NOT-ST14	0.875	0.933333	0.823529
ADAM12 AND NOT-ILDR1 AND KCNK7	0.894737	0.809524	1
PLXND1 AND NOT-ABCB1 AND TAS2R9	0.866667	1	0.764706
DDR2 AND NOT-WNT4 AND ICOS	0.8	0.923077	0.705882
ADAM12 AND NOT-ST14 AND OR10H2	0.85	0.73913	1
ADAM12 AND NOT-ST14 AND TRPA1	0.85	0.73913	1
ADAM12 AND NOT-ST14 AND ICOS	0.809524	0.68	1
ADAM12 AND NOT-ABCB1 AND OR10H2	0.85	0.73913	1
ADAM12 AND GP5 AND NOT-GPR160	0.829268	0.708333	1
ADAM12 AND GP5 AND NOT-CLEC4M	0.809524	0.68	1
ADAM12 AND NOT-SPINT1 AND CDH5	0.971429	0.944444	1
OSMR AND NOT-ACPP AND NOT-GPR160	0.875	0.933333	0.823529
OSMR AND NOT-ACPP AND NOT-FUT1	0.903226	1	0.823529
ADAM12 AND NOT-ST14 AND CATSPER1	0.85	0.73913	1
ADAM12 AND NOT-ACPP AND NOT-FUT1	0.894737	0.809524	1
ADAM12 AND NOT-HPN AND CADM2	1	1	1
ADAM12 AND NOT-ST14 AND KIR2DL3	0.894737	0.809524	1
ADAM12 AND NOT-ACPP AND TAS2R14	0.918919	0.85	1
ADAM12 AND NOT-HPN AND TAS2R14	0.971429	0.944444	1
ADAM12 AND NOT-SPINT1 AND PRRG2	0.888889	0.842105	0.941176
COMPLEX-SFRP1/ST14/OSMR	0.83871	0.928571	0.764706
OSMR AND NOT-ABCB1 AND NOT-CLCA4	0.875	0.933333	0.823529
OSMR AND NOT-ABCB1 AND NOT-FGFR3	0.875	0.933333	0.823529
OSMR AND NOT-ACPP AND NOT-FPR2	0.827586	1	0.705882
ADAM12 AND NOT-SPINT1 AND AQP4	0.909091	0.9375	0.882353
COMPLEX-TM4SF1/ACPP/OSMR	0.827586	1	0.705882
STX2 AND NOT-CXADR AND NOT-ABCB1	0.9375	1	0.882353
ADAM12 AND NOT-MARVELD2 AND KIR2DL3	0.971429	0.944444	1
ADAM12 AND NOT-ST14 AND ESYT3	0.894737	0.809524	1
ADAM12 AND GP5 AND NOT-OR5K1	0.829268	0.708333	1
ADAM12 AND GP5 AND NOT-TLR9	0.829268	0.708333	1
ADAM12 AND NOT-MARVELD2 AND TAS2R9	0.864865	0.8	0.941176
ADAM12 AND GP5 AND NOT-GJB3	0.829268	0.708333	1
OSMR AND NOT-ACPP AND NOT-VSIG2	0.903226	1	0.823529
OSMR AND NOT-ACPP AND NOT-AQP4	0.903226	1	0.823529
OSMR AND NOT-ACPP AND NOT-NPC1L1	0.903226	1	0.823529
ADAM12 AND NOT-SPINT1 AND OR7A17	0.971429	0.944444	1
COMPLEX-ST14/KCNK5/OSMR	0.875	0.933333	0.823529
ADAM12 AND NOT-SPINT1 AND TAS2R9	0.888889	0.842105	0.941176
DDR2 AND NOT-ABCB1 AND ICOS	0.827586	1	0.705882
ADAM12 AND NOT-ST14 AND IL1RAPL1	0.809524	0.68	1
ADAM12 AND NOT-ST14 AND CDH26	0.829268	0.708333	1
ADAM12 AND NOT-SIGLEC6 AND NOT-ST14	0.85	0.73913	1
ADAM12 AND NOT-ST14 AND CHRM4	0.85	0.73913	1
DDR2 AND NOT-ABCB1 AND GP5	0.827586	1	0.705882
ADAM12 AND NOT-ST14 AND CD200R1	0.833333	0.789474	0.882353
ADAM12 AND GP5 AND NOT-FXYD4	0.871795	0.772727	1
ADAM12 AND GP5 AND NOT-OR2K2	0.894737	0.809524	1
ADAM12 AND GP5 AND NOT-OR10C1	0.871795	0.772727	1
OSMR AND NOT-ABCB1 AND CD63	0.83871	0.928571	0.764706

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
NOT-PTPRK AND NOT-GPR171 AND CD37	0.941634	0.920152	0.964143
NOT-PTPRK AND NOT-GPR171 AND CD37	0.941634	0.920152	0.964143
NOT-PTPRK AND CD37 AND NOT-CCR7	0.941406	0.923372	0.960159
NOT-WLS AND HCST AND NOT-CD160	0.941406	0.923372	0.960159
NOT-PODXL AND P2RX1 AND NOT-CD160	0.940952	0.90146	0.984064
NOT-PTPRK AND P2RX1 AND NOT-CD160	0.940727	0.904412	0.98008
NOT-PODXL AND P2RX1 AND NOT-MS4A1	0.940727	0.904412	0.98008
NOT-PERP AND HCST AND NOT-MS4A1	0.940727	0.904412	0.98008
NOT-WLS AND CSF3R AND NOT-TNFRSF17	0.940711	0.933333	0.948207
NOT-WLS AND P2RX1 AND NOT-TNFRSF17	0.940239	0.940239	0.940239
NOT-PTPRK AND CD70 AND NOT-KLRD1	0.943182	0.898917	0.992032
NOT-PTPRK AND CD70 AND NOT-KLRD1	0.943182	0.898917	0.992032
NOT-WLS AND CSF3R AND NOT-CD22	0.939806	0.916667	0.964143
NOT-PODXL AND CSF3R AND NOT-FCRL5	0.939623	0.892473	0.992032
NOT-WLS AND P2RX1 AND NOT-CXCR5	0.939571	0.919847	0.960159
NOT-PTPRK AND CD37 AND NOT-CD28	0.939571	0.919847	0.960159
NOT-PTPRK AND CD37 AND NOT-CD28	0.939571	0.919847	0.960159
NOT-PTPRK AND P2RX1 AND SLC7A5	0.94636	0.911439	0.984064
NOT-PTPRK AND P2RX1 AND SLC7A5	0.94636	0.911439	0.984064
NOT-MTUS1 AND CD46 AND NOT-CD160	0.939335	0.923077	0.956175
NOT-PTPRK AND CD70 AND NOT-CCR7	0.946565	0.908425	0.988048
NOT-PODXL AND NOT-LTB AND CD37	0.938776	0.962343	0.916335
NOT-PODXL AND NOT-LTB AND CD37	0.938776	0.962343	0.916335
NOT-PODXL AND CD37 AND NOT-CLEC2D	0.938462	0.907063	0.972112
NOT-PERP AND HCST AND NOT-CD160	0.938462	0.907063	0.972112
NOT-PTPRK AND NOT-TIGIT AND CD70	0.944762	0.905109	0.988048
NOT-PTPRK AND NOT-TIGIT AND CD70	0.944762	0.905109	0.988048
NOT-MTUS1 AND NOT-IL2RB AND DDX3X	0.93837	0.936508	0.940239
NOT-WLS AND CXCR4 AND NOT-CR2	0.938224	0.910112	0.968127
NOT-PODXL AND NOT-GPR171 AND DDX3X	0.938224	0.910112	0.968127
NOT-PTPRK AND CSF3R AND NOT-FCRL1	0.937984	0.913208	0.964143
NOT-DAG1 AND DDX3X AND NOT-GPR171	0.937984	0.913208	0.964143
NOT-WLS AND P2RX1 AND NOT-CD72	0.937743	0.91635	0.960159
NOT-PTPRK AND CD37 AND NOT-TRAT1	0.937743	0.91635	0.960159
NOT-PTPRK AND NOT-GPR171 AND ANXA1	0.937618	0.892086	0.988048
NOT-PODXL AND TAAR5 AND NOT-CD160	0.941406	0.923372	0.960159
NOT-WLS AND P2RX1 AND NOT-CD79A	0.9375	0.91954	0.956175
NOT-PERP AND HCST AND NOT-HLA-DOB	0.937381	0.894928	0.984064
NOT-PTPRK AND CD37 AND NOT-KLRD1	0.936902	0.900735	0.976096
NOT-PTPRK AND CSF3R AND NOT-FCRL5	0.942748	0.904762	0.984064
NOT-PTPRK AND CSF3R AND NOT-HLA-DOB	0.93633	0.883392	0.996016
NOT-PODXL AND NOT-GPR171 AND CD33	0.954455	0.948819	0.960159
NOT-PODXL AND NOT-GPR171 AND CD33	0.954455	0.948819	0.960159
NOT-PTPRK AND NOT-IL7R AND CD33	0.95069	0.941406	0.960159
NOT-PTPRK AND NOT-IL7R AND CD33	0.95069	0.941406	0.960159
NOT-PODXL AND CD33 AND NOT-BTN3A2	0.949698	0.95935	0.940239
NOT-PODXL AND CD33 AND NOT-KLRD1	0.949416	0.927757	0.972112
NOT-PTPRK AND NOT-BTN3A2 AND CD33	0.94929	0.966942	0.932271
NOT-PTPRK AND NOT-BTN3A2 AND CD33	0.94929	0.966942	0.932271
NOT-PODXL AND CD33 AND NOT-PTGDR	0.949219	0.931034	0.968127
NOT-PTPRK AND NOT-GPR171 AND CD33	0.948413	0.944664	0.952191
NOT-PTPRK AND NOT-GPR171 AND CD33	0.948413	0.944664	0.952191
NOT-PODXL AND CD33 AND NOT-CD8B	0.945313	0.927203	0.964143
NOT-PTPRK AND NOT-KLRF1 AND CD33	0.945098	0.930502	0.960159
NOT-PTPRK AND NOT-CD247 AND CD33	0.945098	0.930502	0.960159
NOT-PTPRK AND NOT-KLRF1 AND CD33	0.945098	0.930502	0.960159
NOT-PTPRK AND NOT-CD247 AND CD33	0.945098	0.930502	0.960159
NOT-PTPRK AND NOT-P2RY10 AND CD33	0.960317	0.956522	0.964143
NOT-PTPRK AND NOT-P2RY10 AND CD33	0.960317	0.956522	0.964143
NOT-PODXL AND CD33 AND NOT-P2RY10	0.943907	0.917293	0.972112
NOT-PODXL AND CD33 AND NOT-SIRPG	0.94347	0.923664	0.964143
NOT-PODXL AND CD33 AND NOT-SIRPG	0.94347	0.923664	0.964143
NOT-PODXL AND CD33 AND NOT-LY9	0.943907	0.917293	0.972112
NOT-PODXL AND CD33 AND NOT-SIT1	0.945525	0.923954	0.968127
NOT-PODXL AND CD33 AND NOT-KLRB1	0.942801	0.933594	0.952191
NOT-PODXL AND NOT-LTB AND CD33	0.942623	0.970464	0.916335
NOT-PODXL AND NOT-LTB AND CD33	0.942623	0.970464	0.916335
NOT-PTPRK AND NOT-KLRB1 AND CD33	0.942116	0.944	0.940239
NOT-PTPRK AND NOT-KLRB1 AND CD33	0.942116	0.944	0.940239
NOT-PTPRK AND NOT-CD8B AND CD33	0.940945	0.929961	0.952191
NOT-PTPRK AND NOT-CD8B AND CD33	0.940945	0.929961	0.952191
NOT-PODXL AND CD33 AND NOT-BTLA	0.947368	0.927481	0.968127
NOT-PODXL AND CD33 AND NOT-CD28	0.942801	0.933594	0.952191
NOT-PODXL AND NOT-AMIGO2 AND CD33	0.940711	0.933333	0.948207
NOT-PODXL AND NOT-AMIGO2 AND CD33	0.940711	0.933333	0.948207
NOT-PTPRK AND NOT-BTLA AND CD33	0.943249	0.926923	0.960159
NOT-PTPRK AND NOT-BTLA AND CD33	0.943249	0.926923	0.960159

Antigen Pair Logic	F1	Precision	Recall
OSMR AND NOT-CXADR AND NOT-ERVFRD-1	0.875	0.933333	0.823529
FAP AND NOT-ABCB1 AND ICOS	0.866667	1	0.764706
FAP AND NOT-ABCB1 AND CORIN	0.866667	1	0.764706
FAP AND NOT-ABCB1 AND OR10H2	0.866667	1	0.764706
FAP AND NOT-SLMAP AND OR10H2	0.866667	1	0.764706
FAP AND NOT-SLMAP AND GPR63	0.866667	1	0.764706
FAP AND NOT-SLMAP AND JPH2	0.866667	1	0.764706
FAP AND NOT-SLMAP AND TAS2R9	0.866667	1	0.764706
FAP AND NOT-SLMAP AND ICOS	0.866667	1	0.764706
FAP AND NOT-SLMAP AND SLC9A3	0.866667	1	0.764706
FAP AND NOT-SLMAP AND CORIN	0.866667	1	0.764706
ADAM12 AND NOT-EPCAM AND NOT-ABCB1	0.918919	0.85	1
FAP AND NOT-SLMAP AND CADM2	0.866667	1	0.764706
FAP AND NOT-ABCB1 AND NOT-CXADR	0.866667	1	0.764706
ADAM12 AND NOT-EPCAM AND SLC39A14	0.918919	0.85	1
FAP AND NOT-ABCB1 AND ST14	0.866667	1	0.764706
FAP AND NOT-SLMAP AND GP5	0.866667	1	0.764706
FAP AND NOT-SLMAP AND CHRM4	0.866667	1	0.764706
FAP AND NOT-SIGLEC6 AND GPR160	0.866667	1	0.764706
FAP AND NOT-SLMAP AND ABCC2	0.866667	1	0.764706
FAP AND NOT-SLMAP AND RTP1	0.866667	1	0.764706
FAP AND NOT-SLMAP AND OR7A17	0.866667	1	0.764706
FAP AND NOT-SLMAP AND SLC26A4	0.866667	1	0.764706
ADAM12 AND NOT-CLDN8 AND CORIN	0.871795	0.772727	1
FAP AND NOT-SLMAP AND NOT-ST14	0.866667	1	0.764706
FAP AND NOT-SLMAP AND NOT-GJB1	0.866667	1	0.764706
FAP AND NOT-SLMAP AND IL20RB	0.866667	1	0.764706
FAP AND NOT-SLMAP AND TAS2R40	0.866667	1	0.764706
FAP AND NOT-ABCB1 AND FGFR3	0.827586	1	0.705882
FAP AND NOT-SIGLEC6 AND IL22RA1	0.827586	1	0.705882
FAP AND NOT-ABCB1 AND TAS2R9	0.827586	1	0.705882
NOT-EPCAM AND SLC2A10 AND NOT-ABCB1	0.827586	1	0.705882
FAP AND NOT-ABCB1 AND GP5	0.827586	1	0.705882
ADAM12 AND NOT-SPINT1 AND NOT-CLDN6	0.918919	0.85	1
ADAM12 AND NOT-CLDN8 AND TAS2R40	0.894737	0.809524	1
FAP AND NOT-SLMAP AND CLEC12A	0.866667	1	0.764706
ADAM12 AND NOT-EPCAM AND GPR82	0.894737	0.809524	1
NOT-EPCAM AND FAM57A AND NOT-ABCB1	0.823529	0.823529	0.823529
FAP AND NOT-SLMAP AND FLT3LG	0.866667	1	0.764706
FAP AND NOT-SLMAP AND NOT-ERVFRD-1	0.866667	1	0.764706
NOT-EPCAM AND FAM57A AND KIR2DL3	0.848485	0.875	0.823529
ADAM12 AND NOT-EPCAM AND NOT-PRLR	0.820513	0.727273	0.941176
FAP AND NOT-SLMAP AND NOT-OR10A4	0.827586	1	0.705882
FAP AND NOT-SLMAP AND GRIN3B	0.827586	1	0.705882
FAP AND NOT-SLMAP AND TAS2R14	0.866667	1	0.764706
NOT-EPCAM AND FAM57A AND NOT-LTA	0.823529	0.823529	0.823529
ADAM12 AND GP5 AND NOT-SLC34A2	0.829268	0.708333	1
FAP AND NOT-HPN AND GPR160	0.866667	1	0.764706
ADAM12 AND NOT-CLDN8 AND AQP4	0.894737	0.809524	1
NOT-EPCAM AND FAM57A AND NOT-TSPAN32	0.823529	0.823529	0.823529
NOT-EPCAM AND FAM57A AND NOT-CORIN	0.823529	0.823529	0.823529
ADAM12 AND NOT-CLDN8 AND TAS2R14	0.894737	0.809524	1
NOT-EPCAM AND FAM57A AND NOT-SLC24A4	0.848485	0.875	0.823529
OSMR AND NOT-ABCB1 AND NOT-CLDN8	0.83871	0.928571	0.764706
NOT-EPCAM AND SLC2A10 AND SLC9A5	0.827586	1	0.705882
OSMR AND NOT-ACPP AND NOT-IL3RA	0.866667	1	0.764706
NOT-EPCAM AND SLC2A10 AND OR10H2	0.827586	1	0.705882
NOT-EPCAM AND SLC2A10 AND CORIN	0.827586	1	0.705882
ADAM12 AND GP5 AND NOT-CLDN7	0.829268	0.708333	1
DDR2 AND NOT-CLDN8 AND MUC15	0.8	0.923077	0.705882
DDR2 AND NOT-CLDN8 AND TAS2R40	0.8	0.923077	0.705882
NOT-EPCAM AND SLC2A10 AND GPR82	0.827586	1	0.705882
DDR2 AND NOT-CLDN8 AND PRLR	0.8	0.923077	0.705882
DDR2 AND NOT-CLDN8 AND CORIN	0.8	0.923077	0.705882
DDR2 AND NOT-CLDN8 AND KIR2DL3	0.8	0.923077	0.705882
OSMR AND NOT-SPINT1 AND NOT-CLDN6	0.875	0.933333	0.823529
DDR2 AND NOT-CLDN8 AND TAS2R9	0.8	0.923077	0.705882
DDR2 AND NOT-CLDN8 AND CD4	0.8	0.923077	0.705882
NOT-EPCAM AND FAM57A AND NOT-OR7A17	0.848485	0.875	0.823529
NOT-EPCAM AND FAM57A AND NOT-TAS2R9	0.8125	0.866667	0.764706
OSMR AND NOT-ABCB1 AND NOT-EPCAM	0.875	0.933333	0.823529
FAP AND NOT-SLMAP AND NOT-GPR160	0.866667	1	0.764706
ADAM12 AND GP5 AND NOT-SSTR5	0.871795	0.772727	1
DDR2 AND NOT-CLDN8 AND TAS2R14	0.8	0.923077	0.705882
DDR2 AND NOT-CLDN8 AND OR7A17	0.8	0.923077	0.705882
DDR2 AND NOT-CLDN8 AND FLT3LG	0.8	0.923077	0.705882
FAP AND NOT-KCNMB2 AND NOT-ACPP	0.866667	1	0.764706

