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(54) Title: ANTIBODIES BINDING CD40 AND USES THEREOF

(57) Abstract: Provided is an isolated monoclonal antibody or an antigen-binding portion thereof that specifically binds human CD40. A nucleic acid molecule encoding the antibody or antigen-binding portion thereof, an expression vector, a host cell and a method for expressing the antibody or the antigen-binding portion thereof are also provided. An immunoconjugate, a bispecific molecule, a chimeric antigen receptor, an oncolytic virus and a pharmaceutical composition comprising the antibody or antigen-binding portion thereof, as well as a treatment method using the same are further provided.



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ANTIBODIES BINDING CD40 AND USES THEREOF

RELATED APPLICATIONS AND INCORPORATION BY REFERENCE

[0001] This application claims priority to US provisional patent application Serial No. 63/001,612 filed on March 30, 2020.

[0002] The foregoing applications, and all documents cited therein or during their prosecution (“appln cited documents”) and all documents cited or referenced herein (including without limitation all literature documents, patents, published patent applications cited herein) (“herein cited documents”), and all documents cited or referenced in herein cited documents, together with any manufacturer’s instructions, descriptions, product specifications, and product sheets for any products mentioned herein or in any document incorporated by reference herein, are hereby incorporated herein by reference, and may be employed in the practice of the invention. More specifically, all referenced documents are incorporated by reference to the same extent as if each individual document was specifically and individually indicated to be incorporated by reference. Any Genbank sequences mentioned in this disclosure are incorporated by reference with the Genbank sequence to be that of the earliest effective filing date of this disclosure.

FIELD OF THE INVENTION

[0003] The present disclosure relates generally to an isolated monoclonal antibody, particularly a mouse, chimeric or humanized monoclonal antibody, or an antigen-binding portion thereof, that specifically binds to human CD40 with high affinity and functionality. A nucleic acid molecule encoding the antibody or antigen-binding portion thereof, an expression vector, a host cell and a method for expressing the antibody or antigen-binding portion thereof are also provided. The present disclosure further provides an immunoconjugate, a bispecific molecule, a chimeric antigen receptor, an oncolytic virus, and a pharmaceutical composition comprising the antibody or antigen-binding portion thereof, as well as a treatment method using the anti-CD40 antibody or antigen-binding portion thereof of the disclosure.

BACKGROUND OF THE INVENTION

[0004] B lymphocyte activation requires antigen receptor-mediated stimulation and co-stimulation, and CD40 is one of the costimulatory molecules participating in the activation process (Jodi L. Karnell *et al.*, (2019) *Advanced Drug Delivery Review* 141:92-103).

[0005] CD40, a type I transmembrane protein, is a member of the TNF receptor superfamily. It was initially characterized on B lymphocytes where it is constitutively expressed and signals to promote B cell activation and proliferation. Later it was found on dendritic cells (DCs), monocytes, macrophages as well as non-hematopoietic cells. The major ligand of CD40 is CD40L, which is primarily expressed by activated T cells and activated B cells and platelets, and is also found on monocytic cells, natural killer cells, and basophils under inflammatory conditions (Jodi L. Karnell *et al.*, (2019) *supra*). The wide distribution of this costimulatory pair indicates the pivotal roles they play in immune processes. For example, the engagement of CD40 with CD40L on DCs promotes cytokine production and costimulatory molecule induction, resulting in T cell activation and differentiation (Quezada SA *et al.*, (2004) *Annu Rev Immunol.* 22:307-328).

[0006] CD40 is also expressed on tumors, including B-cell malignancies, lung, bladder, gastric, breast and ovarian cancers, and has been reported to be involved in pathologies of several inflammatory diseases, including autoimmune diseases, atherothrombosis, cancers, and respiratory diseases (Costello *et al.*, (1999) *Immunol Today* 20(11): 488-493; Tong *et al.*, (2003) *Cancer Gene Ther* 10(1): 1-13; Lee *et al.*, (2014) *Curr Cancer Drug Targets* 14(7): 610-620; Ara A *et al.*, (2018) *supra*; Lee *et al.*, (1999) *Proc Natl Acad Sci USA* 96:9136-9141; Stamenkovic *et al.*, (1989) *EMBO J.* 8:1403-1410). CD40-mediated signaling, on one hand, caused tumor cell growth inhibition and cell death in some B cell-derived tumor lines (Grafton *et al.*, (1997) *Cell. Immunol.* 182:45-56), while, on the other hand, induced increased expression of many factors that protected tumor cells from apoptosis in certain other B-cell malignancies (Lee *et al.*, (1999) *Proc Natl Acad Sci USA* 96:9136-9141).

[0007] Agonistic anti-CD40 antibodies that activate or induce CD40 signaling upon binding CD40, and antagonistic anti-CD40 antibodies that block or inhibit CD40 signaling that may be induced by CD40L engagement, have been developed for disease treatment. Selicrelumab (Pfizer and VLST), an agonistic anti-CD40 antibody, has shown clinical efficacy in a number of settings of patients with advanced cancers (Vonderheide *et al.*, (2013) *Clin Cancer Res.* 19(5):1035-1043). Dacetuzumab (Sett Genetiscs), a weaker CD40 agonist than Selicrelumab, shows anti-tumor activity in diffuse large B cell lymphoma, multiple myeloma and CLL, and has been tested in combination with Rituximab and Gemcitabine in treatment of relapsed or refractory DLBCL (Advani R *et al.*, (2009) *J Clin Oncol.* 27:4371-4377; Furman RR *et al.*, (2010) *Leuk Lymphoma.* 51:228-235; Forero-Torres A *et al.*, (2012) *Leuk Lymphoma* 54(2):277-283). Lucatumumab (Novartis), an antagonistic anti-CD40 antibody, has been tested in clinical trials for treating multiple myeloma and chronic lymphocytic leukemia (Hassan SB *et al.*, (2014) *Immunopharmacol immunotoxicol* 36(2):96-104). Biologics that agonize CD40 signaling also showed efficacy in treating infectious and autoimmune diseases, including HIV-1/AIDS, tuberculosis and malaria (Elizabeth A Thompson, *et al.*, (2015) *J Immunol.* 195(3): 1015-1024).

[0008] There remains a need for more anti-CD40 antibodies with improved pharmaceutical characteristics.

[0009] Citation or identification of any document in this application is not an admission that such document is available as prior art to the present invention.

[0010] SUMMARY OF THE INVENTION

[0011] The present disclosure provides an isolated monoclonal antibody, for example, a mouse, human, chimeric or humanized monoclonal antibody, or an antigen-binding portion thereof, that binds to CD40 (e.g., the human CD40, and monkey CD40) and has comparable, if not higher, binding affinity to CD40, and comparable, if not higher, activity of activating CD40 signaling as compared to prior art anti-CD40 antibodies such as Dacetuzumab and Selicrelumab.

[0012] The antibody or antigen-binding portion thereof of the disclosure can be used for a variety of applications, including detection of the CD40 protein, and treatment and prevention of CD40 associated diseases, such as cancers, infectious diseases and autoimmune diseases.

[0013] Accordingly, in one aspect, the disclosure pertains to an isolated monoclonal antibody (e.g., a mouse, chimeric or humanized antibody), or an antigen-binding portion thereof, that binds CD40, having i) a heavy chain variable region that may comprise a VH CDR1 region, a VH CDR2 region

and a VH CDR3 region, wherein the VH CDR1 region, the VH CDR2 region and the VH CDR3 region may comprise amino acid sequences having at least 85%, 86%, 87%, 88%, 89%, 90%, 91%, 92%, 93%, 94%, 95%, 96%, 97%, 98%, 99% or 100% identity to (1) SEQ ID NOs: 1, 4 and 7, respectively; (2) SEQ ID NOs: 2, 5 and 8, respectively; or (3) SEQ ID NOs: 3, 6 and 9, respectively; and/or ii) a light chain variable region that may comprise a VL CDR1 region, a VL CDR2 region and a VL CDR3 region, wherein the VL CDR1 region, the VL CDR2 region, and the VL CDR3 region may comprise amino acid sequences having at least 85%, 86%, 87%, 88%, 89%, 90%, 91%, 92%, 93%, 94%, 95%, 96%, 97%, 98%, 99% or 100% identity to (1) SEQ ID NOs: 10, 13 and 16, respectively; (2) SEQ ID NOs: 11, 14 and 17, respectively; or (3) SEQ ID NOs: 12, 15 and 18, respectively.

[0014] The antibody or antigen-binding portion thereof of the disclosure may comprise a heavy chain variable region that may comprise a VH CDR1 region, a VH CDR2 region and a VH CDR3 region, and a light chain variable region that may comprise a VL CDR1 region, a VL CDR2 region and a VL CDR3 region, wherein the VH CDR1 region, the VH CDR2 region, the VH CDR3 region, the VL CDR1 region, the VL CDR2 region, and the VL CDR3 region may comprise amino acid sequences having at least 85%, 86%, 87%, 88%, 89%, 90%, 91%, 92%, 93%, 94%, 95%, 96%, 97%, 98%, 99% or 100% identity to (1) SEQ ID NOs: 1, 4, 7, 10, 13 and 16, respectively; (2) SEQ ID NOs: 2, 5, 8, 11, 14 and 17, respectively; or (3) SEQ ID NOs: 3, 6, 9, 12, 15 and 18, respectively, wherein the antibody or antigen-binding fragment thereof binds to CD40.

[0015] The heavy chain variable region of the antibody or antigen-binding portion thereof of the disclosure may comprise an amino acid sequence having at least 85%, 86%, 87%, 88%, 89%, 90%, 91%, 92%, 93%, 94%, 95%, 96%, 97%, 98%, 99% or 100% identity to SEQ ID NOs: 19, 20 (X1=A or S), 21 or 22, wherein the antibody or antigen-binding fragment thereof binds to CD40. The amino acid sequence of SEQ ID NO: 19 may be encoded by nucleotide sequences of SEQ ID NOs: 31 or 32, and the amino acid sequence of SEQ ID NO: 20 (X1=S) may be encoded by the nucleotide sequence of SEQ ID NO: 33.

[0016] The light chain variable region of the antibody or antigen-binding portion thereof of the disclosure may comprise an amino acid sequence having at least 85%, 86%, 87%, 88%, 89%, 90%, 91%, 92%, 93%, 94%, 95%, 96%, 97%, 98%, 99% or 100% identity to SEQ ID NOs: 23, 24 (X1=K, X2=F; or X1=Y, X2=Y), 25 or 26, wherein the antibody or antigen-binding fragment thereof binds to CD40. The amino acid sequence of SEQ ID NO: 23 may be encoded by nucleotide sequences of SEQ ID NOs: 34 or 35. The amino acid sequences of SEQ ID NO: 24 (X1=K, X2=F) may be encoded by the nucleotide sequence of SEQ ID NO: 36.

[0017] The antibody or antigen-binding portion thereof of the disclosure may comprise a heavy chain variable region and a light chain variable region having amino acid sequences having at least 85%, 86%, 87%, 88%, 89%, 90%, 91%, 92%, 93%, 94%, 95%, 96%, 97%, 98%, 99% or 100% identity to (1) SEQ ID NOs: 19 and 23, respectively; (2) SEQ ID NOs: 20 (X1=A) and 24 (X1=K, X2=F), respectively; (3) SEQ ID NOs: 20 (X1=S) and 24 (X1=K, X2=F), respectively; (4) SEQ ID NOs: 20 (X1=A) and 24 (X1=Y, X2=Y), respectively; (5) SEQ ID NOs: 20 (X1=S) and 24 (X1=Y, X2=Y), respectively; (6) SEQ ID NOs: 21 and 25, respectively; or (7) SEQ ID NOs: 22 and 26, respectively, wherein the antibody or antigen-binding fragment thereof binds to CD40.

[0018] The isolated monoclonal antibody, or the antigen-binding portion thereof, of the present disclosure may comprise a heavy chain and a light chain linked by disulfide bonds, the heavy chain may comprise a heavy chain variable region and a heavy chain constant region, the light chain may

comprise a light chain variable region and a light chain constant region, wherein the C terminus of the heavy chain variable region is linked to the N terminus of the heavy chain constant region, and the C terminus of the light chain variable region is linked to the N terminus of the light chain constant region, wherein the heavy chain variable region and the light chain variable region may comprise amino acid sequences described above, and the antibody or antigen-binding portion thereof binds to CD40. The heavy chain constant region may be human IgG2 constant region having the amino acid sequence set forth in e.g., SEQ ID NO.:28, or human IgG1 constant region having the amino acid sequence set forth in e.g., SEQ ID NO: 27, and the light chain constant region may be human kappa constant region having the amino acid sequences set forth in e.g., SEQ ID NO.: 29. The heavy chain constant region, such as the Fc fragment, may be engineered to have reduced or enhanced FcR binding affinity. The amino acid sequences of SEQ ID NOs: 27, 28 and 29 may be encoded by the nucleotide sequences of SEQ ID NOs: 37, 38 and 39, respectively.

[0019] The antibody of the present disclosure in certain embodiments may comprise or consist of two heavy chains and two light chains, wherein each heavy chain may comprise the heavy chain constant region, heavy chain variable region or CDR sequences mentioned above, and each light chain may comprise the light chain constant region, light chain variable region or CDR sequences mentioned above, wherein the antibody binds to CD40. The antibody of the disclosure can be a full-length antibody, for example, of an IgG1, IgG2 or IgG4 isotype. The antibody or the antigen-binding portion thereof of the present disclosure in other embodiments may be a single chain variable fragment (scFv) antibody, or antibody fragments, such as Fab or F(ab')₂ fragments.

[0020] The disclosure also provides a bispecific molecule that may comprise the antibody, or the antigen-binding portion thereof, of the disclosure, linked to a second functional moiety (e.g., a second antibody) having a different binding specificity than said antibody, or antigen-binding portion thereof. The disclosure also provides an immunoconjugate, such as an antibody-drug conjugate, that may comprise an antibody, or antigen-binding portion thereof, of the disclosure, linked to a therapeutic agent, such as a cytotoxin. In another aspect, the antibody or the antigen binding portion thereof of the present disclosure can be made into part of a chimeric antigen receptor (CAR). Also provided is an immune cell that may comprise the antigen chimeric receptor, such as a T cell and a NK cell. The antibody or the antigen binding portion thereof of the present disclosure can also be encoded by or used in conjunction with an oncolytic virus.

[0021] Nucleic acid molecules encoding the antibody, or the antigen-binding portion thereof, of the disclosure are also encompassed by the disclosure, as well as expression vectors that may comprise such nucleic acids and host cells that may comprise such expression vectors. A method for preparing the anti-CD40 antibody or the antigen-binding portion thereof of the disclosure using the host cell is also provided, that may comprise steps of (i) expressing the antibody in the host cell and (ii) isolating the antibody from the host cell or its cell culture.

[0022] Compositions that may comprise the antibody, or the antigen-binding portion thereof, the immunoconjugate, the bispecific molecule, the oncolytic virus, the CAR, the CAR-T cell, the nucleic acid molecule, the expression vector or the host cells of the disclosure, and a pharmaceutically acceptable carrier, are also provided. In certain embodiments, the pharmaceutical composition may further contain a therapeutic agent such as an anti-cancer agent.

[0023] In yet another aspect, the disclosure provides a method of modulating an immune response in a subject comprising administering to the subject the antibody, or antigen-binding portion thereof, of the disclosure such that the immune response in the subject is modulated. Preferably, the antibody

or antigen-binding portion thereof of the disclosure enhances, stimulates or increases the immune response in the subject. In some embodiments, the method comprises administering a bispecific molecule, an immunoconjugate, a CAR-T cell, or an antibody-encoding or antibody-bearing oncolytic virus of the disclosure, or alternatively a nucleic acid molecule capable of expressing the same in the subject.

[0024] In a further aspect, the disclosure provides a method of inhibiting tumor growth in a subject in need thereof, comprising administering to a subject a therapeutically effective amount of the composition of the present disclosure. The tumor may be a solid or non-solid tumor, including, but not limited to, B cell lymphoma, chronic lymphocytic leukemia, multiple myeloma, melanoma, colon adenocarcinoma, pancreas cancer, colon cancer, gastric intestine cancer, prostate cancer, bladder cancer, kidney cancer, ovary cancer, cervix cancer, breast cancer, lung cancer, and nasopharynx cancer. In some embodiments, at least one additional anti-cancer antibody can be administered with the antibody, or an antigen-binding portion thereof, of the disclosure, such as an anti-VISTA antibody, an anti-PD-1 antibody, an anti-PD-L1 antibody, an anti-LAG-3 antibody, an anti-CTLA-4 antibody, an anti-TIM-3 antibody, an anti-STAT3 antibody, and/or an anti-ROR1 antibody. In yet another embodiment, an antibody, or an antigen-binding portion thereof, of the disclosure is administered with a cytokine (e.g., IL-2, IL-21, GM-CSF and/or IL-4), or a costimulatory antibody (e.g., an anti-CD137 and/or anti-GITR antibody). In another embodiment, an antibody, or an antigen-binding portion thereof, of the disclosure is administered with a chemotherapeutic agent, which may be a cytotoxic agent, such as epirubicin, oxaliplatin, and/or 5-fluorouracil (5-FU). The antibody or antigen-binding portion thereof of the present disclosure may be, for example, mouse, human, chimeric or humanized.

[0025] In another aspect, the disclosure provides a method of treating or alleviating an infectious disease in a subject in need thereof, comprising administering to a subject a therapeutically effective amount of the composition of the present disclosure. The infectious disease may be a disease caused by viral, bacterial, fungal or mycoplasma infection. In certain embodiments, the infectious disease is AIDS, tuberculosis or malaria. In certain embodiments, the subject may be further administered with at least one an anti-infective agent, such as an anti-viral agent, an anti-bacterial agent, an anti-fungal agent, or an anti-mycoplasma agent.

[0026] In another aspect, the disclosure provides a method of treating or alleviating an autoimmune disease in a subject in need thereof, comprising administering to the subject a therapeutically effective amount of the composition of the disclosure. In certain embodiments, the subject may be further administered with at least one anti-inflammatory agent.

[0027] Other features and advantages of the instant disclosure will be apparent from the following detailed description and examples, which should not be construed as limiting. The contents of all references, Genbank entries, patents and published patent applications cited throughout this application are expressly incorporated herein by reference.

[0028] Accordingly, it is an object of the invention not to encompass within the invention any previously known product, process of making the product, or method of using the product such that Applicants reserve the right and hereby disclose a disclaimer of any previously known product, process, or method. It is further noted that the invention does not intend to encompass within the scope of the invention any product, process, or making of the product or method of using the product, which does not meet the written description and enablement requirements of the USPTO (35 U.S.C. §112, first paragraph) or the EPO (Article 83 of the EPC), such that Applicants reserve the right and hereby disclose a disclaimer of any previously described product, process of making the product, or

method of using the product. It may be advantageous in the practice of the invention to be in compliance with Art. 53(c) EPC and Rule 28(b) and (c) EPC. All rights to explicitly disclaim any embodiments that are the subject of any granted patent(s) of applicant in the lineage of this application or in any other lineage or in any prior filed application of any third party is explicitly reserved. Nothing herein is to be construed as a promise.

[0029] It is noted that in this disclosure and particularly in the claims and/or paragraphs, terms such as "comprises", "comprised", "comprising" and the like can have the meaning attributed to it in U.S. Patent law; e.g., they can mean "includes", "included", "including", and the like; and that terms such as "consisting essentially of" and "consists essentially of" have the meaning ascribed to them in U.S. Patent law, e.g., they allow for elements not explicitly recited, but exclude elements that are found in the prior art or that affect a basic or novel characteristic of the invention.

BRIEF DESCRIPTION OF THE DRAWINGS

[0030] FIGs. 1A and 1B show the binding capacities of mouse antibodies 1A3 and 1D1 (A), and C1H1 (B) to human CD40 in a capture ELISA.

[0031] FIGs. 2A and 2B show the binding capacities of mouse antibodies 1A3 and 1D1 (A), and C1H1 (B) to 293T cells expressing human CD40 in a cell based binding FACS.

[0032] FIGs. 3A and 3B show the blocking abilities of mouse antibodies 1A3 and 1D1 (A), and C1H1 (B) on human CD40-CD40L binding in a competitive ELISA.

[0033] FIGs. 4A and 4B show the abilities of mouse antibodies 1A3 and 1D1 (A), and C1H1 (B) to block Benchmark -human CD40 binding in a competitive ELISA.

[0034] FIGs. 5A and 5B show the activities of mouse antibodies 1A3 and 1D1 (A), and C1H1 (B) to activate CD40 signaling in a cell based reporter assay.

[0035] FIGs. 6A-6C show the binding capacities of chimeric antibodies 1A3 (A), 1D1 (B), and C1H1 (C) to human CD40 in a capture ELISA.

[0036] FIGs. 7A-7C show the activities of chimeric antibodies 1A3 (A), 1D1 (B), and C1H1 (C) to activate CD40 signaling in a cell based reporter assay.

[0037] FIG. 8 shows the binding capacities of humanized antibodies huC1H1-V1 and huC1H1-V2 to human CD40 in a capture ELISA.

[0038] FIG. 9 shows the binding capacities of humanized antibodies huC1H1-V1 and huC1H1-V2 to 293T cells expressing human CD40 in a cell based binding FACS.

[0039] FIG. 10 shows the blocking abilities of humanized antibodies huC1H1-V1 and huC1H1-V2 on human CD40-CD40L binding in a competitive ELISA.

[0040] FIG. 11 shows the abilities of humanized antibodies huC1H1-V1 and huC1H1-V2 to block Benchmark-human CD40 binding in a competitive ELISA.

[0041] FIG. 12 shows the activities of humanized antibodies huC1H1-V1 and huC1H1-V2 to activate CD40 signaling in a cell based reporter assay.

DETAILED DESCRIPTION OF THE INVENTION

[0042] To ensure that the present disclosure may be more readily understood, certain terms are first defined. Additional definitions are set forth throughout the detailed description.

[0043] The term "CD40" refers to tumor necrosis factor receptor superfamily member 5 (TNFR5). The term "CD40" comprises variants, isoforms, homologs, orthologs and paralogs. For example, an

antibody specific for a human CD40 protein may, in certain cases, cross-react with a CD40 protein from a species other than human, such as monkey. In other embodiments, an antibody specific for a human CD40 protein may be completely specific for the human CD40 protein and exhibit no cross-reactivity to other species or of other types, or may cross-react with CD40 from certain other species but not all other species.

[0044] The term “human CD40” refers to a CD40 protein having an amino acid sequence from a human, such as the amino acid sequence of human CD40 having a Genbank accession number of NP_001241.1 (Amini M *et al.*, (2020) *Life Sci* 254: 117774). The terms “monkey or rhesus CD40” and “mouse CD40” refer to monkey and mouse CD40 sequences, respectively, e.g. those with the amino acid sequences having Genbank Accession Nos. NP_001252791.1 and NP_035741.2, respectively.

