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(54) **Titre : THERAPIES A BASE DE CELLULES EFFECTRICES IMMUNITAIRES DOTEES D'UNE EFFICACITE ACCRUE**
(54) **Title: IMMUNE EFFECTOR CELL THERAPIES WITH ENHANCED EFFICACY**

(57) **Abrégé/Abstract:**
Provided are the use of LSD1 inhibitors in connection with use and manufacture of immune effector cells (e.g., T cells, NK cells), e.g., engineered to express a chimeric antigen receptor (CAR), to treat a subject having a disease, e.g., a disease associated with expression of a tumor antigen.

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(54) **Title**: IMMUNE EFFECTOR CELL THERAPIES WITH ENHANCED EFFICACY

(57) **Abstract**: Provided are the use of LSD1 inhibitors in connection with use and manufacture of immune effector cells (e.g., T cells, NK cells), e.g., engineered to express a chimeric antigen receptor (CAR), to treat a subject having a disease, e.g., a disease associated with expression of a tumor antigen.

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JUMBO APPLICATIONS/PATENTS

THIS SECTION OF THE APPLICATION/PATENT CONTAINS MORE THAN ONE VOLUME

THIS IS VOLUME 1 OF 2
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Immune Effector Cell Therapies with Enhanced Efficacy

RELATED APPLICATIONS

This application claims priority to PCT Patent Application Number PCT/CN2015/099882, filed December 30, 2015, the entire contents of which are incorporated herein by reference.

SEQUENCE LISTING

The instant application contains a Sequence Listing which has been submitted electronically in ASCII format and is hereby incorporated by reference in its entirety. Said ASCII copy, created on December 29, 2016, is named N2067-7105WO3_SL.txt and is 1,118,768 bytes in size.

FIELD OF THE INVENTION

The present invention relates generally to the use LSD1 inhibitors in connection with use and manufacture of immune effector cells (e.g., T cells, NK cells), e.g., engineered to express a chimeric antigen receptor (CAR), to treat a subject having a disease, e.g., a disease associated with expression of a tumor antigen.

BACKGROUND OF THE INVENTION

Adoptive cell transfer (ACT) therapy, for example, with T-cells transduced with Chimeric Antigen Receptors (CARs), has shown promise in cancer trials. There is a medical need for T cell therapies, especially CAR T cell therapies with improved efficacy.

SUMMARY OF THE INVENTION

Methods and compositions disclosed herein are directed to the use of an inhibitor of Lysine-specific demethylase 1 (LSD1) in connection with the use and/or manufacture of immune effector cells (e.g., T cells or NK cells), for example, immune effector cells engineered to express a Chimeric Antigen Receptor (CAR), to treat a disease, e.g., a disease associated with expression of a cancer associated antigen (or tumor marker).

It has been discovered that inhibition of LSD1 is effective in improving the function of immune effector cells, e.g., engineered to express a CAR molecule, e.g., as described herein, and can be combined with T cell, e.g., CAR T cell, therapy and/or manufacturing.

While not wishing to be bound by theory, it is believed that contacting a population of immune effector cells, e.g., engineered to express a CAR molecule, e.g., as described herein, with an LSD1 inhibitor is accompanied by an increase in the proportion of naive T cells (e.g., CD45RA+CD62L+ T cells, e.g., T_{SCM} cells), at least transiently, relative to an uncontacted population, for example, when such cells are stimulated (e.g., with anti-CD3 and/or anti-CD28 stimulation) or induced to proliferate (e.g., in response to antigen recognition, e.g., antigen recognition through a CAR molecule, e.g., as described herein).

While not wishing to be bound by theory, it is believed that contacting a population of immune effector cells, e.g., engineered to express a CAR molecule, e.g., as described herein, with an LSD1 inhibitor is accompanied by an increase in the number of naive T cells (e.g., CD45RA+CD62L+ T cells, e.g., T_{SCM} cells), at least transiently, relative to an uncontacted population, for example, when such cells are stimulated (e.g., with anti-CD3 and/or anti-CD28 stimulation) or induced to proliferate (e.g., in response to antigen recognition, e.g., antigen recognition through a CAR).

While not wishing to be bound by theory, it is believed that contacting a population of immune effector cells, e.g., engineered to express a CAR molecule, e.g., as described herein, with an LSD1 inhibitor is accompanied by a decrease in the number of T_{EM} cells, at least transiently, relative to an uncontacted population, for example, when such cells are stimulated (e.g., with anti-CD3 and/or anti-CD28 stimulation) or induced to proliferate (e.g., in response to antigen recognition, e.g., antigen recognition through a CAR).