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
NOT-PODXL AND CD33 AND NOT-CD8A	0.939335	0.923077	0.956175
NOT-PTPRK AND CD33 AND NOT-CLEC2D	0.945098	0.930502	0.960159
NOT-PODXL AND CD33 AND NOT-CLEC4A	0.955645	0.967347	0.944223
NOT-SGCB AND CD33 AND NOT-KLRF1	0.938614	0.933071	0.944223
NOT-PODXL AND CD33 AND NOT-IL7R	0.938224	0.910112	0.968127
NOT-PTPRK AND NOT-TIGIT AND CD33	0.947368	0.927481	0.968127
NOT-PTPRK AND NOT-TIGIT AND CD33	0.947368	0.927481	0.968127
NOT-PODXL AND CD33 AND NOT-FCRL3	0.947368	0.927481	0.968127
NOT-PTPRK AND NOT-SIRPG AND CD33	0.9375	0.91954	0.956175
NOT-PTPRK AND NOT-SIRPG AND CD33	0.9375	0.91954	0.956175
NOT-PODXL AND CD33 AND NOT-HEG1	0.9375	0.91954	0.956175
NOT-PODXL AND CD33 AND NOT-HEG1	0.9375	0.91954	0.956175
NOT-PTPRK AND NOT-CD8A AND CD33	0.937008	0.92607	0.948207
NOT-PTPRK AND NOT-CD8A AND CD33	0.937008	0.92607	0.948207
NOT-PODXL AND CD33 AND NOT-SLAMF6	0.942085	0.913858	0.972112
NOT-PTPRK AND CD33 AND NOT-STX7	0.965795	0.97561	0.956175
NOT-PODXL AND CD33 AND NOT-KLRF1	0.936508	0.932806	0.940239
NOT-PTPRK AND CD33 AND NOT-CLEC4A	0.959514	0.975309	0.944223
NOT-PODXL AND CD33 AND NOT-CD3G	0.93617	0.909774	0.964143
NOT-PTPRK AND CD33 AND NOT-SIT1	0.941406	0.923372	0.960159
NOT-PTPRK AND CD33 AND NOT-SIT1	0.941406	0.923372	0.960159
NOT-PTPRK AND CD33 AND NOT-PAG1	0.935167	0.922481	0.948207
NOT-PODXL AND CD33 AND NOT-STX7	0.968	0.971888	0.964143
NOT-PODXL AND CD33 AND NOT-CD247	0.934615	0.903346	0.968127
NOT-PODXL AND CD33 AND NOT-TIGIT	0.934615	0.903346	0.968127
NOT-PODXL AND CD33 AND NOT-TIGIT	0.934615	0.903346	0.968127
NOT-PTPRK AND NOT-WLS AND CD33	0.934132	0.936	0.932271
NOT-PTPRK AND NOT-WLS AND CD33	0.934132	0.936	0.932271
NOT-PODXL AND CD33 AND NOT-CLEC2D	0.947368	0.927481	0.968127
NOT-PODXL AND NOT-CD27 AND CD33	0.933868	0.939516	0.928287
NOT-PODXL AND CD33 AND NOT-NPC1	0.934911	0.925781	0.944223
NOT-PODXL AND CD33 AND NOT-ADTRP	0.938462	0.907063	0.972112
NOT-PODXL AND CD33 AND NOT-CD40LG	0.945525	0.923954	0.968127
NOT-PODXL AND CD33 AND NOT-ST3GAL5	0.939335	0.923077	0.956175
NOT-PTPRK AND CD33 AND NOT-CD300A	0.931393	0.973913	0.89243
NOT-PTPRK AND CD33 AND NOT-CCR7	0.946535	0.940945	0.952191
NOT-PODXL AND CD33 AND NOT-CD5	0.945313	0.927203	0.964143
NOT-SGCB AND CD33 AND NOT-SIRPG	0.930966	0.921875	0.940239
NOT-PODXL AND CD33 AND NOT-PAG1	0.930502	0.902622	0.960159
NOT-PTPRK AND CD33 AND NOT-TNFSF10	0.939516	0.95102	0.928287
NOT-PTPRK AND NOT-CD3G AND CD33	0.930233	0.90566	0.956175
NOT-PTPRK AND NOT-CD3G AND CD33	0.930233	0.90566	0.956175
NOT-SGCB AND CD33 AND NOT-CD8B	0.930966	0.921875	0.940239
NOT-PTPRK AND CD33 AND NOT-IL18R1	0.930233	0.90566	0.956175
NOT-PTPRK AND NOT-CD40LG AND CD33	0.946955	0.934109	0.960159
NOT-PTPRK AND NOT-CD40LG AND CD33	0.946955	0.934109	0.960159
NOT-SGCB AND CD33 AND NOT-P2RY10	0.937876	0.943548	0.932271
NOT-PODXL AND NOT-IL2RB AND CD33	0.929134	0.918288	0.940239
NOT-PTPRK AND NOT-NPC1 AND CD33	0.929134	0.918288	0.940239
NOT-PODXL AND NOT-IL2RB AND CD33	0.929134	0.918288	0.940239
NOT-PTPRK AND NOT-NPC1 AND CD33	0.929134	0.918288	0.940239
NOT-PTPRK AND CD33 AND NOT-CD5	0.928709	0.899254	0.960159
NOT-PODXL AND CD33 AND NOT-CCR6	0.929254	0.893382	0.968127
NOT-PODXL AND CD33 AND NOT-ITGAL	0.927875	0.908397	0.948207
NOT-PODXL AND CD33 AND NOT-EPHA4	0.927757	0.887273	0.972112
NOT-PTPRK AND CD33 AND NOT-TGFBR2	0.960317	0.956522	0.964143
NOT-PTPRK AND NOT-CD27 AND CD33	0.931452	0.942857	0.920319
NOT-TMEM47 AND NOT-IL2RB AND CD33	0.927308	0.914729	0.940239
NOT-PODXL AND CD33 AND NOT-HMOX2	0.927203	0.892989	0.964143
NOT-PTPRK AND CD33 AND NOT-DPP4	0.927203	0.892989	0.964143
NOT-PODXL AND CD33 AND NOT-ICOS	0.926733	0.92126	0.932271
NOT-PTPRK AND CD33 AND NOT-ADTRP	0.934363	0.906367	0.964143
NOT-PODXL AND CD33 AND NOT-HVCN1	0.92562	0.961373	0.89243
NOT-PTPRK AND CD33 AND NOT-HMOX2	0.954092	0.956	0.952191
NOT-PTPRK AND NOT-PHLDB2 AND P2RX4	0.96063	0.949416	0.972112
NOT-PHLDB2 AND EMP3 AND NOT-P2RY10	0.967742	0.979592	0.956175
NOT-WLS AND SLC22A16 AND VAMP8	0.957576	0.971311	0.944223
NOT-PHLDB2 AND EMP3 AND TAAR5	0.95723	0.979167	0.936255
NOT-PHLDB2 AND EMP3 AND NOT-SLC44A2	0.96	0.963855	0.956175
NOT-PHLDB2 AND HCST AND NOT-LAT	0.957031	0.938697	0.976096
NOT-PHLDB2 AND NOT-KIAA1324 AND P2RX4	0.956863	0.942085	0.972112
NOT-WLS AND SLC22A16 AND EMP3	0.955466	0.971193	0.940239
NOT-PHLDB2 AND HCST AND P2RX4	0.964427	0.956863	0.972112
NOT-PHLDB2 AND HCST AND P2RX4	0.964427	0.956863	0.972112
NOT-PHLDB2 AND NOT-SLC4A4 AND P2RX4	0.954635	0.945313	0.964143
NOT-PHLDB2 AND NOT-SLC4A4 AND P2RX4	0.954635	0.945313	0.964143
NOT-PHLDB2 AND CXCR4 AND P2RX4	0.96063	0.949416	0.972112

Antigen Pair Logic	F1	Precision	Recall
FAP AND NOT-NPC1L1 AND EDA	0.866667	1	0.764706
ADAM12 AND NOT-ST14 AND FCRL1	0.809524	0.68	1
ADAM12 AND GP5 AND NOT-MUC16	0.918919	0.85	1
FAP AND NOT-LTA AND TAS2R9	0.866667	1	0.764706
FAP AND NOT-LTA AND IL1RAPL1	0.866667	1	0.764706
FAP AND NOT-LTA AND CADM2	0.866667	1	0.764706
NOT-EPCAM AND SLC2A10 AND KIR2DL3	0.827586	1	0.705882
OSMR AND NOT-FOLR1 AND NOT-ST14	0.83871	0.928571	0.764706
OSMR AND NOT-EPCAM AND NOT-PRLR	0.83871	0.928571	0.764706
NOT-EPCAM AND SLC2A10 AND GJD4	0.827586	1	0.705882
OSMR AND NOT-CXADR AND NOT-CLDN6	0.875	0.933333	0.823529
NOT-EPCAM AND SLC2A10 AND TAS2R9	0.827586	1	0.705882
FAP AND NOT-NPC1L1 AND GPR160	0.866667	1	0.764706
FAP AND NOT-NPC1L1 AND ICOS	0.866667	1	0.764706
FAP AND NOT-PTGER3 AND NOT-ACPP	0.827586	1	0.705882
FAP AND NOT-MFSD3 AND CADM2	0.827586	1	0.705882
NOT-EPCAM AND FAM57A AND NOT-MUC15	0.8125	0.866667	0.764706
FAP AND NOT-MFSD3 AND IL1RAPL1	0.827586	1	0.705882
ADAM12 AND NOT-VTCN1 AND CADM2	0.871795	0.772727	1
ADAM12 AND NOT-CLDN8 AND FLT3LG	0.871795	0.772727	1
FAP AND NOT-EPCAM AND CD4	0.866667	1	0.764706
FAP AND NOT-ABCB1 AND NOT-EPCAM	0.866667	1	0.764706
FAP AND NOT-SLMAP AND NOT-CLDN7	0.866667	1	0.764706
FAP AND NOT-SLMAP AND NOT-SLC34A2	0.866667	1	0.764706
FAP AND NOT-CLDN6 AND NOT-SPINT1	0.866667	1	0.764706
FAP AND NOT-CLDN6 AND NOT-ACPP	0.866667	1	0.764706
FAP AND NOT-CLDN8 AND CORIN	0.827586	1	0.705882
FAP AND NOT-CLDN8 AND KIR2DL3	0.827586	1	0.705882
FAP AND NOT-CLDN8 AND CD4	0.827586	1	0.705882
FAP AND NOT-CLDN8 AND AQP4	0.827586	1	0.705882
FAP AND NOT-CLDN8 AND TAS2R40	0.827586	1	0.705882
FAP AND NOT-CLDN6 AND NOT-ST14	0.866667	1	0.764706
FAP AND NOT-CLDN6 AND NOT-CDHR1	0.866667	1	0.764706
FAP AND NOT-CLDN8 AND FLT3LG	0.827586	1	0.705882
NOT-EPCAM AND FAM57A AND NOT-CXCR5	0.823529	0.823529	0.823529
FAP AND NOT-EPCAM AND ITGB2	0.827586	1	0.705882
FAP AND NOT-TSPAN32 AND NOT-EPCAM	0.827586	1	0.705882
DDR2 AND NOT-CLDN8 AND MUC4	0.8	0.923077	0.705882
FAP AND NOT-LTA AND NOT-EPCAM	0.866667	1	0.764706
FAP AND NOT-EPCAM AND NOT-TAS2R14	0.866667	1	0.764706
NOT-EPCAM AND SLC2A10 AND CD34	0.827586	1	0.705882
NOT-EPCAM AND SLC2A10 AND MUC4	0.827586	1	0.705882
ADAM12 AND NOT-CLDN8 AND MUC4	0.85	0.73913	1
FAP AND NOT-NPC1L1 AND MUC4	0.866667	1	0.764706
NOT-EPCAM AND FOLH1 AND CD63	0.823529	0.823529	0.823529
FAP AND NOT-EPCAM AND NOT-OR7A17	0.866667	1	0.764706
FAP AND NOT-EPCAM AND TGFA	0.866667	1	0.764706
FAP AND NOT-EPCAM AND KIR2DL3	0.866667	1	0.764706
FAP AND NOT-EPCAM AND NOT-SLC9A3	0.827586	1	0.705882
FAP AND NOT-EPCAM AND NOT-CDH26	0.827586	1	0.705882
FAP AND NOT-EPCAM AND TAS2R9	0.866667	1	0.764706
FAP AND NOT-EPCAM AND NRG1	0.866667	1	0.764706
FAP AND NOT-EPCAM AND NOT-OR10H2	0.827586	1	0.705882
FAP AND NOT-GRIN3B AND MUC4	0.866667	1	0.764706
ADAM12 AND NOT-CLDN7 AND NOT-MUC16	0.871795	0.772727	1
FAP AND NOT-TSPAN32 AND CLDN7	0.827586	1	0.705882
VMP1 AND NOT-EPCAM AND CD34	0.8125	0.866667	0.764706
FAP AND NOT-EPCAM AND TLR4	0.827586	1	0.705882
ADAM12 AND NOT-CLDN7 AND MUC4	0.829268	0.708333	1
FAP AND NOT-TAS2R14 AND NOT-SLC34A2	0.866667	1	0.764706
FAP AND NOT-OR7A17 AND NOT-VTCN1	0.866667	1	0.764706
FAP AND NOT-SSTR5 AND TAS2R9	0.866667	1	0.764706
FAP AND NOT-TAS2R14 AND NOT-CLDN7	0.866667	1	0.764706
FAP AND NOT-CXCR5 AND ICOS	0.866667	1	0.764706
FAP AND NOT-CXCR5 AND OR10H2	0.866667	1	0.764706
FAP AND NOT-SSTR5 AND GP5	0.866667	1	0.764706
FAP AND NOT-TAS2R14 AND NOT-VTCN1	0.827586	1	0.705882
FAP AND NOT-CXCR5 AND GPR63	0.866667	1	0.764706
FAP AND NOT-TAS2R14 AND MUC16	0.866667	1	0.764706
SLC38A6 AND NOT-EPCAM AND MUC4	0.8125	0.866667	0.764706
FAP AND NOT-CXCR5 AND GP5	0.866667	1	0.764706
FAP AND NOT-CXCR5 AND IL1RAPL1	0.866667	1	0.764706
NOT-EPCAM AND GPNMB AND CD4	0.882353	0.882353	0.882353
FAP AND NOT-VTCN1 AND GPR63	0.827586	1	0.705882
FAP AND NOT-VTCN1 AND CADM2	0.827586	1	0.705882
FAP AND NOT-TAS2R14 AND CXCR5	0.827586	1	0.705882
FAP AND NOT-MUC16 AND ICOS	0.866667	1	0.764706

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
NOT-PHLDB2 AND CXCR4 AND P2RX4	0.96063	0.949416	0.972112
NOT-PHLDB2 AND EMP3 AND NOT-BTN3A2	0.953722	0.963415	0.944223
NOT-WLS AND SLC22A16 AND LAPTM5	0.953722	0.963415	0.944223
NOT-PHLDB2 AND HCST AND SLC39A8	0.96146	0.979339	0.944223
NOT-PHLDB2 AND NOT-ABHD6 AND P2RX4	0.953125	0.934866	0.972112
NOT-PHLDB2 AND NOT-ABHD6 AND P2RX4	0.953125	0.934866	0.972112
SLC22A16 AND NOT-PHLDB2 AND FXYD5	0.952577	0.987179	0.920319
FLT3 AND EMP3 AND NOT-ATP8B1	0.952381	0.991379	0.916335
FLT3 AND EMP3 AND NOT-GHR	0.952381	0.991379	0.916335
NOT-PHLDB2 AND NOT-SLC25A4 AND P2RX4	0.956863	0.942085	0.972112
NOT-PHLDB2 AND NOT-SLC25A4 AND P2RX4	0.956863	0.942085	0.972112
NOT-PHLDB2 AND HCST AND SLC30A1	0.952015	0.918519	0.988048
NOT-PHLDB2 AND HCST AND SLC30A1	0.952015	0.918519	0.988048
NOT-PHLDB2 AND CXCR4 AND NOT-SEMA4D	0.95183	0.921642	0.984064
P2RY8 AND NOT-PHLDB2 AND P2RX4	0.956175	0.956175	0.956175
P2RY8 AND NOT-PHLDB2 AND P2RX4	0.956175	0.956175	0.956175
SLC22A16 AND HCST AND NOT-PHLDB2	0.950617	0.982979	0.920319
COMPLEX-FLT3/ATP8B1/CD44	0.950617	0.982979	0.920319
SLC22A16 AND NOT-PHLDB2 AND CD69	0.950413	0.987124	0.916335
NOT-PTPRK AND CD44 AND NOT-BTN3A1	0.950192	0.915129	0.988048
NOT-PHLDB2 AND ITGA5 AND P2RX4	0.951417	0.967078	0.936255
NOT-PTPRK AND NOT-LAT AND SLC30A1	0.957529	0.928839	0.988048
NOT-PTPRK AND NOT-LAT AND SLC30A1	0.957529	0.928839	0.988048
NOT-PHLDB2 AND NOT-KCNK1 AND P2RX4	0.9501	0.952	0.948207
NOT-PHLDB2 AND P2RX4 AND NOT-TSPAN15	0.949416	0.927757	0.972112
NOT-PHLDB2 AND NOT-MRAP2 AND P2RX4	0.949416	0.927757	0.972112
NOT-PHLDB2 AND NOT-MRAP2 AND P2RX4	0.949416	0.927757	0.972112
NOT-PHLDB2 AND P2RX4 AND NOT-TSPAN15	0.949416	0.927757	0.972112
NOT-PHLDB2 AND HCST AND NOT-BTN3A1	0.966862	0.946565	0.988048
SLC22A16 AND NOT-LAMP3 AND NOT-GHR	0.94929	0.966942	0.932271
SLC22A16 AND NOT-TSPAN1 AND NOT-PHLDB2	0.948665	0.978814	0.920319
SLC22A16 AND P2RY8 AND NOT-PHLDB2	0.948665	0.978814	0.920319
SLC22A16 AND NOT-SLC6A16 AND NOT-PHLDB2	0.948665	0.978814	0.920319
SLC22A16 AND NOT-LAMP3 AND NOT-PHLDB2	0.948665	0.978814	0.920319
SLC22A16 AND ICAM3 AND NOT-PHLDB2	0.948665	0.978814	0.920319
NOT-PTPRK AND NOT-PHLDB2 AND CD44	0.948375	0.911765	0.988048
NOT-PHLDB2 AND CXCR4 AND NOT-BTN3A1	0.948375	0.911765	0.988048
NOT-PTPRK AND NOT-PHLDB2 AND CD44	0.948375	0.911765	0.988048
NOT-PHLDB2 AND NOT-KIAA1324 AND VAMP8	0.961089	0.939163	0.984064
NOT-PHLDB2 AND NOT-KIAA1324 AND VAMP8	0.961089	0.939163	0.984064
NOT-PHLDB2 AND CXCR4 AND NOT-SEMA4B	0.948177	0.914815	0.984064
NOT-PTPRK AND NOT-LAT AND P2RX4	0.947791	0.955466	0.940239
SLC22A16 AND NOT-ATP8B1 AND NOT-EPHA4	0.947154	0.966805	0.928287
SLC22A16 AND NOT-TMEM231 AND NOT-PHLDB2	0.946721	0.974684	0.920319
SLC22A16 AND NOT-TSPAN6 AND NOT-PHLDB2	0.946721	0.974684	0.920319
SLC22A16 AND NOT-LSR AND NOT-PHLDB2	0.946721	0.974684	0.920319
SLC22A16 AND NOT-TUSC3 AND NOT-PHLDB2	0.946721	0.974684	0.920319
NOT-PHLDB2 AND HCST AND VAMP8	0.946565	0.908425	0.988048
NOT-PHLDB2 AND HCST AND VAMP8	0.946565	0.908425	0.988048
P2RY8 AND NOT-PHLDB2 AND SLC39A8	0.946535	0.940945	0.952191
SLC22A16 AND CXCR4 AND NOT-PHLDB2	0.946502	0.978723	0.916335
SLC22A16 AND EVI2B AND NOT-PHLDB2	0.946502	0.978723	0.916335
NOT-ATP8B1 AND CD44 AND NOT-KLRB1	0.946322	0.944444	0.948207
NOT-PHLDB2 AND TAAR5 AND NOT-FAT1	0.9499	0.955645	0.944223
NOT-PHLDB2 AND GLIPR1 AND P2RX4	0.945892	0.951613	0.940239
NOT-PHLDB2 AND SELL AND P2RX4	0.950298	0.948413	0.952191
NOT-PHLDB2 AND SELL AND P2RX4	0.950298	0.948413	0.952191
P2RY8 AND NOT-PHLDB2 AND SLC30A1	0.945736	0.920755	0.972112
P2RY8 AND NOT-PHLDB2 AND SLC30A1	0.945736	0.920755	0.972112
NOT-ATP8B1 AND CD44 AND NOT-GPR171	0.945736	0.920755	0.972112
NOT-PHLDB2 AND NOT-PTPRF AND P2RX4	0.945736	0.920755	0.972112
NOT-PHLDB2 AND NOT-PTPRF AND P2RX4	0.945736	0.920755	0.972112
NOT-PHLDB2 AND EVI2B AND SLC30A1	0.945946	0.917603	0.976096
NOT-PHLDB2 AND EVI2B AND SLC30A1	0.945946	0.917603	0.976096
NOT-PHLDB2 AND TAAR5 AND P2RX4	0.945892	0.951613	0.940239
NOT-SGCE AND EMP3 AND NOT-BTN3A1	0.951456	0.92803	0.976096
NOT-PHLDB2 AND NOT-LRIG3 AND LTBR	0.945098	0.930502	0.960159
NOT-PHLDB2 AND NOT-LRIG3 AND LTBR	0.945098	0.930502	0.960159
NOT-PHLDB2 AND EMP3 AND NOT-LY9	0.945098	0.930502	0.960159
NOT-PHLDB2 AND NOT-LRIG3 AND LTBR	0.945098	0.930502	0.960159
NOT-PHLDB2 AND FXYD5 AND P2RX4	0.964427	0.956863	0.972112
NOT-PHLDB2 AND FXYD5 AND P2RX4	0.964427	0.956863	0.972112
NOT-PHLDB2 AND EMP3 AND NOT-SIRPG	0.944882	0.933852	0.956175
NOT-PHLDB2 AND EMP3 AND NOT-CD27	0.944882	0.933852	0.956175
NOT-PHLDB2 AND FXYD5 AND SLC30A1	0.955684	0.925373	0.988048
NOT-PHLDB2 AND FXYD5 AND SLC30A1	0.955684	0.925373	0.988048
SLC22A16 AND NOT-PSENEN AND NOT-PHLDB2	0.944785	0.970588	0.920319