[0045] The term “immune response” refers to the action of, for example, lymphocytes, antigen presenting cells, phagocytic cells, granulocytes, and soluble macromolecules produced by the above cells or the liver (including antibodies, cytokines, and complement) that results in selective damage to, destruction of, or elimination from the human body of invading pathogens, cells or tissues infected with pathogens, cancerous cells, or, in cases of autoimmunity or pathological inflammation, normal human cells or tissues.

[0046] The term “antibody” as used herein refers to an immunoglobulin molecule that recognizes and specifically binds a target, such as CD40, through at least one antigen-binding site wherein the antigen-binding site is usually within the variable region of the immunoglobulin molecule. As used herein, the term encompasses intact polyclonal antibodies, intact monoclonal antibodies, single-chain Fv (scFv) antibodies, heavy chain antibodies (HCAs), light chain antibodies (LCAs), multispecific antibodies, bispecific antibodies, monospecific antibodies, monovalent antibodies, fusion proteins comprising an antigen-binding site of an antibody, and any other modified immunoglobulin molecule comprising an antigen-binding site (e.g., dual variable domain immunoglobulin molecules) as long as the antibodies exhibit the desired biological activity. Antibodies also include, but are not limited to, mouse antibodies, chimeric antibodies, humanized antibodies, and human antibodies. An antibody can be any of the five major classes of immunoglobulins: IgA, IgD, IgE, IgG, and IgM, or subclasses (isotypes) thereof (e.g., IgG1, IgG2, IgG3, IgG4, IgA1 and IgA2), based on the identity of their heavy-chain constant domains referred to as alpha, delta, epsilon, gamma, and mu, respectively. The different classes of immunoglobulins have different and well-known subunit structures and three-dimensional configurations. Antibodies can be naked or conjugated to other molecules, including but not limited to, toxins and radioisotopes. Unless expressly indicated otherwise, the term “antibody” as used herein include “antigen-binding portion” of the intact antibodies. An IgG is a glycoprotein which may comprise two heavy (H) chains and two light (L) chains inter-connected by disulfide bonds. Each heavy chain may be comprised of a heavy chain variable region (abbreviated herein as V_H) and a heavy chain constant region. The heavy chain constant region may be comprised of three domains, C_{H1}, C_{H2} and C_{H3}. Each light chain may be comprised of a light chain variable region (abbreviated herein as V_L) and a light chain constant region. The light chain constant region may be comprised of one domain, C_L. The V_H and V_L regions can be further subdivided into regions of hypervariability, termed complementarity determining regions (CDR), interspersed with regions that are more conserved, termed framework regions (FR). Each V_H and V_L is composed of three CDRs and four FRs, arranged from amino-terminus to carboxy-terminus in the following order: FR1, CDR1, FR2, CDR2, FR3, CDR3, FR4. The variable regions of the heavy and light chains contain a binding

domain that interacts with an antigen. The constant regions of the antibodies can mediate the binding of the immunoglobulin to host tissues or factors, including various cells of the immune system (e.g., effector cells) and the first component (C1q) of the classical complement system.

[0047] The term “antigen-binding portion” of an antibody (or simply “antibody portion”), as used herein, refers to one or more fragments of an antibody that retain the ability to specifically bind to an antigen (e.g., a CD40 protein). It has been shown that the antigen-binding function of an antibody can be performed by fragments of a full-length antibody. Examples of binding fragments encompassed within the term “antigen-binding portion” of an antibody include (i) a Fab fragment, a monovalent fragment consisting of the V_L , V_H , C_L and C_{H1} domains; (ii) a $F(ab')_2$ fragment, a bivalent fragment which may comprise two Fab fragments linked by a disulfide bridge at the hinge region; (iii) a Fd fragment consisting of the V_H and C_{H1} domains; (iv) a Fv fragment consisting of the V_L and V_H domains of a single arm of an antibody, (v) a dAb fragment (Ward *et al.*, (1989) *Nature* 341:544-546), which consists of a V_H domain; (vi) an isolated complementarity determining region (CDR); and (viii) a nanobody, a heavy chain variable region containing a single variable domain and two constant domains. Furthermore, although the two domains of the Fv fragment, V_L and V_H , are coded by separate genes, they can be joined, using recombinant methods, by a synthetic linker that enables them to be made as a single protein chain in which the V_L and V_H regions pair to form monovalent molecules (known as single chain Fv (scFv); see e.g., Bird *et al.*, (1988) *Science* 242:423-426; and Huston *et al.*, (1988) *Proc. Natl. Acad. Sci. USA* 85:5879-5883). Such single chain antibodies are also intended to be encompassed within the term “antigen-binding portion” of an antibody. These antibody fragments are obtained using conventional techniques known to those with skill in the art, and the fragments are screened for utility in the same manner as are intact antibodies.

[0048] An “isolated antibody”, as used herein, is intended to refer to an antibody that is substantially free of other antibodies having different antigenic specificities (e.g., an isolated antibody that specifically binds a CD40 protein is substantially free of antibodies that specifically bind antigens other than CD40 proteins). An isolated antibody that specifically binds a human CD40 protein may, however, have cross-reactivity to other antigens, such as CD40 proteins from other species. Moreover, an isolated antibody can be substantially free of other cellular material and/or chemicals.

[0049] The terms “monoclonal antibody” or “monoclonal antibody composition” as used herein refer to a preparation of antibody molecules of single molecular composition. A monoclonal antibody composition displays a single binding specificity and affinity for a particular epitope.

[0050] The term “mouse antibody”, as used herein, is intended to include antibodies having variable regions in which both the framework and CDR regions are derived from mouse germline immunoglobulin sequences. Furthermore, if the antibody contains a constant region, the constant region also is derived from mouse germline immunoglobulin sequences. The mouse antibodies of the disclosure can include amino acid residues not encoded by mouse germline immunoglobulin sequences (e.g., mutations introduced by random or site-specific mutagenesis *in vitro* or by somatic mutation *in vivo*). However, the term “mouse antibody”, as used herein, is not intended to include antibodies in which CDR sequences derived from the germline of another mammalian species have been grafted onto mouse framework sequences.

[0051] The term “chimeric antibody” refers to an antibody made by combining genetic material from a nonhuman source with genetic material from a human being. Or more generally, a chimeric antibody is an antibody having genetic material from a certain species with genetic material from another species.

[0052] The term “humanized antibody”, as used herein, refers to an antibody from non-human species whose protein sequences have been modified to increase similarity to antibody variants produced naturally in humans.

[0053] The phrases “an antibody recognizing an antigen” and “an antibody specific for an antigen” are used interchangeably herein with the term “an antibody which binds specifically to an antigen.”

[0054] As used herein, an antibody or antigen-binding portion thereof that “specifically binds to human CD40” is intended to refer to an antibody that binds to human CD40 protein (and possibly a CD40 protein from one or more non-human species) but does not substantially bind to non-CD40 proteins. Preferably, the antibody binds to human CD40 protein with “high affinity”, namely with a K_D of 5.0×10^{-8} M or less, and more preferably 1.0×10^{-8} M or less.

[0055] The term “does not substantially bind” to a protein or cells, as used herein, means does not bind or does not bind with a high affinity to the protein or cells, i.e. binds to the protein or cells with a K_D of 1.0×10^{-6} M or more, more preferably 1.0×10^{-5} M or more, more preferably 1.0×10^{-4} M or more, more preferably 1.0×10^{-3} M or more, even more preferably 1.0×10^{-2} M or more.

[0056] The term “high affinity” for an IgG antibody refers to an antibody having a K_D of 1.0×10^{-6} M or less, more preferably 9.0×10^{-9} M or less, more preferably 5.0×10^{-9} M or less, even more preferably 1.0×10^{-9} M or less, and even more preferably 5.0×10^{-10} M or less for a target antigen. However, “high affinity” binding can vary for other antibody isotypes. For example, “high affinity” binding for an IgM isotype refers to an antibody having a K_D of 10^{-6} M or less, more preferably 10^{-7} M or less, even more preferably 10^{-8} M or less.

[0057] The term “ K_{assoc} ” or “ K_a ”, as used herein, is intended to refer to the association rate of a particular antibody-antigen interaction, whereas the term “ K_{dis} ” or “ K_d ”, as used herein, is intended to refer to the dissociation rate of a particular antibody-antigen interaction. The term “ K_D ”, as used herein, is intended to refer to the dissociation constant, which is obtained from the ratio of K_d to K_a (i.e., K_d/K_a) and is expressed as a molar concentration (M). K_D values for antibodies can be determined using methods well established in the art. A preferred method for determining the K_D of an antibody is by using surface plasmon resonance, preferably using a biosensor system such as a Biacore™ system.

[0058] The term “ EC_{50} ”, also known as half maximal effective concentration, refers to the concentration of an antibody which induces a response halfway between the baseline and maximum after a specified exposure time.

[0059] The term “ IC_{50} ”, also known as half maximal inhibitory concentration, refers to the concentration of an antibody which inhibits a specific biological or biochemical function by 50% relative to the absence of the antibody.

[0060] The term “subject” includes any human or nonhuman animal. The term “nonhuman animal” includes all vertebrates, e.g., mammals and non-mammals, such as non-human primates, sheep, dogs, cats, cows, horses, chickens, amphibians, and reptiles, although mammals are preferred, such as non-human primates, sheep, dogs, cats, cows and horses.

[0061] The term “therapeutically effective amount” means an amount of the antibody or antigen-binding portion thereof of the present disclosure sufficient to prevent or ameliorate the symptoms associated with a disease or condition (such as a cancer) and/or lessen the severity of the disease or condition. A therapeutically effective amount is understood to be in context to the condition being treated, where the actual effective amount is readily discerned by those of skill in the art.

[0062] The term “agonistic CD40 antibody” or “agonistic anti-CD40 antibody” refers to an anti-CD40 antibody that binds to CD40 and activates or induces CD40 signaling to e.g., promote immune cell activation and proliferation as well as cytokine and chemokine production. While the term “antagonistic CD40 antibody” refers to an anti-CD40 antibody that blocks or inhibits CD40 signaling that may be induced by CD40L engagement. The agonistic CD40 antibody may promote a tumor-bearing subject’s innate and adaptive immune response to tumors, via elevated antigen presenting ability of APCs, activation of tumor specific CD4+ and CD8+ T cells, secretion of cytokines and chemokines by lymphocytes and monocytes, enhanced tumor cell killings by cytotoxic lymphocytes and NK cells, etc.

[0063] The percent “identity” as used herein in the context of two or more nucleic acids or polypeptides, refers to two or more sequences or subsequences that are the same or have a specified percentage of nucleotides or amino acid residues that are the same, when compared and aligned (introducing gaps, if necessary) for maximum correspondence, considering or not considering conservative amino acid substitutions as part of the sequence identity. The percent identity can be measured using sequence comparison software or algorithms or by visual inspection. Various algorithms and software that can be used to obtain alignments of amino acid or nucleotide sequences are well-known in the art. These include, but are not limited to, BLAST, ALIGN, Megalign, BestFit, GCG Wisconsin Package, and variants thereof. In some embodiments, two nucleic acids or polypeptides of the invention are substantially identical, meaning they have at least 70%, at least 75%, at least 80%, at least 85%, at least 90%, and in some embodiments at least 95%, 96%, 97%, 98%, 99% nucleotide or amino acid residue identity, when compared and aligned for maximum correspondence, as measured using a sequence comparison algorithm or by visual inspection.

[0064] Various aspects of the disclosure are described in further detail in the following subsections.

[0065] The antibody, or the antigen-binding portion thereof, of the disclosure specifically binds to human CD40 with comparable, if not better, binding affinity as compared to previously described anti-CD40 antibodies, such as Dacetuzumab and Selicrelumab.

[0066] Additional functional properties include the capacity to block CD40-CD40L binding, and to activate CD40 signaling.

[0067] Preferred antibodies of the disclosure are humanized monoclonal antibodies. Additionally or alternatively, the antibodies can be, for example, chimeric monoclonal antibodies.

Table 1. Amino acid sequence ID numbers of heavy/light chain variable regions

Antibody	V _H CDR1	V _H CDR2	V _H CDR3	V _H	V _L CDR1	V _L CDR2	V _L CDR3	V _L
C1H1	1	4	7	19	10	13	16	23
huC1H1-V1	1	4	7	20, X1=A	10	13	16	24, X1=K, X2=F
huC1H1-V2	1	4	7	20, X1=S	10	13	16	24, X1=K, X2=F
hu-C1H1-V3	1	4	7	20, X1=A	10	13	16	24, X1=Y, X2=Y
huC1H1-V4	1	4	7	20, X1=S	10	13	16	24, X1=Y, X2=Y
1A3	2	5	8	21	11	14	17	25
1D1	3	6	9	22	12	15	18	26

[0068] The exemplary antibody or antigen-binding portion thereof of the disclosure is structurally and chemically characterized as described below and in the following Examples. The amino acid sequence ID numbers of the heavy/light chain variable regions of the antibodies are summarized in

Table 1 below, some antibodies sharing the same V_H or V_L . The heavy chain constant region for the antibodies may be human IgG1 or IgG2 heavy chain constant region having an amino acid sequence set forth in, e.g., SEQ ID NOS: 27 and 28, respectively, and the light chain constant region for the antibodies may be human kappa constant region having an amino acid sequence set forth in, e.g., SEQ ID NO: 29. These antibodies may also contain mouse IgG1 or IgG2 heavy chain constant region, and/or mouse kappa constant region.

[0069] The heavy chain variable region CDRs and the light chain variable region CDRs in Table 1 have been defined by the Kabat numbering system. However, as is well known in the art, CDR regions can also be determined by other systems such as Chothia, and IMGT, AbM, or Contact numbering system/method, based on heavy chain/light chain variable region sequences.

[0070] The V_H and V_L sequences (or CDR sequences) of other anti-CD40 antibodies which bind to human CD40 can be “mixed and matched” with the V_H and V_L sequences (or CDR sequences) of the anti-CD40 antibody of the present disclosure. Preferably, when V_H and V_L chains (or the CDRs within such chains) are mixed and matched, a V_H sequence from a particular V_H/V_L pairing is replaced with a structurally similar V_H sequence. Likewise, preferably a V_L sequence from a particular V_H/V_L pairing is replaced with a structurally similar V_L sequence.

[0071] Accordingly, in one embodiment, an antibody of the disclosure, or an antigen binding portion thereof, comprises:

- (a) a heavy chain variable region comprising an amino acid sequence listed above in Table 1; and
- (b) a light chain variable region comprising an amino acid sequence listed above in Table 1, or the V_L of another anti-CD40 antibody, wherein the antibody specifically binds human CD40.

[0072] In another embodiment, an antibody of the disclosure, or an antigen binding portion thereof, comprises:

- (a) the CDR1, CDR2, and CDR3 regions of the heavy chain variable region listed above in Table 1; and
- (b) the CDR1, CDR2, and CDR3 regions of the light chain variable region listed above in Table 1 or the CDRs of another anti-CD40 antibody, wherein the antibody specifically binds human CD40.

[0073] In yet another embodiment, the antibody, or antigen binding portion thereof, includes the heavy chain variable CDR2 region of anti-CD40 antibody combined with CDRs of other antibodies which bind human CD40, e.g., CDR1 and/or CDR3 from the heavy chain variable region, and/or CDR1, CDR2, and/or CDR3 from the light chain variable region of a different anti-CD40 antibody.

[0074] In addition, it is well known in the art that the CDR3 domain, independently from the CDR1 and/or CDR2 domain(s), alone can determine the binding specificity of an antibody for a cognate antigen and that multiple antibodies can predictably be generated having the same binding specificity based on a common CDR3 sequence. See, e.g., Klimka *et al.*, *British J. of Cancer* 83(2):252-260 (2000); Beiboer *et al.*, *J. Mol. Biol.* 296:833-849 (2000); Rader *et al.*, *Proc. Natl. Acad. Sci. U.S.A.* 95:8910-8915 (1998); Barbas *et al.*, *J. Am. Chem. Soc.* 116:2161-2162 (1994); Barbas *et al.*, *Proc. Natl. Acad. Sci. U.S.A.* 92:2529-2533 (1995); Ditzel *et al.*, *J. Immunol.* 157:739-749 (1996); Berezov *et al.*, *BIAjournal 8: Scientific Review* 8 (2001); Igarashi *et al.*, *J. Biochem (Tokyo)* 117:452-7 (1995); Bourgeois *et al.*, *J. Virol* 72:807-10 (1998); Levi *et al.*, *Proc. Natl. Acad. Sci. U.S.A.* 90:4374-8 (1993); Polymenis and Stoller, *J. Immunol.* 152:5218-5329 (1994) and Xu and Davis, *Immunity* 13:37-45 (2000). See also, U.S. Pat. Nos. 6,951,646; 6,914,128; 6,090,382; 6,818,216; 6,156,313;

6,827,925; 5,833,943; 5,762,905 and 5,760,185. Each of these references is hereby incorporated by reference in its entirety.

[0075] Accordingly, in another embodiment, antibodies of the disclosure comprise the CDR2 of the heavy chain variable region of the anti-CD40 antibody and at least the CDR3 of the heavy and/or light chain variable region of the anti-CD40 antibody, or the CDR3 of the heavy and/or light chain variable region of another anti-CD40 antibody, wherein the antibody is capable of specifically binding to human CD40. These antibodies preferably (a) compete for binding with CD40; (b) retain the functional characteristics; (c) bind to the same epitope; and/or (d) have a similar binding affinity as the anti-CD40 antibody of the present disclosure. In yet another embodiment, the antibodies further may comprise the CDR2 of the light chain variable region of the anti-CD40 antibody, or the CDR2 of the light chain variable region of another anti-CD40 antibody, wherein the antibody is capable of specifically binding to human CD40. In another embodiment, the antibodies of the disclosure may further include the CDR1 of the heavy and/or light chain variable region of the anti-CD40 antibody, or the CDR1 of the heavy and/or light chain variable region of another anti-CD40 antibody, wherein the antibody is capable of specifically binding to human CD40.

[0076] In another embodiment, an antibody of the disclosure comprises a heavy and/or light chain variable region sequences of CDR1, CDR2 and CDR3 sequences which differ from those of the anti-CD40 antibodies of the present disclosure by one or more conservative modifications. It is understood in the art that certain conservative sequence modification can be made which do not remove antigen binding. See, e.g., Brummell *et al.*, (1993) *Biochem* 32:1180-8; de Wildt *et al.*, (1997) *Prot. Eng.* 10:835-41; Komissarov *et al.*, (1997) *J. Biol. Chem.* 272:26864-26870; Hall *et al.*, (1992) *J. Immunol.* 149:1605-12; Kelley and O'Connell (1993) *Biochem.*32:6862-35; Adib-Conquy *et al.*, (1998) *Int. Immunol.*10:341-6 and Beers *et al.*, (2000) *Clin. Can. Res.* 6:2835-43.

[0077] Accordingly, in one embodiment, the antibody comprises a heavy chain variable region comprising CDR1, CDR2, and CDR3 sequences and/or a light chain variable region comprising CDR1, CDR2, and CDR3 sequences, wherein:

- (a) the heavy chain variable region CDR1 sequence comprises a sequence listed in Table 1 above, and/or conservative modifications thereof; and/or
- (b) the heavy chain variable region CDR2 sequence comprises a sequence listed in Table 1 above, and/or conservative modifications thereof; and/or
- (c) the heavy chain variable region CDR3 sequence comprises a sequence listed in Table 1 above, and/or conservative modifications thereof; and/or
- (d) the light chain variable region CDR1, and/or CDR2, and/or CDR3 sequences comprise the sequence(s) listed in Table 1 above; and/or conservative modifications thereof; and
- (e) the antibody specifically binds human CD40.

[0078] The antibody of the present disclosure possesses one or more of the following functional properties described above, such as high affinity binding to human CD40, and the ability to activate CD40 signaling in CD40-expressing cells.

[0079] In various embodiments, the antibody can be, for example, a mouse, human, humanized or chimeric antibody.

[0080] As used herein, the term “conservative sequence modifications” is intended to refer to amino acid modifications that do not significantly affect or alter the binding characteristics of the antibody containing the amino acid sequence. Such conservative modifications include amino acid

substitutions, additions and deletions. Modifications can be introduced into an antibody of the disclosure by standard techniques known in the art, such as site-directed mutagenesis and PCR-mediated mutagenesis. Conservative amino acid substitutions are ones in which the amino acid residue is replaced with an amino acid residue having a similar side chain. Families of amino acid residues having similar side chains have been defined in the art. These families include amino acids with basic side chains (e.g., lysine, arginine, histidine), acidic side chains (e.g., aspartic acid, glutamic acid), uncharged polar side chains (e.g., glycine, asparagine, glutamine, serine, threonine, tyrosine, cysteine, tryptophan), nonpolar side chains (e.g., alanine, valine, leucine, isoleucine, proline, phenylalanine, methionine), beta-branched side chains (e.g., threonine, valine, isoleucine) and aromatic side chains (e.g., tyrosine, phenylalanine, tryptophan, histidine). Thus, one or more amino acid residues within the CDR regions of an antibody of the disclosure can be replaced with other amino acid residues from the same side chain family and the altered antibody can be tested for retained function (i.e., the functions set forth above) using the functional assays described herein.

[0081] Antibodies of the disclosure can be prepared using an antibody having one or more of the V_H/V_L sequences of the anti-CD40 antibody of the present disclosure as starting material to engineer a modified antibody. An antibody can be engineered by modifying one or more residues within one or both variable regions (i.e., V_H and/or V_L), for example within one or more CDR regions and/or within one or more framework regions. Additionally or alternatively, an antibody can be engineered by modifying residues within the constant region(s), for example to alter the effector function(s) of the antibody.

[0082] In certain embodiments, CDR grafting can be used to engineer variable regions of antibodies. Antibodies interact with target antigens predominantly through amino acid residues that are located in the six heavy and light chain complementarity determining regions (CDRs). For this reason, the amino acid sequences within CDRs are more diverse between individual antibodies than sequences outside of CDRs. Because CDR sequences are responsible for most antibody-antigen interactions, it is possible to express recombinant antibodies that mimic the properties of specific naturally occurring antibodies by constructing expression vectors that include CDR sequences from the specific naturally occurring antibody grafted onto framework sequences from a different antibody with different properties (see, e.g., Riechmann *et al.*, (1998) *Nature* 332:323-327; Jones *et al.*, (1986) *Nature* 321:522-525; Queen *et al.*, (1989) *Proc. Natl. Acad.* See also U.S.A. 86:10029-10033; U.S. Pat. Nos. 5,225,539; 5,530,101; 5,585,089; 5,693,762 and 6,180,370).