While not wishing to be bound by theory, it is believed that contacting a population of immune effector cells, e.g., engineered to express a CAR molecule, e.g., as described herein, with an LSD1 inhibitor is accompanied by a decrease in the proportion of T_{EM} cells, at least transiently, relative to an uncontacted population, for example, when such cells are stimulated (e.g., with anti-CD3 and/or anti-CD28 stimulation) or induced to proliferate (e.g., in response to antigen recognition, e.g., antigen recognition through a CAR molecule, e.g., as described herein).

While not wishing to be bound by theory, it is believed that contacting a population of immune effector cells, e.g., engineered to express a CAR molecule, e.g., as described herein, with an LSD1 inhibitor is accompanied by an increase in the production of cytokines (e.g., IFNg and/or IL-2) from said population of immune effector cells, at least transiently, relative to an uncontacted population, for example, when such cells are stimulated (e.g., with anti-CD3 and/or anti-CD28 stimulation) or induced to

proliferate (e.g., in response to antigen recognition, e.g., antigen recognition through a CAR molecule, e.g., as described herein).

While not wishing to be bound by theory, it is believed that contacting a population of immune effector cells, e.g., engineered to express a CAR molecule, e.g., as described herein, with an LSD1 inhibitor is accompanied by a decrease in the proportion of PD-1 positive immune effector cells, at least transiently, relative to an uncontacted population, for example, when such cells are stimulated (e.g., with anti-CD3 and/or anti-CD28 stimulation) or induced to proliferate (e.g., in response to antigen recognition, e.g., antigen recognition through a CAR molecule, e.g., as described herein).

While not wishing to be bound by theory, it is believed that contacting a population of immune effector cells, e.g., engineered to express a CAR molecule, e.g., as described herein, with an LSD1 inhibitor is accompanied by an increase in the ratio of PD-1 negative immune effector cells / PD-1 positive immune effector cells, at least transiently, relative to an uncontacted population, for example, when such cells are stimulated (e.g., with anti-CD3 and/or anti-CD28 stimulation) or induced to proliferate (e.g., in response to antigen recognition, e.g., antigen recognition through a CAR molecule, e.g., as described herein).

While not wishing to be bound by theory, it is believed that contacting a population of immune effector cells, e.g., engineered to express a CAR molecule, e.g., as described herein, with an LSD1 inhibitor is accompanied by a decrease in the proportion of PD-1+/Lag3+/Tim3+ immune effector cells, at least transiently, relative to an uncontacted population, for example, when such cells are stimulated (e.g., with anti-CD3 and/or anti-CD28 stimulation) or induced to proliferate (e.g., in response to antigen recognition, e.g., antigen recognition through a CAR molecule, e.g., as described herein).

While not wishing to be bound by theory, it is believed that contacting a population of immune effector cells, e.g., engineered to express a CAR molecule, e.g., as described herein, with an LSD1 inhibitor is accompanied by an increase in the ratio of PD-1-/Lag3-/Tim3- immune effector cells to PD-1+/Lag3+/Tim3+ immune effector cells, at least transiently, relative to an uncontacted population, for example, when such cells are stimulated (e.g., with anti-CD3 and/or anti-CD28 stimulation) or induced to proliferate (e.g., in response to antigen recognition, e.g., antigen recognition through a CAR molecule, e.g., as described herein).

While not wishing to be bound by theory, it is believed that contacting a population of immune effector cells, e.g., engineered to express a CAR molecule, e.g., as described herein, with an LSD1 inhibitor is accompanied by an increase in the proliferation of the immune effector cells, at least transiently, relative to an uncontacted population, for example, when such cells are stimulated (e.g., with anti-CD3 and/or anti-CD28 stimulation) or induced to proliferate (e.g., in response to antigen recognition, e.g., antigen recognition through a CAR molecule, e.g., as described herein). Again, without being bound by theory, it is believed that T cells can be exhausted by, for example, stimulation with CD3/CD28 stimulation or antigen stimulation (e.g., by induced signaling through a CAR). Such exhaustion can lead to decreased efficacy or function (e.g., decreased proliferation, persistence, and/or anti-tumor efficacy) of such immune effector cells. As described herein, the inventors have discovered that inhibiting LSD1 increases the proliferation and/or survival of more naive T cells, e.g., T_{SCM} cells, which in turn have better efficacy and function. Thus, embodiments of the invention are based, at least in part, on the recognition that LSD1 inhibition, is associated with improved T cell function and/or phenotype.