Antigen Pair Logic	F1	Precision	Recall
ROR1 AND NOT-ABCB1 AND NOT-EPCAM	0.866667	1	0.764706
ROR1 AND NOT-CLDN6 AND NOT-CXADR	0.866667	1	0.764706
ROR1 AND NOT-CLDN6 AND NOT-SPINT1	0.866667	1	0.764706
ROR1 AND NOT-CLDN6 AND NOT-ACPP	0.866667	1	0.764706
FAP AND NOT-CXCR5 AND NOT-ST14	0.866667	1	0.764706
FAP AND NOT-SLC34A2 AND NOT-TAS2R9	0.827586	1	0.705882
FAP AND NOT-CXCR5 AND CHRM4	0.866667	1	0.764706
FAP AND NOT-AGTR2 AND PCDHGC4	0.866667	1	0.764706
FAP AND NOT-GUCY2F AND FXYD7	0.827586	1	0.705882
FAP AND NOT-AGTR2 AND PCDHAC2	0.866667	1	0.764706
FAP AND NOT-OR8D1 AND LRRC52	0.866667	1	0.764706
FAP AND NOT-OR8D1 AND SLC34A1	0.866667	1	0.764706
FAP AND NOT-OR8D1 AND PCDHAC2	0.866667	1	0.764706
FAP AND NOT-OR8D1 AND EPHA10	0.866667	1	0.764706
FAP AND NOT-OR8D1 AND GJA10	0.866667	1	0.764706
FAP AND NOT-TSPAN16 AND PCDHAC2	0.866667	1	0.764706
FAP AND NOT-OR8D1 AND SLC5A8	0.827586	1	0.705882
TGFBI AND NOT-EPCAM AND NOT-ACSL6	0.909091	0.9375	0.882353
FAP AND NOT-CRB2 AND HTR6	0.827586	1	0.705882
FAP AND NOT-SLC39A2 AND GJA10	0.866667	1	0.764706
FAP AND NOT-SLC39A2 AND FXYD7	0.866667	1	0.764706
FAP AND NOT-SLC39A2 AND PCDHGC4	0.866667	1	0.764706
NOT-EPCAM AND SLC31A1 AND NOT-ACSL6	0.833333	0.789474	0.882353
FAP AND NOT-NPBWR1 AND SV2C	0.866667	1	0.764706
FAP AND NOT-SLC39A2 AND PLVAP	0.866667	1	0.764706
FAP AND NOT-NPBWR1 AND GHSR	0.866667	1	0.764706
FAP AND NOT-NPBWR1 AND CALN1	0.866667	1	0.764706
FAP AND NOT-NPBWR1 AND PCDHGC4	0.866667	1	0.764706
FAP AND NOT-HTR6 AND PCDHGC4	0.827586	1	0.705882
FAP AND NOT-HTR6 AND SV2C	0.827586	1	0.705882
FAP AND NOT-HTR6 AND CALN1	0.827586	1	0.705882
ROR1 AND NOT-AGTR2 AND PLVAP	0.866667	1	0.764706
FAP AND NOT-NPBWR1 AND ADCY10	0.866667	1	0.764706
FAP AND NOT-HTR6 AND GJA10	0.827586	1	0.705882
ROR1 AND NOT-OR8D1 AND PLVAP	0.866667	1	0.764706
FAP AND NOT-HTR6 AND PLVAP	0.827586	1	0.705882
FAP AND NOT-NPBWR1 AND ROS1	0.866667	1	0.764706
ROR1 AND NOT-SLC22A6 AND PLVAP	0.866667	1	0.764706
ROR1 AND NOT-GPR78 AND PLVAP	0.866667	1	0.764706
ROR1 AND PLVAP AND NOT-SLC39A2	0.866667	1	0.764706
ROR1 AND NOT-CCR9 AND PLVAP	0.866667	1	0.764706
ROR1 AND NOT-AGTR2 AND SV2C	0.827586	1	0.705882
ROR1 AND NOT-OR8D1 AND LRRC52	0.866667	1	0.764706
FAP AND NOT-AGER AND ROS1	0.866667	1	0.764706
STAB1 AND FSHR AND NOT-CXCR5	0.827586	1	0.705882
ROR1 AND NOT-AGTR2 AND PCDHGC4	0.866667	1	0.764706
ROR1 AND NOT-OR8D1 AND EPHA10	0.866667	1	0.764706
ROR1 AND NOT-AGTR2 AND PCDHAC2	0.866667	1	0.764706
ROR1 AND NOT-OR8D1 AND SLC34A1	0.866667	1	0.764706
ROR1 AND PLVAP AND NOT-HTR6	0.866667	1	0.764706
FAP AND NOT-CLDN8 AND CD93	0.827586	1	0.705882
FAP AND NOT-CLDN8 AND CD74	0.827586	1	0.705882
FAP AND NOT-CLDN8 AND LCT	0.827586	1	0.705882
FAP AND NOT-CLDN8 AND GJA10	0.827586	1	0.705882
FAP AND NOT-CLDN8 AND GUCY2F	0.827586	1	0.705882
FAP AND NOT-GUCY2F AND NOT-EPCAM	0.827586	1	0.705882
FAP AND VAMP8 AND NOT-CLDN7	0.866667	1	0.764706
NOT-EPCAM AND SLC31A1 AND CD34	0.882353	0.882353	0.882353
TGFBI AND NOT-EPCAM AND CD34	0.9375	1	0.882353
FAP AND NOT-SSTR5 AND GHSR	0.866667	1	0.764706
FAP AND NOT-SSTR5 AND PCDHAC2	0.866667	1	0.764706
FAP AND NOT-SSTR5 AND GJA10	0.866667	1	0.764706
FAP AND NOT-SSTR5 AND SV2C	0.866667	1	0.764706
FAP AND NOT-SSTR5 AND ADCY10	0.866667	1	0.764706
FAP AND NOT-SSTR5 AND GRIK4	0.866667	1	0.764706
FAP AND NOT-SSTR5 AND PCDHGC4	0.827586	1	0.705882
FAP AND NOT-CXCR5 AND SV2C	0.866667	1	0.764706
FAP AND NOT-CXCR5 AND PCDHGC4	0.827586	1	0.705882
FAP AND NOT-CXCR5 AND GRIK4	0.866667	1	0.764706
FAP AND NOT-SLC39A2 AND NOT-CLDN7	0.866667	1	0.764706
FAP AND NOT-CXCR5 AND ROS1	0.866667	1	0.764706
FAP AND NOT-SLC34A2 AND ACSL6	0.827586	1	0.705882
FAP AND NOT-SLC34A2 AND PCDHGC4	0.866667	1	0.764706
ROR1 AND NOT-CLDN6 AND PLVAP	0.866667	1	0.764706
ROR1 AND NOT-CLDN8 AND CD93	0.866667	1	0.764706
ROR1 AND NOT-CLDN8 AND CD74	0.866667	1	0.764706
FAP AND NOT-MUC16 AND GHSR	0.866667	1	0.764706

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
SLC22A16 AND NOT-TSPAN15 AND NOT-PHLDB2	0.944785	0.970588	0.920319
SLC22A16 AND NOT-SLC25A4 AND NOT-PHLDB2	0.944785	0.970588	0.920319
NOT-PHLDB2 AND CXCR4 AND SLC30A1	0.944762	0.905109	0.988048
NOT-PHLDB2 AND HCST AND NOT-GABRB1	0.944762	0.905109	0.988048
NOT-PHLDB2 AND HCST AND ALCAM	0.944762	0.905109	0.988048
NOT-PHLDB2 AND CXCR4 AND SLC30A1	0.944762	0.905109	0.988048
NOT-PHLDB2 AND HCST AND GYPC	0.944762	0.905109	0.988048
SLC22A16 AND NOT-SLC7A2 AND NOT-PHLDB2	0.944559	0.974576	0.916335
SLC22A16 AND NOT-KIAA1324 AND NOT-PHLDB2	0.944559	0.974576	0.916335
SLC22A16 AND NOT-ENPP5 AND NOT-PHLDB2	0.944559	0.974576	0.916335
NOT-PHLDB2 AND EMP3 AND NOT-BTN3A1	0.962076	0.964	0.960159
NOT-PHLDB2 AND EMP3 AND SLC39A8	0.954918	0.983122	0.928287
SLC22A16 AND NOT-SLC4A8 AND NOT-PHLDB2	0.946721	0.974684	0.920319
NOT-PHLDB2 AND NOT-FAT1 AND P2RX4	0.946322	0.944444	0.948207
NOT-PHLDB2 AND NOT-FAT1 AND P2RX4	0.946322	0.944444	0.948207
SLC22A16 AND EMP3 AND NOT-PHLDB2	0.944099	0.982759	0.908367
SLC22A16 AND NOT-KIAA0319 AND NOT-PHLDB2	0.942857	0.966527	0.920319
SLC22A16 AND NOT-OXTR AND NOT-PHLDB2	0.942857	0.966527	0.920319
SLC22A16 AND NOT-GRM7 AND NOT-PHLDB2	0.942857	0.966527	0.920319
SLC22A16 AND NOT-SMPD2 AND NOT-PHLDB2	0.942387	0.974468	0.912351
SLC22A16 AND NOT-LRIG3 AND NOT-PHLDB2	0.940937	0.9625	0.920319
SLC22A16 AND NOT-PHLDB2 AND NOT-SLC22A5	0.940937	0.9625	0.920319
SLC22A16 AND NOT-PTPRT AND NOT-PHLDB2	0.940937	0.9625	0.920319
NOT-PHLDB2 AND NOT-LRIG3 AND P2RX4	0.94027	0.910448	0.972112
NOT-PHLDB2 AND NOT-LRIG3 AND P2RX4	0.94027	0.910448	0.972112
NOT-ATP8B1 AND CD44 AND NOT-BTN3A1	0.944123	0.914179	0.976096
SLC22A16 AND NOT-ATP8B1 AND NOT-PHLDB2	0.938776	0.962343	0.916335
NOT-PHLDB2 AND P2RX4 AND NOT-SLC22A5	0.940039	0.913534	0.968127
NOT-PHLDB2 AND P2RX4 AND NOT-SLC22A5	0.940039	0.913534	0.968127
SLC22A16 AND NOT-SLC31A1 AND NOT-PHLDB2	0.936605	0.962185	0.912351
NOT-PHLDB2 AND CD44 AND SLCO1B3	0.940711	0.933333	0.948207
NOT-PHLDB2 AND CD44 AND P2RX4	0.958084	0.96	0.956175
NOT-PHLDB2 AND NOT-ATP8B1 AND P2RX4	0.934615	0.903346	0.968127
NOT-PHLDB2 AND NOT-ATP8B1 AND P2RX4	0.934615	0.903346	0.968127
SLC22A16 AND NOT-FAT1 AND NOT-PHLDB2	0.933884	0.969957	0.900398
SLC22A16 AND NOT-ATP8B1 AND NOT-SEMA4D	0.933602	0.943089	0.924303
SLC22A16 AND EMP3 AND NOT-CDH11	0.933333	0.946721	0.920319
NOT-PHLDB2 AND P2RX4 AND NOT-OXTR	0.933078	0.897059	0.972112
NOT-PHLDB2 AND P2RX4 AND NOT-OXTR	0.933078	0.897059	0.972112
NOT-PHLDB2 AND P2RX4 AND NOT-SLC4A8	0.945736	0.920755	0.972112
NOT-PHLDB2 AND P2RX4 AND NOT-SLC4A8	0.945736	0.920755	0.972112
NOT-GHR AND EMP3 AND NOT-BTN3A1	0.932039	0.909091	0.956175
SLC22A16 AND EMP3 AND NOT-GHR	0.931452	0.942857	0.920319
NOT-PHLDB2 AND SLC39A8 AND NOT-SLC22A5	0.931393	0.973913	0.89243
NOT-PHLDB2 AND SLC39A8 AND NOT-LRIG3	0.931393	0.973913	0.89243
NOT-PHLDB2 AND SLC39A8 AND NOT-ATP8B1	0.931393	0.973913	0.89243
NOT-ATP8B1 AND CD44 AND NOT-BTN3A3	0.931034	0.896679	0.968127
NOT-PHLDB2 AND EMP3 AND NOT-PIK3IP1	0.930966	0.921875	0.940239
SLC22A16 AND NOT-SLC31A1 AND NOT-GHR	0.929577	0.939024	0.920319
NOT-PHLDB2 AND NOT-PTPRT AND P2RX4	0.929524	0.890511	0.972112
NOT-PHLDB2 AND NOT-PTPRT AND P2RX4	0.929524	0.890511	0.972112
SLC22A16 AND NOT-ATP8B1 AND NOT-IFNAR2	0.929006	0.946281	0.912351
NOT-CDH11 AND EMP3 AND NOT-BTN3A1	0.940039	0.913534	0.968127
NOT-PHLDB2 AND SLC39A8 AND NOT-KIAA0319	0.927536	0.965517	0.89243
NOT-PHLDB2 AND SLC39A8 AND NOT-PTPRT	0.927536	0.965517	0.89243
NOT-ATP8B1 AND CD44 AND NOT-LAT	0.927481	0.89011	0.968127
SLC22A16 AND CD44 AND NOT-MRGPRF	0.930894	0.950207	0.912351
NOT-PHLDB2 AND SLC39A8 AND NOT-FAT1	0.926931	0.973684	0.884462
NOT-PHLDB2 AND NOT-ATP8B1 AND VAMP8	0.926829	0.875887	0.984064
NOT-PHLDB2 AND NOT-ATP8B1 AND VAMP8	0.926829	0.875887	0.984064
NOT-PHLDB2 AND P2RX4 AND NOT-SEMA4B	0.930502	0.902622	0.960159
NOT-GHR AND EMP3 AND NOT-BTN3A3	0.926357	0.901887	0.952191
NOT-PHLDB2 AND P2RX4 AND NOT-SLC31A1	0.92543	0.889706	0.964143
NOT-PHLDB2 AND P2RX4 AND NOT-SLC31A1	0.92543	0.889706	0.964143
SLC22A16 AND NOT-ATP8B1 AND NOT-SYT11	0.924644	0.945833	0.904382
SLC22A16 AND NOT-ATP8B1 AND NOT-PAQR8	0.923695	0.931174	0.916335
NOT-PHLDB2 AND SLC39A8 AND NOT-SLC31A1	0.923077	0.965217	0.884462
NOT-PHLDB2 AND VAMP8 AND NOT-SLC22A5	0.92365	0.867133	0.988048
NOT-PHLDB2 AND VAMP8 AND NOT-SLC22A5	0.92365	0.867133	0.988048
NOT-ATP8B1 AND SLC30A1 AND NOT-S1PR1	0.922481	0.898113	0.948207
NOT-ATP8B1 AND VAMP8 AND NOT-S1PR1	0.922481	0.898113	0.948207
NOT-ATP8B1 AND VAMP8 AND NOT-LAT	0.933837	0.888489	0.984064
NOT-ATP8B1 AND SLC30A1 AND NOT-LAT	0.923364	0.869718	0.984064
NOT-CDH11 AND EMP3 AND NOT-BTN3A3	0.93617	0.909774	0.964143
NOT-ATP8B1 AND VAMP8 AND NOT-BTN3A1	0.943182	0.898917	0.992032
NOT-ATP8B1 AND VAMP8 AND NOT-BTN3A3	0.937381	0.894928	0.984064
NOT-PHLDB2 AND P2RX4 AND NOT-FUT3	0.93666	0.903704	0.972112