[0083] Accordingly, another embodiment of the disclosure pertains to an isolated monoclonal antibody, or antigen binding portion thereof, comprising a heavy chain variable region comprising CDR1, CDR2, and CDR3 sequences comprising the sequences of the present disclosure, as described above, and/or a light chain variable region comprising CDR1, CDR2, and CDR3 sequences comprising the sequences of the present disclosure, as described above. While these antibodies contain the V_H and V_L CDR sequences of the monoclonal antibody of the present disclosure, they can contain different framework sequences.

[0084] Such framework sequences can be obtained from public DNA databases or published references that include germline antibody gene sequences. For example, germline DNA sequences for human heavy and light chain variable region genes can be found in the "VBase" human germline sequence database (available on the Internet at www.mrc-cpe.cam.ac.uk/vbase), as well as in Kabat *et al.*, (1991), cited supra; Tomlinson *et al.*, (1992) *J. Mol. Biol.* 227:776-798; and Cox *et al.*, (1994) *Eur. J. Immunol.* 24:827-836; the contents of each of which are expressly incorporated herein by reference.

As another example, the germline DNA sequences for human heavy and light chain variable region genes can be found in the Genbank database. For example, the following heavy chain germline sequences found in the HCo7 HuMAb mouse are available in the accompanying Genbank Accession Nos.: 1-69 (NG--0010109, NT--024637 & BC070333), 3-33 (NG--0010109 & NT--024637) and 3-7 (NG--0010109 & NT--024637). As another example, the following heavy chain germline sequences found in the HCo12 HuMAb mouse are available in the accompanying Genbank Accession Nos.: 1-69 (NG--0010109, NT--024637 & BC070333), 5-51 (NG--0010109 & NT--024637), 4-34 (NG--0010109 & NT--024637), 3-30.3 (CAJ556644) & 3-23 (AJ406678).

[0085] Antibody protein sequences are compared against a compiled protein sequence database using one of the sequence similarity searching methods called the Gapped BLAST (Altschul *et al.*, (1997), *supra*), which is well known to those skilled in the art.

[0086] Preferred framework sequences for use in the antibodies of the disclosure are those that are structurally similar to the framework sequences used by antibodies of the disclosure. The V_H CDR1, CDR2, and CDR3 sequences can be grafted onto framework regions that have the identical sequence as that found in the germline immunoglobulin gene from which the framework sequence derives, or the CDR sequences can be grafted onto framework regions that contain one or more mutations as compared to the germline sequences. For example, it has been found that in certain instances it is beneficial to mutate residues within the framework regions to maintain or enhance the antigen binding ability of the antibody (see e.g., U.S. Pat. Nos. 5,530,101; 5,585,089; 5,693,762 and 6,180,370).

[0087] Another type of variable region modification is to mutate amino acid residues within the V_H and/or V_L CDR1, CDR2 and/or CDR3 regions to thereby improve one or more binding properties (e.g., affinity) of the antibody of interest. Site-directed mutagenesis or PCR-mediated mutagenesis can be performed to introduce the mutation(s) and the effect on antibody binding, or other functional property of interest, can be evaluated in *in vitro* or *in vivo* assays as known in the art. Preferably conservative modifications (as known in the art) are introduced. The mutations can be amino acid substitutions, additions or deletions, but are preferably substitutions. Moreover, typically no more than one, two, three, four or five residues within a CDR region are altered.

[0088] Accordingly, in another embodiment, the disclosure provides isolated anti-CD40 monoclonal antibodies, or antigen binding portions thereof, comprising a heavy chain variable region comprising: (a) a V_H CDR1 region comprising the sequence of the present disclosure, or an amino acid sequence having one, two, three, four or five amino acid substitutions, deletions or additions; (b) a V_H CDR2 region comprising the sequence of the present disclosure, or an amino acid sequence having one, two, three, four or five amino acid substitutions, deletions or additions; (c) a V_H CDR3 region comprising the sequence of the present disclosure, or an amino acid sequence having one, two, three, four or five amino acid substitutions, deletions or additions; (d) a V_L CDR1 region comprising the sequence of the present disclosure, or an amino acid sequence having one, two, three, four or five amino acid substitutions, deletions or additions; (e) a V_L CDR2 region comprising the sequence of the present disclosure, or an amino acid sequence having one, two, three, four or five amino acid substitutions, deletions or additions; and (f) a V_L CDR3 region comprising the sequence of the present disclosure, or an amino acid sequence having one, two, three, four or five amino acid substitutions, deletions or additions.

[0089] Engineered antibodies of the disclosure include those in which modifications have been made to framework residues within V_H and/or V_L, e.g. to improve the properties of the antibody. Typically, such framework modifications are made to decrease the immunogenicity of the antibody.

For example, one approach is to “backmutate” one or more framework residues to the corresponding germline sequence. More specifically, an antibody that has undergone somatic mutation can contain framework residues that differ from the germline sequence from which the antibody is derived. Such residues can be identified by comparing the antibody framework sequences to the germline sequences from which the antibody is derived.

[0090] Another type of framework modification involves mutating one or more residues within the framework region, or even within one or more CDR regions, to remove T cell epitopes to thereby reduce the potential immunogenicity of the antibody. This approach is also referred to as “deimmunization” and is described in further detail in U.S. Patent Publication No. 20030153043.

[0091] In addition, or as an alternative to modifications made within the framework or CDR regions, antibodies of the disclosure can be engineered to include modifications within the Fc region, typically to alter one or more functional properties of the antibody, such as serum half-life, complement fixation, Fc receptor binding, and/or antigen-dependent cellular cytotoxicity. Furthermore, an antibody of the disclosure can be chemically modified (e.g., one or more chemical moieties can be attached to the antibody) or be modified to alter its glycosylation, again to alter one or more functional properties of the antibody.

[0092] In one embodiment, the hinge region between the CH1 and CH2 regions is modified in such that the number of cysteine residues in the hinge region is altered, e.g., increased or decreased. This approach is described further in U.S. Pat. No. 5,677,425. The number of cysteine residues in the hinge region is altered to, for example, facilitate assembly of the light and heavy chains or to increase or decrease the stability of the antibody.

[0093] In another embodiment, the Fc hinge region of an antibody is mutated to increase or decrease the biological half-life of the antibody. More specifically, one or more amino acid mutations are introduced into the C_{H2}-C_{H3} domain interface region of the Fc-hinge fragment such that the antibody has impaired Staphylococcal protein A (SpA) binding relative to native Fc-hinge domain SpA binding. This approach is described in further detail in U.S. Pat. No. 6,165,745.

[0094] In still another embodiment, the glycosylation of an antibody is modified. For example, a glycosylated antibody can be made (i.e., the antibody lacks glycosylation). Glycosylation can be altered to, for example, increase the affinity of the antibody for antigen. Such carbohydrate modifications can be accomplished by, for example, altering one or more sites of glycosylation within the antibody sequence. For example, one or more amino acid substitutions can be made that result in elimination of one or more variable region framework glycosylation sites to thereby eliminate glycosylation at that site. Such a glycosylation may increase the affinity of the antibody for antigen. See, e.g., U.S. Pat.Nos. 5,714,350 and 6,350,861.

[0095] Additionally or alternatively, an antibody can be made that has an altered type of glycosylation, such as a hypofucosylated antibody having reduced amounts of fucosyl residues or an antibody having increased bisecting GlcNAc structures. Such altered glycosylation patterns have been demonstrated to increase the ADCC ability of antibodies. Such carbohydrate modifications can be accomplished by, for example, expressing the antibody in a host cell with altered glycosylation machinery. Cells with altered glycosylation machinery have been described in the art and can be used as host cells in which to express recombinant antibodies of the disclosure to thereby produce an antibody with altered glycosylation. For example, the cell lines Ms704, Ms705, and Ms709 lack the fucosyl transferase gene, FUT8 (α (1, 6)-fucosyl transferase), such that antibodies expressed in the

Ms704, Ms705, and Ms709 cell lines lack fucose on their carbohydrates. The Ms704, Ms705, and Ms709 FUT8^{-/-} cell lines were created by the targeted disruption of the FUT8 gene in CHO/DG44 cells using two replacement vectors (see U.S. Patent Publication No. 20040110704 and Yamane-Ohnuki *et al.*, (2004) *Biotechnol Bioeng* 87:614-22). As another example, EP 1,176,195 describes a cell line with a functionally disrupted FUT8 gene, which encodes a fucosyl transferase, such that antibodies expressed in such a cell line exhibit hypofucosylation by reducing or eliminating the α -1, 6 bond-related enzyme. EP 1,176,195 also describes cell lines which have a low enzyme activity for adding fucose to the N-acetylglucosamine that binds to the Fc region of the antibody or does not have the enzyme activity, for example the rat myeloma cell line YB2/0 (ATCC CRL 1662). PCT Publication WO 03/035835 describes a variant CHO cell line, Lec13 cells, with reduced ability to attach fucose to Asn(297)-linked carbohydrates, also resulting in hypofucosylation of antibodies expressed in that host cell (see also Shields *et al.*, (2002) *J. Biol. Chem.* 277:26733-26740). Antibodies with a modified glycosylation profile can also be produced in chicken eggs, as described in PCT Publication WO 06/089231. Alternatively, antibodies with a modified glycosylation profile can be produced in plant cells, such as Lemna. Methods for production of antibodies in a plant system are disclosed in the U.S. patent application corresponding to Alston & Bird LLP attorney docket No. 040989/314911, filed on Aug. 11, 2006. PCT Publication WO 99/54342 describes cell lines engineered to express glycoprotein-modifying glycosyl transferases (e.g., β (1,4)-N-acetylglucosaminyltransferase III (GnTIII)) such that antibodies expressed in the engineered cell lines exhibit increased bisecting GlcNAc structures which results in increased ADCC activity of the antibodies (see also Umana *et al.*, (1999) *Nat. Biotech.* 17:176-180). Alternatively, the fucose residues of the antibody can be cleaved off using a fucosidase enzyme; e.g., the fucosidase α -L-fucosidase removes fucosyl residues from antibodies (Tarentino *et al.*, (1975) *Biochem.* 14:5516-23).

[0096] Another modification of the antibodies herein that is contemplated by this disclosure is pegylation. An antibody can be pegylated to, for example, increase the biological (e.g., serum) half-life of the antibody. To pegylate an antibody, the antibody, or fragment thereof, typically is reacted with polyethylene glycol (PEG), such as a reactive ester or aldehyde derivative of PEG, under conditions in which one or more PEG groups become attached to the antibody or antibody fragment. Preferably, the pegylation is carried out via an acylation reaction or an alkylation reaction with a reactive PEG molecule (or an analogous reactive water-soluble polymer). As used herein, the term “polyethylene glycol” is intended to encompass any of the forms of PEG that have been used to derivatize other proteins, such as mono (C₁-C₁₀) alkoxy- or aryloxy-polyethylene glycol or polyethylene glycol-maleimide. In certain embodiments, the antibody to be pegylated is an aglycosylated antibody. Methods for pegylating proteins are known in the art and can be applied to the antibodies of the disclosure. See, e.g., EPO 154 316 and EP 0 401 384.

[0097] Antibodies of the disclosure can be characterized by their various physical properties, to detect and/or differentiate different classes thereof.

[0098] For example, antibodies can contain one or more glycosylation sites in either the light or heavy chain variable region. Such glycosylation sites may result in increased immunogenicity of the antibody or an alteration of the pK of the antibody due to altered antigen binding (Marshall et al (1972) *Annu Rev Biochem* 41:673-702; Gala and Morrison (2004) *J Immunol* 172:5489-94; Wallick et al (1988) *J Exp Med* 168:1099-109; Spiro (2002) *Glycobiology* 12:43R-56R; Parekh et al (1985) *Nature* 316:452-7; Mimura *et al.*, (2000) *Mol Immunol* 37:697-706). Glycosylation has been known to occur at motifs containing an N-X-S/T sequence. In some instances, it is preferred to have an anti-CD40

antibody that does not contain variable region glycosylation. This can be achieved either by selecting antibodies that do not contain the glycosylation motif in the variable region or by mutating residues within the glycosylation region.

[0099] In a preferred embodiment, the antibodies do not contain asparagine isomerism sites. The deamidation of asparagine may occur on N-G or D-G sequences and result in the creation of an isoaspartic acid residue that introduces a link into the polypeptide chain and decreases its stability (isoaspartic acid effect).

[00100] Each antibody will have a unique isoelectric point (pI), which generally falls in the pH range between 6 and 9.5. The pI for an IgG1 antibody typically falls within the pH range of 7-9.5 and the pI for an IgG4 antibody typically falls within the pH range of 6-8. There is speculation that antibodies with a pI outside the normal range may have some unfolding and instability under *in vivo* conditions. Thus, it is preferred to have an anti-CD40 antibody that contains a pI value that falls in the normal range. This can be achieved either by selecting antibodies with a pI in the normal range or by mutating charged surface residues.

[00101] In another aspect, the disclosure provides nucleic acid molecules that encode heavy and/or light chain variable regions, or CDRs, of the antibodies of the disclosure. The nucleic acids can be present in whole cells, in a cell lysate, or in a partially purified or substantially pure form. A nucleic acid is "isolated" or "rendered substantially pure" when purified away from other cellular components or other contaminants, e.g., other cellular nucleic acids or proteins, by standard techniques. A nucleic acid of the disclosure can be, e.g., DNA or RNA and may or may not contain intronic sequences. In a preferred embodiment, the nucleic acid is a cDNA molecule.

[00102] Nucleic acids of the disclosure can be obtained using standard molecular biology techniques. For antibodies expressed by hybridomas (e.g., hybridomas prepared from transgenic mice carrying human immunoglobulin genes as described further below), cDNAs encoding the light and heavy chains of the antibody made by the hybridoma can be obtained by standard PCR amplification or cDNA cloning techniques. For antibodies obtained from an immunoglobulin gene library (e.g., using phage display techniques), a nucleic acid encoding such antibodies can be recovered from the gene library.

[00103] Preferred nucleic acids molecules of the disclosure include those encoding the V_H and V_L sequences of the CD40 monoclonal antibody or the CDRs. Once DNA fragments encoding V_H and V_L segments are obtained, these DNA fragments can be further manipulated by standard recombinant DNA techniques, for example to convert the variable region genes to full-length antibody chain genes, to Fab fragment genes or to a scFv gene. In these manipulations, a V_L - or V_H -encoding DNA fragment is operatively linked to another DNA fragment encoding another protein, such as an antibody constant region or a flexible linker. The term "operatively linked", as used in this context, is intended to mean that the two DNA fragments are joined such that the amino acid sequences encoded by the two DNA fragments remain in-frame.

[00104] The isolated DNA encoding the V_H region can be converted to a full-length heavy chain gene by operatively linking the V_H -encoding DNA to another DNA molecule encoding heavy chain constant regions (C_{H1} , C_{H2} and C_{H3}). The sequences of human heavy chain constant region genes are known in the art and DNA fragments encompassing these regions can be obtained by standard PCR amplification. The heavy chain constant region can be an IgG1, IgG2, IgG3, IgG4, IgA, IgE, IgM or IgD constant region, but most preferably is an IgG1 or IgG2 constant region. For a Fab fragment

heavy chain gene, the V_H-encoding DNA can be operatively linked to another DNA molecule encoding only the heavy chain C_{H1} constant region.

[00105] The isolated DNA encoding the V_L region can be converted to a full-length light chain gene (as well as a Fab light chain gene) by operatively linking the V_L-encoding DNA to another DNA molecule encoding the light chain constant region, C_L. The sequences of human light chain constant region genes are known in the art and DNA fragments encompassing these regions can be obtained by standard PCR amplification. In preferred embodiments, the light chain constant region can be a kappa or lambda constant region.

[00106] To create a scFv gene, the V_H- and V_L-encoding DNA fragments are operatively linked to another fragment encoding a flexible linker, e.g., encoding the amino acid sequence (Gly4-Ser)₃, such that the V_H and V_L sequences can be expressed as a contiguous single-chain protein, with the V_L and V_H regions joined by the flexible linker (see e.g., Bird *et al.*, (1988) *Science* 242:423-426; Huston *et al.*, (1988) *Proc. Natl. Acad. Sci. USA* 85:5879-5883; McCafferty *et al.*, (1990) *Nature* 348:552-554).

[00107] Monoclonal antibodies (mAbs) of the present disclosure can be produced using the well-known somatic cell hybridization (hybridoma) technique of Kohler and Milstein (1975) *Nature* 256:495. Other embodiments for producing monoclonal antibodies include viral or oncogenic transformation of B lymphocytes and phage display techniques. Chimeric or humanized antibodies are also well known in the art. See e.g., U.S. Pat. Nos. 4,816,567; 5,225,539; 5,530,101; 5,585,089; 5,693,762 and 6,180,370, the contents of which are specifically incorporated herein by reference in their entirety.

[00108] Antibodies of the disclosure also can be produced in a host cell transfectoma using, for example, a combination of recombinant DNA techniques and gene transfection methods as is well known in the art (e.g., Morrison, S. (1985) *Science* 229:1202). In one embodiment, DNA encoding partial or full-length light and heavy chains obtained by standard molecular biology techniques is inserted into one or more expression vectors such that the genes are operatively linked to transcriptional and translational regulatory sequences. In this context, the term “operatively linked” is intended to mean that an antibody gene is ligated into a vector such that transcriptional and translational control sequences within the vector serve their intended function of regulating the transcription and translation of the antibody gene.

[00109] The term “regulatory sequence” is intended to include promoters, enhancers and other expression control elements (e.g., polyadenylation signals) that control the transcription or translation of the antibody genes. Such regulatory sequences are described, e.g., in Goeddel (Gene Expression Technology. Methods in Enzymology 185, Academic Press, San Diego, Calif. (1990)). Preferred regulatory sequences for mammalian host cell expression include viral elements that direct high levels of protein expression in mammalian cells, such as promoters and/or enhancers derived from cytomegalovirus (CMV), Simian Virus 40 (SV40), adenovirus, e.g., the adenovirus major late promoter (AdMLP) and polyoma. Alternatively, non-viral regulatory sequences can be used, such as the ubiquitin promoter or β -globin promoter. Still further, regulatory elements composed of sequences from different sources, such as the SR α promoter system, which contains sequences from the SV40 early promoter and the long terminal repeat of human T cell leukemia virus type 1 (Takebe *et al.*, (1988) *Mol. Cell. Biol.* 8:466-472). The expression vector and expression control sequences are chosen to be compatible with the expression host cell used.

[00110] The antibody light chain gene and the antibody heavy chain gene can be inserted into the same or separate expression vectors. In preferred embodiments, the variable regions are used to create full-length antibody genes of any antibody isotype by inserting them into expression vectors already encoding heavy chain constant and light chain constant regions of the desired isotype such that the V_H segment is operatively linked to the C_H segment(s) within the vector and the V_L segment is operatively linked to the C_L segment within the vector. Additionally or alternatively, the recombinant expression vector can encode a signal peptide that facilitates secretion of the antibody chain from a host cell. The antibody chain gene can be cloned into the vector such that the signal peptide is linked in-frame to the amino terminus of the antibody chain gene. The signal peptide can be an immunoglobulin signal peptide or a heterologous signal peptide (i.e., a signal peptide from a non-immunoglobulin protein).

[00111] In addition to the antibody chain genes and regulatory sequences, the recombinant expression vectors of the disclosure can carry additional sequences, such as sequences that regulate replication of the vector in host cells (e.g., origins of replication) and selectable marker genes. The selectable marker gene facilitates selection of host cells into which the vector has been introduced (see, e.g., U.S. Pat. Nos. 4,399,216; 4,634,665 and 5,179,017). For example, typically the selectable marker gene confers resistance to drugs, such as G418, hygromycin or methotrexate, on a host cell into which the vector has been introduced. Preferred selectable marker genes include the dihydrofolate reductase (DHFR) gene (for use in dhfr-host cells with methotrexate selection/amplification) and the neo gene (for G418 selection).

[00112] For expression of the light and heavy chains, the expression vector(s) encoding the heavy and light chains is transfected into a host cell by standard techniques. The various forms of the term “transfection” are intended to encompass a wide variety of techniques commonly used for the introduction of exogenous DNA into a prokaryotic or eukaryotic host cell, e.g., electroporation, calcium-phosphate precipitation, DEAE-dextran transfection and the like. Although it is theoretically possible to express the antibodies of the disclosure in either prokaryotic or eukaryotic host cells, expression of antibodies in eukaryotic cells, and most preferably mammalian host cells, is the most preferred because such eukaryotic cells, and in particular mammalian cells, are more likely than prokaryotic cells to assemble and secrete a properly folded and immunologically active antibody.

[00113] Preferred mammalian host cells for expressing the recombinant antibodies of the disclosure include Chinese Hamster Ovary (CHO cells) (including dhfr- CHO cells, described in Urlaub and Chasin, (1980) *Proc. Natl. Acad. Sci. USA* 77:4216-4220, used with a DHFR selectable marker, e.g., as described in R. J. Kaufman and P. A. Sharp (1982) *J. Mol. Biol.* 159:601-621), NSO myeloma cells, COS cells and SP2 cells. In particular for use with NSO myeloma cells, another preferred expression system is the GS gene expression system disclosed in WO 87/04462, WO 89/01036 and EP 338,841. When recombinant expression vectors encoding antibody genes are introduced into mammalian host cells, the antibodies are produced by culturing the host cells for a period of time sufficient to allow for expression of the antibody in the host cells or, more preferably, secretion of the antibody into the culture medium in which the host cells are grown. Antibodies can be recovered from the culture medium using standard protein purification methods.

[00114] Antibodies of the disclosure can be conjugated to a therapeutic agent to form an immunoconjugate such as an antibody-drug conjugate (ADC). Suitable therapeutic agents include cytotoxins, alkylating agents, DNA minor groove binders, DNA intercalators, DNA crosslinkers, histone deacetylase inhibitors, nuclear export inhibitors, proteasome inhibitors, topoisomerase I or II

inhibitors, heat shock protein inhibitors, tyrosine kinase inhibitors, antibiotics, and anti-mitotic agents. In the ADC, the antibody and therapeutic agent preferably are conjugated via a linker cleavable such as a peptidyl, disulfide, or hydrazone linker. More preferably, the linker is a peptidyl linker such as Val-Cit, Ala-Val, Val-Ala-Val, Lys-Lys, Pro-Val-Gly-Val-Val, Ala-Asn-Val, Val-Leu-Lys, Ala-Ala-Asn, Cit-Cit, Val-Lys, Lys, Cit, Ser, or Glu. The ADCs can be prepared as described in U.S. Pat. Nos. 7,087,600; 6,989,452; and 7,129,261; PCT Publications WO 02/096910; WO 07/038,658; WO 07/051,081; WO 07/059,404; WO 08/083,312; and WO 08/103,693; U.S. Patent Publications 20060024317; 20060004081; and 20060247295; the disclosures of which are incorporated herein by reference.

[00115] In another aspect, the present disclosure features bispecific molecules comprising one or more antibodies of the disclosure linked to at least one other functional molecule, e.g., another peptide or protein (e.g., another antibody or ligand for a receptor) to generate a bispecific molecule that binds to at least two different binding sites or target molecules. Thus, as used herein, “bispecific molecule” includes molecules that have three or more specificities.