In an embodiment these approaches can be used to optimize the performance of immune effector cells, e.g., T cells, in the subject. While not wishing to be bound by theory, it is believed that, in an embodiment, the performance of endogenous, non-modified immune effector cells, e.g., T cells, is improved. While not wishing to be bound by theory, it is believed that, in an embodiment, the performance of immune effector cells, e.g., T cells, harvested (e.g., from a subject administered an LSD1 inhibitor) and engineered to express a CAR molecule, e.g., as described herein, is improved. In other embodiments, a population of immune effector cells, e.g., T cells, which have been, or will be engineered to express a CAR molecule, e.g., as described herein, can be treated ex vivo by contact with an amount of an LSD1 inhibitor that improves the number or ratio of naive T cells, e.g., T_{SCM} cells, and/or improves the number or ratio of PD-1 negative, e.g., PD-1-/Tim3-/Lag3- T cells, relative to an uncontacted population.

In an embodiment, the LSD1 inhibitor is administered for an amount of time sufficient to decrease the proportion of PD-1 positive T cells, increase the proportion of PD-1 negative T cells, or increase the ratio of PD-1 negative T cells/PD-1 positive T cells, in the peripheral blood of the subject (or in a preparation of T cells isolated from the subject).

In an embodiment, the method of treating, e.g., promoting an immune response in, a subject, e.g., a human subject, comprises inhibiting a negative immune response mediated by the engagement of PD-1 with PD-L1 or PD-L2, e.g., relative to a T cell not contacted with an LSD1 inhibitor.

In an embodiment, the method of treating, e.g., promoting an immune response in, a subject, e.g., a human subject, comprises increasing the number of T cells capable of proliferation, e.g., relative to a T cell not contacted with an LSD1 inhibitor.

In an embodiment, the method of treating, e.g., promoting an immune response in, a subject, e.g., a human subject, comprises increasing the number of T cells capable of cytotoxic function, secreting cytokines, or activation, e.g., relative to a T cell not contacted with an LSD1 inhibitor.

In an embodiment, the method of treating, e.g., promoting an immune response in, a subject, e.g., a human subject, comprises increasing the amount of cytokine secretion (e.g., interferon gamma (IFN-g) and/or interleukin 2 (IL-2)) in response to stimulation and/or activation of the T cell, e.g., relative to a T cell not contacted with an LSD1 inhibitor.

In an embodiment, the LSD1 inhibitor is administered (in vivo or ex vivo) prior to administration of immune effector cells, e.g., T cells to be engineered to express a CAR molecule, e.g., as described herein, (e.g., prior to or after harvest of the immune effector cells) for an amount of time sufficient for one or more of the following to occur:

1) an increase in the proportion of naive T cells, e.g., T_{SCM} cells;

2) an increase in the number of naive T cells, e.g., T_{SCM} cells;

3) a decrease in the number of T_{EM} cells;

4) a decrease in the proportion of T_{EM} cells;

5) an increase in the proportion of CD45RA+CD62L+ T cells;

6) an increase in the number of CD45RA+CD62L+ T cells;

7) an increase in the proportion of CD45RA+CCR7+ T cells;

8) an increase in the number of CD45RA+CCR7+ T cells;

9) a decrease in the proportion of PD-1 positive immune effector cells;

10) an increase in the ratio of PD-1 negative immune effector cells / PD-1 positive immune effector cells;

11) a decrease in the proportion of PD-1+/Lag3+/Tim3+ immune effector cells;

12) an increase in the ratio of PD-1-/Lag3-/Tim3- immune effector cells to PD-1+/Lag3+/Tim3+ immune effector cells;

13) an increase in the proliferation of the immune effector cells;

14 an increase in the production of cytokines (e.g., IFNg and/or IL-2) from said population of immune effector cells; or

15) a combination of two, three, four, five, six, seven, eight, nine, ten, eleven, twelve, thirteen, or more (e.g., all) of the above;

and wherein 1), 2), 3), 4), 5), 6), 7), 8), 9), 10), 11), 12) 13), 14) or 15) occurs e.g., at least transiently, e.g., permanently, e.g., as compared to a non-treated subject. In an embodiment, the immune effector cell, e.g., T cell, to be engineered to express a CAR molecule, e.g., as described herein, is harvested at least 5, 10, 15, 20, 25, 30, 35, 40, 45, 50, 55, or 60 days after initiation, or completion, of dosing with the LSD1 inhibitor.