Antigen Pair Logic	F1	Precision	Recall
FAP AND NOT-SLC34A2 AND GJA10	0.866667	1	0.764706
FAP AND NOT-VTCN1 AND CRB1	0.827586	1	0.705882
FAP AND NOT-CXCR5 AND FSHR	0.866667	1	0.764706
FAP AND NOT-MUC16 AND PCDHGC4	0.866667	1	0.764706
FAP AND NOT-MUC16 AND GRIK4	0.866667	1	0.764706
ROR1 AND NOT-EPCAM AND NOT-GUCY2F	0.866667	1	0.764706
ROR1 AND NOT-CLDN8 AND OR4N4	0.866667	1	0.764706
FAP AND NOT-CLDN7 AND SLC22A8	0.866667	1	0.764706
FAP AND NOT-CLDN7 AND ACSL6	0.827586	1	0.705882
ROR1 AND NOT-CLDN8 AND ACSL6	0.827586	1	0.705882
ROR1 AND NOT-CLDN8 AND GUCY2F	0.866667	1	0.764706
ROR1 AND NOT-EPCAM AND NOT-PCDHAC2	0.827586	1	0.705882
FAP AND NOT-CLDN7 AND GHSR	0.866667	1	0.764706
FAP AND NOT-CLDN7 AND CASR	0.866667	1	0.764706
ROR1 AND NOT-CLDN8 AND LCT	0.866667	1	0.764706
ROR1 AND NOT-CLDN8 AND PCDHGC4	0.866667	1	0.764706
ROR1 AND NOT-EPCAM AND NOT-GRIK4	0.827586	1	0.705882
COMPLEX-OR4N4/EPCAM/ROR1	0.83871	0.928571	0.764706
ROR1 AND NOT-SSTR5 AND SV2C	0.827586	1	0.705882
ROR1 AND NOT-CXCR5 AND SV2C	0.827586	1	0.705882
ROR1 AND NOT-CXCR5 AND OR4N4	0.866667	1	0.764706
ROR1 AND NOT-CLDN7 AND VAMP8	0.83871	0.928571	0.764706
ROR1 AND NOT-CRB2 AND CXCR5	0.827586	1	0.705882
ROR1 AND NOT-SSTR5 AND PCDHAC2	0.866667	1	0.764706
ROR1 AND NOT-CLDN8 AND GJA10	0.827586	1	0.705882
FAP AND NOT-CLDN8 AND MUC4	0.827586	1	0.705882
FAP AND NOT-EPCAM AND NOT-CXCR5	0.866667	1	0.764706
FAP AND NOT-CXCR5 AND NOT-CLDN7	0.866667	1	0.764706
ROR1 AND NOT-EPCAM AND NOT-CXCR5	0.866667	1	0.764706
ROR1 AND NOT-EPCAM AND CD34	0.83871	0.928571	0.764706
ROR1 AND NOT-CLDN8 AND MUC4	0.866667	1	0.764706
ROR1 AND NOT-CLDN7 AND CD34	0.83871	0.928571	0.764706
NOT-EPCAM AND IL13RA1 AND CD34	0.827586	1	0.705882
ATP8B2 AND NOT-SLC39A2 AND PLVAP	0.827586	1	0.705882
EMP3 AND PLVAP AND NOT-AGTR2	0.827586	1	0.705882
ATP8B2 AND NOT-GPR78 AND PLVAP	0.827586	1	0.705882
EMP3 AND NOT-ACSL6 AND GJA10	0.827586	1	0.705882
EMP3 AND PLVAP AND NOT-GPR78	0.827586	1	0.705882
EMP3 AND NOT-CCR9 AND PLVAP	0.827586	1	0.705882
EMP3 AND PLVAP AND NOT-SLC22A6	0.827586	1	0.705882
EMP3 AND PLVAP AND NOT-OR8D1	0.827586	1	0.705882
Stomach Cancer			
GPRC5A AND NOT-SCNN1B AND NOT-SGCG	0.941176	0.952381	0.930233
GPRC5A AND NOT-SCNN1B AND NOT-ITGB5	0.953488	0.953488	0.953488
ST14 AND NOT-SCNN1B AND NOT-SGCG	0.91954	0.909091	0.930233
ST14 AND NOT-SCNN1B AND ITGB5	0.883721	0.883721	0.883721
LSR AND NOT-SLC16A9 AND NOT-LAMP2	0.9	0.972973	0.837209
LSR AND NOT-SLC16A9 AND LRP8	0.897436	1	0.813953
GPRC5A AND NOT-SLC16A9 AND NOT-HEG1	0.8	0.864865	0.744186
PTPRH AND NOT-AQP8 AND SPINT1	0.842105	0.969697	0.744186
LSR AND NOT-SLC16A9 AND XPR1	0.9	0.972973	0.837209
FXYD3 AND NOT-SCNN1B AND PLXND1	0.8	0.730769	0.883721
SMAGP AND NOT-SCNN1B AND NOT-ITGB5	0.952381	0.97561	0.930233
TSPAN8 AND PLXND1 AND NOT-FXYD3	0.831169	0.941176	0.744186
FXYD3 AND NOT-SCNN1B AND PDGFRB	0.8	0.730769	0.883721
TSPAN8 AND PDGFRB AND NOT-FXYD3	0.842105	0.969697	0.744186
FXYD3 AND NOT-SCNN1B AND PANX1	0.878049	0.923077	0.837209
LSR AND NOT-SLC16A9 AND HEG1	0.91358	0.973684	0.860465
LSR AND NOT-SLC16A9 AND RELL1	0.925	1	0.860465
PTPRH AND PDGFRB AND FXYD3	0.883721	0.883721	0.883721
PTPRH AND PDGFRB AND SPINT1	0.941176	0.952381	0.930233
FXYD3 AND NOT-SCNN1B AND FADS2	0.853933	0.826087	0.883721
LSR AND NOT-SLC16A9 AND PCDHB7	0.925	1	0.860465
LSR AND NOT-SLC16A9 AND NOT-SLC12A1	0.91358	0.973684	0.860465
LSR AND NOT-FXYD3 AND NOT-SLCO4C1	0.823529	0.833333	0.813953
LSR AND NOT-SLC16A9 AND STX7	0.9	0.972973	0.837209
CLDN18 AND NOT-SLCO4C1 AND SLC39A10	0.857143	0.878049	0.837209
CLDN18 AND NOT-SGCG AND LRP11	0.886076	0.972222	0.813953
CLDN18 AND NOT-SLCO4C1 AND LRP11	0.888889	0.947368	0.837209
PTPRH AND NOT-SLC26A3 AND EPCAM	0.976744	0.976744	0.976744
CLDN18 AND NOT-SCNN1B AND NOT-SGCG	0.833333	0.853659	0.813953
CLDN18 AND NOT-SLCO4C1 AND ZDHHC5	0.911392	1	0.837209
CLDN18 AND NOT-EVA1A AND BCAP31	0.878049	0.923077	0.837209
CLDN18 AND NOT-SCNN1B AND ITGB5	0.827586	0.818182	0.837209
CLDN18 AND NOT-SLCO4C1 AND NPC1	0.878049	0.923077	0.837209
EPCAM AND GJB2 AND NOT-SLC26A3	0.966292	0.934783	1
CLDN18 AND NOT-SLCO4C1 AND S100A10	0.847059	0.857143	0.837209

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
NOT-ATP8B1 AND CD44 AND NOT-S1PR1	0.91945	0.906977	0.932271
NOT-PHLDB2 AND P2RX4 AND NOT-SMPD2	0.919231	0.888476	0.952191
NOT-PHLDB2 AND P2RX4 AND NOT-SMPD2	0.919231	0.888476	0.952191
NOT-GHR AND EMP3 AND NOT-STEAP4	0.918489	0.916667	0.920319
NOT-GHR AND EMP3 AND NOT-STEAP4	0.918489	0.916667	0.920319
NOT-PHLDB2 AND NOT-FAT1 AND VAMP8	0.918406	0.876812	0.964143
NOT-PHLDB2 AND NOT-FAT1 AND SLC30A1	0.918406	0.876812	0.964143
NOT-PHLDB2 AND NOT-FAT1 AND SLC30A1	0.918406	0.876812	0.964143
NOT-PHLDB2 AND VAMP8 AND NOT-SEMA4B	0.938931	0.901099	0.98008
NOT-ATP8B1 AND SLC30A1 AND NOT-BTN3A1	0.929368	0.87108	0.996016
NOT-CDH11 AND EMP3 AND NOT-S1PR1	0.917323	0.906615	0.928287
NOT-ATP8B1 AND SLC30A1 AND NOT-BTN3A3	0.920518	0.858621	0.992032
NOT-PHLDB2 AND SLC39A8 AND NOT-SMPD2	0.916318	0.964758	0.87251
NOT-PHLDB2 AND P2RX4 AND NOT-CACNG6	0.929524	0.890511	0.972112
NOT-PHLDB2 AND CD36 AND P2RX4	0.922432	0.973451	0.876494
SLC22A16 AND NOT-ATP8B1 AND NOT-FLVCR1	0.915811	0.944915	0.888446
NOT-ATP8B1 AND CD44 AND NOT-IFNAR2	0.915572	0.865248	0.972112
NOT-ATP8B1 AND CD44 AND NOT-IFNAR2	0.915572	0.865248	0.972112
NOT-PHLDB2 AND EMP3 AND NOT-S1PR1	0.91498	0.930041	0.900398
NOT-GHR AND EMP3 AND NOT-LAT	0.912959	0.887218	0.940239
NOT-CDH11 AND EMP3 AND NOT-LAT	0.912879	0.870036	0.960159
NOT-PHLDB2 AND NOT-SLC31A1 AND VAMP8	0.912801	0.854167	0.98008
NOT-PHLDB2 AND NOT-SLC31A1 AND VAMP8	0.912801	0.854167	0.98008
NOT-ATP8B1 AND VAMP8 AND NOT-IFNAR2	0.913444	0.849315	0.988048
NOT-ATP8B1 AND VAMP8 AND NOT-IFNAR2	0.913444	0.849315	0.988048
NOT-PHLDB2 AND LAPTM5 AND NOT-LRIG3	0.915129	0.852234	0.988048
NOT-PHLDB2 AND LAPTM5 AND NOT-ATP8B1	0.913444	0.849315	0.988048
NOT-PHLDB2 AND LAPTM5 AND NOT-ATP8B1	0.913444	0.849315	0.988048
NOT-PHLDB2 AND SLC39A8 AND SLCO1B3	0.911765	0.964444	0.864542
NOT-PHLDB2 AND LAPTM5 AND NOT-PTPRT	0.911765	0.846416	0.988048
NOT-PHLDB2 AND LAPTM5 AND NOT-GRM7	0.911765	0.846416	0.988048
NOT-STEAP4 AND EMP3 AND NOT-CDH11	0.916667	0.913043	0.920319
NOT-PHLDB2 AND P2RX4 AND NOT-PPAPDC1B	0.910506	0.889734	0.932271
NOT-PHLDB2 AND EMP3 AND SLC7A5	0.962076	0.964	0.960159
NOT-PHLDB2 AND EMP3 AND SLC7A5	0.962076	0.964	0.960159
NOT-PHLDB2 AND SLC7A5 AND VAMP8	0.948177	0.914815	0.984064
NOT-PHLDB2 AND SLC7A5 AND VAMP8	0.948177	0.914815	0.984064
NOT-PHLDB2 AND NOT-ERBB3 AND P2RX4	0.947573	0.924242	0.972112
NOT-PHLDB2 AND EMP3 AND NOT-CR2	0.945098	0.930502	0.960159
NOT-PHLDB2 AND SLC7A5 AND PTGER4	0.957364	0.932075	0.984064
NOT-PHLDB2 AND NOT-ALDH1A1 AND P2RX4	0.943026	0.930233	0.956175
SLC22A16 AND NOT-ERBB3 AND NOT-PHLDB2	0.942857	0.966527	0.920319
SLC22A16 AND NOT-CLDN8 AND NOT-PHLDB2	0.942857	0.966527	0.920319
SLC22A16 AND NOT-STEAP2 AND NOT-PHLDB2	0.942857	0.966527	0.920319
SLC22A16 AND NOT-ERBB4 AND NOT-PHLDB2	0.942857	0.966527	0.920319
SLC22A16 AND NOT-BMPR1B AND NOT-PHLDB2	0.942623	0.970464	0.916335
NOT-PHLDB2 AND EMP3 AND NOT-MS4A1	0.941176	0.926641	0.956175
SLC22A16 AND NOT-CLDN7 AND NOT-PHLDB2	0.940937	0.9625	0.920319
SLC22A16 AND SLC7A5 AND NOT-PHLDB2	0.940937	0.9625	0.920319
SLC22A16 AND NOT-ERBB2 AND NOT-PHLDB2	0.940937	0.9625	0.920319
SLC22A16 AND NOT-VTCN1 AND NOT-PHLDB2	0.940937	0.9625	0.920319
SLC22A16 AND NOT-SDC1 AND NOT-PHLDB2	0.940937	0.9625	0.920319
SLC22A16 AND NOT-RNF43 AND NOT-PHLDB2	0.940937	0.9625	0.920319
NOT-PHLDB2 AND SLC7A5 AND P2RX4	0.940039	0.913534	0.968127
NOT-PHLDB2 AND EMP3 AND NOT-HLA-DOB	0.939571	0.919847	0.960159
NOT-PHLDB2 AND CD44 AND SLC7A5	0.96063	0.949416	0.972112
SLC22A16 AND NOT-EPCAM AND NOT-PHLDB2	0.938776	0.962343	0.916335
SLC22A16 AND NOT-MET AND NOT-PHLDB2	0.938525	0.966245	0.912351
NOT-PHLDB2 AND EMP3 AND NOT-FCRL1	0.9375	0.91954	0.956175
NOT-PHLDB2 AND EMP3 AND NOT-CD22	0.9375	0.91954	0.956175
NOT-PHLDB2 AND EMP3 AND NOT-FCRL5	0.9375	0.91954	0.956175
NOT-PHLDB2 AND SLC7A5 AND NOT-GRM3	0.937381	0.894928	0.984064
NOT-PHLDB2 AND SLC7A5 AND NOT-DTNA	0.937381	0.894928	0.984064
NOT-PHLDB2 AND NOT-CEACAM5 AND P2RX4	0.93666	0.903704	0.972112
NOT-PHLDB2 AND P2RX4 AND NOT-CLDN3	0.93666	0.903704	0.972112
SLC22A16 AND NOT-STEAP1 AND NOT-PHLDB2	0.936345	0.966102	0.908367
SLC22A16 AND NOT-FOLH1 AND NOT-PHLDB2	0.936345	0.966102	0.908367
NOT-PHLDB2 AND NOT-ERBB3 AND SLC30A1	0.935849	0.888889	0.988048
NOT-PHLDB2 AND NOT-ERBB3 AND SLC30A1	0.935849	0.888889	0.988048
NOT-PHLDB2 AND EMP3 AND NOT-FCRL2	0.935673	0.916031	0.956175
NOT-PHLDB2 AND SLC7A5 AND LAPTM5	0.935606	0.891697	0.984064
NOT-PHLDB2 AND SLC7A5 AND NOT-CSPG5	0.935606	0.891697	0.984064
NOT-PHLDB2 AND SLC7A5 AND NOT-SLC1A2	0.935606	0.891697	0.984064
NOT-PHLDB2 AND EMP3 AND NOT-CD79A	0.935421	0.919231	0.952191
SLC22A16 AND EMP3 AND NOT-KDR	0.935223	0.950617	0.920319
NOT-PHLDB2 AND NOT-CEACAM5 AND VAMP8	0.935849	0.888889	0.988048
NOT-PHLDB2 AND NOT-CLDN7 AND P2RX4	0.934866	0.900369	0.972112