[00116] In an embodiment, a bispecific molecule has, in addition to an anti-Fc binding specificity and an anti-CD40 binding specificity, a third specificity. The third specificity can be for an anti-enhancement factor (EF), e.g., a molecule that binds to a surface protein involved in cytotoxic activity and thereby increases the immune response against the target cell. For example, the anti-enhancement factor can bind a cytotoxic T-cell (e.g. via CD2, CD3, CD8, CD28, CD4, CD40, or ICAM-1) or other immune cell, resulting in an increased immune response against the target cell.

[00117] Bispecific molecules may be in many different formats and sizes. At one end of the size spectrum, a bispecific molecule retains the traditional antibody format, except that, instead of having two binding arms of identical specificity, it has two binding arms each having a different specificity. At the other extreme are bispecific molecules consisting of two single-chain antibody fragments (scFv's) linked by a peptide chain, a so-called Bs(scFv) 2 construct. Intermediate-sized bispecific molecules include two different F(ab) fragments linked by a peptidyl linker. Bispecific molecules of these and other formats can be prepared by genetic engineering, somatic hybridization, or chemical methods. See, e.g., Kufer et al, cited supra; Cao and Suresh, *Bioconjugate Chemistry*, 9 (6), 635-644 (1998); and van Spriël *et al.*, *Immunology Today*, 21 (8), 391-397 (2000), and the references cited therein.

[00118] Also provided herein is an oncolytic virus that preferentially infects and kills cancer cells. Antibodies of the present disclosure can be used in conjunction with oncolytic viruses. Alternatively, oncolytic viruses encoding antibodies of the present disclosure can be introduced into human body.

[00119] Also provided herein are a chimeric antigen receptor (CAR) containing an anti-CD40 scFv, the anti- CD40 scFv comprising CDRs and heavy/light chain variable regions described herein.

[00120] The anti-CD40 CAR may comprise (a) an extracellular antigen binding domain comprising an anti-CD40 scFv; (b) a transmembrane domain; and (c) an intracellular signaling domain.

[00121] The CAR may contain a signal peptide at the N-terminus of the extracellular antigen binding domain that directs the nascent receptor into the endoplasmic reticulum, and a hinge peptide at the N-terminus of the extracellular antigen binding domain that makes the receptor more available for binding. The CAR preferably comprises, at the intracellular signaling domain, a primary intracellular signaling domain and one or more co-stimulatory signaling domains. The mainly used and most effective primary intracellular signaling domain is CD3-zeta cytoplasmic domain which contains

ITAMs, the phosphorylation of which results in T cell activation. The costimulatory signaling domain may be derived from the co-stimulatory proteins such as CD28, CD137 and OX40.

[00122] The CARs may further add factors that enhance T cell expansion, persistence, and anti-tumor activity, such as cytokines, and co-stimulatory ligands.

[00123] Also provided are engineered immune effector cells, comprising the CAR provided herein. In some embodiments, the immune effector cell is a T cell, an NK cell, a peripheral blood mononuclear cell (PBMC), a hematopoietic stem cell, a pluripotent stem cell, or an embryonic stem cell. In some embodiments, the immune effector cell is a T cell.

[00124] In another aspect, the present disclosure provides a pharmaceutical composition which may comprise one or more antibodies or antigen-binding portions thereof, the bispecifics, CAR-T cells, oncolytic viruses, immunoconjugates, nucleic acid molecules, expression vectors, or host cells of the present disclosure formulated together with a pharmaceutically acceptable carrier. The antibodies or antigen-binding portion thereof, the bispecifics, CAR-T cells, oncolytic viruses, immunoconjugates, nucleic acid molecules, expression vectors, or host cells can be dosed separately when the composition contains more than one antibody (or antigen-binding portion thereof, the bispecifics, CAR-T cells, oncolytic viruses, immunoconjugates, nucleic acid molecules, expression vectors, or host cells). The composition may optionally contain one or more additional pharmaceutically active ingredients, such as another antibody or a drug, such as an anti-tumor drug.

[00125] The pharmaceutical composition can comprise any number of excipients. Excipients that can be used include carriers, surface active agents, thickening or emulsifying agents, solid binders, dispersion or suspension aids, solubilizers, colorants, flavoring agents, coatings, disintegrating agents, lubricants, sweeteners, preservatives, isotonic agents, and combinations thereof. The selection and use of suitable excipients is taught in Gennaro, ed., Remington: *The Science and Practice of Pharmacy*, 20th Ed. (Lippincott Williams & Wilkins 2003), the disclosure of which is incorporated herein by reference.

[00126] Preferably, the pharmaceutical composition is suitable for intravenous, intramuscular, subcutaneous, parenteral, spinal or epidermal administration (e.g., by injection or infusion). Depending on the route of administration, the active ingredient can be coated in a material to protect it from the action of acids and other natural conditions that may inactivate it. The phrase "parenteral administration" as used herein means modes of administration other than enteral and topical administration, usually by injection, and includes, without limitation, intravenous, intramuscular, intraarterial, intrathecal, intracapsular, intraorbital, intracardiac, intradermal, intraperitoneal, transtracheal, subcutaneous, subcuticular, intraarticular, subcapsular, subarachnoid, intraspinal, epidural and intrasternal injection and infusion. Alternatively, the pharmaceutical composition of the disclosure can be administered via a non-parenteral route, such as a topical, epidermal or mucosal route of administration, e.g., intranasally, orally, vaginally, rectally, sublingually or topically.

[00127] Pharmaceutical compositions can be in the form of sterile aqueous solutions or dispersions. They can also be formulated in a microemulsion, liposome, or other ordered structure suitable to high drug concentration.

[00128] The amount of active ingredient which can be combined with a carrier material to produce a single dosage form will vary depending upon the subject being treated and the particular mode of administration and will generally be that amount of the composition which produces a therapeutic

effect. Generally, out of one hundred percent, this amount will range from about 0.01% to about 99% of active ingredient in combination with a pharmaceutically acceptable carrier.

[00129] Dosage regimens are adjusted to provide the optimum desired response (e.g., a therapeutic response). For example, a single bolus can be administered, several divided doses can be administered over time or the dose can be proportionally reduced or increased as indicated by the exigencies of the therapeutic situation. It is especially advantageous to formulate parenteral compositions in dosage unit form for ease of administration and uniformity of dosage. Dosage unit form as used herein refers to physically discrete units suited as unitary dosages for the subjects to be treated; each unit contains a predetermined quantity of active ingredient calculated to produce the desired therapeutic effect in association with the required pharmaceutical carrier. Alternatively, antibody can be administered as a sustained release formulation, in which case less frequent administration is required.

[00130] For administration of the antibody, the dosage may range from about 0.0001 to 100 mg/kg.

[00131] A “therapeutically effective dosage” of an anti-CD40 antibody or antigen-binding portion thereof of the disclosure preferably results in a decrease in severity of disease symptoms, an increase in frequency and duration of disease symptom-free periods, or a prevention of impairment or disability due to the disease affliction. For example, for the treatment of tumor-bearing subjects, a “therapeutically effective dosage” preferably inhibits tumor growth by at least about 20%, more preferably by at least about 40%, even more preferably by at least about 60%, and still more preferably by at least about 80% relative to untreated subjects. A therapeutically effective amount of a therapeutic antibody can decrease tumor size, or otherwise ameliorate symptoms in a subject, which is typically a human or can be another mammal.

[00132] The pharmaceutical composition can be a controlled release formulation, including implants, transdermal patches, and microencapsulated delivery systems. Biodegradable, biocompatible polymers can be used, such as ethylene vinyl acetate, polyanhydrides, polyglycolic acid, collagen, polyorthoesters, and polylactic acid. See, e.g., Sustained and Controlled Release Drug Delivery Systems, J. R. Robinson, ed., Marcel Dekker, Inc., New York, 1978.

[00133] Therapeutic compositions can be administered via medical devices such as (1) needleless hypodermic injection devices (e.g., U.S. Pat. Nos. 5,399,163; 5,383,851; 5,312,335; 5,064,413; 4,941,880; 4,790,824; and 4,596,556); (2) micro-infusion pumps (U.S. Pat. No. 4,487,603); (3) transdermal devices (U.S. Pat.No. 4,486,194); (4) infusion apparatuses (U.S. Pat.Nos. 4,447,233 and 4,447,224); and (5) osmotic devices (U.S. Pat. Nos. 4,439,196 and 4,475,196); the disclosures of which are incorporated herein by reference.

[00134] In certain embodiments, the monoclonal antibodies of the disclosure can be formulated to ensure proper distribution *in vivo*. For example, to ensure that the therapeutic antibody of the disclosure cross the blood-brain barrier, they can be formulated in liposomes, which may additionally comprise targeting moieties to enhance selective transport to specific cells or organs. See, e.g. U.S. Pat. Nos. 4,522,811; 5,374,548; 5,416,016; and 5,399,331; V. V. Ranade (1989) *J. Clin.Pharmacol.*29:685; Umezawa *et al.*, (1988) *Biochem. Biophys. Res. Commun.* 153:1038; Bloeman *et al.*, (1995) *FEBS Lett.*357:140; M. Owais *et al.*, (1995) *Antimicrob. Agents Chemother.* 39:180; Briscoe *et al.*, (1995) *Am. J. Physiol.* 1233:134; Schreier *et al.*, (1994) *J. Biol. Chem.* 269:9090; Keinanen and Laukkanen (1994) *FEBS Lett.* 346:123; and Killion and Fidler (1994) *Immunomethods* 4:273.

[00135] The pharmaceutical compositions of the present disclosure have numerous *in vitro* and *in vivo* utilities, including, for example, treatment and/or prevention of cancers, or more generally immune response enhancement in patients with cancers. The pharmaceutical compositions can be administered to human subjects, e.g., *in vivo*, to inhibit tumor growth, or to treat or alleviate an infectious disease or an autoimmune disease.

[00136] Given the ability of the pharmaceutical compositions of the disclosure to inhibit proliferation and survival of cancer cells, the disclosure provides methods for inhibiting growth of tumor cells in a subject in need thereof comprising administering to the subject a pharmaceutical composition of the disclosure such that growth of the tumor is inhibited in the subject. Non-limiting examples of tumors that can be treated by the pharmaceutical compositions of the disclosure include, but not limited to, B cell lymphoma, chronic lymphocytic leukemia, multiple myeloma, melanoma, colon adenocarcinoma, pancreas cancer, colon cancer, gastric intestine cancer, prostate cancer, bladder cancer, kidney cancer, ovary cancer, cervix cancer, breast cancer, lung cancer, and nasopharynx cancer. Additionally, the pharmaceutical compositions of the disclosure may also apply to refractory or recurrent malignancies whose growth may be inhibited by the compositions of the disclosure.

[00137] These and other methods of the disclosure are discussed in further detail below.

[00138] In another aspect, the disclosure provides methods of combination therapy in which a pharmaceutical composition of the present disclosure is co-administered with one or more additional antibodies or non-antibody agents that are effective in inhibiting tumor growth in a subject. In one embodiment, the disclosure provides a method for inhibiting tumor growth in a subject comprising administering to the subject a pharmaceutical composition of the disclosure and one or more additional antibodies, such as an anti-TIM3 antibody, an anti- an anti-PD-L1 antibody, and anti-PD-1 antibody and/or an anti-CTLA-4 antibody. In certain embodiments, the subject is human. In certain embodiments, the pharmaceutical composition of the disclosure may be further combined with standard cancer treatments. For example, CD40 signaling activation by the pharmaceutical composition of the disclosure can be combined with CTLA-4 and/or PD-1 blockade and also chemotherapeutic regimens. For example, a chemotherapeutic agent can be administered with the pharmaceutical composition of the disclosure, which may be a cytotoxic agent. For example, epirubicin, oxaliplatin, and 5-FU are administered to patients receiving anti-CD40 therapy. Other therapies that may be combined with anti-CD40 therapy includes, but not limited to, interleukin-2 (IL-2) administration, radiation, surgery, or hormone deprivation.

[00139] The combination of therapeutic agents discussed herein can be administered concurrently as a single composition in a pharmaceutically acceptable carrier, or concurrently as separate compositions with each agent in a pharmaceutically acceptable carrier. In another embodiment, the combination of therapeutic agents can be administered sequentially.

[00140] Furthermore, if more than one dose of the combination therapy is administered sequentially, the order of the sequential administration can be reversed or kept in the same order at each time point of administration, sequential administrations can be combined with concurrent administrations, or any combination thereof.

[00141] The present disclosure is further illustrated by the following examples, which should not be construed as further limiting. The contents of all figures and all references, Genbank sequences, patents and published patent applications cited throughout this application are expressly incorporated herein by reference.

Examples

Example 1 Generation of Mouse Anti-CD40 Monoclonal Antibodies Using Hybridoma Technology *Immunization*

[00142] Mice were immunized according to the method as described in E Harlow, D. Lane, *Antibody: A Laboratory Manual*, Cold Spring Harbor Laboratory Press, Cold Spring Harbor, N.Y., 1998. Recombinant human CD40 protein (AA region 21-193 of Uniprot Number #P25942, amino acid residue 21-193 of SEQ ID NO.: 30) with human IgG1 Fc tag (SEQ ID NO.: 27) at the C-terminus was used as the immunogen. Human CD40-his protein (Acro biosystems, Cat#CD0-H5228) was used for determining anti-sera titer and for screening hybridomas secreting antigen-specific antibodies. Immunizing dosages contained 25 µg human CD40-Fc protein/mouse/injection for both primary and boost immunizations. To increase immune response, the complete Freud's adjuvant and incomplete Freud's adjuvant (Sigma, St. Louis, Mo., USA) were used respectively for primary and boost immunizations. Briefly, adjuvant-antigen mixture was prepared by first gently mixing the adjuvant in a vial using a vortex. The desired amount of adjuvant was transferred to an autoclaved 1.5 mL micro-centrifuge tube. The antigen was prepared in PBS or saline with the concentration ranging from 0.5-1.0 mg/ml. The calculated amount of antigen was then added to the micro-centrifuge tube with the adjuvant, and the resulting mixture was mixed by gently vortexing for 2 minutes to generate water-in-oil emulsions. The adjuvant-antigen emulsion was then drawn into the proper syringe for animal injection. A total of 25µg of antigen was injected in a volume of 50-100 µl. Each animal was immunized, and then boosted for 2 to 3 times depending on the anti-sera titer. Animals with good titers were given a final boost by intraperitoneal injection before fusion.

Hybridoma fusion and screening

[00143] Cells of murine myeloma cell line (SP2/0-Ag14, ATCC#CRL-1581) were cultured to reach the log phase stage right before fusion. Spleen cells from immunized mice were prepared sterilely and fused with myeloma cells according to the method as described in Kohler G, and Milstein C, "Continuous cultures of fused cells secreting antibody of predefined specificity," *Nature*, 256: 495-497(1975). Fused "hybrid cells" were subsequently dispensed into 96-well plates in DMEM/20% FCS/HAT media. Surviving hybridoma colonies were observed under the microscope seven to ten days post fusion. After two weeks, the supernatant from each well was subjected to ELISA-based screening using recombinant human CD40-his protein, and cell-based binding FACS using 293T-CD40 cells expressing human CD40 protein (uniprot #P25942-1, SEQ ID NO.: 30) on cell membrane. Hybridomas secreting antibodies that bound to human CD40-his protein and showed high specific binding to the 293T-CD40 cells, namely Clone 1A3, 1D1 and C1H1, were subcloned by limiting dilution to ensure the clonality of the cell line, and then monoclonal antibodies were purified. Briefly, Protein A sepharose column (from bestchrom (Shanghai) Biosciences, Cat#AA0273) was washed using PBS buffer in 5 to 10 column volumes. Cell supernatants were passed through the columns, and then the columns were washed using PBS buffer until the absorbance for protein reached the baseline. The columns were eluted with elution buffer (0.1 M Glycine-HCl, pH 2.7), and immediately collected into 1.5 ml tubes with neutralizing buffer (1 M Tris-HCl, pH 9.0). Fractions containing immunoglobulins were pooled and dialyzed in PBS overnight at 4°C. Subsequently, the *in vitro* functional activities of purified monoclonal antibodies were characterized as follows.

Example 2 Affinity Determination of Mouse Anti-CD40 Monoclonal Antibodies Using BIACORE Surface Plasmon Resonance Technology

[00144] The purified anti-CD40 mouse monoclonal antibodies (mAbs) generated in Example 1 were characterized for affinity and binding kinetics by Biacore T200 system (GE healthcare, Pittsburgh, PA, USA).

[00145] Briefly, goat anti-mouse IgG (GE healthcare, Cat#BR100838, Mouse Antibody Capture Kit) was covalently linked to a CM5 chip (carboxy methyl dextran coated chip from GE healthcare #BR100530) via primary amines, using a standard amine coupling kit provided by Biacore (GE healthcare, Pittsburgh, PA, USA), and a Protein G chip (GE healthcare, Cat#29-1793-15) was used for the benchmark's affinity determination. Un-reacted moieties on the biosensor surface were blocked with ethanolamine. Then, purified anti-CD40 antibodies of the disclosure and two benchmark antibodies, BM1 (Dacetuzumab, Genentech Inc., also referred to as CD40-BM1, in house made with heavy chain and light chain amino acid sequences set forth in SEQ ID NOs: 40 and 41, respectively) and BM2 (Selicrelumab, Abgenix Inc., also referred to as CD40-BM2, in house made with heavy chain and light chain amino acid sequences set forth in SEQ ID NOs: 42 and 43, respectively), at the concentration of 66.7 nM were respectively flowed onto the chip at a flow rate of 10 μ L/min. Then, serially diluted recombinant human CD40-his (Acro biosystems, Cat#CD0-H5228, starting at 80 nM with a 2-fold serial dilution) or cynomolgus monkey CD40-his protein (Acro biosystems, Cat#CD0-C52H6, starting at 80 nM with a 2-fold serial dilution) in HBS-EP⁺ buffer (provided by Biacore) was flowed onto the chip at a flow rate of 30 μ L/min. The antigen-antibody association kinetics was followed for 2 minutes and the dissociation kinetics was followed for 10 minutes. The association and dissociation curves were fit to a 1:1 Langmuir binding model using BIAcore evaluation software. The K_D , K_a and K_d values were determined and summarized in Table 2 below.

Table 2. Binding affinity of mouse anti-CD40 antibodies

Mouse mAb	Kinetics on Biacore					
	Human CD40-his			Cynomolgus CD40-his		
	K_a ($M^{-1}s^{-1}$)	K_d (s^{-1})	K_D (M)	K_a ($M^{-1}s^{-1}$)	K_d (s^{-1})	K_D (M)
C1H1	2.10E+06	8.97E-04	4.28E-10	2.29E+06	8.63E-04	3.77E-10
BM1	8.03E+05	7.562E-03	9.42E-09	7.64E+05	7.397E-03	9.68E-09
BM2	2.06E+05	1.616E-03	7.84E-09	1.73E+05	1.508E-03	8.73E-09

[00146] The mouse antibody C1H1 of the disclosure specifically bound to human and monkey CD40s with a higher binding affinity than BM1 and BM2.

Example 3 Binding Activity of Mouse Anti-CD40 Monoclonal Antibodies

[00147] The binding activities of mouse anti-CD40 antibodies were determined by Capture ELISA and Flow Cytometry (FACS).

[00148] For the capture ELISA, 96-well micro plates were coated with 2 μ g/ml AffiniPure_Goat Anti-Mouse IgG, F(ab')₂ fragment specific (Jackson Immuno Research, Cat#115-005-072) in PBS, 100 μ l/well, and incubated overnight at 4°C. Plates were washed once with wash buffer (PBS+0.05% v/v Tween-20, PBST) and then blocked with 200 μ l/well blocking buffer (5% w/v non-fatty milk in PBST) for 2 hours at 37°C. Plates were washed 4 times and incubated with 100 μ l/well serially diluted anti-CD40 antibodies of the disclosure, BM1, BM2 and negative control hIgG (human immunoglobulin (pH4) for intravenous injection, Hualan Biological Engineering Inc.) (5-fold serial dilution in 2.5% w/v non-fatty milk in PBST, starting at 66.7 nM) for 40 minutes at 37°C, and then washed 4 times again. Plates containing captured anti-CD40 antibodies were incubated with 100 μ l

biotin-labeled human CD40-Fc protein (amino acid residue 21-193 of SEQ ID NO: 30 linked to N-terminal amino acid residue 99-330 of SEQ ID NO.:28, 1.15 nM in 2.5% w/v non-fatty milk in PBST) for 40 minutes at 37°C, washed 4 times, and incubated with streptavidin conjugated HRP (1:10000 dilution in PBST, Jackson Immuno Research, Cat#016-030-084, 100 µl/well) for 40 minutes at 37°C. After a final wash, plates were incubated with 100 µl/well ELISA substrate TMB (Innoreagents, Cat#TMB-S-002) at room temperature. The reaction was stopped in 3-10 minutes with 50 µl/well 1M H₂SO₄, and the absorbance of each well was read on a microplate reader using dual wavelength mode with 450 nm for TMB and 630 nm as the reference wavelength, then the OD (450-630) values were plotted against antibody concentration. Data was analyzed using Graphpad Prism software and EC₅₀ values were reported.

[00149] For binding of anti-CD40 antibodies to 293T-CD40 cells tested by flow cytometry (FACS), Biosion in-house prepared 293T-CD40 cells stably expressing full length human CD40 (uniprot#P25942-1, SEQ ID NO.: 30) on cell membrane were used. The 293T-CD40 cells were prepared by transfecting 293T cells with pCMV-T-P plasmids inserted with CD40 coding sequence between *EcoRI* and *XbaI* sites, following the instruction of lipofectamine 3000 transfection reagent (Thermo Fisher). In specific, the 293T-CD40 cells were harvested from cell culture flasks, washed twice and resuspended in phosphate buffered saline (PBS) containing 2% v/v Fetal Bovine Serum (FACS buffer). 2 x 10⁵ cells were incubated in each well of 96 well-plates in 100 µl anti-CD40 antibodies of the disclosure or controls of various concentrations (starting at 80 nM with a 4-fold serial dilution) in FACS buffer for 40 minutes on ice. Cells were washed twice with FACS buffer, and 100 µL/well R-Phycoerythrin Affini Pure F(ab')₂ Fragment Goat Anti-Mouse IgG (H+L) (1:1000 dilution in FACS buffer, Jackson Immuno Research, Cat#115-116-146) was added. Following an incubation of 40 minutes at 4°C in dark, cells were washed three times and resuspended in FACS buffer. Fluorescence was measured using a Becton Dickinson FACS Canto II-HTS equipment and plotted against antibody concentration. Data was analyzed using Graphpad Prism software and EC₅₀ values were reported.