In an embodiment, the LSD1 inhibitor is administered to a subject prior to harvest of immune effector cells, e.g., T cells to be engineered to express an CAR molecule, e.g., as described herein, for an amount of time sufficient for one or more of the following to occur e.g., to occur in the harvested cells or in the engineered cells (or in non-harvested cells, or in both):

1) an increase in the proportion of naive T cells, e.g., T_{SCM} cells;

2) an increase in the number of naive T cells, e.g., T_{SCM} cells;

3) a decrease in the number of T_{EM} cells;

4) a decrease in the proportion of T_{EM} cells;

5) an increase in the proportion of CD45RA+CD62L+ T cells;

6) an increase in the number of CD45RA+CD62L+ T cells;

7) an increase in the proportion of CD45RA+CCR7+ T cells;

8) an increase in the number of CD45RA+CCR7+ T cells;

9) a decrease in the proportion of PD-1 positive immune effector cells;

10) an increase in the ratio of PD-1 negative immune effector cells / PD-1 positive immune effector cells;

11) a decrease in the proportion of PD-1+/Lag3+/Tim3+ immune effector cells;

12) an increase in the ratio of PD-1-/Lag3-/Tim3- immune effector cells to PD-1+/Lag3+/Tim3+ immune effector cells;

13) an increase in the proliferation of the immune effector cells;

14 an increase in the production of cytokines (e.g., IFNg and/or IL-2) from said population of immune effector cells; or

15) a combination of two, three, four, five, six, seven, eight, nine, ten, eleven, twelve, thirteen, or more (e.g., all) of the above;

and wherein 1), 2), 3), 4), 5), 6), 7), 8), 9), 10), 11), 12) 13), 14) or 15) occurs e.g., at least transiently, e.g., permanently, e.g., as compared to a non-treated subject. In an embodiment, the immune effector cell, e.g., T cell, to be engineered to express a CAR molecule, e.g., as described herein, is harvested at least 5, 10, 15, 20, 25, 30, 35, 40, 45, 50, 55, or 60 days after initiation, or completion, of dosing with the LSD1 inhibitor.

In an embodiment, the LSD1 inhibitor is administered after harvest of immune effector cells, e.g., T cells to be engineered to express an CAR molecule, e.g., as described herein, for an amount of time sufficient for one or more of the following to occur:

1) an increase in the proportion of naive T cells, e.g., T_{SCM} cells;

2) an increase in the number of naive T cells, e.g., T_{SCM} cells;

3) a decrease in the number of T_{EM} cells;

4) a decrease in the proportion of T_{EM} cells;

5) an increase in the proportion of CD45RA+CD62L+ T cells;

6) an increase in the number of CD45RA+CD62L+ T cells;

7) an increase in the proportion of CD45RA+CCR7+ T cells;

8) an increase in the number of CD45RA+CCR7+ T cells;

9) a decrease in the proportion of PD-1 positive immune effector cells;

10) an increase in the ratio of PD-1 negative immune effector cells / PD-1 positive immune effector cells;

11) a decrease in the proportion of PD-1+/Lag3+/Tim3+ immune effector cells;

12) an increase in the ratio of PD-1-/Lag3-/Tim3- immune effector cells to PD-1+/Lag3+/Tim3+ immune effector cells;

13) an increase in the proliferation of the immune effector cells;

14 an increase in the production of cytokines (e.g., IFNg and/or IL-2) from said population of immune effector cells; or

15) a combination of two, three, four, five, six, seven, eight, nine, ten, eleven, twelve, thirteen, or more (e.g., all) of the above;

and wherein 1), 2), 3), 4), 5), 6), 7), 8), 9), 10), 11), 12) 13), 14) or 15) occurs e.g., at least transiently, e.g., permanently, e.g., as compared to a non-treated subject.

In an embodiment, the LSD1 inhibitor is administered after administration of immune effector cells, e.g., T cells to be engineered to express an CAR molecule, e.g., as described herein, for an amount of time sufficient for one or more of the following to occur:

1) an increase in the proportion of naive T cells, e.g., T_{SCM} cells;

2) an increase in the number of naive T cells, e.g., T_{SCM} cells;

3) a decrease in the number of T_{EM} cells;

4) a decrease in the proportion of T_{EM} cells;

5) an increase in the proportion of CD45RA+CD62L+ T cells;

6) an increase in the number of CD45RA+CD62L+ T cells;

7) an increase in the proportion of CD45RA+CCR7+ T cells;

8) an increase in the number of CD45RA+CCR7+ T cells;

9) a decrease in the proportion of PD-1 positive immune effector cells;

10) an increase in the ratio of PD-1 negative immune effector cells / PD-1 positive immune effector cells;