Antigen Pair Logic	F1	Precision	Recall
CLDN18 AND NOT-EVA1A AND SLMAP	0.837209	0.837209	0.837209
CLDN18 AND NOT-SLCO4C1 AND RYK	0.819277	0.85	0.790698
CLDN18 AND NOT-SLCO4C1 AND ITGB5	0.835443	0.916667	0.767442
CLDN18 AND NOT-SLCO4C1 AND CD99	0.814815	0.868421	0.767442
CLDN18 AND NOT-SGCG AND ZDHHC5	0.897436	1	0.813953
CLDN18 AND NOT-EVA1A AND RYK	0.819277	0.85	0.790698
EPCAM AND SLC16A1 AND NOT-SLC26A3	0.945055	0.895833	1
CLDN18 AND NOT-SLCO4C1 AND CLSTN1	0.809524	0.829268	0.790698
CLDN18 AND NOT-SLCO4C1 AND LRP8	0.809524	0.829268	0.790698
CLDN18 AND NOT-SLCO4C1 AND FADS2	0.808989	0.782609	0.837209
MUC13 AND NOT-SLC51B AND MMD	0.868421	1	0.767442
CLDN18 AND NOT-SLCO4C1 AND ITGA2	0.897436	1	0.813953
CLDN18 AND NOT-SLCO4C1 AND SLC3A2	0.878049	0.923077	0.837209
CLDN18 AND NOT-SLCO4C1 AND MTDH	0.9	0.972973	0.837209
CLDN18 AND NOT-SGCG AND ITGB5	0.883117	1	0.790698
EPCAM AND NOT-SCNN1B AND CYBA	1	1	1
TM4SF5 AND NOT-SLC26A3 AND EPCAM	0.888889	0.851064	0.930233
CLDN18 AND NOT-SGCG AND PIGU	0.868421	1	0.767442
CLDN18 AND NOT-SGCG AND S100A10	0.897436	1	0.813953
CLDN18 AND NOT-SLCO4C1 AND PLGRKT	0.888889	0.947368	0.837209
CLDN18 AND NOT-SLCO4C1 AND NOT-SGCG	0.823529	0.833333	0.813953
CLDN18 AND SLC4A11 AND NOT-CADM3	0.875	0.945946	0.813953
CLDN18 AND NOT-KLRF1 AND ZDHHC5	0.911392	1	0.837209
CLDN18 AND NOT-SLCO4C1 AND ATRAID	0.875	0.945946	0.813953
MUC13 AND NOT-SLC16A9 AND LRP8	0.897436	1	0.813953
CLDN18 AND NOT-SLCO4C1 AND TCIRG1	0.847059	0.857143	0.837209
EPCAM AND SLC12A2 AND NOT-SLC26A3	0.952381	0.97561	0.930233
CLDN18 AND NOT-EVA1A AND ATRAID	0.875	0.945946	0.813953
EPCAM AND NOT-SCNN1B AND PIGU	0.914894	0.843137	1
EPCAM AND NOT-SLCO4C1 AND NOT-SLC26A3	0.886598	0.796296	1
EPCAM AND NOT-SCNN1B AND ZDHHC5	1	1	1
CLDN18 AND NOT-SGCG AND SLC3A2	0.897436	1	0.813953
CLDN18 AND NOT-SLCO4C1 AND MGST2	0.9	0.972973	0.837209
CLDN18 AND NOT-EVA1A AND PLGRKT	0.888889	0.947368	0.837209
CLDN18 AND NOT-ECSCR AND ATP13A1	0.837838	1	0.72093
EPCAM AND SEMA4G AND NOT-SLC26A3	0.826923	0.704918	1
MUC13 AND PLXND1 AND NOT-FXYD3	0.888889	0.947368	0.837209
MUC13 AND NOT-SLC16A9 AND HEG1	0.9	0.972973	0.837209
EPCAM AND NOT-SCNN1B AND TMEM63B	0.895833	0.811321	1
CLDN18 AND NOT-SGCG AND PLGRKT	0.897436	1	0.813953
CLDN18 AND NOT-SGCG AND ZNRF3	0.823529	0.833333	0.813953
MUC13 AND PDGFRB AND NOT-FXYD3	0.902439	0.948718	0.860465
CLDN18 AND NOT-SLCO4C1 AND SPINT1	0.911392	1	0.837209
EPCAM AND NOT-SCNN1B AND NOT-PKHD1	0.886598	0.796296	1
EPCAM AND KCNN4 AND NOT-SLC26A3	0.930233	0.930233	0.930233
CLDN18 AND NOT-EVA1A AND PSENEN	0.847059	0.857143	0.837209
CLDN18 AND NOT-CXCR2 AND SLC39A10	0.804878	0.846154	0.767442
CLDN18 AND NOT-SLCO4C1 AND BMPR2	0.818182	0.8	0.837209
CLDN18 AND NOT-SLCO4C1 AND STX7	0.883117	1	0.790698
CLDN18 AND NOT-SLCO4C1 AND VAMP3	0.875	0.945946	0.813953
CLDN18 AND LRP11 AND NOT-STX1B	0.9	0.972973	0.837209
MUC13 AND NOT-SLCO4C1 AND FADS2	0.865979	0.777778	0.976744
EPCAM AND NOT-SCNN1B AND VNN3	0.886598	0.796296	1
MUC13 AND NOT-ANO10 AND NOT-SLCO4C1	0.836735	0.745455	0.953488
EPCAM AND NOT-CA4 AND NOT-CDHR3	0.837209	0.837209	0.837209
EPCAM AND CDH3 AND NOT-CDHR3	0.833333	0.754717	0.930233
CLDN18 AND SLC4A11 AND NOT-PTPRN	0.911392	1	0.837209
CLDN18 AND NOT-SLCO4C1 AND NOT-CLSTN2	0.819277	0.85	0.790698
EPCAM AND NOT-SCNN1B AND ADAM10	0.955556	0.914894	1
MUC13 AND NOT-SLCO4C1 AND NOT-FXYD3	0.901099	0.854167	0.953488
CLDN18 AND SLC4A11 AND NOT-SLC47A2	0.835443	0.916667	0.767442
EPCAM AND NOT-SCNN1B AND MCOLN2	0.905263	0.826923	1
MUC13 AND NOT-SLCO4C1 AND HEG1	0.891304	0.836735	0.953488
EPCAM AND GPR35 AND NOT-SLC26A3	0.895833	0.811321	1
CLDN18 AND NOT-CX3CL1 AND ESYT2	0.835443	0.916667	0.767442
GPRC5A AND NOT-SCNN1B AND CD44	0.953488	0.953488	0.953488
CDH17 AND NOT-SLC51B AND ZDHHC5	0.860465	0.860465	0.860465
CDH17 AND NOT-SLC51B AND SLC25A5	0.870588	0.880952	0.860465
B3GNT3 AND NOT-SCNN1B AND CD44	0.945055	0.895833	1
CDH17 AND NOT-AQP8 AND NOT-SPINT1	0.853333	1	0.744186
ST14 AND NOT-SCNN1B AND CD44	0.942529	0.931818	0.953488
GPRC5A AND NOT-SCNN1B AND TGFBI	0.941176	0.952381	0.930233
TSPAN1 AND NOT-SLCO4C1 AND CD93	0.903226	0.84	0.976744
ST14 AND NOT-SCNN1B AND TGFBI	0.930233	0.930233	0.930233
LSR AND NOT-SCNN1B AND CD44	0.938272	1	0.883721
TSPAN1 AND NOT-SCNN1B AND CD44	0.875	0.792453	0.976744
FXYD3 AND NOT-SCNN1B AND CATSPERD	0.844444	0.808511	0.883721

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
NOT-PHLDB2 AND P2RX4 AND NOT-ERBB2	0.934866	0.900369	0.972112
NOT-PHLDB2 AND NOT-CLDN8 AND P2RX4	0.934866	0.900369	0.972112
NOT-PHLDB2 AND P2RX4 AND NOT-RNF43	0.934866	0.900369	0.972112
NOT-PHLDB2 AND NOT-CLDN8 AND P2RX4	0.934866	0.900369	0.972112
NOT-PHLDB2 AND NOT-SDC1 AND P2RX4	0.934866	0.900369	0.972112
NOT-PHLDB2 AND P2RX4 AND NOT-RNF43	0.934866	0.900369	0.972112
NOT-PHLDB2 AND NOT-SDC1 AND P2RX4	0.934866	0.900369	0.972112
NOT-PHLDB2 AND NOT-CLDN7 AND P2RX4	0.934866	0.900369	0.972112
NOT-PHLDB2 AND P2RX4 AND NOT-ERBB2	0.934866	0.900369	0.972112
NOT-PHLDB2 AND P2RX4 AND NOT-EPCAM	0.934615	0.903346	0.968127
NOT-PHLDB2 AND P2RX4 AND NOT-EPCAM	0.934615	0.903346	0.968127
NOT-PHLDB2 AND NOT-ERBB3 AND VAMP8	0.934087	0.885714	0.988048
NOT-PHLDB2 AND NOT-ERBB3 AND VAMP8	0.934087	0.885714	0.988048
NOT-PHLDB2 AND SLC7A5 AND NOT-SLC24A2	0.933837	0.888489	0.984064
NOT-PHLDB2 AND SLC39A8 AND NOT-SDC1	0.933333	0.978166	0.89243
NOT-PHLDB2 AND P2RX4 AND NOT-CLDN23	0.932821	0.9	0.968127
SLC22A16 AND NOT-ERBB2 AND CD44	0.933063	0.950413	0.916335
NOT-PHLDB2 AND SLC7A5 AND NOT-KCNJ10	0.937381	0.894928	0.984064
NOT-PHLDB2 AND P2RX4 AND NOT-FOLH1	0.937743	0.91635	0.960159
NOT-PHLDB2 AND SLC39A8 AND NOT-ERBB3	0.931393	0.973913	0.89243
NOT-PHLDB2 AND SLC39A8 AND NOT-ERBB2	0.931393	0.973913	0.89243
NOT-PHLDB2 AND SLC39A8 AND NOT-STEAP2	0.931393	0.973913	0.89243
NOT-PHLDB2 AND SLC39A8 AND SLC7A5	0.931393	0.973913	0.89243
NOT-PHLDB2 AND SLC39A8 AND NOT-CLDN8	0.931393	0.973913	0.89243
NOT-PHLDB2 AND SLC39A8 AND NOT-CLDN7	0.931393	0.973913	0.89243
COMPLEX-VTCN1/PHLDB2/CD44	0.93666	0.903704	0.972112
NOT-PHLDB2 AND NOT-STEAP1 AND P2RX4	0.930502	0.902622	0.960159
NOT-PHLDB2 AND NOT-STEAP1 AND P2RX4	0.930502	0.902622	0.960159
SLC22A16 AND NOT-CLDN8 AND NOT-GHR	0.930417	0.928571	0.932271
NOT-PHLDB2 AND SLC7A5 AND PPAPDC1B	0.929524	0.890511	0.972112
NOT-PHLDB2 AND P2RX4 AND NOT-VTCN1	0.929524	0.890511	0.972112
NOT-PHLDB2 AND P2RX4 AND NOT-VTCN1	0.929524	0.890511	0.972112
NOT-PHLDB2 AND SLC39A8 AND NOT-RNF43	0.929461	0.969697	0.89243
NOT-PHLDB2 AND P2RX4 AND NOT-MUC13	0.93665	0.903704	0.972112
NOT-ATP8B1 AND CD44 AND NOT-CD160	0.929254	0.893382	0.968127
NOT-PHLDB2 AND SLC39A8 AND NOT-EPCAM	0.929167	0.973799	0.888446
SLC22A16 AND EMP3 AND NOT-AXL	0.929006	0.946281	0.912351
NOT-PHLDB2 AND DDX3X AND NOT-SLC4A8	0.928983	0.896296	0.964143
SLC22A16 AND SLC7A5 AND NOT-GHR	0.928	0.931727	0.924303
SLC22A16 AND NOT-CLDN8 AND CD44	0.931174	0.946502	0.916335
NOT-PHLDB2 AND SLC39A8 AND NOT-VTCN1	0.927536	0.965517	0.89243
NOT-PHLDB2 AND P2RX4 AND NOT-MOK	0.927481	0.89011	0.968127
NOT-PHLDB2 AND SLC7A5 AND NOT-SYT11	0.927203	0.892989	0.964143
NOT-PHLDB2 AND SLC39A8 AND NOT-MUC1	0.926931	0.973684	0.884462
NOT-PHLDB2 AND NOT-FAT1 AND DDX3X	0.926923	0.895911	0.960159
NOT-PHLDB2 AND NOT-MET AND P2RX4	0.926641	0.898876	0.956175
NOT-PHLDB2 AND NOT-MET AND P2RX4	0.926641	0.898876	0.956175
SLC22A16 AND NOT-ERBB2 AND EMP3	0.926441	0.924603	0.928287
NOT-AXL AND EMP3 AND NOT-BTN3A1	0.931298	0.893773	0.972112
NOT-ATP8B1 AND CD44 AND NOT-MS4A1	0.925714	0.886861	0.968127
SLC22A16 AND NOT-MUC1 AND NOT-PHLDB2	0.92562	0.961373	0.89243
NOT-PHLDB2 AND SLC7A5 AND NOT-GABRB1	0.925094	0.872792	0.984064
NOT-PHLDB2 AND SLC39A8 AND NOT-STEAP1	0.924686	0.973568	0.880478
SLC22A16 AND EMP3 AND NOT-EDNRB	0.924644	0.945833	0.904382
NOT-ATP8B1 AND VAMP8 AND NOT-HLA-DOB	0.924214	0.862069	0.996016
NOT-ATP8B1 AND VAMP8 AND NOT-MS4A1	0.935606	0.891697	0.984064
NOT-PHLDB2 AND CD33 AND SLCO1B3	0.941414	0.954918	0.928287
NOT-PHLDB2 AND CD33 AND SLC30A1	0.948413	0.944664	0.952191
NOT-PHLDB2 AND CD33 AND NOT-ATP8B1	0.939096	0.926357	0.952191
NOT-PHLDB2 AND CD33 AND NOT-LRIG3	0.939096	0.926357	0.952191
NOT-PHLDB2 AND CD33 AND P2RX4	0.959016	0.987342	0.932271
NOT-PHLDB2 AND CD33 AND NOT-SLC22A5	0.939096	0.926357	0.952191
NOT-PHLDB2 AND CD33 AND NOT-SLC31A1	0.934911	0.925781	0.944223
NOT-PHLDB2 AND CD33 AND NOT-SMPD2	0.932806	0.92549	0.940239
CD33 AND NOT-ATP8B1 AND NOT-SEMA4D	0.910156	0.89272	0.928287
CD33 AND NOT-ATP8B1 AND NOT-PIK3IP1	0.888889	0.901639	0.876494
CD33 AND NOT-IFNAR2 AND NOT-ATP8B1	0.91453	0.986175	0.85259
CD33 AND NOT-ATP8B1 AND NOT-PAQR8	0.865784	0.823741	0.912351
CD33 AND NOT-ATP8B1 AND NOT-SYT11	0.865385	0.836431	0.896414
CD33 AND NOT-ATP8B1 AND NOT-FLVCR1	0.859345	0.83209	0.888446
CD33 AND NOT-ATP8B1 AND NOT-PPAPDC1B	0.825263	0.875	0.780876
CD33 AND NOT-ATP8B1 AND NOT-SLCO1B3	0.800693	0.708589	0.920319
NOT-PHLDB2 AND HCST AND SLC7A5	0.966862	0.946565	0.988048
NOT-PHLDB2 AND HCST AND SLC7A5	0.966862	0.946565	0.988048
NOT-PHLDB2 AND SLC7A5 AND PTGER2	0.966601	0.953488	0.98008
NOT-PHLDB2 AND CXCR4 AND SLC7A5	0.957529	0.928839	0.988048
NOT-PHLDB2 AND CXCR4 AND SLC7A5	0.957529	0.928839	0.988048

Antigen Pair Logic	F1	Precision	Recall
PPAP2C AND NOT-SCNN1B AND TGFBI	0.931818	0.911111	0.953488
TSPAN8 AND NOT-SCNN1B AND CD44	0.883117	1	0.790698
CDH17 AND NOT-SLC16A9 AND SLC25A5	0.888889	0.947368	0.837209
TSPAN1 AND NOT-SCNN1B AND TGFBI	0.904762	0.926829	0.883721
CDH17 AND NOT-SPPL2A AND SLC25A5	0.8	0.765957	0.837209
PTPRH AND NOT-SLC30A10 AND SPINT1	0.965517	0.954545	0.976744
PIGR AND NOT-SLCO4C1 AND CD93	0.83871	0.78	0.906977
CDH17 AND NOT-SCNN1B AND ITGB5	0.876404	0.847826	0.906977
TM4SF5 AND NOT-SLC30A10 AND ESYT2	0.879121	0.833333	0.930233
TSPAN8 AND NOT-SCNN1B AND TGFBI	0.857143	0.970588	0.767442
PTPRH AND CD44 AND NOT-SCNN1B	0.869565	0.816327	0.930233
IYD AND NOT-SCNN1B AND CD44	0.883117	1	0.790698
KCNE3 AND NOT-SLCO4C1 AND CD44	0.803738	0.671875	1
FXYD3 AND NOT-SCNN1B AND CD44	0.850575	0.840909	0.860465
FXYD3 AND NOT-SCNN1B AND CD93	0.835165	0.791667	0.883721
CDH17 AND PODXL AND SLC25A5	0.878049	0.923077	0.837209
TMPRSS2 AND NOT-SCNN1B AND TGFBI	0.928571	0.95122	0.906977
KCNE3 AND NOT-SLCO4C1 AND CD93	0.865979	0.777778	0.976744
B3GNT3 AND NOT-SCNN1B AND TGFBI	0.95122	1	0.906977
CDH17 AND NOT-FXYD3 AND ABCA1	0.818182	0.8	0.837209
DSG2 AND NOT-SCNN1B AND CD44	0.934783	0.877551	1
TM4SF5 AND NOT-SLC30A10 AND LRP8	0.842105	0.769231	0.930233
FXYD3 AND NOT-SCNN1B AND TGFBI	0.850575	0.840909	0.860465
CDH17 AND NOT-SLC16A9 AND LITAF	0.857143	0.878049	0.837209
ABCC3 AND NOT-SCNN1B AND CD44	0.886598	0.796296	1
FZD5 AND NOT-SCNN1B AND CD44	0.91358	0.973684	0.860465
TMPRSS2 AND NOT-SCNN1B AND CD44	0.906977	0.906977	0.906977
CDH17 AND NOT-FXYD3 AND NRP1	0.804598	0.795455	0.813953
PIGR AND NOT-SCNN1B AND CD44	0.863636	0.844444	0.883721
LSR AND NOT-SLC30A10 AND PRRG1	0.904762	0.926829	0.883721
PPAP2C AND NOT-SCNN1B AND CD44	0.930233	0.930233	0.930233
LSR AND NOT-SLC16A9 AND NOT-SLC31A1	0.857143	0.970588	0.767442
TNFRSF21 AND NOT-SCNN1B AND TGFBI	0.850575	0.840909	0.860465
ST14 AND NOT-SLC16A9 AND NOT-CD1A	0.808989	0.782609	0.837209
TM4SF5 AND CD44 AND NOT-SCNN1B	0.95122	1	0.906977
ITGA2 AND NOT-SCNN1B AND CD44	0.813187	0.770833	0.860465
TM4SF5 AND NOT-SLC30A10 AND CD99	0.870588	0.880952	0.860465
SLC12A2 AND NOT-SCNN1B AND CD44	0.909091	0.888889	0.930233
DSG2 AND NOT-SCNN1B AND TGFBI	0.903226	0.84	0.976744
LSR AND NOT-SLCO4C1 AND CD44	0.835165	0.791667	0.883721
PIGR AND NOT-SCNN1B AND TGFBI	0.844444	0.808511	0.883721
COMPLEX-ALCAM/SLC16A9/LSR	0.91358	0.973684	0.860465
TM4SF5 AND NOT-SLC30A10 AND PCDHB7	0.835165	0.791667	0.883721
PVRL3 AND NOT-SCNN1B AND CD44	0.871795	0.971429	0.790698
SEMA4G AND NOT-SCNN1B AND CD44	0.913043	0.857143	0.976744
CLDN18 AND NOT-SCNN1B AND CD44	0.847059	0.857143	0.837209
CDH17 AND NOT-SLC26A3 AND EPCAM	0.953488	0.953488	0.953488
CLDN18 AND NOT-SLCO4C1 AND CD44	0.837209	0.837209	0.837209
CLDN18 AND NOT-SLCO4C1 AND CD93	0.827586	0.818182	0.837209
CLDN18 AND LRP11 AND NOT-AOC3	0.9	0.972973	0.837209
CLDN18 AND NOT-SCNN1B AND TGFBI	0.846154	0.942857	0.767442
CLDN18 AND NOT-SLCO4C1 AND VANGL1	0.875	0.945946	0.813953
EPCAM AND NOT-SCNN1B AND TGFBI	0.965517	0.954545	0.976744
MUC13 AND NOT-SCNN1B AND CD44	0.988235	1	0.976744
CEACAM5 AND NOT-SCNN1B AND CD44	0.888889	0.947368	0.837209
MUC13 AND NOT-SCNN1B AND TGFBI	0.938272	1	0.883721
EPCAM AND FXYD5 AND NOT-S1PR1	0.965517	0.954545	0.976744
CLDN18 AND NOT-ECSCR AND CD58	0.842105	0.969697	0.744186
CLDN7 AND NOT-SCNN1B AND CD44	0.930233	0.930233	0.930233
CLDN18 AND NOT-EVA1A AND CD58	0.9	0.972973	0.837209
CDH17 AND NOT-SLC16A9 AND ANXA1	0.857143	0.878049	0.837209
EPCAM AND NOT-SLCO4C1 AND CD93	0.94382	0.913043	0.976744
EPCAM AND NOT-SCNN1B AND CD44	0.976744	0.976744	0.976744
CLDN18 AND NOT-EVA1A AND SLC31A1	0.853333	1	0.744186
MST1R AND NOT-SCNN1B AND CD44	0.942529	0.931818	0.953488
EPCAM AND CDH3 AND NOT-CATSPERD	0.888889	0.851064	0.930233
EPCAM AND FXYD5 AND NOT-C8B	0.977273	0.955556	1
CLDN18 AND SLC2A1 AND NOT-HEG1	0.814815	0.868421	0.767442
CLDN18 AND SLC2A1 AND NOT-SLC6A1	0.864198	0.921053	0.813953
CLDN18 AND SLC4A11 AND NOT-SLC22A9	0.888889	0.947368	0.837209
EPCAM AND NOT-CA4 AND NOT-CATSPERD	0.837209	0.837209	0.837209
EPCAM AND NOT-CA4 AND NOT-OXTR	0.809524	0.829268	0.790698
EPCAM AND NOT-SCNN1B AND SEMA4B	0.914894	0.843137	1
MST1R AND NOT-SCNN1B AND TGFBI	0.941176	0.952381	0.930233
EPCAM AND NOT-SCNN1B AND CATSPERD	0.886598	0.796296	1
CLDN18 AND NOT-SLCO4C1 AND SEMA4B	0.886076	0.972222	0.813953
CLDN18 AND NOT-SGCG AND SLC31A1	0.837838	1	0.72093