[00150] The results were shown in FIGs. 1A-1B and FIGs. 2A-2B.

[00151] The results indicated that the mouse antibodies of the disclosure bound to human CD40 specifically, with 1A3, 1D1, and C1H1 having lower EC₅₀ values than either BM1 or BM2, or both, suggesting that they more efficiently bound to human CD40 protein. Further, as can be seen from FIGs. 1A-1B and FIGs. 2A-2B, the maximum binding of the mouse antibodies 1A3/C1H1 was comparable to that of BM1 or BM2.

Example 4 Blocking Activity of Mouse Anti-CD40 Antibodies on CD40-CD40L or CD40-Benchmark Binding

4.1 Ligand Blocking ELISA

[00152] The abilities of anti-CD40 antibodies to block CD40-CD40L binding were measured in a competitive ELISA assay. Briefly, 100 µl human CD40-Fc protein (amino acid residue 21-193 of SEQ ID NO: 30 linked to N-terminus of amino acid residue 99-330 of SEQ ID NO.:28) was coated on 96-well micro plates at 2 µg/mL in coating buffer (carbonate/bicarbonate buffer) and incubated overnight at 4°C. The next day, plates were washed once with wash buffer (PBS+0.05% v/v Tween-20, PBST), and blocked with 5% w/v non-fatty milk in PBST for 2 hours at 37°C. Plates were then washed 4 times using wash buffer.

[00153] Serially diluted anti-CD40 antibodies of the disclosure or controls (starting at 200 nM with a five-fold serial dilution) in PBST with 2.5% w/v non-fatty milk were added to the CD40-Fc bound plates, 100 μ l per well, and incubated at 37°C for 40 minutes. Plates were washed 4 times again using wash buffer, and then added and incubated for 40 minutes at 37°C with 100 μ l/well 95 ng/mL biotin-labeled human CD40L-his protein (Sino biological Inc., Cat#10239-H08E). Plates were washed again using wash buffer. Thereafter, the plates were added with 100 μ l/well of streptavidin conjugated HRP (1:10000 dilution in PBST buffer, Jackson Immunoresearch, Cat#016-030-084) and incubated for 40 minutes at 37°C. Plates were washed again using wash buffer. Finally, TMB was added and the reaction was stopped using 1M H₂SO₄, and the absorbance of each well was read on a microplate reader using dual wavelength mode with 450 nm for TMB and 630 nm as the reference wavelength, then the OD (450-630) values were plotted against antibody concentration. Data was analyzed using Graphpad Prism software and IC₅₀ values were reported.

4.2 Benchmark Blocking ELISA

[00154] The abilities of the anti-CD40 antibodies of the disclosure to block Benchmark -human CD40 binding was measured in a competitive ELISA assay. Briefly, the BM2 antibody was coated on 96-well micro plates at 1 μ g/mL in PBS, 100 μ l per well, and incubated overnight at 4°C. The next day, plates were washed once with wash buffer, and blocked with blocking buffer (5% w/v non-fatty milk in PBST) for 2 hours at 37°C. While blocking, the anti-CD40 antibodies of the disclosure or controls were diluted with biotin labeled human CD40-Fc protein (amino acid residue 21-193 of SEQ ID NO: 30 linked to N-terminus of amino acid residue 99-330 of SEQ ID NO.:28, 0.23 nM in 2.5% v/v non-fatty milk in PBST), starting at 66.7 nM with a 5-fold serial dilution, and incubated at room temperature for 40 minutes. After plate washing 4 times, the antibody/human CD40-Fc-biotin mixtures were added to BM2 coated plates, 100 μ l/well. After incubation at 37°C for 40 minutes, plates were washed 4 times again using wash buffer. Then the plates were added and incubated with 100 μ l/well streptavidin conjugated HRP for 40 minutes at 37°C to detect biotin-labeled human CD40-Fc bound to plates. Plates were washed 4 times again using wash buffer. Finally, TMB was added and the reaction was stopped using 1M H₂SO₄, and the absorbance was read on a microplate reader using dual wavelength mode with 450 nm for TMB and 630 nm as the reference wavelength, then the OD (450-630) values were plotted against antibody concentration. Data was analyzed using Graphpad Prism software and IC₅₀ values were reported.

[00155] The results of the two assays were shown in FIGs. 3A-3B and 4A-4B.

[00156] It can be seen from FIG. 3B that the mouse antibody C1H1 was capable of blocking human CD40-human CD40L binding at an IC₅₀ value similar or even lower as compared to BM1 and BM2. Further, as shown in FIG. 3B, the mouse antibody C1H1 showed higher maximum blocking than BM1 and BM2.

[00157] FIGs. 4A and 4B showed that the mouse antibody C1H1 was able to block human CD40-BM2 binding, suggesting that it bound to the same or similar epitope as BM2 did. The mouse antibodies 1A3 and 1D1 showing no blocking may bind to different epitopes.

Example 5 Cell Based Reporter Assay of Mouse Anti-CD40 Antibodies

[00158] The anti-CD40 antibodies of the disclosure were further tested for their agonistic activities using a CD40-expressing reporter cell line 293T-NF- κ B-Luc-CD40 which stably expressed full length human CD40 (uniprot No. P25942-1, SEQ ID NO.: 30). The 293T-NF- κ B-Luc-CD40 cells were prepared, following the instruction of lipofectamine 3000 transfection reagent (Thermo Fisher), by

transfecting 293T cells with pGL4.32[luc2P NF- κ B-RE Hygro] vectors (Promega, GenBank® Accession Number: EU581860) and later pCMV-T-P plasmids inserted with CD40 coding sequence between *EcoRI* and *XbaI* sites. When CD40 agonists were added to these cells, CD40 signaling was activated and luciferase expression was up-regulated which can be measured in a luminescence assay.

[00159] Briefly, 5×10^3 293T-NF- κ B-Luc-CD40 cells at the log phase stage in 20 μ L DMEM medium (Gibco Inc., Cat# 10566-016) supplemented with 10% FBS (Gibco, Cat#10099-141) were plated into 384-well cell culture plates (Corning, Cat#3707). Then, the plates were added with 20 μ l/well serially diluted anti-CD40 antibodies of the disclosure or controls (including an in house made anti-CD22 antibody as a negative control) (starting from 200 nM, 3-fold serial dilution in the culture medium), and incubated at 37°C for 6 hours. Then, the plates were added with the reagents of ONE-Glo™ Luciferase Assay System (Promega, Cat#E6120, 30 μ l/well) and incubated for 5 minutes at room temperature. Chemiluminescence was measured using a Tecan Infiniit® 200 Pro equipment. Data was analyzed using Graphpad Prism software and EC₅₀ values were reported.

[00160] The results were shown in FIGs. 5A-5B.

[00161] It can be seen that the mouse antibodies 1A3, 1D1 and C1H1 had comparable agonistic activities compared to BM1 but not as good as that of BM2.

Example 6 Generation and Characterization of Chimeric Antibodies

[00162] The variable domains of the heavy and light chain of the anti-CD40 mouse mAbs were sequenced and summarized in Table 1.

[00163] The variable domains of the heavy and light chain of the anti-CD40 mouse mAbs C1H1, 1D1 and 1A3 were cloned in frame to human IgG1 heavy-chain (SEQ ID NO.: 27) and human kappa light-chain constant regions (SEQ ID NO.: 29), respectively, wherein the C terminus of the variable region was linked to the N terminus of respective constant region.

[00164] The vectors each containing a nucleotide encoding a heavy chain variable region linked to human IgG1 heavy-chain constant region, and the vectors each containing a nucleotide encoding a light chain variable region linked to human kappa light-chain constant region were transiently transfected into 50 ml of 293F suspension cell cultures in a ratio of 1.1:1 light to heavy chain construct, with 1 mg/mL PEI.

[00165] Cell supernatants were harvested after six days in shaking flasks, spun down to pellet cells, and then chimeric antibodies were purified from cell supernatants as described above. The purified chimeric antibodies were tested in the capture ELISA, BIAcore affinity test and cell-based reporter assays, following the protocols in the foregoing Examples with or without modifications and protocols described below.

[00166] For the BIAcore, goat anti-human IgG (GE healthcare, Cat#BR100839, Human Antibody Capture Kit) was covalently linked to a CM5 chip instead of goat anti-mouse IgG, and a CM5 chip was used for BM1 and BM2 instead of a Protein G chip. The results were shown in Table 3.

[00167] For the capture ELISA, AffiniPure Goat Anti-Human IgG, F(ab')₂ fragment specific (Jackson Immuno Research, Cat#109-005-097) was used instead of AffiniPure Goat Anti-Mouse IgG, F(ab')₂ fragment specific, 100 μ l/well.

[00168] The results were shown in Table 3, FIGs. 6A-6C and FIGs. 7A-7C.

[00169] The data showed that the chimeric antibodies had similar binding capacities and agonistic activities to their parent antibodies. In particular, the chimeric C1H1 antibody showed higher binding affinity and capacity to human CD40, and higher binding affinity to Cynomolgus CD40 than BM1.

Table 3. Binding affinity of chimeric antibodies

Clone ID#	Kinetics on Biacore					
	Human CD40-his			Cyno CD40-his		
	K_a ($M^{-1}s^{-1}$)	K_d (s^{-1})	K_D (M)	K_a ($M^{-1}s^{-1}$)	K_d (s^{-1})	K_D (M)
mouse C1H1	1.79E+06	0.001227	6.84E-10	1.39E+06	0.001184	8.52E-10
chimeric C1H1	1.82E+06	0.001567	8.61E-10	1.39E+06	0.001478	1.06E-09
BM1	7.39E+05	0.006556	8.87E-09	6.23E+05	0.006164	9.89E-09

Example 7 Humanization of Anti-CD40 Mouse Monoclonal Antibody C1H1

[00170] The mouse anti-CD40 antibody C1H1 was selected for humanization and further investigations. Humanization of this antibody was conducted using the well-established CDR-grafting method as described in detail below.

[00171] To select acceptor frameworks for humanization of mouse antibody C1H1, the light and heavy chain variable region sequences of mouse C1H1 were blasted against the human immunoglobulin gene database. The human germlines with the highest homology to mouse C1H1 were selected as the acceptor frameworks for humanization. The mouse antibody heavy/light chain variable region CDRs were inserted into the selected frameworks, and the residue(s) in the frameworks was/were further mutated to obtain more candidate heavy chain/light chain variable regions. A total of 4 humanized C1H1 antibodies, namely from huC1H1-V1 to huC1H1-V4 were obtained whose heavy/light chain variable region sequences were in Table 1.

[00172] The vectors each containing a nucleotide encoding a the heavy chain variable region linked to human IgG2 heavy-chain constant (SEQ ID NOs: 28), and the vectors each containing a nucleotide encoding a humanized light chain variable region linked to human kappa light-chain constant regions (SEQ ID NOs: 29) were transiently transfected into 50 ml 293F suspension cell cultures in a ratio of 60% to 40% light to heavy chain construct, with 1 mg/ml PEI.

Example 8 Characterization of Humanized Antibodies

[00173] Cell supernatants containing humanized antibodies huC1H1-V1 to huC1H1-V4 were harvested after six days in shaking flasks and tested for binding affinity to human CD40 by Octect using the following protocol below, using Octet system (ForteBio, Octet RED 96). Briefly, AHC biosensors (anti-human IgG Fc capture, from ForteBio) were presoaked with 10 mM glycine (pH 1.5) for 3 seconds, and then dipped in a well with running buffer (0.5% w/v BSA in PBST) for 3 seconds, the soaking and dipping steps were repeated for three times. Then, the sensors were dipped in a well with cell supernatant containing humanized anti-CD40 antibodies, the chimeric C1H1 antibody in HBS-EP⁺ at 5 μ g/ml, or the benchmark in HBS-EP⁺ at 5 μ g/ml for 120 seconds, and then immersed in a well with running buffer for 5 minutes. A new baseline was run for 180 seconds in another well with running buffer. Then the sensors were dipped in a well with serially diluted human CD40-his protein (Acro biosystems, Cat#CD0-H5228, starting at 40 nM with a two-fold serial dilution) in running buffer for 120 seconds, and then immersed in a baseline well for 10 minutes. Finally, sensors were presoaked with 10 mM glycine (pH 1.5) for 3 seconds, and then dipped in a well with running buffer for 3 seconds, the soaking and dipping steps repeated for three times. The association and dissociation curves were fit to a 1:1 Langmuir binding model using ForteBio Data Analysis 8.1. The K_a , K_d and K_D values were determined and summarized in Table 4 below.

[00174] The results indicated that huC1H1-V1 and huC1H1-V2 as tested had similar human CD40 binding affinity, compared to the chimeric antibody C1H1, which was better than that of BM2.

Table 4. Affinity of Humanized C1H1 mAbs

Clone ID#	Octet Kinetics of Humanized C1H1 mAbs Binding to Human CD40		
	Kinetics on Octet		
	Human CD40-his		
	K_a ($M^{-1}s^{-1}$)	K_d (s^{-1})	K_D (M)
chimeric C1H1	1.47E+06	9.29E-04	6.33E-10
huC1H1-V1	1.41E+06	1.28E-03	9.04E-10
huC1H1-V2	1.65E+06	9.81E-04	5.96E-10
huC1H1-V3	5.23E+05	2.83E-03	5.41E-09
huC1H1-V4	4.24E+05	3.67E-03	8.66E-09
BM2	8.74E+05	9.79E-04	1.12E-09

[00175] The humanized antibodies huC1H1-V1 and huC1H1-V2 were purified as described above and tested in Biacore, Capture ELISA, Cell-based binding FACS, Competitive ELISA, Cell-based reporter assay and Protein thermal shift assay, following the protocols in the foregoing Examples with or without modifications and protocols described below.

[00176] For the capture ELISA, AffiniPure Goat Anti-Human IgG, F(ab')₂ fragment specific, (Jackson Immuno Research, Cat#109-005-097) was used instead of AffiniPure Goat Anti-Mouse IgG, F(ab')₂ fragment specific, 100 µl/well.

[00177] For the BIAcore, goat anti-human IgG (GE healthcare, Cat#BR100839, Human Antibody Capture Kit) was covalently linked to a CM5 chip instead of goat anti-mouse IgG, and a CM5 chip was used for BM1 and BM2 instead of a Protein G chip.

[00178] For the cell-based binding FACS, R-Phycoerythrin AffiniPure Goat Anti-human IgG Fcγ fragment specific (Jackson Immuno Research, Cat#109-115-098) was used instead of R-Phycoerythrin AffiniPure F(ab')₂ Fragment Goat Anti-Mouse IgG (H+L), 100 µl/well.

[00179] The humanized antibody huC1H1-V2 was also tested for the thermal stability assay. A protein thermal shift assay was used to determine T_m (melting temperature) using a GloMelt™ Thermal Shift Protein Stability Kit (Biotium, Cat# 33022-T). Briefly, the GloMelt™ dye was allowed to thaw and reach room temperature. The vial containing the dye was vortexed and centrifuged. Then, 10x dye was prepared by adding 5 µL 200x dye to 95 µL PBS. 2 µL 10x dye and 10 µg humanized antibodies were added, and PBS was added to a total reaction volume of 20 µL. The tubes containing the dye and antibodies were briefly spun and placed in real-time PCR thermocycler (Roche, LightCycler 480 II) set up with a melt curve program having the parameters in Table 5.

Table 5. Parameters for Melt Curve Program

Profile step	Temperature	Ramp rate	Holding Time
Initial hold	25°C	NA	30 s
Melt curve	25-99°C	0.1°C/s	NA

[00180] The assay results were shown in Table 6 and FIGs. 8-12.

[00181] It can be seen from Table 6 that the humanized antibodies huC1H1-V1 and huC1H1-V2 showed comparable binding affinities to human and cynomolgus CD40 compared to the chimeric antibody C1H1. In other words, the huC1H1-V1 and huC1H1-V2's binding affinities to human and cynomolgus CD40 were higher than those of BM1 and BM2.

[00182] FIG. 10 showed that the humanized antibodies huC1H1-V1 and huC1H1-V2 of the disclosure were capable of blocking CD40-CD40L binding, and the blocking activities were comparable or a bit lower when compared to BM1 and BM2.

[00183] According to FIG. 11, the humanized antibodies huC1H1-V1 and huC1H1-V2 of the disclosure were able to block human CD40-BM2 binding, suggesting that the antibodies huC1H1-V1 and huC1H1-V2 of the disclosure might bind to a similar epitope as BM2 did.

[00184] As shown in FIG. 12, the humanized antibodies huC1H1-V1 and huC1H1-V2 of the disclosure had higher agonistic activities in the cell-based reporter assay when compared to the BM1 and BM2.

Table 6. Binding and Functional activities of Humanized C1H1 antibodies

Clone ID#	Kinetics on Biacore						T _m (melting temperature) °C
	Human CD40-his			Cynomolgus CD40-his			
	K _a (M ⁻¹ s ⁻¹)	K _d (s ⁻¹)	K _D (M)	K _a (M ⁻¹ s ⁻¹)	K _d (s ⁻¹)	K _D (M)	
mouse C1H1	2.10E+06	8.97E-04	4.28E-10	2.29E+06	8.63E-04	3.77E-10	*
chimeric C1H1	2.51E+06	0.002221	8.84E-10	2.54E+06	0.002147	8.46E-10	*
huC1H1-V1	2.25E+06	2.48E-03	1.10E-09	2.14E+06	0.002371	1.11E-09	*
huC1H1-V2	2.31E+06	0.002704	1.17E-09	2.56E+06	0.002735	1.07E-09	70
BM1	8.03E+05	0.007562	9.42E-09	7.64E+05	0.007397	9.68E-09	*
BM2	2.06E+05	0.001616	7.84E-09	1.73E+05	0.001508	8.73E-09	*

*Not tested.

Example 9 *In Vivo* Anti-tumor Efficacy of huC1H1-V2 Antibody in Human B-cell Lymphoma Ramos Xenograft Model

[00185] The *in vivo* anti-tumor activity of huC1H1-V2 antibody was tested in NOD-SCID mice. Briefly, NOD-SCID mice were subcutaneously injected with 3×10^7 Human B-cell Lymphoma Ramos cells (Chinese Academy of Sciences) at the right axilla. Tumor volumes were measured using electronic caliper and calculated as $(\text{length} \times \text{width}^2)/2$. When tumors reached an average volume of about 100-150 mm³, 40 tumor-bearing mice were selected and randomized into 4 groups (10 mice/group), and the day doing the animal grouping was designated as Day 0. The animals were intravenously injected at the tail vein with isotype control Ab (anti-HEL-Human IgG2 Isotype-control, also referred to as IgG2, Biointron Inc) or huC1H1-V2 antibody from Day 0 according to dosing regimen shown in Table 7.

Table 7. Study Design of Human B-cell Lymphoma Ramos Xenograft Model

Group No.	N	Compound	Dose (mg/kg)	Dose Route	Dosing Volume (ml/kg)	Dosing Schedule
1	10	isotype control Ab (IgG2)	15	i.v.	10	Q2D×8 times
2	10	huC1H1-V2	1.5	i.v.	10	Q2D×8 times
3	10	huC1H1-V2	5	i.v.	10	Q2D×8 times
4	10	huC1H1-V2	15	i.v.	10	Q2D×8 times

Note: N: animal number per group; Q2D: every 2 days.

[00186] The data was summarized in Table 8.

[00187] All treatments were well tolerated by the tumor-bearing animals and there was no significant weight loss or other symptoms. The huC1H1-V2 antibody treatment at 5 mg/kg exhibited strongest anti-tumor activity, resulting in mean tumor size of 230.2 mm³ on Day 14 with tumor growth inhibition (TGI) of 96%. The mean tumor volume in Group 3 at Day 14 was statistically smaller than that of the control group. However, the huC1H1-V2 antibody treatment at 15 mg/kg did not show further enhanced efficacy, i.e., the tumor volume of Group 4 was not significantly different from that of Group 3.

Table 8. Anti-Tumor Efficacy of huC1H1-V2 antibody in Human B-cell Lymphoma Ramos Xenograft Model

Treatment	Tumor Volume (mm ³) ^a on Day 0 of treatment	Tumor Volume (mm ³) ^{a, b} on Day 14 of treatment	T/C (%) ^c	TGI (%) ^d	P value
IgG2, 15 mg/kg, Q2D	114.7±1.4	2830.4±235.7	-	-	-
huC1H1-V2, 1.5 mg/kg, Q2D	111.1±2.0	431.9±58.0**	12	88	<0.001
huC1H1-V2, 5 mg/kg, Q2D	114.5±1.5	230.2±36.3**	4	96	<0.001
huC1H1-V2, 15 mg/kg, Q2D	114.3±1.3	242.5±22.1**	5	95	<0.001

Note: a. Mean ± SEM; b. Tumor volume at Day 14, compared with control group by student's t test, *P<0.05 and **P<0.01; c. T/C(%)=(T-T₀)/(C-C₀) × 100%, where T and C referred to the mean tumor volume of the treated and control groups on Day 14, respectively, and T₀ and C₀ referred to the mean tumor volume of the treated and control groups on Day 0, respectively; d. TGI (%)=(1- T/C) × 100%.

Example 10 *In Vivo* Anti-tumor Efficacy of huC1H1-V2 Antibody in mouse colon cancer MC38 Xenograft Model

[00188] The *in vivo* anti-tumor activity of huC1H1-V2 antibody was further tested in CD40 humanized transgenic mice (also referred to as hCD40 mice). Briefly, hCD40 mice were subcutaneously injected with 1 × 10⁶ mouse colon cancer MC38 cells (Shanghai Lanli Biological Technology Co., Ltd.) at the right flank. Tumor volumes were measured using electronic caliper and calculated as (length × width²)/2. When tumors reached an average volume of about 100 mm³, 18 tumor-bearing mice were selected and randomized into 3 groups, 6 mice per group, and the day doing the animal grouping was designated as Day 0. The animals were intravenously injected at the tail vein with vehicle (Phosphate-buffered saline, also referred to as PBS) or huC1H1-V2 antibody from Day 1 according to regimen shown in Table 9.

Table 9. Study Design of MC38 Xenograft Model

Group No.	N	Compound	Dose (mg/kg)	Dose Route	Dosing Volume (ml/kg)	Dosing Schedule
1	6	Vehicle (PBS)	-	i.v.	10	Q2D×7 times
2	6	huC1H1-V2	3	i.v.	10	Q2D×7 times
3	6	huC1H1-V2	10	i.v.	10	Q2D×7 times

Note: N: animal number per group; Q2D: every 2 days.

[00189] The data was summarized in Table 10.

[00190] All treatments were well tolerated by the tumor-bearing animals and there was no significant weight loss or symptoms. The huC1H1-V2 antibody treatment at 3 mg/kg exhibited strong anti-tumor activity, resulting in mean tumor size of 1169 mm³ on day 14 with a TGI of 64%. The tumor size of

Group 2 at Day 14 was statistically smaller than that of the vehicle group. The huC1H1-V2 antibody treatment at 10 mg/kg also exhibited strong anti-tumor activity, resulting in mean tumor size of 1446 mm³ on Day 14 with TGI of 56%. The mean tumor size of Group 3 on Day 14 was statistically smaller than that of the vehicle group.