11) a decrease in the proportion of PD-1+/Lag3+/Tim3+ immune effector cells;

12) an increase in the ratio of PD-1-/Lag3-/Tim3- immune effector cells to PD-1+/Lag3+/Tim3+ immune effector cells;

13) an increase in the proliferation of the immune effector cells;

14 an increase in the production of cytokines (e.g., IFNg and/or IL-2) from said population of immune effector cells; or

15) a combination of two, three, four, five, six, seven, eight, nine, ten, eleven, twelve, thirteen, or more (e.g., all) of the above;

and wherein 1), 2), 3), 4), 5), 6), 7), 8), 9), 10), 11), 12) 13), 14) or 15) occurs e.g., at least transiently, e.g., permanently, e.g., as compared to a non-treated subject.

In an embodiment, LSD1 inhibitor is administered to immune effector cells, e.g., T cells, which have, or will be engineered to express a CAR molecule, e.g., as described herein, ex vivo for an amount of time sufficient for one or more of the following to occur:

1) an increase in the proportion of naive T cells, e.g., T_{SCM} cells;

2) an increase in the number of naive T cells, e.g., T_{SCM} cells;

3) a decrease in the number of T_{EM} cells;

4) a decrease in the proportion of T_{EM} cells;

5) an increase in the proportion of CD45RA+CD62L+ T cells;

6) an increase in the number of CD45RA+CD62L+ T cells;

7) an increase in the proportion of CD45RA+CCR7+ T cells;

8) an increase in the number of CD45RA+CCR7+ T cells;

9) a decrease in the proportion of PD-1 positive immune effector cells;

10) an increase in the ratio of PD-1 negative immune effector cells / PD-1 positive immune effector cells;

11) a decrease in the proportion of PD-1+/Lag3+/Tim3+ immune effector cells;

12) an increase in the ratio of PD-1-/Lag3-/Tim3- immune effector cells to PD-1+/Lag3+/Tim3+ immune effector cells;

13) an increase in the proliferation of the immune effector cells;

14 an increase in the production of cytokines (e.g., IFNg and/or IL-2) from said population of immune effector cells; or

15) a combination of two, three, four, five, six, seven, eight, nine, ten, eleven, twelve, thirteen, or more (e.g., all) of the above;

and wherein 1), 2), 3), 4), 5), 6), 7), 8), 9), 10), 11), 12) 13), 14) or 15) occurs e.g., at least transiently, e.g., permanently, e.g., as compared to a non-treated cell.

Without being bound by theory, it is believed that LSD1 may also directly demethylate p53 (Nature Reviews Molecular Cell Biology 13, 297-311 (May 2012) doi:10.1038/nrm3327). Thus, in an embodiment, the compounds and methods disclosed herein may be used to inhibit demethylation of p53.

In an embodiment, the subject has cancer and the method comprises promoting the subject's immune response to the cancer. In an embodiment, the subject was selected on the basis of having cancer. In an embodiment, a cell of the cancer expresses PD-L1 or PD-L2. In an embodiment, a cell in the cancer microenvironment expresses PD-L1 or PD-L2.

In an embodiment, the cancer comprises a solid tumor. In an embodiment, the cancer is a hematological cancer. In an embodiment, the cancer is a leukemia. In an embodiment, the cancer is a chronic lymphocytic leukemia (CLL). In an embodiment, the cancer is CLL and wherein the antigen binding domain of the CAR targets CD19. In an embodiment, the cancer is melanoma.

In an embodiment, the method further comprises administering an additional treatment, e.g., a chemotherapeutic, radiation, a cellular therapy, or bone marrow transplant to the subject. In an embodiment, the method further comprises administering an additional treatment that kills T cells, e.g., radiation or cytotoxic chemotherapy. In an embodiment, the method further comprises administering to the subject an mTOR pathway inhibitor, such as vitamin E, vitamin A, an antibacterial antibiotic, an antioxidant, L- carnitine, lipoic acid, metformin, resveratrol, leptine, a non-steroid anti- inflammatory drug, or a COX inhibitor. In an embodiment, the method further comprises administering metformin to the subject. In an embodiment, the LSD1 inhibitor is administered prior to or after the initiation of the additional treatment. In an embodiment, the method further comprises administering an additional treatment for the cancer.

In an embodiment, the method further comprises administering the immune effector cell, e.g., T cell, engineered to express a CAR molecule, e.g., as described herein, in combination with another agent (in addition to the LSD1 inhibitor). In one embodiment, the agent can be a kinase inhibitor, e.g., a CDK4/6 inhibitor, a BTK inhibitor, an mTOR inhibitor, a MNK inhibitor, or a dual mTOR/PI3K kinase inhibitor, and combinations thereof.