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
NOT-PHLDB2 AND SLC7A5 AND LTBR	0.965795	0.97561	0.956175
NOT-PHLDB2 AND SLC7A5 AND LTBR	0.965795	0.97561	0.956175
NOT-PHLDB2 AND CLEC12A AND SLC7A5	0.96371	0.97551	0.952191
FLT3 AND EMP3 AND NOT-EDNRB	0.952381	0.991379	0.916335
NOT-PHLDB2 AND NOT-KIAA1324 AND ANXA1	0.959223	0.935606	0.984064
NOT-PHLDB2 AND NOT-KIAA1324 AND DDX3X	0.962376	0.956693	0.968127
NOT-PHLDB2 AND HCST AND NOT-HLA-DOB	0.952015	0.918519	0.988048
NOT-PHLDB2 AND SLC7A5 AND PLP2	0.955513	0.928571	0.984064
NOT-PHLDB2 AND SLC7A5 AND PLP2	0.955513	0.928571	0.984064
NOT-PHLDB2 AND HCST AND NOT-CR2	0.96124	0.935849	0.988048
COMPLEX-FLT3/GHR/CLDN8	0.950617	0.982979	0.920319
NOT-PHLDB2 AND SLC7A5 AND TLR2	0.954813	0.94186	0.968127
NOT-PHLDB2 AND HCST AND DDX3X	0.95	0.918216	0.984064
NOT-PHLDB2 AND EVI2B AND SLC7A5	0.949612	0.924528	0.976096
NOT-PHLDB2 AND EVI2B AND SLC7A5	0.949612	0.924528	0.976096
NOT-PHLDB2 AND NOT-ERBB3 AND LTBR	0.949416	0.927757	0.972112
NOT-PHLDB2 AND NOT-ERBB3 AND LTBR	0.949416	0.927757	0.972112
P2RY8 AND NOT-PHLDB2 AND NOT-CR2	0.95499	0.938462	0.972112
P2RY8 AND NOT-PHLDB2 AND NOT-CR2	0.95499	0.938462	0.972112
NOT-PHLDB2 AND HCST AND CD70	0.95534	0.931818	0.98008
NOT-PHLDB2 AND CXCR4 AND NOT-CR2	0.948375	0.911765	0.988048
NOT-PHLDB2 AND TAAR5 AND NOT-STEAP2	0.957916	0.96371	0.952191
FLT3 AND NOT-AXL AND EMP3	0.948025	0.991304	0.908367
NOT-PHLDB2 AND HCST AND SLC39A6	0.947977	0.91791	0.98008
NOT-PHLDB2 AND HCST AND NOT-MS4A1	0.95183	0.921642	0.984064
NOT-PHLDB2 AND HCST AND NOT-FCRL5	0.947977	0.91791	0.98008
NOT-PHLDB2 AND TAAR5 AND NOT-ERBB3	0.954092	0.956	0.952191
P2RY8 AND NOT-PHLDB2 AND SLC7A5	0.951267	0.931298	0.972112
P2RY8 AND NOT-PHLDB2 AND SLC7A5	0.951267	0.931298	0.972112
NOT-PHLDB2 AND CYSLTR1 AND SLC7A5	0.946502	0.978723	0.916335
NOT-PHLDB2 AND PIEZO1 AND NOT-ERBB3	0.946322	0.944444	0.948207
P2RY8 AND NOT-PHLDB2 AND NOT-HLA-DOB	0.949416	0.927757	0.972112
NOT-PHLDB2 AND SLC7A5 AND ESYT2	0.954092	0.956	0.952191
SLC22A16 AND NOT-ABHD6 AND NOT-KDR	0.945455	0.959016	0.932271
NOT-PHLDB2 AND FXYD5 AND SLC7A5	0.964981	0.942966	0.988048
NOT-PHLDB2 AND FXYD5 AND SLC7A5	0.964981	0.942966	0.988048
CSF3R AND NOT-PHLDB2 AND SLC7A5	0.952965	0.978992	0.928287
NOT-PHLDB2 AND GLIPR1 AND SLC7A5	0.944882	0.933852	0.956175
NOT-PHLDB2 AND HCST AND NOT-CEACAM5	0.944762	0.905109	0.988048
NOT-PHLDB2 AND TAAR5 AND NOT-CLDN8	0.944664	0.937255	0.952191
SLC22A16 AND NOT-LAMP3 AND NOT-EDNRB	0.944559	0.974576	0.916335
NOT-PHLDB2 AND HCST AND NOT-FCRL1	0.944551	0.908088	0.984064
NOT-PHLDB2 AND SLC7A5 AND CD46	0.953125	0.934866	0.972112
NOT-PHLDB2 AND SLC7A5 AND CD46	0.953125	0.934866	0.972112
NOT-PHLDB2 AND SLC7A5 AND ICAM3	0.948177	0.914815	0.984064
NOT-PHLDB2 AND SLC7A5 AND ICAM3	0.948177	0.914815	0.984064
NOT-PHLDB2 AND HCST AND NOT-CD79A	0.944338	0.911111	0.98008
NOT-PTPRK AND NOT-PHLDB2 AND CD70	0.944338	0.911111	0.98008
NOT-PTPRK AND NOT-PHLDB2 AND CD70	0.944338	0.911111	0.98008
NOT-PTPRK AND SLC22A16 AND CD37	0.94433	0.978632	0.912351
NOT-PHLDB2 AND HCST AND NOT-CD160	0.944123	0.914179	0.976096
NOT-PHLDB2 AND FXYD5 AND DDX3X	0.948177	0.914815	0.984064
NOT-PHLDB2 AND SLC7A5 AND LILRB2	0.951456	0.92803	0.976096
NOT-PHLDB2 AND SLC7A5 AND TCIRG1	0.944338	0.911111	0.98008
NOT-PHLDB2 AND NOT-CEACAM5 AND LTBR	0.943249	0.926923	0.960159
NOT-PHLDB2 AND CXCR4 AND NOT-CXCR5	0.942966	0.901818	0.988048
NOT-PHLDB2 AND TAAR5 AND SLC7A5	0.942801	0.933594	0.952191
NOT-PHLDB2 AND TAAR5 AND NOT-RNF43	0.942801	0.933594	0.952191
NOT-PHLDB2 AND TAAR5 AND NOT-CLDN7	0.942801	0.933594	0.952191
NOT-PHLDB2 AND SLC7A5 AND ATP2A3	0.942801	0.933594	0.952191
NOT-PHLDB2 AND SLC7A5 AND NOT-FAIM2	0.942748	0.904762	0.984064
NOT-PHLDB2 AND SLC7A5 AND NOT-PLP1	0.942748	0.904762	0.984064
NOT-PHLDB2 AND SLC7A5 AND NOT-GPRC5B	0.942748	0.904762	0.984064
NOT-PHLDB2 AND SLC7A5 AND SLC10A3	0.946154	0.914498	0.98008
SLC22A16 AND NOT-ABHD6 AND NOT-AXL	0.944785	0.970588	0.920319
NOT-PHLDB2 AND EVI2B AND NOT-CR2	0.942308	0.910781	0.976096
NOT-PHLDB2 AND EVI2B AND DDX3X	0.942085	0.913858	0.972112
NOT-PHLDB2 AND HCST AND HSPA5	0.94186	0.916981	0.968127
P2RY8 AND NOT-PHLDB2 AND NOT-FCRL5	0.941634	0.920152	0.964143
P2RY8 AND NOT-PHLDB2 AND NOT-FCRL5	0.941634	0.920152	0.964143
SLC22A16 AND P2RY8 AND NOT-CR2	0.945233	0.96281	0.928287
NOT-PHLDB2 AND SLC7A5 AND NOT-ATP1A2	0.940952	0.90146	0.984064
NOT-PHLDB2 AND CXCR4 AND DDX3X	0.940952	0.90146	0.984064
NOT-PHLDB2 AND CD300LF AND SLC7A5	0.940937	0.9625	0.920319
P2RY8 AND NOT-PHLDB2 AND DDX3X	0.940039	0.913534	0.968127
NOT-SGCE AND EMP3 AND NOT-CD160	0.940039	0.913534	0.968127
NOT-PHLDB2 AND NKG7 AND SLC7A5	0.939759	0.947368	0.932271

Antigen Pair Logic	F1	Precision	Recall
CLDN18 AND SLC2A1 AND NOT-GPR18	0.847059	0.857143	0.837209
CLDN18 AND SLC2A1 AND SLC25A5	0.9	0.972973	0.837209
EPCAM AND FAT1 AND NOT-SLC26A3	0.903226	0.84	0.976744
MUC13 AND NOT-SLCO4C1 AND CD44	0.875	0.792453	0.976744
EPCAM AND S1PR1 AND NOT-CA4	0.831169	0.941176	0.744186
CEACAM5 AND NOT-SCNN1B AND TGFBI	0.883117	1	0.790698
EPCAM AND CDH3 AND NOT-OXTR	0.808511	0.745098	0.883721
CLDN18 AND SLC2A1 AND NOT-ZNRF3	0.833333	0.853659	0.813953
EPCAM AND NOT-SLCO4C1 AND CD44	0.933333	0.893617	0.976744
CLDN18 AND SLC2A1 AND NOT-PLGRKT	0.833333	0.853659	0.813953
ERBB3 AND NOT-SCNN1B AND CD44	0.857143	0.763636	0.976744
CLDN18 AND SLC2A1 AND STX7	0.886076	0.972222	0.813953
EPCAM AND F2R AND NOT-SLC26A3	0.977273	0.955556	1
CLDN18 AND NOT-KLRF1 AND SLC31A1	0.853333	1	0.744186
MUC13 AND NOT-SLC30A10 AND PRRG1	0.94382	0.913043	0.976744
CLDN18 AND NOT-S1PR1 AND ITGA2	0.883117	1	0.790698
CLDN18 AND SLC2A1 AND PCDHB7	0.847059	0.857143	0.837209
EPCAM AND FXYD5 AND NOT-CATSPERD	0.988506	0.977273	1
EPCAM AND TTYH3 AND NOT-SLC26A3	0.848485	0.75	0.976744
EPCAM AND NOT-SCNN1B AND S1PR1	0.886364	0.866667	0.906977
CLDN18 AND LRP11 AND NOT-SLC39A2	0.888889	0.947368	0.837209
CLDN18 AND NOT-CD83 AND CD93	0.819277	0.85	0.790698
EPCAM AND NOT-SLCO4C1 AND CATSPERD	0.803738	0.671875	1
EPCAM AND FXYD5 AND SEMA4B	0.965517	0.954545	0.976744
CLDN18 AND SLC4A11 AND NOT-SLC31A1	0.847059	0.857143	0.837209
CLDN18 AND SLC4A11 AND NOT-ANTXR2	0.814815	0.868421	0.767442
CLDN18 AND NOT-CSF3R AND CD58	0.871795	0.971429	0.790698
EPCAM AND F2R AND NOT-C1orf210	0.934783	0.877551	1
CLDN18 AND NOT-S1PR1 AND SDC4	0.833333	0.853659	0.813953
CLDN18 AND SLC4A11 AND CD93	0.878049	0.923077	0.837209
CLDN18 AND SLC2A1 AND MTDH	0.9	0.972973	0.837209
CLDN18 AND LRP11 AND CD1A	0.875	0.945946	0.813953
EPCAM AND S1PR1 AND CDH3	0.888889	0.851064	0.930233
CLDN18 AND CACNG6 AND ESYT2	0.878049	0.923077	0.837209
CLDN18 AND NOT-FXYD6 AND SLC25A3	0.810811	0.967742	0.697674
EPCAM AND NOT-CATSPERD AND NOT-PARM1	0.814815	0.868421	0.767442
EPCAM AND PDPN AND NOT-S1PR1	0.888889	0.851064	0.930233
EPCAM AND THY1 AND NOT-S1PR1	0.816327	0.727273	0.930233
EPCAM AND THY1 AND NOT-C8B	0.816327	0.727273	0.930233
EPCAM AND THY1 AND SEMA4B	0.860215	0.8	0.930233
EPCAM AND THY1 AND NOT-OXTR	0.816327	0.727273	0.930233
EPCAM AND THY1 AND CATSPERD	0.824742	0.740741	0.930233
CLDN18 AND NOT-FOLR1 AND SLC31A1	0.842105	0.969697	0.744186
CLDN18 AND NOT-SLC34A2 AND SLC30A1	0.818182	0.8	0.837209
CLDN18 AND EPCAM AND NOT-S1PR1	0.888889	0.947368	0.837209
EPCAM AND SPON2 AND NOT-CATSPERD	0.931818	0.911111	0.953488
CLDN18 AND NOT-SLC34A2 AND SLC31A1	0.853333	1	0.744186
EPCAM AND NOT-ENPP3 AND SEMA4B	0.901099	0.854167	0.953488
CLDN18 AND EPCAM AND NOT-CATSPERD	0.888889	0.947368	0.837209
EPCAM AND SPON2 AND NOT-OXTR	0.906977	0.906977	0.906977
EPCAM AND SLC7A5 AND NOT-C8B	0.917647	0.928571	0.906977
EPCAM AND SLC7A5 AND NOT-S1PR1	0.917647	0.928571	0.906977
EPCAM AND FAP AND NOT-S1PR1	0.817204	0.76	0.883721
EPCAM AND SPON2 AND S1PR1	0.942529	0.931818	0.953488
EPCAM AND NOT-CLDN8 AND SEMA4B	0.842105	0.969697	0.744186
EPCAM AND FAP AND NOT-C8B	0.835165	0.791667	0.883721
EPCAM AND NOT-ENPP3 AND NOT-S1PR1	0.803922	0.694915	0.953488
EPCAM AND FAP AND NOT-CATSPERD	0.873563	0.863636	0.883721
EPCAM AND SLC7A5 AND CATSPERD	0.917647	0.928571	0.906977
EPCAM AND NOT-ENPP3 AND NOT-C8B	0.815534	0.7	0.976744
EPCAM AND NOT-ENPP3 AND NOT-CATSPERD	0.815534	0.7	0.976744
EPCAM AND NOT-CATSPERD AND CLDN1	0.955556	0.914894	1
CLDN18 AND RNF43 AND NOT-SLC22A9	0.86747	0.9	0.837209
EPCAM AND SPON2 AND NOT-C8B	0.931818	0.911111	0.953488
MUC13 AND CD93 AND NOT-CD33	0.821918	1	0.697674
MUC17 AND SLC7A5 AND NOT-PTK7	0.853333	1	0.744186
CLDN18 AND RNF43 AND NOT-KCND2	0.809524	0.829268	0.790698
EPCAM AND NOT-OXTR AND CLDN1	0.945055	0.895833	1
CLDN18 AND RNF43 AND NOT-ANTXR2	0.804878	0.846154	0.767442
ATP8B1 AND SLC7A5 AND PTK7	0.897436	1	0.813953
MUC13 AND S1PR1 AND EPCAM	0.826667	0.96875	0.72093
EPCAM AND NOT-BMPR1B AND SLC6A6	0.848485	0.75	0.976744
EPCAM AND S1PR1 AND CLDN1	0.945055	0.895833	1
CLDN18 AND RNF43 AND CD58	0.911392	1	0.837209
EPCAM AND VCAM1 AND EPHA2	0.839506	0.894737	0.790698
EPCAM AND NOT-BMPR1B AND EPHA2	0.839506	0.894737	0.790698
EPCAM AND VCAM1 AND SLC6A6	0.84	0.736842	0.976744