Table 10. Anti-Tumor Efficacy of huC1H1-V2 antibody in MC38 Xenograft Model

Treatment	Tumor Volume (mm ³) ^a on Day 0 of treatment	Tumor Volume (mm ³) ^{a, b} on Day 14 of treatment	T/C (%) ^c	TGI (%) ^d	P value
Vehicle (DPBS), Q2D	112±9	3241±383	-	-	-
huC1H1-V2, 3 mg/kg, Q2D	111±5	1169±217**	36	64	<0.01
huC1H1-V2, 10 mg/kg, Q2D	113±8	1446±467*	44	56	0.014

Note: a. Mean ± SEM; b. Tumor volume at Day 14, compared with vehicle control group by student's t test, *P≤0.05 and **P<0.01; c. T/C(%)=T_{RTV}/C_{RTV} × 100%, where T_{RTV} and C_{RTV} referred to the mean RTVs of the treatment and control groups, respectively, RTV=V_t/V₀, where V_t referred to the mean tumor volume on Day 14, and V₀ referred to the mean tumor volume on Day 0; d. TGI (%)=(1-T/C) × 100%.

[00191] While the disclosure has been described above in connection with one or more embodiments, it should be understood that the disclosure is not limited to those embodiments, and the description is intended to cover all alternatives, modifications, and equivalents, as may be included within the spirit and scope of the appended claims. All referenced cited herein are further incorporated by reference in their entirety.

[00192] Sequences in the present application are summarized below.

Description/ Sequence/SEQ ID NO.
VH-CDR1 for mouse, chimeric and humanized C1H1 antibodies NYGIS (SEQ ID NO: 1)
VH-CDR2 for mouse, chimeric and humanized C1H1 antibodies SISSGGDNTYYPDNVKG (SEQ ID NO: 4)
VH-CDR3 for mouse, chimeric and humanized C1H1 antibodies AGEKAMDY (SEQ ID NO: 7)
VL-CDR1 for mouse, chimeric and humanized C1H1 antibodies RASQTINNNLH (SEQ ID NO: 10)
VL-CDR2 for mouse, chimeric and humanized C1H1 antibody YASQIS (SEQ ID NO: 13)
VL-CDR3 for mouse, chimeric and humanized C1H1 antibodies QQFSSWPLT (SEQ ID NO: 16)
VH for mouse and chimeric C1H1 antibodies EVKLVESGGGLVKPGASLKLSCAASGFTFSNYGISWVRQTS DKRLEWVASISSGGDNTYYPD NVKGRFTISRENAKNTLYLQMSSLKSEDTALYYCARAGEKAMDYWGQGTSVTVSS (SEQ ID NO: 19)
VH for mouse C1H1 GAAGTGAACTGGTGGAGTCTGGGGGAGGCTTAGTGAAGCCTGGAGCGTCTCTGAACT CTCCTGTGCAGCCTCTGGATTCACTTTCAGTAACTATGGCATACTTGGGTTCCGACACT TCAGACAAGAGGCTGGAGTGGGTCGCATCCATTAGTAGTGGTGGTGATAACACCTACTA TCCAGACAATGTAAAGGGCCGATTCACCATCTCCAGAGAGAATGCCAAGAACACCCTAT ACCTACAAATGAGTAGTCTGAAGTCTGAGGACACGGCCTTGTAATTACTGTGCAAGAGCTG GGGAGAAGGCTATGGACTACTGGGGTCAAGGAACCTCAGTCACCGTCTCCTCA (SEQ ID NO: 31)
VH for chimeric C1H1 GAGGTGAAGCTGGTGGAGTCCGGCGGCGGCCTGGTGAAGCCTGGAGCTAGCCTGAAGCT

<p>GAGCTGCGCCGCCAGCGGCTTCACCTTTTCCAACACTACGGCATCAGCTGGGTGAGGCAGAC AAGCGATAAGAGGCTGGAGTGGGTGGCCAGCATCAGCAGCGGCGGCATAACACATACT ACCCTGACAACGTGAAGGGCAGATTCACCATCAGCAGGGAGAACGCCAAGAATACCCTG TACCTGCAGATGAGCAGCCTGAAGAGCGAGGACACCGCCCTGTACTACTGTGCCAGGGC CGGCGAGAAGGCCATGGACTACTGGGGCCAGGGCACCTCCGTGACCGTGAGCTCC (SEQ ID NO.:32)</p>
<p>VH for huC1H1-V1 and huC1H1-V3 EVQLVESGGGLVKPGGSLRLSCAASGFTFSNYGISWVRQAPGKGLEWVXI SSSGGDNTYYP DNVKGRFTISRDNKNSLYLQMNSLRAEDTAVYYCARAGEKAMDYWGQGLVTVSS (SEQ ID NO: 20, X1=A) EVQLVESGGGLVKPGGSLRLSCAASGFTFSNYGISWVRQAPGKGLEWVASISSGGDNTYYPD NVKGRFTISRDNKNSLYLQMNSLRAEDTAVYYCARAGEKAMDYWGQGLVTVSS</p>
<p>VH for huC1H1-V2 and huC1H1-V4 EVQLVESGGGLVKPGGSLRLSCAASGFTFSNYGISWVRQAPGKGLEWVXI SSSGGDNTYYP DNVKGRFTISRDNKNSLYLQMNSLRAEDTAVYYCARAGEKAMDYWGQGLVTVSS (SEQ ID NO: 20, X1=S) EVQLVESGGGLVKPGGSLRLSCAASGFTFSNYGISWVRQAPGKGLEWVSSISSGGDNTYYPD NVKGRFTISRDNKNSLYLQMNSLRAEDTAVYYCARAGEKAMDYWGQGLVTVSS GAGGTGCAGCTGGTGGAGAGCGGCGGCGGACTGGTGAAGCCTGGCGGAAGCCTGAGAC TGAGCTGCGCCGCTCCGGCTTCACCTTCTCCAACACTACGGCATCAGCTGGGTGAGGCAGG CCCCCGAAAGGGCCTGGAGTGGGTGAGCAGCATCAGCAGCGGCGGCGACAATACCTAC TACCCTGACAACGTGAAGGGCAGGTTACCATCAGCAGAGACAATGCCAAGAATTCCCT GTACCTGCAGATGAACAGCCTGAGAGCCGAGGATACAGCCGTGTACTACTGTGCCAGAG CCGGCGAGAAGGCCATGGATTACTGGGGCCAGGGCACCTGGTGACCGTGTCTCTCC (SEQ ID NO: 33)</p>
<p>VL for mouse and chimeric C1H1 antibodies DIVLTQSPVTLSPGDSVLSLSCRASQTINNNLHWYQQKSHESPRLLIKYASQSIGIPSRFSGS GSGTDFTLTSSVETEDFGIYFCQQFSSWPLTFGAGTKLELK (SEQ ID NO: 23) VL for mouse C1H1 GATATTGTAATACTCAGTCTCCAGTCACCCGTCTGTGACTCCAGGAGATAGCGTCAGT CTTCTCTGCAGGGCCAGCCAACTATTAACAACAACCTACACTGGTATCAACAAAATCA CATGAGTCTCAAGGCTTCTCATCAAGTATGCTTCCCAGTCCATCTCTGGGATCCCCTCA GGTTCAGTGGCAGTGGATCAGGGACAGATTTCACTCTCAGTATCAACAGTGTGGAGACT GAAGATTTTGAATATATTTCTGTCAACAGTTTAGCAGCTGGCCTCTTACGTTCCGGTGCTG GGACTAAGCTGGAGCTGAAA (SEQ ID NO: 34) VL for chimeric C1H1 GATATCGTGCTGACCCAGAGCCCCGTGACCCTGAGCGTGACACCCGGCGACAGCGTGAG CCTGTCTGCAGAGCCAGCCAGCCATCAACAACAATCTGCACTGGTATCAACAGAAGA GCCACGAGAGCCCCAGGCTGCTGATCAAGTACGCCAGCCAGAGCATCTCCGGCATCCCT AGCAGATTCAGCGGCTCCGGCTCCGGCACAGACTTTACCCTGAGCATCAACAGCGTGGA GACCGAGGATTTCCGGCATCTACTTTTGCCAGCAGTTTTCCTCCTGGCCTCTGACATTCGGC GCCGGCACAAAGCTGGAGCTGAAG (SEQ ID NO: 35)</p>
<p>VL for huC1H1-V1 and huC1H1-V2 EIVLTQSPATLSLSPGERATLSCRASQTINNNLHWYQQKPGQAPRLLIXIYASQSIGIPARFSG SGSGTDFTLTISSLEPEDFAVYX2CQQFSSWPLTFGGGKVEIK (SEQ ID NO: 24, X1=K, X2=F) EIVLTQSPATLSLSPGERATLSCRASQTINNNLHWYQQKPGQAPRLLIKYASQSIGIPARFSGS GSGTDFTLTISSLEPEDFAVYFCQQFSSWPLTFGGGKVEIK GAGATCGTGCTGACCCAGAGCCCCGCCACCCTGAGCCTGAGCCCTGGAGAGAGGGCCAC CCTGTCTGCAGGGCCTCCAGACAATCAATAATAATCTGCACTGGTATCAACAGAAGCC CGGCCAGGCCCCAGGCTGCTGATCAAGTACGCCAGCCAGTCCATCAGCGGCATCCCTG CCAGGTTCTCCGGCAGCGGCAGCGGAACAGACTTACCCTGACCATCTCCTCCCTGGAGC CTGAGGACTTTGCCGTGTACTTTTGCCAGCAGTTCTCCAGCTGGCCTCTGACCTTTGGCG CGGCACCAAGGTGGAGATCAAG (SEQ ID NO:36)</p>

<p>VL for huC1H1-V3 and huC1H1-V4 EIVLTQSPATLSLSPGERATLSCRASQTINNNLHWYQQKPGQAPRLLIX1YASQSIGIPARFSG SSGTDFLTISSLEPEDFAVYX2CQQFSSWPLTFGGGKVEIK (SEQ ID NO: 24, X1=Y, X2=Y) EIVLTQSPATLSLSPGERATLSCRASQTINNNLHWYQQKPGQAPRLLIYYASQSIGIPARFSGS GSGTDFLTISSLEPEDFAVYYCQQFSSWPLTFGGGKVEIK</p>
<p>VH-CDR1 for mouse and chimeric 1A3 antibodies GYYLH (SEQ ID NO: 2)</p>
<p>VH-CDR2 for mouse and chimeric 1A3 antibodies YISCHDGTIIYNQKFKG (SEQ ID NO: 5)</p>
<p>VH-CDR3 for mouse and chimeric 1A3 antibodies FLNYYGSNYAMDY (SEQ ID NO: 8)</p>
<p>VL-CDR1 for mouse and chimeric 1A3 antibodies KASQDVGPAVA (SEQ ID NO: 11)</p>
<p>VL-CDR2 for mouse and chimeric 1A3 antibodies WASTRHT (SEQ ID NO: 14)</p>
<p>VL-CDR3 for mouse and chimeric 1A3 antibodies QQYFTYPLT (SEQ ID NO: 17)</p>
<p>VH for mouse and chimeric 1A3 antibodies EVQLQQSGPELVKGTGASVKISCKASGYSGFYGLHWVKQSLGKGLEWIGYISCHDGTIIYNQK FKGKATFTLDTSSSTAYMQFSSLTSEDSAVYFCARFLNYYGSNYAMDYWGQGTSTVTVSS (SEQ ID NO: 21)</p>
<p>VL for mouse and chimeric 1A3 antibodies DIVMTQSHKFMSTSVGDRVSITCKASQDVGPAAVWYQQKPGQSPKLLIYWASTRHTGVPDR FTGSGYGTDFLTINNVQSEDLADYFCQQYFTYPLTFGGGKLEIK (SEQ ID NO: 25)</p>
<p>VH-CDR1 for mouse and chimeric 1D1 antibodies DYVMH (SEQ ID NO: 3)</p>
<p>VH-CDR2 for mouse and chimeric 1D1 antibodies YINPYNDGTTYNEKFKG (SEQ ID NO: 6)</p>
<p>VH-CDR3 for mouse and chimeric 1D1 antibodies GFLRESWFGY (SEQ ID NO: 9)</p>
<p>VL-CDR1 for mouse and chimeric 1D1 antibodies RSSQNIVHSNGNTYLD (SEQ ID NO: 12)</p>
<p>VL-CDR2 for mouse and chimeric 1D1 antibodies KVSNRFS (SEQ ID NO: 15)</p>
<p>VL-CDR3 for mouse and chimeric 1D1 antibodies FQGSHPPT (SEQ ID NO: 18)</p>
<p>VH for mouse and chimeric 1D1 antibodies EVQLQQSGPELVKPGASVKMSCKASGYTFTDYVMHWVKQKPGQGLEICIGYINPYNDGTTY NEKFKGKATLTSKSSNAAYLESSLTSEDSAVYYCARGFLRESWFGYWGQGTSLTVSA (SEQ ID NO: 22)</p>
<p>VL for mouse and chimeric 1D1 antibodies DVLLTQTPLSLPVSLGDQASISCRSSQNIVHSNGNTYLDWYLQKPGQSPKLLIYKVSNRFSGV PDRFSGSGSGTDFLTKISRVEAEDLGVYYCFQGSHPPTFGGGKLELK (SEQ ID NO:26)</p>
<p>Heavy chain constant region (for chimeric antibodies) ASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAVLQSSGLY SLSSVTVPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPAPELLGGPSVFLFP PKPKDTLMISRTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYNSTYRVVSV LTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSREEMTKNQVSLTC LVKGFYPSDIAVEWESNGQPENNYKTPPVLDSDGSFFLYSKLTVDKSRWQQGNVDFCSVM HEALHNHYTQKSLSLSPGK (SEQ ID NO.: 27) GCTAGCACCAAGGGCCATCGGTCTTCCCCCTGGCACCTCCTCCAAGAGCACCTCTGGG GGCACAGCGCCCTGGGCTGCCTGGTCAAGGACTACTTCCCCGAACCGGTGACGGTGTC GTGGAACCTCAGGCGCCCTGACCAGCGGCGTGCACACCTTCCCGGCTGTCTACAGTCCTC</p>

AGGACTCTACTCCCTCAGCAGCGTGGTGACCGTGCCCTCCAGCAGCTTGGGGACCCAGAC
 CTACATCTGCAACGTGAATCACAAGCCAGCAACACCAAGGTGGACAAGAAAGTTGAGC
 CCAAATCTTGTGACAAAACCTCACACATGCCACCGTGCCAGCACCTGAACTCCTGGGGG
 GACCGTCAGTCTTCTCTTCCCCCAAACCCAAGGACACCCTCATGATCTCCCGGACCC
 CTGAGGTCACATGCGTGGTGGTGGACGTGAGCCACGAAGACCCTGAGGTCAAGTTCAAC
 TGGTACGTGGACGGCGTGGAGGTGCATAATGCCAAGACAAAGCCGCGGGAGGAGCAGT
 ACAACAGCACGTACCGTGTGGTACGCGTCTCACCGTCTGCACCAGGACTGGCTGAATG
 GCAAGGAGTACAAGTGCAAGGTCTCCAACAAAGCCCTCCAGCCCCATCGAGAAAACC
 ATCTCAAAGCCAAAGGGCAGCCCCGAGAACCACAGGTGTACACCCTGCCCCATCCCG
 GGAGGAGATGACCAAGAACCAGGTGAGCCTGACCTGCCTGGTCAAAGGCTTCTATCCCA
 GCGACATCGCCGTGGAGTGGGAGAGCAATGGGCAGCCGGAGAACAATAAGACCAC
 GCCTCCCGTGCTGGACTCCGACGGCTCCTTCTTCTCTACAGCAAGCTCACCGTGGACAA
 GAGCAGGTGGCAGCAGGGGAACGTCTTCTCATGCTCCGTGATGCATGAGGCTCTGCACA
 ACCACTACACGCAGAAGAGCCTCTCCCTGTCTCCGGGTAAATGA (SEQ ID NO:37)

Heavy chain constant region (for humanized antibodies)
 ASTKGPSVFPLAPCSRSTSESTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAVLQSSGLY
 SLSSVVTVPSSNFGTQTYTCNVDHKPSNTKVDKTKVERKCCVECPPCPAPPVAGPSVFLFPPKP
 KDTLMISRTPEVTCVVVDVSHEDPEVQFNWYVDGVEVHNAKTKPREEQFNSTFRVVSIVLTV
 VHQDWLNGKEYKCKVSNKGLPAPIEKTIKTKGQPREPQVYTLPPSREEMTKNQVSLTCLVK
 GFYPSDIAVEWESNGQPENNYKTTTPMLDSDGSFFLYSKLTVDKSRWQQGNVFCFSVMHEA
 LHNHYTQKSLSLSPGK (SEQ ID NO.:28)
 GCCTCCACCAAGGGCCATCGGTCTTCCCCCTGGCGCCCTGCTCCAGGAGCACCTCTGAG
 AGCACAGCGGCCCTGGGCTGCCTGGTCAAGGACTACTTCCCCGAACCGGTGACGGTGTC
 GTGGAACTCAGGCGCTCTGACCAGCGGCGTGCACACCTTCCAGCTGTCTTACAGTCTCTC
 AGGACTCTACTCCCTCAGCAGCGTGGTGACCGTGCCCTCCAGCAACTTCGGCACCCAGAC
 CTACACCTGCAACGTAGATCACAAGCCAGCAACACCAAGGTGGACAAGACAGTTGAGC
 GCAAATGTTGTGTCGAGTGCCACCGTGCCAGCACACCTGTGGCAGGACCGTCACTCT
 TCTCTTCCCCCAAACCCAAGGACACCCTCATGATCTCCCGGACCCCTGAGGTACAGT
 GCGTGGTGGTGGACGTGAGCCACGAAGACCCCGAGGTCCAGTTCAACTGGTACGTGGAC
 GGCGTGGAGGTGCATAATGCCAAGACAAAGCCACGGGAGGAGCAGTTCAACAGCACGTT
 CCGTGTGGTCAGCGTCTCACCGTTGTGCACCAGGACTGGCTGAACGGCAAGGAGTACA
 AGTGCAAGGTCTCCAACAAAGGCCTCCAGCCCCATCGAGAAAACCATCTCCAAAACC
 AAAGGGCAGCCCCGAGAACCACAGGTGTACACCCTGCCCCATCCCGGGAGGAGATGAC
 CAAGAACCAGGTCAGCCTGACCTGCCTGGTCAAAGGCTTCTACCCAGCGACATCGCCGT
 GGAGTGGGAGAGCAATGGGCAGCCGGAGAACAATAAGACCACACCTCCCATGCTG
 GACTCCGACGGCTCCTTCTTCTCTACAGCAAGCTCACCGTGGACAAGAGCAGGTGGCAG
 CAGGGGAACGTCTTCTCATGCTCCGTGATGCATGAGGCTCTGCACAACCACTACACGCAG
 AAGAGCCTCTCCCTGTCTCCGGGTAAATGA (SEQ ID NO:38)

Light chain constant region for chimeric and humanized antibodies
 RTVAAPSVFIFPPSDEQLKSGTASVIVCLLNNFYPREAKVQWKVDNALQSGNSQESVTEQDSK
 DSTYLSSTLTLSKADYEKHKVYACEVTHQGLSSPVTKSFNRGEC (SEQ ID NO.: 29)
 CGTACGGTGGCGGCGCCATCTGTCTTCTATCTTCCCGCCATCTGATGAGCAGTTGAAATCT
 GGAACTGCCTCTGTTGTGTGCCTGCTGAATAACTTCTATCCAGAGAGGCCAAAGTACAG
 TGGAAGGTGGATAACGCCCTCCAATCGGGTAACTCCAGGAGAGTGTACAGAGCAGGA
 CAGCAAGGACAGCACCTACAGCCTCAGCAGCACCTGACGCTGAGCAAAGCAGACTACG
 AGAAACACAAAGTCTACGCCTGCGAAGTCAACCATCAGGGCCTGAGCTCGCCGTCACA
 AAGAGCTTCAACAGGGGAGAGTGTGA (SEQ ID NO:39)

Human CD40, uniprot #P25942-1
 MVRLPLQCVLWGCLLTAHVHPEPPTACREKQYLINSQCCSLCQPGQKLVSDCTEFTETECLPC
 GESEFLDTWNRETHCHQHKYCDPNLGLRVQKGTSETDTICTCEEGWHCTSEACESVLHRS
 CSPGFVKQIATGVSDTICEPCPVGFFSNVSSAFEKCHPWTSCETKDLVVQQAGTNKTDVVC
 GPQDRRLRALVVIPIIFGILFAILLVLFIKKVAKKPTNKAPHPKQEPQEINFDDLPGSNTAAPV
 QETLHGCQPVTQEDGKESRISVQERQ (SEQ ID NO:30)

Heavy chain of Dacetuzumab

<p>EVQLVESGGGLVQPGGSLRSLCAASGYSFTGYIHWVRQAPGKGLEWVARVIPNAGGTSYN QKFKGRFTLSVDNSKNTAYLQMNSLRAEDTAVYYCAREGIYWWGQGLTVTVSSASTKGPS VFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSQVHTFPAVLQSSGLYSLSSVVT VPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPAPELLGGPSVFLFPPKPKDTL MISRTPEVTCVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYNSTYRVVSVLTVLHQ DWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSREEMTKNQVSLTCLVKGFY PSDIAVEWESNGQPENNYKTPPVLDSDGSFFLYSKLTVDKSRWQQGNVFSCSVMHEALHN HYTQKSLSLSPGK (SEQ ID NO:40)</p>
<p>Light chain of Dacetuzumab DIQMTQSPSSLSASVGDRVTITCRSSQSLVHSNGNTFLHWYQQKPGKAPKLLIYTVSNRFSGV PSRFSGSGSGTDFLTISLQPEDFATYFCSQTHVPTWTFGGTKVEIKRTVAAPSVFIFPPSDE QLKSGTASVVCLLNNFYPREAKVQWKVDNALQSGNSQESVTEQDSKDSTYLSSTLTLSKA DYEKHKVYACEVTHQGLSSPVTKSFNRGEC (SEQ ID NO:41)</p>
<p>Heavy chain of Selicrelumab QVQLVQSGAEVKKPGASVKVSCKASGYTFTGYIMHWVRQAPGQGLEWMGWINPDSGGTN YAQKFQGRVTMTRDTSISTAYMELNRLRSDDTAVYYCARDQPLGYCTNGVCSYFDYWGQG TLTVTVSSASTKGPSVFPLAPCSRSTSESTAALGCLVKDYFPEPVTVSWNSGALTSQVHTFPAV LQSSGLYSLSSVVTVPSSNFGTQTYTCNVDHKPSNTKVDKTKVERKCCVECPCPAPPVAGPSV FLFPPKPKDTLMISRTPEVTCVVDVSHEDPEVQFNWYVDGVEVHNAKTKPREEQFNSTFRV VSVLTVVHQDWLNGKEYKCKVSNKGLPAPIEKTISKTKGQPREPQVYTLPPSREEMTKNQVS LTCLVKGFYPSDIAVEWESNGQPENNYKTPPMLDSDGSFFLYSKLTVDKSRWQQGNVFSCS VMHEALHNHYTQKSLSLSPGK (SEQ ID NO:42)</p>
<p>Light chain of Selicrelumab DIQMTQSPSSVSASVGDRVTITCRASQGIYSWLAWYQQKPGKAPNLLIYTASTLQSGVPSRFS GSGSGTDFLTISLQPEDFATYYCQANIFPLTFGGGKVEIKRTVAAPSVFIFPPSDEQLKSG TASVVCLLNNFYPREAKVQWKVDNALQSGNSQESVTEQDSKDSTYLSSTLTLSKADYEK KVYACEVTHQGLSSPVTKSFNRGEC (SEQ ID NO:43)</p>

* * *

[00193] Having thus described in detail preferred embodiments of the present invention, it is to be understood that the invention defined by the above paragraphs is not to be limited to particular details set forth in the above description as many apparent variations thereof are possible without departing from the spirit or scope of the present invention.