In an embodiment, the method comprises providing an anti-tumor immunity in a mammal. In one embodiment, the cell is an autologous T cell or an autologous NK cell. In one embodiment, the cell is an allogeneic T cell or an allogeneic NK cell. In one embodiment, the mammal is a human.

In an embodiment the method comprises treating a mammal having a disease associated with expression of a cancer associated antigen or tumor marker.

In one embodiment, the method comprises administering an agent that increases the efficacy of the immune effector cell, e.g., T cell or NK cell, engineered to express a CAR molecule, e.g., as described herein, e.g., an agent described herein.

In one embodiment, the method comprises administering an agent that ameliorates one or more side effect associated with administration of a cell expressing a CAR molecule, e.g., as described herein, the immune effector cell, e.g., T cell or NK cell, engineered to express a CAR molecule, e.g., as described herein, e.g., an agent described herein.

In one embodiment, the method comprises administering an agent that treats the disease associated with a cancer associated antigen as described herein, e.g., an agent described herein.

In one embodiment, the immune effector cell, e.g., T cell or NK cell, engineered to express a CAR molecule, e.g., as described herein, expresses two or more CAR molecules and, e.g., is administered to a subject in need thereof to treat cancer.

In one embodiment, the CAR molecule is introduced into immune effector cells (e.g., T cells, NK cells), e.g., using in vitro transcription, and the subject (e.g., human) receives an initial administration of cells comprising a CAR molecule, and one or more subsequent administrations of cells comprising a CAR molecule, wherein the one or more subsequent administrations are administered less than 15 days, e.g., 14, 13, 12, 11, 10, 9, 8, 7, 6, 5, 4, 3, or 2 days after the previous administration. In one embodiment, more than one administration of cells comprising a CAR molecule are administered to the subject (e.g., human) per week, e.g., 2, 3, or 4 administrations of cells comprising a CAR molecule are administered per week. In one embodiment, the subject (e.g., human subject) receives more than one administration of cells comprising a CAR molecule per week (e.g., 2, 3 or 4 administrations per week) (also referred to herein as a cycle), followed by a week of no administration of cells comprising a CAR molecule, and then one or more additional administrations of cells comprising a CAR molecule (e.g., more than one administration of the cells comprising a CAR molecule per week) are administered to the subject. In another embodiment, the subject (e.g., human subject) receives more than one cycle of cells comprising a CAR molecule, and the time between each cycle is less than 10, 9, 8, 7, 6, 5, 4, or 3 days. In one embodiment, the cells comprising a CAR molecule are administered every other day for 3 administrations per week. In one embodiment, the cells comprising a CAR molecule are administered for at least two, three, four, five, six, seven, eight or more weeks.

In one embodiment, the immune effector cell, e.g., T cell or NK cell, engineered to express a CAR, e.g., a CAR molecule described herein, is administered as a first line treatment for the disease, e.g., the cancer, e.g., the cancer described herein. In another embodiment, the immune effector cell, e.g., T cell, engineered to express a CAR, e.g., a CAR molecule described herein, is administered as a second, third, fourth line treatment for the disease, e.g., the cancer, e.g., the cancer described herein.

In one embodiment, a population of cells described herein is administered.

In one embodiment, the LSD1 inhibitor and the immune effector cell, e.g., a T cell, engineered to express a CAR molecule, e.g., as described herein, are present in a single composition, e.g., are administered as a single composition. In one embodiment, LSD1 inhibitor and the immune effector cell, e.g., a T cell, engineered to express a CAR molecule, e.g., as described herein, are present in separate compositions, e.g., are administered as separate compositions.

In certain aspects, the disclosure provides an LSD1 inhibitor for use in treating a subject, wherein said LSD1 inhibitor enhances an immune response of said subject, and wherein said subject has received, is receiving or is about to receive an immune effector cell engineered to express a CAR molecule, e.g., as described herein.

In certain aspects, the disclosure provides an immune effector cell engineered to express a CAR molecule, e.g., as described herein for use in treating a subject, wherein said subject has received, is receiving, or is about to receive, an LSD1 inhibitor, e.g., one that enhances an immune response of said subject.

In certain aspects, the disclosure provides an immune effector cell engineered to express a CAR molecule, e.g., as described herein for use in treating a subject, wherein said immune effector cell engineered to express a CAR molecule, e.g., as described herein has been contacted with an LSD1 inhibitor, e.g., contacted ex vivo with an LSD1 inhibitor.