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
NOT-PHLDB2 AND NOT-ERBB3 AND TYROBP	0.944762	0.905109	0.988048
NOT-PHLDB2 AND NOT-ERBB3 AND TYROBP	0.944762	0.905109	0.988048
SLC22A16 AND P2RY8 AND NOT-FCRL5	0.944785	0.970588	0.920319
NOT-PHLDB2 AND LTBR AND NOT-ERBB2	0.939571	0.919847	0.960159
NOT-PHLDB2 AND LTBR AND NOT-ERBB2	0.939571	0.919847	0.960159
NOT-PHLDB2 AND NOT-SDC1 AND LTBR	0.939571	0.919847	0.960159
NOT-PHLDB2 AND NOT-SDC1 AND LTBR	0.939571	0.919847	0.960159
NOT-PHLDB2 AND LTBR AND NOT-CLDN3	0.939571	0.919847	0.960159
SLC22A16 AND P2RY8 AND NOT-MS4A1	0.939516	0.95102	0.928287
NOT-PHLDB2 AND NOT-ERBB3 AND TMEM219	0.939394	0.895307	0.988048
NOT-PHLDB2 AND SLC7A5 AND NOT-PTPRZ1	0.939163	0.898182	0.984064
NOT-PHLDB2 AND CXCR4 AND NOT-MS4A1	0.939163	0.898182	0.984064
NOT-PHLDB2 AND SLC7A5 AND CD69	0.939163	0.898182	0.984064
NOT-PHLDB2 AND SLC7A5 AND NOT-AQP4	0.939163	0.898182	0.984064
NOT-PHLDB2 AND SLC7A5 AND NOT-PTPRZ1	0.939163	0.898182	0.984064
NOT-PHLDB2 AND SLC7A5 AND CD69	0.939163	0.898182	0.984064
NOT-PTPRK AND NOT-PHLDB2 AND DDX3X	0.939163	0.898182	0.984064
NOT-PHLDB2 AND NOT-PTPRF AND DDX3X	0.939163	0.898182	0.984064
NOT-PHLDB2 AND CD33 AND NOT-KIAA1324	0.961771	0.971545	0.952191
NOT-PHLDB2 AND CD33 AND NOT-LRRC4	0.95122	0.970954	0.932271
NOT-PHLDB2 AND CD33 AND NOT-SLC25A4	0.944664	0.937255	0.952191
NOT-PODXL AND CD33 AND NOT-BTN3A3	0.954813	0.94186	0.968127
NOT-PODXL AND NOT-PHLDB2 AND CD33	0.945098	0.930502	0.960159
NOT-PODXL AND NOT-PHLDB2 AND CD33	0.945098	0.930502	0.960159
NOT-PHLDB2 AND CD33 AND NOT-PSENEN	0.961771	0.971545	0.952191
NOT-PHLDB2 AND CD33 AND NOT-PTPRZ1	0.944664	0.937255	0.952191
NOT-PHLDB2 AND CD33 AND NOT-ABHD6	0.942801	0.933594	0.952191
NOT-PHLDB2 AND CD33 AND NOT-LAMP3	0.942801	0.933594	0.952191
NOT-PHLDB2 AND CD33 AND NOT-STIM2	0.942801	0.933594	0.952191
NOT-PHLDB2 AND HCST AND CD33	0.942748	0.904762	0.984064
NOT-PHLDB2 AND CD33 AND EVI2B	0.942574	0.937008	0.948207
NOT-PHLDB2 AND CD33 AND NOT-UGT8	0.942801	0.933594	0.952191
NOT-PHLDB2 AND CD33 AND OR6A2	0.953535	0.967213	0.940239
COMPLEX-SLC4A1/PHLDB2/CD33	0.949612	0.924528	0.976096
NOT-PHLDB2 AND CD33 AND NOT-SLC22A4	0.94929	0.966942	0.932271
NOT-PTPRK AND NOT-PHLDB2 AND CD33	0.94186	0.916981	0.968127
NOT-PTPRK AND NOT-PHLDB2 AND CD33	0.94186	0.916981	0.968127
NOT-PHLDB2 AND CD33 AND NOT-ABCC5	0.946322	0.944444	0.948207
NOT-PHLDB2 AND CD33 AND NOT-DAG1	0.940945	0.929961	0.952191
NOT-PHLDB2 AND CD33 AND NOT-SLC6A16	0.940945	0.929961	0.952191
NOT-PHLDB2 AND CD33 AND NOT-TSPAN1	0.940945	0.929961	0.952191
NOT-PHLDB2 AND CD33 AND ITGB2	0.940945	0.929961	0.952191
NOT-PHLDB2 AND CD33 AND NOT-PPAP2C	0.940945	0.929961	0.952191
NOT-PHLDB2 AND CD33 AND CD53	0.940476	0.936759	0.944223
NOT-PODXL AND CD33 AND NOT-BTN3A1	0.970179	0.968254	0.972112
NOT-PODXL AND NOT-LAT AND CD33	0.939571	0.919847	0.960159
NOT-PODXL AND NOT-LAT AND CD33	0.939571	0.919847	0.960159
NOT-PTPRK AND CD33 AND NOT-BTN3A1	0.96994	0.975806	0.964143
NOT-PHLDB2 AND CD33 AND NOT-SLC2A10	0.939096	0.926357	0.952191
NOT-PHLDB2 AND CD33 AND NOT-CACFD1	0.939096	0.926357	0.952191
NOT-PHLDB2 AND CD33 AND NOT-SCNN1A	0.939096	0.926357	0.952191
NOT-PHLDB2 AND CD33 AND NOT-CFTR	0.939096	0.926357	0.952191
NOT-PHLDB2 AND CD33 AND NOT-GOLM1	0.939096	0.926357	0.952191
NOT-PHLDB2 AND CD33 AND NOT-C1orf210	0.939096	0.926357	0.952191
NOT-PHLDB2 AND CD33 AND NOT-ITPR3	0.939096	0.926357	0.952191
NOT-PHLDB2 AND CD33 AND NOT-TSPAN3	0.939096	0.926357	0.952191
NOT-PHLDB2 AND CD33 AND NOT-PERP	0.939096	0.926357	0.952191
NOT-PHLDB2 AND CD33 AND NOT-EMP2	0.939096	0.926357	0.952191
NOT-PHLDB2 AND CD33 AND NOT-ST6GALNAC6	0.939096	0.926357	0.952191
NOT-PHLDB2 AND CD33 AND NOT-SLC38A6	0.939096	0.926357	0.952191
NOT-PHLDB2 AND CD33 AND NOT-TMPRSS2	0.939096	0.926357	0.952191
NOT-PHLDB2 AND CD33 AND NOT-WDR19	0.939096	0.926357	0.952191
NOT-PHLDB2 AND CD33 AND NOT-STOML3	0.939096	0.926357	0.952191
NOT-PHLDB2 AND CD33 AND NOT-SEMA5A	0.939096	0.926357	0.952191
NOT-PHLDB2 AND CD33 AND NOT-SLC17A5	0.939096	0.926357	0.952191
NOT-PHLDB2 AND CD33 AND NOT-PROS1	0.939096	0.926357	0.952191
NOT-PHLDB2 AND CD33 AND NOT-LSR	0.939096	0.926357	0.952191
NOT-PHLDB2 AND CD33 AND NOT-PON2	0.939096	0.926357	0.952191
NOT-PHLDB2 AND CD33 AND NOT-TSPAN6	0.939096	0.926357	0.952191
NOT-PTPRK AND CD33 AND NOT-IFNAR2	0.944882	0.933852	0.956175
NOT-PHLDB2 AND CD33 AND NOT-GPRC5A	0.937008	0.92607	0.948207
NOT-PHLDB2 AND CD33 AND NOT-AQP3	0.937008	0.92607	0.948207
NOT-PHLDB2 AND CD33 AND NOT-ATP2C2	0.937008	0.92607	0.948207
P2RY8 AND NOT-PHLDB2 AND CD33	0.93666	0.903704	0.972112
NOT-PHLDB2 AND CD33 AND CD69	0.9499	0.955645	0.944223
COMPLEX-FXYD5/PHLDB2/CD33	0.935849	0.888889	0.988048
NOT-PHLDB2 AND CXCR4 AND CD33	0.935849	0.888889	0.988048

Antigen Pair Logic	F1	Precision	Recall
EPCAM AND BST2 AND VCAM1	0.9	0.972973	0.837209
EPCAM AND S1PR1 AND NOT-GPA33	0.807692	0.688525	0.976744
EPCAM AND VCAM1 AND NOT-BEST2	0.82	0.719298	0.953488
EPCAM AND NOT-SLC22A5 AND VCAM1	0.850575	0.840909	0.860465
EPCAM AND NOT-BMPR1B AND NOT-BEST2	0.815534	0.7	0.976744
MST1R AND NOT-TYR AND CD93	0.837838	1	0.72093
EPCAM AND NOT-CATSPERD AND MOK	0.804598	0.795455	0.813953
EPCAM AND VCAM1 AND NOT-SLC30A10	0.803738	0.671875	1
CLDN18 AND SLC2A1 AND CD93	0.878049	0.923077	0.837209
CLDN18 AND NOT-AOC3 AND SLC31A1	0.805556	1	0.674419
CLDN18 AND NOT-GYPC AND SLC31A1	0.805556	1	0.674419
EPCAM AND BST2 AND NOT-S1PR1	0.886076	0.972222	0.813953
EPCAM AND BST2 AND NOT-CATSPERD	0.9	0.972973	0.837209
EPCAM AND F2R AND NOT-CATSPERD	0.886598	0.796296	1
EPCAM AND F2R AND NOT-S1PR1	0.86	0.754386	1
EPCAM AND BST2 AND NOT-C8B	0.9	0.972973	0.837209
EPCAM AND NOT-CATSPERD AND SLC6A6	0.875	0.792453	0.976744
EPCAM AND NOT-CATSPERD AND EPHA2	0.894118	0.904762	0.883721
EPCAM AND NOT-SLC22A5 AND NOT-CATSPERD	0.804348	0.755102	0.860465
EPCAM AND BST2 AND SEMA4B	0.886076	0.972222	0.813953
EPCAM AND BST2 AND NOT-OXTR	0.9	0.972973	0.837209
EPCAM AND CDH20 AND S1PR1	0.8	0.765957	0.837209
EPCAM AND TTYH3 AND S1PR1	0.844444	0.808511	0.883721
ATP8B1 AND ENG AND NOT-SLC31A1	0.857143	0.878049	0.837209
EPCAM AND NOT-OXTR AND SLC6A6	0.831683	0.724138	0.976744
THY1 AND NOT-SLC39A2 AND CD58	0.804124	0.722222	0.906977
EPCAM AND S1PR1 AND EPHA2	0.860759	0.944444	0.790698
EPCAM AND S1PR1 AND NOT-BEST2	0.803922	0.694915	0.953488
EPCAM AND NOT-SLC22A5 AND S1PR1	0.822222	0.787234	0.860465
CLDN18 AND SLC30A10 AND CD58	0.853659	0.897436	0.813953
EPCAM AND NOT-OXTR AND EPHA2	0.839506	0.894737	0.790698
MST1R AND NOT-SLC39A2 AND CD44	0.921348	0.891304	0.953488
MUC17 AND S1PR1 AND NOT-RNF43	0.826667	0.96875	0.72093
EPCAM AND CXCL16 AND NOT-CATSPERD	0.853333	1	0.744186
EPCAM AND S1PR1 AND SLC6A6	0.875	0.792453	0.976744
EPCAM AND S1PR1 AND NOT-SLC30A10	0.811321	0.68254	1
EPCAM AND S1PR1 AND LRFN2	0.82	0.719298	0.953488
EPCAM AND NOT-SLC22A5 AND NOT-OXTR	0.804348	0.755102	0.860465
EPCAM AND NOT-CATSPERD AND ICAM1	0.826087	0.77551	0.883721
EPCAM AND S1PR1 AND MLNR	0.8125	0.735849	0.906977
CLDN18 AND NOT-SLC34A2 AND ESYT2	0.878049	0.923077	0.837209
CLDN18 AND NOT-SLC34A2 AND S100A10	0.878049	0.923077	0.837209
EPCAM AND THY1 AND SLC25A5	0.860215	0.8	0.930233
EPCAM AND THY1 AND CYBA	0.898876	0.869565	0.930233
CLDN18 AND NOT-SLC34A2 AND SLMAP	0.837209	0.837209	0.837209
EPCAM AND THY1 AND NOT-PKHD1	0.91954	0.909091	0.930233
CLDN18 AND NOT-SLCO4C1 AND SLC39A6	0.857143	0.970588	0.767442
CLDN18 AND NOT-SLCO4C1 AND ANXA1	0.853659	0.897436	0.813953
CLDN18 AND NOT-SLC34A2 AND STX7	0.886076	0.972222	0.813953
CLDN18 AND NOT-SGCG AND ANXA1	0.897436	1	0.813953
EPCAM AND THY1 AND NCSTN	0.816327	0.727273	0.930233
MUC13 AND NOT-SLC16A9 AND ANXA1	0.91358	0.973684	0.860465
EPCAM AND BIRC5 AND NOT-SLC26A3	0.963855	1	0.930233
EPCAM AND THY1 AND PIGU	0.842105	0.769231	0.930233
EPCAM AND THY1 AND VNN3	0.833333	0.754717	0.930233
EPCAM AND THY1 AND TSPAN13	0.857143	0.970588	0.767442
CLDN18 AND NOT-SLC34A2 AND SLC3A2	0.9	0.972973	0.837209
EPCAM AND THY1 AND C1orf210	0.8	0.730769	0.883721
MUC13 AND NOT-SLCO4C1 AND SLC39A6	0.876404	0.847826	0.906977
CLDN18 AND NOT-SLC34A2 AND ATRAID	0.875	0.945946	0.813953
CLDN18 AND NOT-SLC34A2 AND PLGRKT	0.888889	0.947368	0.837209
CLDN18 AND NOT-SLC34A2 AND PAM	0.804598	0.795455	0.813953
EPCAM AND EPHB2 AND NOT-SLC26A3	0.901099	0.854167	0.953488
CLDN18 AND NOT-SLC34A2 AND SPINT1	0.911392	1	0.837209
CLDN18 AND NOT-SLC34A2 AND FXYD3	0.837209	0.837209	0.837209
EPCAM AND NOT-CA4 AND NOT-BMPR1B	0.857143	0.878049	0.837209
EPCAM AND THY1 AND CDHR3	0.816327	0.727273	0.930233
CLDN18 AND SLC4A11 AND NOT-TYR	0.864198	0.921053	0.813953
CLDN18 AND NOT-SLC34A2 AND MTDH	0.9	0.972973	0.837209
EPCAM AND MST1R AND NOT-SLC26A3	0.953488	0.953488	0.953488
EPCAM AND VCAM1 AND NOT-CA4	0.831169	0.941176	0.744186
CLDN18 AND NOT-CD33 AND ATP13A1	0.86747	0.9	0.837209
EPCAM AND NOT-ENPP3 AND PIGU	0.91954	0.909091	0.930233
EPCAM AND SPON2 AND NOT-CDHR3	0.931818	0.911111	0.953488
EPCAM AND CDH3 AND NOT-BMPR1B	0.869565	0.816327	0.930233
EPCAM AND SLC7A5 AND LRP11	0.939759	0.975	0.906977
EPCAM AND NOT-SLCO4C1 AND SLC39A6	0.824742	0.740741	0.930233

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
COMPLEX-LPAR1/PHLDB2/CD33	0.939096	0.926357	0.952191
NOT-PHLDB2 AND CD33 AND NOT-AMIGO2	0.934911	0.925781	0.944223
NOT-PHLDB2 AND CD33 AND SELL	0.934132	0.936	0.932271
NOT-PHLDB2 AND CD33 AND LTBR	0.933602	0.943089	0.924303
NOT-PODXL AND CD33 AND NOT-IFNAR2	0.941634	0.920152	0.964143
NOT-PHLDB2 AND CD33 AND NOT-ST14	0.930693	0.925197	0.936255
NOT-PTPRK AND NOT-S1PR1 AND CD33	0.930502	0.902622	0.960159
NOT-PTPRK AND CD33 AND NOT-SEMA4D	0.930502	0.902622	0.960159
NOT-PTPRK AND NOT-S1PR1 AND CD33	0.930502	0.902622	0.960159
NOT-PHLDB2 AND CD33 AND NOT-LRP11	0.930417	0.928571	0.932271
NOT-PTPRK AND CD33 AND NOT-BTN3A3	0.956	0.959839	0.952191
NOT-PHLDB2 AND CD33 AND NOT-TSPAN31	0.928571	0.924901	0.932271
NOT-PODXL AND CD33 AND NOT-S1PR1	0.925253	0.938525	0.912351
NOT-PHLDB2 AND CD33 AND NOT-CD46	0.925	0.969432	0.884462
NOT-PODXL AND CD33 AND NOT-STEAP4	0.953157	0.975	0.932271
NOT-PODXL AND CD33 AND NOT-SEMA4D	0.923954	0.883636	0.968127
NOT-PTPRK AND CD33 AND NOT-PIK3IP1	0.91498	0.930041	0.900398
CD33 AND NOT-ATP8B1 AND NOT-CD27	0.926148	0.928	0.924303
NOT-PTPRK AND CD33 AND NOT-KCNK6	0.907721	0.860714	0.960159
NOT-PTPRK AND CD33 AND NOT-FMNL1	0.907692	0.877323	0.940239
NOT-PODXL AND CD33 AND NOT-TGFBI	0.907258	0.918367	0.896414
NOT-PTPRK AND CD33 AND NOT-SLC6A6	0.906367	0.855124	0.964143
NOT-PTPRK AND CD33 AND NOT-TGFBI	0.906122	0.92887	0.884462
NOT-PTPRK AND CD33 AND NOT-AOC3	0.904315	0.85461	0.960159
NOT-PODXL AND CD33 AND NOT-PIK3IP1	0.904315	0.85461	0.960159
NOT-PTPRK AND CD33 AND NOT-IL17RA	0.904215	0.870849	0.940239
CD33 AND NOT-AOC3 AND NOT-EPHA4	0.932271	0.932271	0.932271
CD33 AND NOT-STX7 AND NOT-GHR	0.956701	0.991453	0.924303
CD33 AND NOT-STX7 AND NOT-ATP8B1	0.954733	0.987234	0.924303
NOT-PODXL AND NOT-GPR137B AND CD33	0.898969	0.931624	0.868526
NOT-PTPRK AND NOT-GPR137B AND CD33	0.898969	0.931624	0.868526
CD33 AND NOT-LAT AND NOT-DAG1	0.910931	0.925926	0.896414
CD33 AND NOT-ATP8B1 AND NOT-HVCN1	0.91453	0.986175	0.85259
CD33 AND NOT-ATP8B1 AND NOT-PAG1	0.927126	0.942387	0.912351
COMPLEX-SLC4A1/STEAP4/CD33	0.896679	0.835052	0.968127
NOT-PHLDB2 AND CD33 AND NOT-ADIPOR1	0.892704	0.967442	0.828685
CD33 AND NOT-SEMA4D AND NOT-DAG1	0.903101	0.879245	0.928287
COMPLEX-SLC4A1/TGFBI/CD33	0.891945	0.879845	0.904382
CD33 AND NOT-EPHA4 AND NOT-GHR	0.915187	0.90625	0.924303
CD33 AND NOT-BTN3A1 AND NOT-DAG1	0.945233	0.96281	0.928287
COMPLEX-BTN3A1/SLC4A1/CD33	0.931818	0.888087	0.98008
NOT-PTPRK AND CD70 AND NOT-HLA-DOB	0.927374	0.870629	0.992032
NOT-PTPRK AND CD70 AND NOT-SLAMF7	0.932331	0.882562	0.988048
NOT-PTPRK AND CD70 AND NOT-FCRL5	0.925094	0.872792	0.984064
NOT-PTPRK AND CD70 AND NOT-MS4A1	0.930057	0.884892	0.98008
P2RY8 AND NOT-HLA-DOB AND NOT-AXL	0.922787	0.875	0.976096
NOT-PTPRK AND SLC7A5 AND DDX3X	0.922495	0.877698	0.972112
NOT-PTPRK AND CD37 AND NOT-CD160	0.921905	0.883212	0.964143
P2RY8 AND NOT-ST8SIA1 AND NOT-MS4A1	0.935673	0.916031	0.956175
P2RY8 AND NOT-MS4A1 AND NOT-KDR	0.927757	0.887273	0.972112
NOT-PTPRK AND CD70 AND NOT-SPON2	0.923933	0.864583	0.992032
P2RY8 AND NOT-HLA-DOB AND NOT-CLDN8	0.926829	0.875887	0.984064
P2RY8 AND NOT-ST8SIA1 AND NOT-FCRL1	0.924855	0.895522	0.956175
NOT-PTPRK AND CD37 AND NOT-SLAMF7	0.919021	0.871429	0.972112
NOT-PTPRK AND SLC7A5 AND NOT-SLAMF7	0.918519	0.858131	0.988048
NOT-PODXL AND CD37 AND NOT-CD160	0.918406	0.876812	0.964143
NOT-PODXL AND CD37 AND NOT-MS4A1	0.918406	0.876812	0.964143
NOT-PTPRK AND CD37 AND NOT-MS4A1	0.918406	0.876812	0.964143
P2RY8 AND NOT-ST8SIA1 AND NOT-FCRL5	0.930769	0.899628	0.964143
P2RY8 AND NOT-MS4A1 AND NOT-AXL	0.920152	0.88	0.964143
P2RY8 AND NOT-MS4A1 AND NOT-AXL	0.920152	0.88	0.964143
P2RY8 AND NOT-PCYT1A AND NOT-CR2	0.936902	0.900735	0.976096
COMPLEX-MST1R/PTPRK/SLC7A5	0.917293	0.868327	0.972112
P2RY8 AND NOT-HLA-DOB AND NOT-EDNRB	0.916667	0.873646	0.964143
COMPLEX-PTPRK/SSTR4/SLC7A5	0.915572	0.865248	0.972112
NOT-PTPRK AND SLC7A5 AND NOT-HLA-DOB	0.915441	0.849829	0.992032
P2RY8 AND NOT-MS4A1 AND NOT-EDNRB	0.915709	0.881919	0.952191
NOT-PTPRK AND CD70 AND NOT-FCRL1	0.914498	0.857143	0.98008
NOT-PTPRK AND SLC7A5 AND NOT-FCRL5	0.914179	0.859649	0.976096
NOT-PTPRK AND CD37 AND NOT-FCRL5	0.913208	0.867384	0.964143
NOT-PTPRK AND CD70 AND NOT-CD22	0.918216	0.860627	0.984064
P2RY8 AND NOT-MS4A1 AND NOT-TNC	0.927757	0.887273	0.972112
P2RY8 AND NOT-ST8SIA1 AND NOT-FCRL2	0.927203	0.892989	0.964143
NOT-PTPRK AND SLC7A5 AND NOT-CD160	0.911439	0.848797	0.984064
P2RY8 AND NOT-ST8SIA1 AND NOT-CR2	0.918406	0.876812	0.964143
COMPLEX-PTPRK/VTCN1/SLC7A5	0.910448	0.85614	0.972112
NOT-PODXL AND CD37 AND NOT-FCRL5	0.910448	0.85614	0.972112