We claim:

1. An isolated monoclonal antibody or an antigen-binding portion thereof, binding to CD40, comprising (i) a heavy chain variable region comprising a VH CDR1 region, a VH CDR2 region and a VH CDR3 region, wherein the VH CDR1 region, the VH CDR2 region and the VH CDR3 region comprise amino acid sequences having at least 85%, 86%, 87%, 88%, 89%, 90%, 91%, 92%, 93%, 94%, 95%, 96%, 97%, 98%, 99% or 100% identity to (1) SEQ ID NOs: 1, 4 and 7, respectively; (2) SEQ ID NOs: 2, 5 and 8, respectively; or (3) SEQ ID NOs: 3, 6 and 9, respectively; and/or (ii) a light chain variable region comprising a VL CDR1 region, a VL CDR2 region and a VL CDR3 region, wherein the VL CDR1 region, the VL CDR2 region and the VL CDR3 region comprise amino acid sequences having at least 85%, 86%, 87%, 88%, 89%, 90%, 91%, 92%, 93%, 94%, 95%, 96%, 97%, 98%, 99% or 100% identity to (1) SEQ ID NOs: 10, 13 and 16, respectively; (2) SEQ ID NOs: 11, 14 and 17, respectively; or (3) SEQ ID NOs: 12, 15 and 18, respectively.
2. The isolated monoclonal antibody or the antigen-binding portion thereof of claim 1, wherein the heavy chain variable region comprises an amino acid sequence having at least 85%, 86%, 87%, 88%, 89%, 90%, 91%, 92%, 93%, 94%, 95%, 96%, 97%, 98%, 99% or 100% identity to SEQ ID NOs: 19, 20 (X1=A or S), 21 or 22.
3. The isolated monoclonal antibody or the antigen-binding portion thereof of claim 1, wherein the light chain variable region comprises an amino acid sequence having at least 85%, 86%, 87%, 88%, 89%, 90%, 91%, 92%, 93%, 94%, 95%, 96%, 97%, 98%, 99% or 100% identity to SEQ ID NO: 23, 24 (X1=K, X2=F; or X1=Y, X2=Y), 25 or 26.
4. The isolated monoclonal antibody or the antigen-binding portion thereof of claim 1, wherein the heavy chain variable region and the light chain variable region comprise amino acid sequences having at least 85%, 86%, 87%, 88%, 89%, 90%, 91%, 92%, 93%, 94%, 95%, 96%, 97%, 98%, 99% or 100% identity to (1) SEQ ID NOs: 19 and 23, respectively; (2) SEQ ID NOs: 20 (X1=A) and 24 (X1=K, X2=F), respectively; (3) SEQ ID NOs: 20 (X1=S) and 24 (X1=K, X2=F), respectively; (4) SEQ ID NOs: 20 (X1=A) and 24 (X1=Y, X2=Y), respectively; (5) SEQ ID NOs: 20 (X1=S) and 24 (X1=Y, X2=Y), respectively; (6) SEQ ID NOs: 21 and 25, respectively; or (7) SEQ ID NOs: 22 and 26, respectively.
5. The isolated monoclonal antibody or the antigen-binding portion thereof of claim 1, comprising a heavy chain constant region having an amino acid sequence of SEQ ID NOs: 27 or 28, linked to the heavy chain variable region, and a light chain constant region having an amino acid sequence of SEQ ID NO: 29, linked to the light chain variable region.
6. The isolated monoclonal antibody or the antigen-binding portion thereof of claim 1, which (a) binds human CD40; (b) binds monkey CD40; (c) blocks CD40-CD40L interaction; and/or (c) activates CD40 signaling.
7. The isolated monoclonal antibody or the antigen-binding portion thereof of claim 1, which is a mouse, chimeric or humanized antibody.
8. The isolated monoclonal antibody or the antigen-binding portion thereof of claim 1, which is an IgG1, IgG2 or IgG4 isotype.

9. A nucleotide encoding the isolated monoclonal antibody or the antigen-binding portion thereof of any one of claims 1 to 8.
10. An expression vector containing the nucleotide of claim 9.
11. A host cell containing the nucleotide of claim 9 or the expression vector of claim 10.
12. A pharmaceutical composition comprising the antibody or the antigen-binding portion thereof of any one of claims 1 to 8, and a pharmaceutically acceptable carrier.
13. The pharmaceutical composition of claim 12, further comprising an anti-tumor agent and/or a cytokine.
14. A method for treating a cancer in a subject in need thereof, comprising administering to the subject a therapeutically effective amount of the pharmaceutical composition of claim 12 or 13.
15. The method of claim 14, wherein the cancer is a solid or non-solid tumor.
16. The method of claim 15, wherein the cancer disease is selected from the group consisting of B cell lymphoma, chronic lymphocytic leukemia, multiple myeloma, melanoma, colon adenocarcinoma, pancreas cancer, colon cancer, gastric intestine cancer, prostate cancer, bladder cancer, kidney cancer, ovary cancer, cervix cancer, breast cancer, lung cancer, and nasopharynx cancer.
17. A method for treating an infectious disease in a subject in need thereof, comprising administering to the subject a therapeutically effective amount of the pharmaceutical composition of claim 12.
18. A method for treating an autoimmune disease in a subject in need thereof, comprising administering to the subject a therapeutically effective amount of the pharmaceutical composition of claim 12.

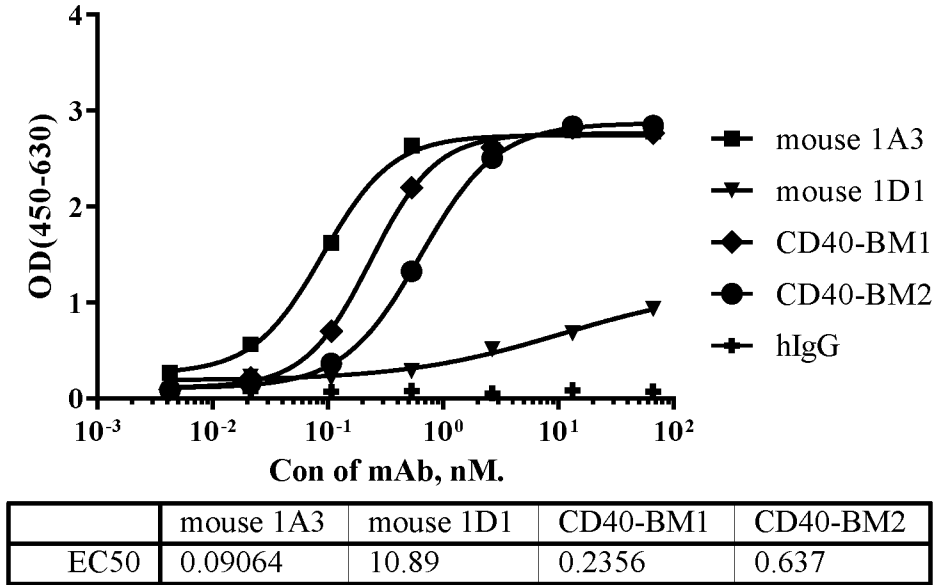


FIG. 1A

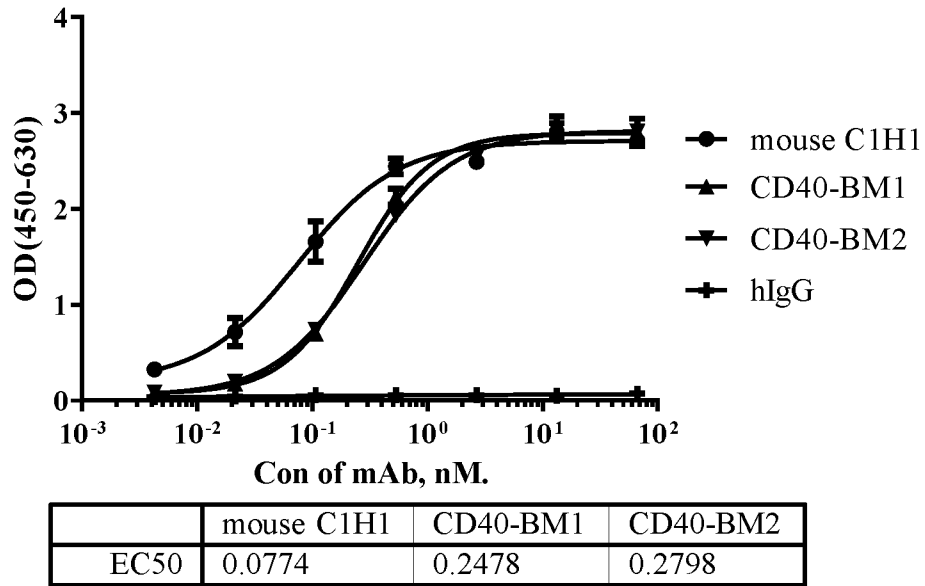


FIG. 1B

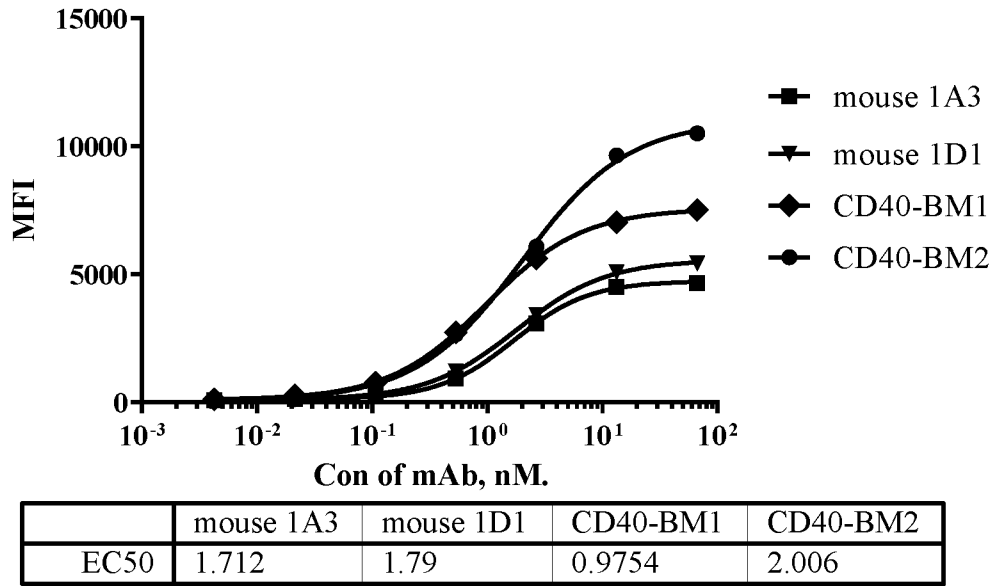


FIG. 2A

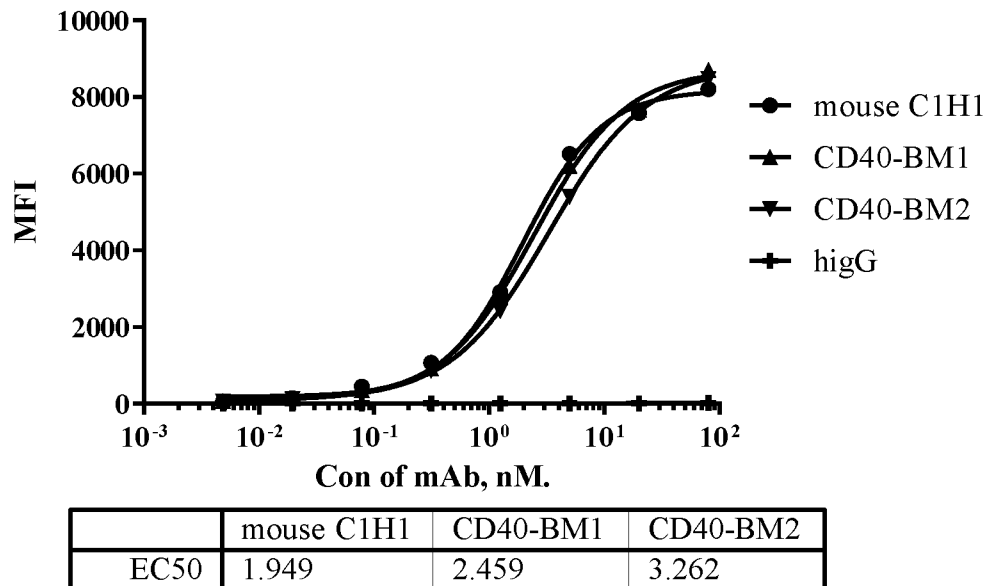
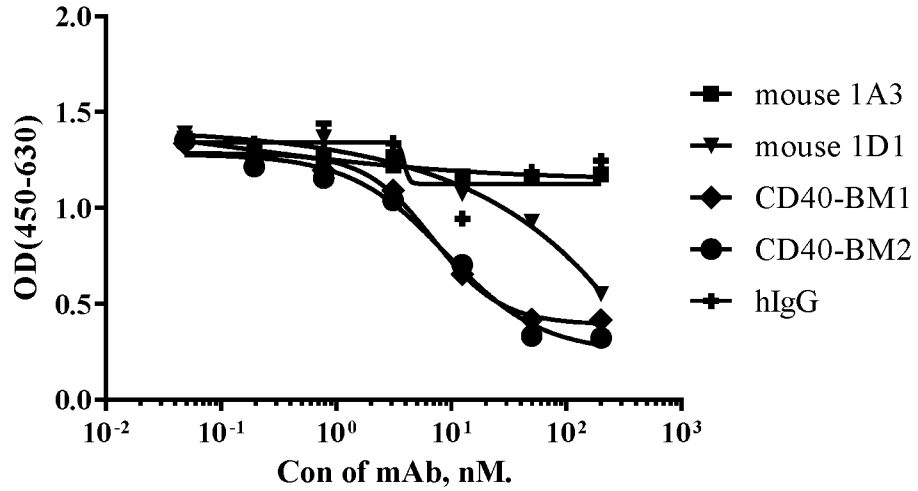
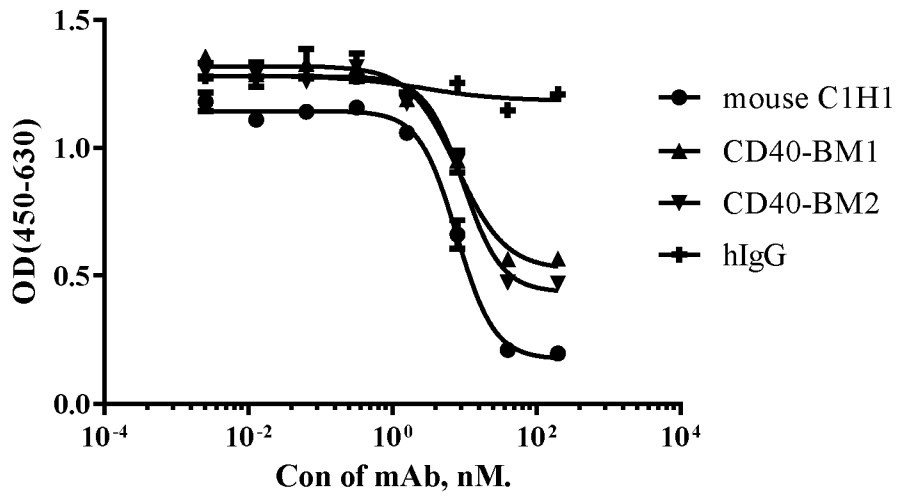