In one embodiment, the invention the population of autologous or allogeneic immune effector cells are transfected or transduced with a vector comprising a nucleic acid molecule encoding a CAR molecule, e.g., as described herein. In one embodiment, the vector is a retroviral vector. In one embodiment, the vector is a self-inactivating lentiviral vector as described elsewhere herein. In one embodiment, the vector is delivered (e.g., by transfecting or electroporating) to a cell, e.g., a T cell or a NK cell, wherein the vector comprises a nucleic acid molecule encoding a CAR molecule, e.g., as described herein, which is transcribed as an mRNA molecule, and the CAR molecule is translated from the RNA molecule and expressed on the surface of the cell.

In an embodiment, a population of CAR-expressing cells, e.g., CAR-expressing T cells (CART cells) or CAR-expressing NK cells, is administered. In some embodiments, the population of CAR-expressing cells comprises a mixture of cells expressing different CARs. For example, in one embodiment, the population of CAR-expressing cells can include a first cell expressing a CAR having an antigen binding domain that binds to a first tumor marker as described herein, and a second cell expressing a CAR having a different antigen binding domain that binds to a second tumor marker as described herein. As another example, the population of CAR-expressing cells can include a first cell expressing a CAR that includes an antigen binding domain that binds to a tumor marker as described herein, and a second cell expressing a CAR that includes an antigen binding domain to a target other than a tumor marker as described herein. In one embodiment, the population of CAR-expressing cells includes, e.g., a first cell

expressing a CAR that includes a primary intracellular signaling domain, and a second cell expressing a CAR that includes a secondary signaling domain.

In one aspect, the invention features a method of treating a subject (e.g., a subject suffering from a disease, e.g., a disease associated with expression of a tumor antigen, e.g., a cancer), that includes administering an LSD1 inhibitor and a population of immune effector cells engineered to express a chimeric antigen receptor (CAR). In embodiments, the method includes administering the LSD1 inhibitor before the population of immune effector cells. In embodiments, the method includes administering the LSD1 inhibitor concurrently with the population of immune effector cells. In embodiments, the method includes administering the LSD1 inhibitor after the population of immune effector cells. In embodiments, the method includes administering the LSD1 inhibitor (e.g., for an interval) before and after the population of immune effector cells is administered.

In one aspect, the invention features a method of treating a subject (e.g., a subject suffering from a disease, e.g., a disease associated with expression of a tumor antigen, e.g., a cancer), that includes administering an LSD1 inhibitor to the subject, wherein said subject has received, is receiving or is about to receive a population of immune effector cells engineered to express a chimeric antigen receptor (CAR). In embodiments, the method includes administering to a subject an LSD1 inhibitor and a population of immune effector cells engineered to express a CAR molecule, e.g., as described herein. In embodiments, the LSD1 inhibitor is administered before the population of immune effector cells engineered to express a CAR molecule, e.g., as described herein, and wherein said administration of the LSD1 inhibitor is continued for a period of time after the administration of the population of immune effector cells engineered to express a CAR molecule, e.g., as described herein. In other embodiments, the administration of the LSD1 inhibitor after the administration of the population of immune effector cells engineered to express a CAR molecule, e.g., as described herein is in an amount sufficient to increase an anti-tumor effect of the population of immune effector cells engineered to express a CAR molecule, e.g., as described herein relative to an equivalent population of immune effector cells engineered to express a CAR molecule, e.g., as described herein administered in the absence of said LSD1 inhibitor.

In another aspect, the invention features a method of increasing the therapeutic efficacy in a subject of a population of immune effector cells engineered to express a CAR molecule, e.g., as described herein, e.g., a CAR19 (e.g., CTL019), including a step of decreasing the activity or expression of LSD1 in said cell, at least transiently. In embodiments, the step of decreasing the activity or expression

of LSD1 in said cell includes contacting the cell with an LSD1 inhibitor. In embodiments, the contacting is done ex vivo. In embodiments, the contacting is done in vivo (e.g., the population of immune effector cells and the LSD1 inhibitor are coadministered to the subject).