Antigen Pair Logic	F1	Precision	Recall
EPCAM AND SPON2 AND EMP1	0.964706	0.97619	0.953488
EPCAM AND NOT-ENPP3 AND CLSTN1	0.831683	0.724138	0.976744
EPCAM AND NOT-CLDN8 AND PIGU	0.810811	0.967742	0.697674
EPCAM AND SPON2 AND NOT-VNN3	0.931818	0.911111	0.953488
EPCAM AND SPON2 AND HEPH	0.931818	0.911111	0.953488
EPCAM AND SPON2 AND PRRG1	0.931818	0.911111	0.953488
EPCAM AND CDH3 AND VCAM1	0.833333	0.754717	0.930233
EPCAM AND FAP AND SLC25A5	0.853933	0.826087	0.883721
CLDN18 AND EPCAM AND NOT-CDHR3	0.875	0.945946	0.813953
CLDN18 AND NOT-SLC34A2 AND F11R	0.883117	1	0.790698
EPCAM AND SLC7A5 AND CYBA	0.95122	1	0.906977
CLDN18 AND EPCAM AND NOT-OR2S2	0.883117	1	0.790698
EPCAM AND SLC7A5 AND NOT-FXYD3	0.917647	0.928571	0.906977
EPCAM AND BIRC5 AND NOT-C1orf210	0.925	1	0.860465
EPCAM AND NOT-SCNN1B AND VCAM1	0.934783	0.877551	1
EPCAM AND NOT-ENPP3 AND ZDHHC5	0.965517	0.954545	0.976744
EPCAM AND NOT-CLDN8 AND ZDHHC5	0.857143	0.970588	0.767442
EPCAM AND SLC7A5 AND C1orf210	0.891566	0.925	0.860465
EPCAM AND FAP AND CYBA	0.853933	0.826087	0.883721
EPCAM AND SLC7A5 AND VNN3	0.917647	0.928571	0.906977
EPCAM AND FAP AND NOT-OR2S2	0.886076	0.972222	0.813953
EPCAM AND NOT-ENPP3 AND NOT-PKHD1	0.875	0.792453	0.976744
CLDN18 AND EPCAM AND VNN3	0.888889	0.947368	0.837209
EPCAM AND SLC7A5 AND NOT-SLC26A3	0.904762	0.926829	0.883721
EPCAM AND SLC7A5 AND OR2S2	0.904762	0.926829	0.883721
EPCAM AND SLC7A5 AND NOT-EMP1	0.835443	0.916667	0.767442
MUC13 AND NOT-SPPL2A AND NOT-BMPR1B	0.88172	0.82	0.953488
CLDN18 AND NOT-SLC34A2 AND SPPL2A	0.883117	1	0.790698
MUC13 AND SLC7A5 AND NOT-FXYD3	0.897436	1	0.813953
EPCAM AND NOT-SCNN1B AND NOT-BMPR1B	0.886598	0.796296	1
CLDN18 AND EPCAM AND PIGU	0.883117	1	0.790698
EPCAM AND NOT-CLDN8 AND TMEM63B	0.804878	0.846154	0.767442
EPCAM AND NOT-ENPP3 AND CYBA	0.954545	0.933333	0.976744
EPCAM AND NOT-CLDN8 AND CYBA	0.857143	0.970588	0.767442
EPCAM AND NOT-CLDN8 AND ADAM10	0.835443	0.916667	0.767442
CDH17 AND NOT-SCNN1B AND CD44	0.911111	0.87234	0.953488
CDH17 AND NOT-SCNN1B AND TGFBI	0.870588	0.880952	0.860465
CDH17 AND NOT-SLC30A10 AND SDC4	0.953488	0.953488	0.953488
CDH17 AND NOT-FXYD3 AND CD93	0.857143	0.8125	0.906977
CDH17 AND NOT-SLC30A10 AND SPINT1	0.931818	0.911111	0.953488
CDH17 AND NOT-SLC30A10 AND SLC25A5	0.901099	0.854167	0.953488
CDH17 AND NOT-SLC30A10 AND PCDHB7	0.811881	0.706897	0.953488
CDH17 AND NOT-SLC30A10 AND F11R	0.911111	0.87234	0.953488
CDH17 AND NOT-SLC30A10 AND MTDH	0.931818	0.911111	0.953488
PTPRH AND NOT-SLC30A10 AND CD58	0.87234	0.803922	0.953488
FUT3 AND NOT-SCNN1B AND CD44	0.869565	0.816327	0.930233
MUC17 AND NOT-SLC30A10 AND LRP8	0.853333	1	0.744186
MUC17 AND NOT-SLC30A10 AND ESYT2	0.853333	1	0.744186
MUC17 AND NOT-SPPL2A AND NOT-GPR22	0.842105	0.969697	0.744186
CDH17 AND NOT-SLC30A10 AND CD99	0.823529	0.833333	0.813953
ATP8B1 AND CD93 AND NOT-SLC6A1	0.826667	0.96875	0.72093
ATP8B1 AND PLXND1 AND NOT-SLC31A1	0.871795	0.971429	0.790698
PVRL3 AND NOT-SLC30A10 AND CD58	0.839506	0.894737	0.790698
SYT13 AND NOT-SLC30A10 AND CD58	0.909091	0.888889	0.930233
COMPLEX-MUC17/CD93/LRP8	0.853333	1	0.744186
ATP8B1 AND CD93 AND NOT-STX7	0.871795	0.971429	0.790698
COMPLEX-MUC17/CD93/SPINT1	0.826667	0.96875	0.72093
MUC17 AND NOT-SCNN1B AND CD44	0.853333	1	0.744186
FUT3 AND NOT-SCNN1B AND TGFBI	0.886076	0.972222	0.813953
ATP8B1 AND TGFBI AND NOT-SCNN1B	0.883117	1	0.790698
ATP8B1 AND NOT-SCNN1B AND CD44	0.91358	0.973684	0.860465
MUC17 AND NOT-GPR22 AND PCDHB7	0.842105	0.969697	0.744186
MUC17 AND NOT-SPPL2A AND NOT-MRAP	0.842105	0.969697	0.744186
MUC17 AND NOT-SPPL2A AND NOT-CD1A	0.842105	0.969697	0.744186
MUC17 AND NOT-SPPL2A AND NOT-KCND2	0.842105	0.969697	0.744186
MUC17 AND SLC39A10 AND NOT-CD1A	0.831169	0.941176	0.744186
MUC17 AND SLC39A10 AND NOT-OMG	0.831169	0.941176	0.744186
MUC17 AND SLC39A10 AND NOT-SLCO1C1	0.831169	0.941176	0.744186
TSPAN8 AND CD93 AND SLC2A1	0.826667	0.96875	0.72093
MUC17 AND NOT-SLCO4C1 AND GYPC	0.815789	0.939394	0.72093
TSPAN8 AND CD93 AND SLC31A1	0.826667	0.96875	0.72093
MUC17 AND LRP11 AND NOT-AOC3	0.831169	0.941176	0.744186
MUC17 AND NOT-SLCO4C1 AND PTGIS	0.826667	0.96875	0.72093
MUC17 AND SYT11 AND NOT-ZDHHC5	0.810811	0.967742	0.697674
MUC17 AND LRP11 AND NOT-SLC39A2	0.831169	0.941176	0.744186
MUC17 AND HEG1 AND NOT-GPR6	0.831169	0.941176	0.744186
MUC17 AND NOT-SLCO4C1 AND VANGL1	0.831169	0.941176	0.744186

FIG. 15 (cont.)

Antigen Pair Logic	F1	Precision	Recall
CXCR4 AND NOT-SLAMF7 AND NOT-KDR	0.913958	0.878676	0.952191
P2RY8 AND NOT-GPNMB AND NOT-MS4A1	0.909441	0.880597	0.940239
P2RY8 AND NOT-GPNMB AND NOT-MS4A1	0.909441	0.880597	0.940239
P2RY8 AND NOT-HLA-DOB AND NOT-CLDN5	0.926829	0.875887	0.984064
NOT-PTPRK AND SLC7A5 AND NOT-ALDH1A1	0.907721	0.860714	0.960159
P2RY8 AND NOT-MS4A1 AND NOT-CLDN8	0.920755	0.874552	0.972112
P2RY8 AND NOT-MS4A1 AND NOT-CLDN8	0.920755	0.874552	0.972112
P2RY8 AND NOT-MS4A1 AND NOT-CLDN11	0.924242	0.880866	0.972112
NOT-PODXL AND CD37 AND NOT-HLA-DOB	0.907407	0.847751	0.976096
NOT-PTPRK AND CD37 AND NOT-HLA-DOB	0.907407	0.847751	0.976096
P2RY8 AND NOT-MS4A1 AND NOT-STEAP1	0.919847	0.882784	0.960159
P2RY8 AND NOT-MS4A1 AND NOT-PCYT1A	0.938224	0.910112	0.968127
NOT-PTPRK AND CD37 AND NOT-FCRL1	0.90631	0.871324	0.944223
P2RY8 AND NOT-MS4A1 AND NOT-CEACAM5	0.924242	0.880866	0.972112
P2RY8 AND NOT-MS4A1 AND NOT-RNF43	0.922495	0.877698	0.972112
P2RY8 AND NOT-MS4A1 AND NOT-ITGAV	0.905222	0.879699	0.932271
P2RY8 AND NOT-MS4A1 AND NOT-FOLR2	0.904854	0.882576	0.928287
P2RY8 AND NOT-HLA-DOB AND NOT-GPNMB	0.904215	0.870849	0.940239
CXCR4 AND NOT-SLAMF7 AND NOT-SDC1	0.903592	0.859712	0.952191
P2RY8 AND NOT-MS4A1 AND NOT-ROR1	0.920755	0.874552	0.972112
P2RY8 AND NOT-MS4A1 AND NOT-GPC3	0.920755	0.874552	0.972112
P2RY8 AND NOT-FCRL1 AND NOT-TNC	0.903346	0.84669	0.968127
NOT-PTPRK AND CD70 AND NOT-CD79A	0.903108	0.834459	0.984064
CXCR4 AND NOT-HLA-DOB AND NOT-KDR	0.902033	0.841379	0.972112
NOT-PODXL AND CD37 AND NOT-CR2	0.902033	0.841379	0.972112
P2RY8 AND NOT-MS4A1 AND NOT-CLDN23	0.920455	0.877256	0.968127
P2RY8 AND NOT-PCYT1A AND NOT-FCRL1	0.920455	0.877256	0.968127
NOT-EDNRB AND ITGB2 AND NOT-HLA-DOB	0.900735	0.836177	0.976096
CXCR4 AND NOT-HLA-DOB AND NOT-SDC1	0.900369	0.838488	0.972112
CD37 AND NOT-MS4A1 AND P2RY8	0.900196	0.884615	0.916335

(cont. at top)

Antigen Pair Logic	F1	Precision	Recall
MUC17 AND PCDHB7 AND NOT-ATP13A5	0.842105	0.969697	0.744186
MUC17 AND CSF2RB AND NOT-ATP13A5	0.831169	0.941176	0.744186
COMPLEX-MUC17/ATP13A5/SPPL2A	0.842105	0.969697	0.744186
COMPLEX-MUC17/CACNG6/SPPL2A	0.842105	0.969697	0.744186
EPCAM AND SPON2 AND NOT-BMPR1B	0.942529	0.931818	0.953488
EPCAM AND SPON2 AND VCAM1	0.942529	0.931818	0.953488
MUC13 AND SLC7A5 AND NOT-PTK7	0.938272	1	0.883721
EPCAM AND NOT-CLDN8 AND VCAM1	0.835443	0.916667	0.767442
CLDN18 AND RNF43 AND NOT-TYR	0.843373	0.875	0.813953
CEACAM5 AND SLC7A5 AND PTK7	0.814815	0.868421	0.767442
EPCAM AND NOT-BMPR1B AND CLDN1	0.955556	0.914894	1
EPCAM AND VCAM1 AND CLDN1	0.945055	0.895833	1
EPCAM AND NOT-BMPR1B AND NOT-GPA33	0.807692	0.688525	0.976744
EPCAM AND VCAM1 AND NOT-GPA33	0.823529	0.711864	0.976744
EPCAM AND TNC AND NOT-BMPR1B	0.808511	0.745098	0.883721
EPCAM AND VCAM1 AND NOT-SSTR2	0.824742	0.740741	0.930233
EPCAM AND FCRL5 AND BMPR1B	0.836735	0.745455	0.953488
EPCAM AND VCAM1 AND MOK	0.818182	0.8	0.837209
CDH17 AND NOT-SLC30A10 AND CD58	0.909091	0.888889	0.930233
COMPLEX-SLC31A1/MUC17/CD1A	0.815789	0.939394	0.72093
MUC17 AND CD44 AND NOT-SLC39A2	0.842105	0.969697	0.744186
ATP8B1 AND AOC3 AND NOT-SLC31A1	0.847059	0.857143	0.837209
ATP8B1 AND SYT11 AND NOT-SLC31A1	0.829268	0.871795	0.790698
ATP8B1 AND CD44 AND NOT-SLC39A2	0.888889	0.947368	0.837209
FAT1 AND NOT-SLC39A2 AND CD44	0.828283	0.732143	0.953488
ATP8B1 AND ANTXR2 AND TMPRSS11D	0.823529	0.833333	0.813953
ATP8B1 AND ANTXR2 AND SLCO1C1	0.853659	0.897436	0.813953
ATP8B1 AND ANTXR2 AND NKAIN2	0.875	0.945946	0.813953
FAT1 AND NOT-SLC30A10 AND CD58	0.863158	0.788462	0.953488
TTYH3 AND NOT-CD1A AND AOC3	0.8	0.809524	0.790698

INTERNATIONAL SEARCH REPORT

International application No.
PCT/US17/31389

A. CLASSIFICATION OF SUBJECT MATTER
IPC - A61K 35/14; C12N 5/0783; C07K 14/725 (2017.01)
CPC - A61K 35/17, C12N 5/0636, 5/0638, 9/16; C07K 14/4703, 14/7051, 14/7155, 16/2803, 16/2866

According to International Patent Classification (IPC) or to both national classification and IPC

B. FIELDS SEARCHED

Minimum documentation searched (classification system followed by classification symbols)
See Search History document

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched
See Search History document

Electronic data base consulted during the international search (name of data base and, where practicable, search terms used)
See Search History document

C. DOCUMENTS CONSIDERED TO BE RELEVANT

Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
X --- Y	WO 2015/124715 A1 (CELLECTIS) August 27, 2015; abstract; page 3, lines 10-13; page 10, lines 2-4; page 19, lines 25-27; page 22, lines 8-9; page 24, lines 5-6; page 25, lines 18-30; claims 1, 9, 13, 16, 38	1-2, 3/1-2, 19, 23, 25/19, 25/23, 26/19, 26/23, 27, 30 --- 20-22, 24 25/20-22, 25/24, 26/20-22, 26/24, 28-29
X --- Y	WO 2015/123642 A1 (BOARD OF REGENTS, THE UNIVERSITY OF TEXAS SYSTEM) August 20, 2015; paragraphs [0009], [0021], [0025], [0096]	46-48, 49/46-48 --- 24, 25/24, 26/24, 28
Y	MORSUT et al. 'Engineering Customized Cell Sensing and Response Behaviors Using Synthetic Notch Receptors' Cell, February 11, 2016, Vol. 164, No 4, pp 1-25; abstract; page 3, paragraph 4; page 6, paragraph 3. DOI:10.1016/j.cell.2016.01.012.	20-22, 25/20-22, 26/20-22
Y	WO 2016/033331 A1 (BIOATLA, LLC) August 27, 2015; abstract; paragraph [00040]	29
A	US 2010/0304410 A1 (KIJANKA et al.) December 2, 2010; paragraphs [0012], [0053]	35-36, 37/35-36
A	WO 2015/105995 A2 (THE BOARD OF TRUSTEES OF THE LELAND STANFORD JUNIOR UNIVERSITY) November 12, 2015; paragraph [00102]	35-36, 37/35-36

☐ Further documents are listed in the continuation of Box C. ☐ See patent family annex.

* Special categories of cited documents:
"A" document defining the general state of the art which is not considered to be of particular relevance
"E" earlier application or patent but published on or after the international filing date
"L" document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified)
"O" document referring to an oral disclosure, use, exhibition or other means
"P" document published prior to the international filing date but later than the priority date claimed
"T" later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention
"X" document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step when the document is taken alone
"Y" document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents, such combination being obvious to a person skilled in the art
"&" document member of the same patent family

Date of the actual completion of the international search	Date of mailing of the international search report
17 July 2017 (17.07.2017)	07 AUG 2017
Name and mailing address of the ISA/ Mail Stop PCT, Attn: ISA/US, Commissioner for Patents P.O. Box 1450, Alexandria, Virginia 22313-1450 Facsimile No. 571-273-8300	Authorized officer Shane Thomas PCT Helpdesk: 571-272-4300 PCT OSP: 571-272-7774

Form PCT/ISA/210 (second sheet) (January 2015)

INTERNATIONAL SEARCH REPORT

International application No.
PCT/US17/31389

Box No. II Observations where certain claims were found unsearchable (Continuation of item 2 of first sheet)

This international search report has not been established in respect of certain claims under Article 17(2)(a) for the following reasons:

1. ☐ Claims Nos.:
because they relate to subject matter not required to be searched by this Authority, namely:

2. ☐ Claims Nos.:
because they relate to parts of the international application that do not comply with the prescribed requirements to such an extent that no meaningful international search can be carried out, specifically:

3. ☒ Claims Nos.: 4-18, 31-34, 38-45
because they are dependent claims and are not drafted in accordance with the second and third sentences of Rule 6.4(a).

Box No. III Observations where unity of invention is lacking (Continuation of item 3 of first sheet)

This International Searching Authority found multiple inventions in this international application, as follows:

1. ☐ As all required additional search fees were timely paid by the applicant, this international search report covers all searchable claims.

2. ☐ As all searchable claims could be searched without effort justifying additional fees, this Authority did not invite payment of additional fees.

3. ☐ As only some of the required additional search fees were timely paid by the applicant, this international search report covers only those claims for which fees were paid, specifically claims Nos.:

4. ☐ No required additional search fees were timely paid by the applicant. Consequently, this international search report is restricted to the invention first mentioned in the claims; it is covered by claims Nos.:

Remark on Protest

☐ The additional search fees were accompanied by the applicant's protest and, where applicable, the payment of a protest fee.

☐ The additional search fees were accompanied by the applicant's protest but the applicable protest fee was not paid within the time limit specified in the invitation.

☐ No protest accompanied the payment of additional search fees.