FIG. 2B



	mouse 1A3	mouse 1D1	CD40-BM1	CD40-BM2
IC50	0.1425	~ 362008128	6.946	9

FIG. 3A



	mouse C1H1	CD40-BM1	CD40-BM2
IC50	7.829	7.908	9.698

FIG. 3B

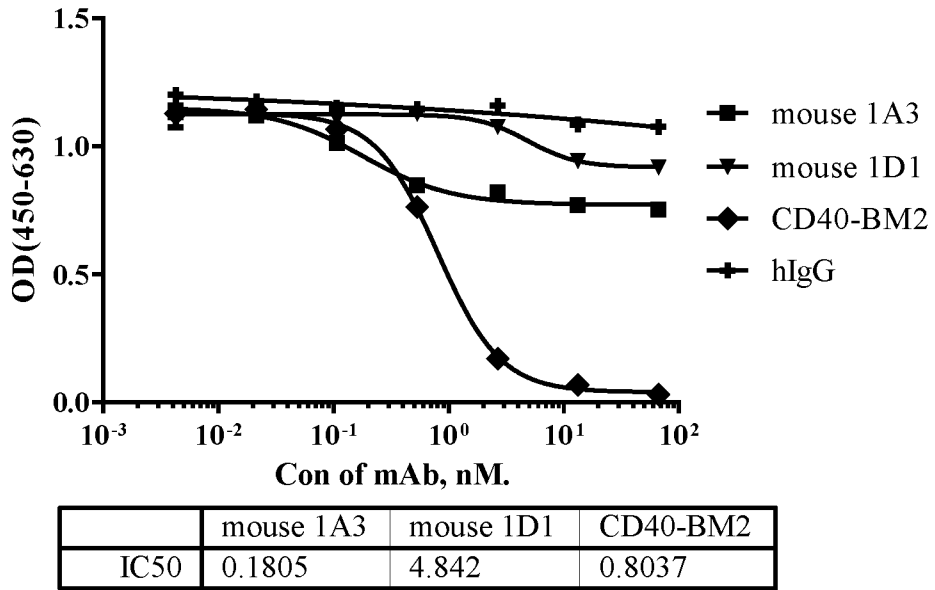


FIG. 4A

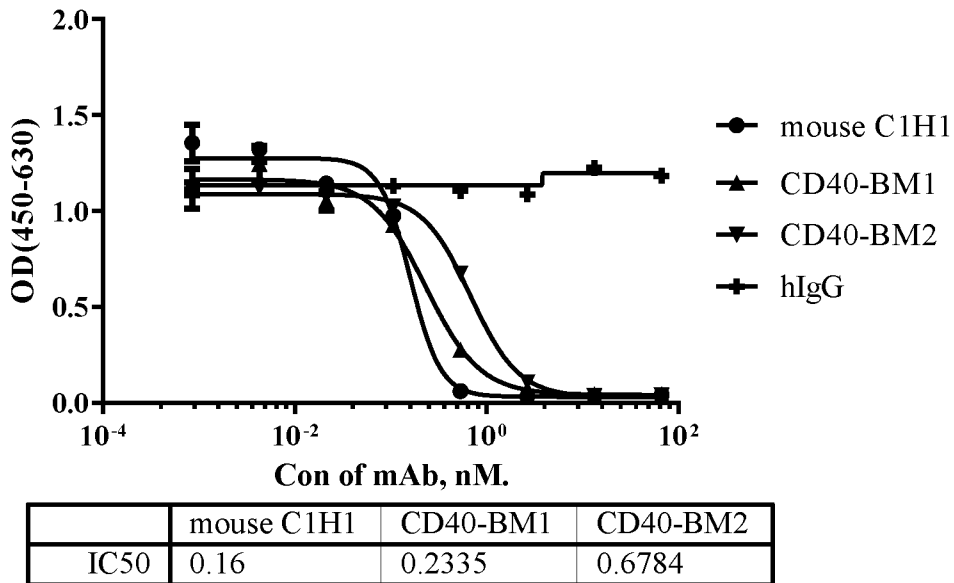


FIG. 4B

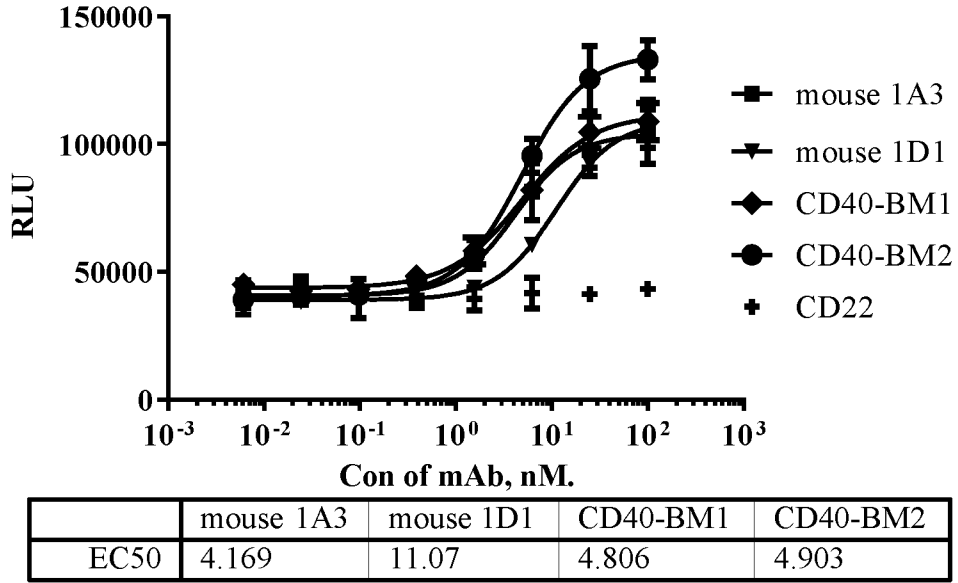


FIG. 5A

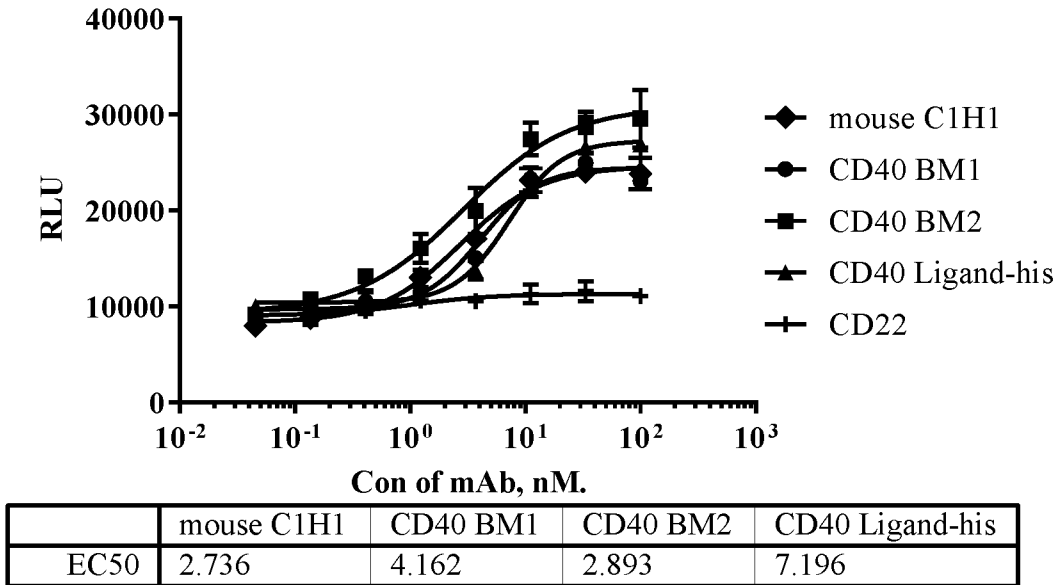
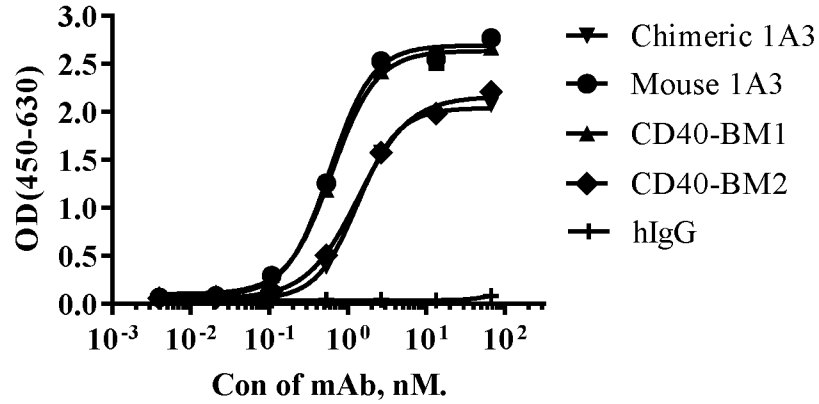
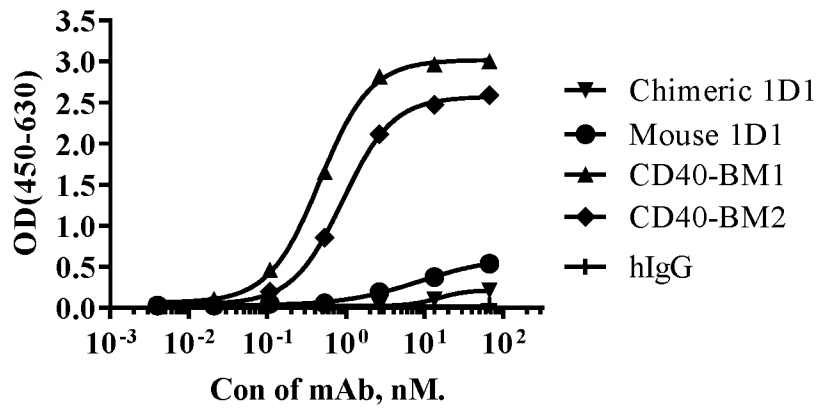


FIG. 5B



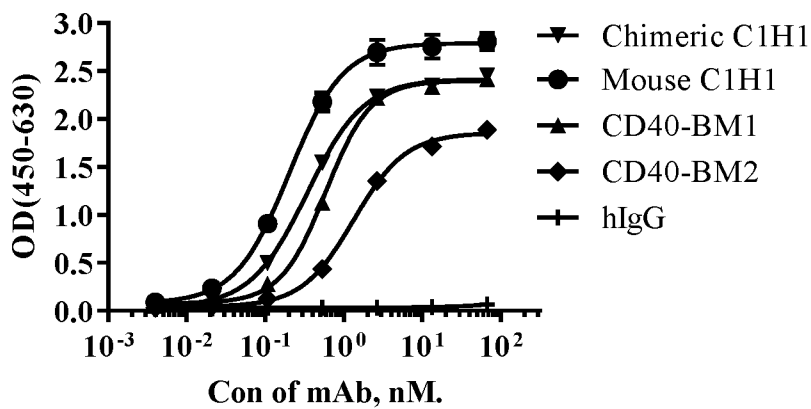
	Chimeric 1A3	Mouse 1A3	CD40-BM1	CD40-BM2
EC50	1.332	0.5917	0.6279	1.362

FIG. 6A



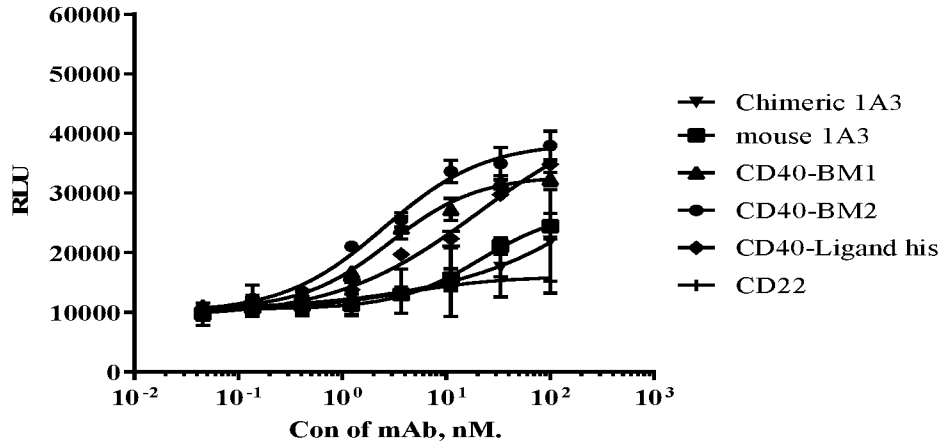
	Chimeric 1D1	Mouse 1D1	CD40-BM1	CD40-BM2
EC50	14.97	8.703	0.463	0.9092

FIG. 6B



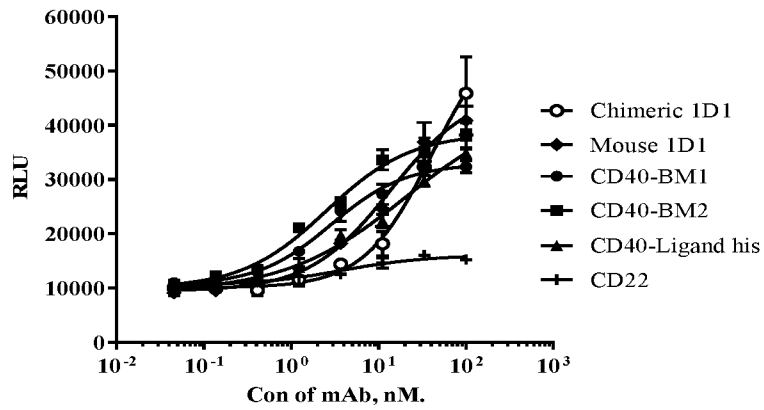
	Chimeric C1H1	Mouse C1H1	CD40-BM1	CD40-BM2
EC50	0.3426	0.2029	0.59	1.344

FIG. 6C



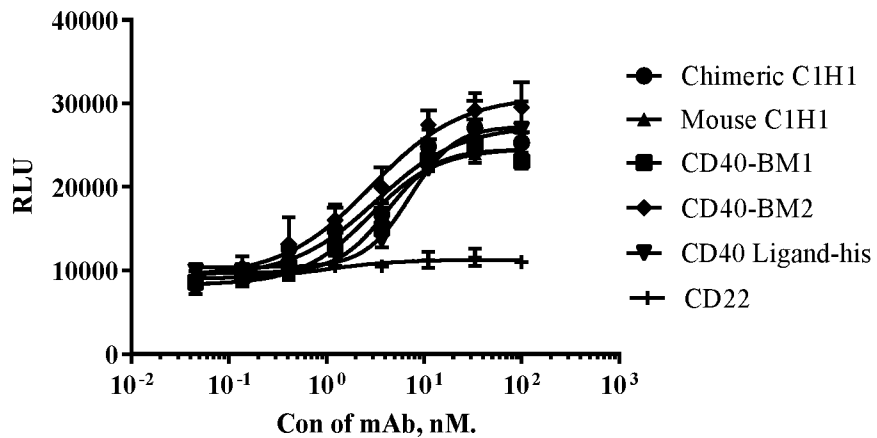
	mouse 1A3	CD40-BM1	CD40-BM2	CD40-Ligand his
EC50	21.46	2.868	2.535	20.56

FIG.7A



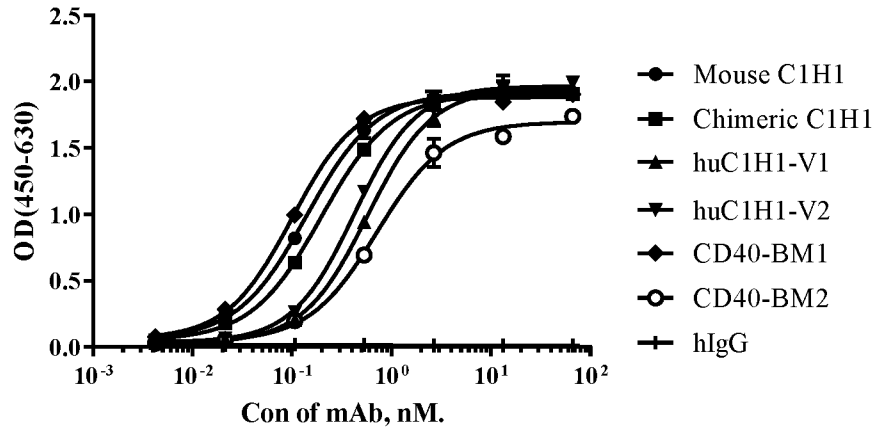
	Chimeric 1D1	Mouse 1D1	CD40-BM1	CD40-BM2	CD40-Ligand his
EC50	40.63	13.65	2.868	2.535	20.56

FIG.7B



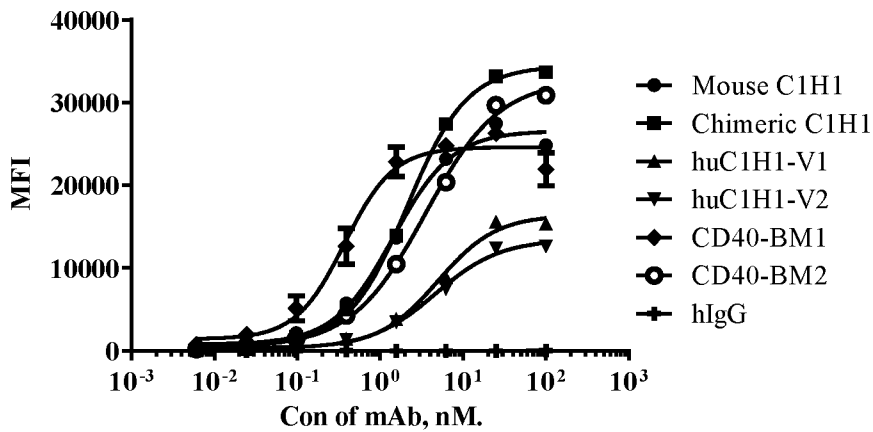
	Chimeric C1H1	Mouse C1H1	CD40-BM1	CD40-BM2	CD40 Ligand-his
EC50	3.222	2.736	4.162	2.893	7.196

FIG.7C



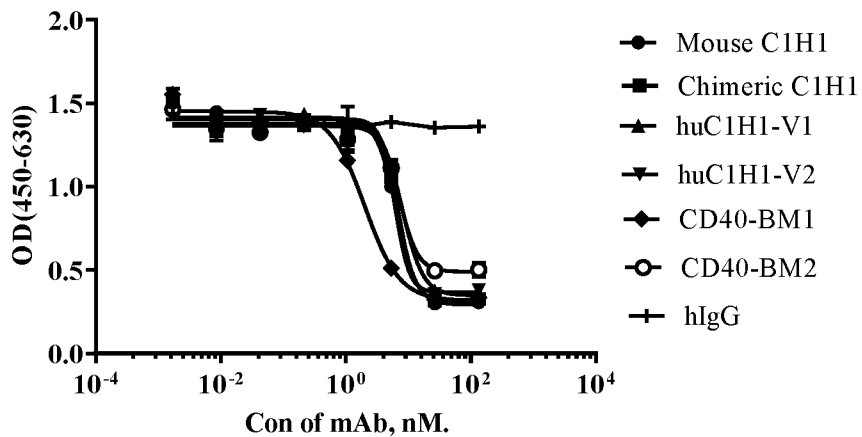
	Mouse C1H1	Chimeric C1H1	huC1H1-V1	huC1H1-V2	CD40-BM1	CD40-BM2
EC50	0.1407	0.1977	0.5742	0.4245	0.1005	0.7196

FIG. 8



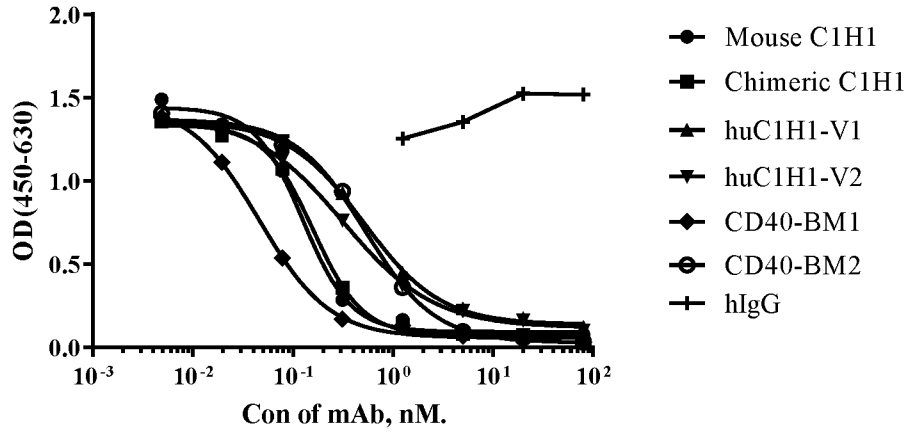
	Mouse C1H1	Chimeric C1H1	huC1H1-V1	huC1H1-V2	CD40-BM1	CD40-BM2
EC50	1.437	2.131	4.95	4.637	0.3749	3.517

FIG. 9



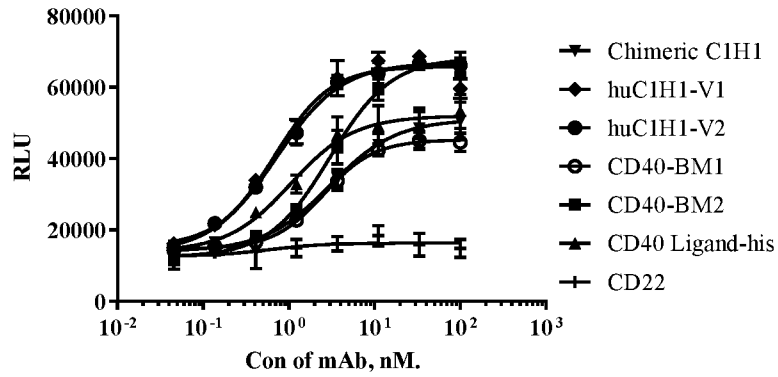
	Mouse C1H1	Chimeric C1H1	huC1H1-V1	huC1H1-V2	CD40-BM1	CD40-BM2
IC50	6.465	6.968	7.91	6.102	2.009	7.218

FIG. 10



	Mouse C1H1	Chimeric C1H1	huC1H1-V1	huC1H1-V2	CD40-BM1	CD40-BM2
IC50	0.1252	0.1596	0.5131	0.3177	0.04612	0.5291

FIG. 11



	Chimeric C1H1	huC1H1-V1	huC1H1-V2	CD40-BM1	CD40-BM2	CD40 Ligand-his
EC50	3.044	0.679	0.7168	2.546	3.025	1.054

FIG. 12

INTERNATIONAL SEARCH REPORT

International application No.

PCT/CN2021/084013

A. CLASSIFICATION OF SUBJECT MATTER		
A61K 39/00(2006.01)i; C07K 16/28(2006.01)i		
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)		
A61K,C07K		
Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched		
Electronic data base consulted during the international search (name of data base and, where practicable, search terms used)		
STN, Genbank, EMBL, VEN, CNKI, CNABS, PubMed, ISI Web of Knowledge,;anti-CD40 antibody, antibody, CD40, CD40L,blockade,SEQ ID NO:1-29		
C. DOCUMENTS CONSIDERED TO BE RELEVANT		
Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
A	US 2018066053 A1 (CELLDEX THERAPEUTICS, INC.) 08 March 2018 (2018-03-08) the whole document	1-18
A	WO 2018088850 A2 (DINONA) 17 May 2018 (2018-05-17) the whole document	1-18
A	WO 2018222019 A1 (SEOUL NATIONAL UNIVERSITY R&DB FOUNDATION) 06 December 2018 (2018-12-06) the whole document	1-18
A	WO 2018219327 A1 (JIANGSU HENGRUI MEDICINE CO., LTD.et al.) 06 December 2018 (2018-12-06) the whole document	1-18
A	VISVANATHAN, S. et al. "Effects of BI 655064, an antagonistic anti-CD40 antibody, on clinical and biomarker variables in patients with active rheumatoid arthritis: a randomised, double-blind, placebo-controlled, phase IIa study" <i>Ann Rheum Dis</i> , Vol. 78, 22 March 2019 (2019-03-22), pages754-760	1-18
<input checked="" type="checkbox"/> Further documents are listed in the continuation of Box C. <input checked="" type="checkbox"/> 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
08 June 2021		02 July 2021
Name and mailing address of the ISA/CN		Authorized officer
National Intellectual Property Administration, PRC 6, Xitucheng Rd., Jimen Bridge, Haidian District, Beijing 100088 China		LIU,Xinlei
Facsimile No. (86-10)62019451		Telephone No. 86-(10)-53962097

INTERNATIONAL SEARCH REPORT

International application No.

PCT/CN2021/084013

C. DOCUMENTS CONSIDERED TO BE RELEVANT		
Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
A	KOSAKA, A. et al. "Combination of an agonistic anti-CD40 monoclonal antibody and the COX-2 inhibitor celecoxib induces anti-glioma effects by promotion of type-1 immunity in myeloid cells and T-cells" <i>Cancer Immunol Immunother</i> , Vol. 63, No. 8, 31 August 2014 (2014-08-31), pages847-857	1-18
.....		

Box No. I Nucleotide and/or amino acid sequence(s) (Continuation of item 1.c of the first sheet)

1. With regard to any nucleotide and/or amino acid sequence disclosed in the international application, the international search was carried out on the basis of a sequence listing:
 - a. forming part of the international application as filed:
 - in the form of an Annex C/ST.25 text file.
 - on paper or in the form of an image file.
 - b. furnished together with the international application under PCT Rule 13ter.1(a) for the purposes of international search only in the form of an Annex C/ST.25 text file.
 - c. furnished subsequent to the international filing date for the purposes of international search only:
 - in the form of an Annex C/ST.25 text file (Rule 13ter.1(a)).
 - on paper or in the form of an image file (Rule 13ter.1(b) and Administrative Instructions, Section 713).
2. In addition, in the case that more than one version or copy of a sequence listing has been filed or furnished, the required statements that the information in the subsequent or additional copies is identical to that forming part of the application as filed or does not go beyond the application as filed, as appropriate, were furnished.
3. Additional comments:

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.: **14-18**
because they relate to subject matter not required to be searched by this Authority, namely:
 - [1] Claims 14-18 direct to a method of treating diseases, and therefore do not warrant an international search according to the criteria set out in PCT Rule 39.1(iv). An international search is still carried out on the basis on the use of the isolated monoclonal antibody for the manufacturing of a medicament for the treatment of diseases.
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.:
because they are dependent claims and are not drafted in accordance with the second and third sentences of Rule 6.4(a).

INTERNATIONAL SEARCH REPORT
Information on patent family members

International application No.

PCT/CN2021/084013

Patent document cited in search report			Publication date (day/month/year)	Patent family member(s)			Publication date (day/month/year)
US	2018066053	A1	08 March 2018	SG	11201808821W	A	29 November 2018
				KR	20180133493	A	14 December 2018
				EA	201892362	A1	30 April 2019
				US	2019322743	A1	24 October 2019
				EP	3445783	A2	27 February 2019
				SG	10201913248V	A	27 February 2020
				AU	2017252527	A1	08 November 2018
				CN	109071665	A	21 December 2018
				US	10865244	B2	15 December 2020
				JP	2019521645	A	08 August 2019
				MX	2018012757	A	24 June 2019
				US	2020377606	A1	03 December 2020
				CA	3021328	A1	26 October 2017
				IL	262269	D0	29 November 2018
				WO	2017184619	A2	26 October 2017
				BR	112018071307	A2	26 February 2019
				US	2020369768	A1	26 November 2020
US	10633444	B2	28 April 2020				
WO	2017184619	A3	21 December 2017				
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WO	2018088850	A2	17 May 2018	KR	102019033	B1	06 September 2019
				KR	20180053255	A	21 May 2018
				CN	110546164	A	06 December 2019
				WO	2018088850	A3	26 July 2018
<hr/>							
WO	2018222019	A1	06 December 2018	US	2020291123	A1	17 September 2020
				KR	20180131989	A	11 December 2018
				CN	111344304	A	26 June 2020
				EP	3632933	A4	03 March 2021
				KR	102198998	B1	07 January 2021
				EP	3632933	A1	08 April 2020
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WO	2018219327	A1	06 December 2018	CA	3064298	A1	06 December 2018
				JP	2020521504	A	27 July 2020
				TW	201902925	A	16 January 2019
				US	2020148778	A1	14 May 2020
				AU	2018278051	A1	30 January 2020
				TW	I699376	B	21 July 2020
				BR	112019025048	A2	30 June 2020
				EP	3632932	A1	08 April 2020
				MX	2019014375	A	23 January 2020
				KR	20200012920	A	05 February 2020
				CN	110267989	A	20 September 2019