In embodiments of any of the foregoing aspect, the administration or the contacting of the LSD1 inhibitor results in:

1) an increase in the proportion of naive T cells, e.g., T_{SCM} cells;

2) an increase in the number of naive T cells, e.g., T_{SCM} cells;

3) a decrease in the number of T_{EM} cells;

4) a decrease in the proportion of T_{EM} cells;

5) an increase in the proportion of CD45RA+CD62L+ T cells;

6) an increase in the number of CD45RA+CD62L+ T cells;

7) an increase in the proportion of CD45RA+CCR7+ T cells;

8) an increase in the number of CD45RA+CCR7+ T cells;

9) a decrease in the proportion of PD-1 positive immune effector cells;

10) an increase in the ratio of PD-1 negative immune effector cells / PD-1 positive immune effector cells;

11) a decrease in the proportion of PD-1+/Lag3+/Tim3+ immune effector cells;

12) an increase in the ratio of PD-1-/Lag3-/Tim3- immune effector cells to PD-1+/Lag3+/Tim3+ immune effector cells;

13) an increase in the proliferation of the immune effector cells;

14 an increase in the production of cytokines (e.g., IFNg and/or IL-2) from said population of immune effector cells; or

15) a combination of two, three, four, five, six, seven, eight, nine, ten, eleven, twelve, thirteen, or more (e.g., all) of the above.

In embodiments, the effect is transient. In embodiments, the effect is permanent. In embodiments, the effect is as compared to cells not contacted with the LSD1 inhibitor. In embodiments, the effect is as compared to cells of the same subject not contacted with the LSD1 inhibitor.

In another aspect, the invention provides a method of treating a subject that includes:

a) administering an LSD1 inhibitor to a subject;

b) collecting a population of immune effector cells from the subject of a), after said administration of the LSD1 inhibitor;

c) providing said population of immune effector cells *ex vivo*;

d) contacting said *ex vivo* population of immune effector cells with the LSD1 inhibitor, wherein the contacting with the LSD1 inhibitor causes one or more of the following to occur:

1) an increase in the proportion of naive T cells, e.g., T_{SCM} cells;

2) an increase in the number of naive T cells, e.g., T_{SCM} cells;

3) a decrease in the number of T_{EM} cells;

4) a decrease in the proportion of T_{EM} cells;

5) an increase in the proportion of CD45RA+CD62L+ T cells;

6) an increase in the number of CD45RA+CD62L+ T cells;

7) an increase in the proportion of CD45RA+CCR7+ T cells;

8) an increase in the number of CD45RA+CCR7+ T cells;

9) a decrease in the proportion of PD-1 positive immune effector cells;

10) an increase in the ratio of PD-1 negative immune effector cells / PD-1 positive immune effector cells;

11) a decrease in the proportion of PD-1+/Lag3+/Tim3+ immune effector cells;

12) an increase in the ratio of PD-1-/Lag3-/Tim3- immune effector cells to PD-1+/Lag3+/Tim3+ immune effector cells;

13) an increase in the proliferation of the immune effector cells;

14 an increase in the production of cytokines (e.g., IFNg and/or IL-2) from said population of immune effector cells; or

15) a combination of two, three, four, five, six, seven, eight, nine, ten, eleven, twelve, thirteen, or more (e.g., all) of the above;

and e) administering the population of immune effector cells to a subject.

In embodiments, the effect of d) is transient. In embodiments, the effect of d) is permanent. In embodiments, the effect of d) is as compared to cells not contacted with the LSD1 inhibitor. In embodiments, the administering of step e) is to the same subject as the subject of step b) (e.g., relates to a method of treatment using a population of autologous immune effector cells). In embodiments, the administering of step e) is to a different subject, e.g., of the same species, as the subject of step b) (e.g., relates to a method of treatment using a population of allogeneic immune effector cells).

In embodiments, step of e) further includes administering the LSD1 inhibitor to the subject. In embodiments, the method further includes the step of inserting nucleic acid that encodes a CAR molecule, e.g., as described herein into cells of the *ex vivo* population of immune effector cells.

In another aspect, the invention features the use of LSD1 inhibitors in the manufacture of a population of immune effector cells, e.g., engineered to express a CAR molecule, e.g., as described herein. In one aspect, the invention provides a method of making a population of immune effector cells, which is optionally a population of T cells, including the steps of:

a) contacting a population of immune effector cells with an LSD1 inhibitor;

thereby making a population of immune effector cells, which is optionally a population of T cells, wherein the contacting with the LSD1 inhibitor causes one or more of the following to occur:

1) an increase in the proportion of naive T cells, e.g., T_{SCM} cells;

2) an increase in the number of naive T cells, e.g., T_{SCM} cells;

3) a decrease in the number of T_{EM} cells;

4) a decrease in the proportion of T_{EM} cells;

5) an increase in the proportion of CD45RA+CD62L+ T cells;

6) an increase in the number of CD45RA+CD62L+ T cells;

7) an increase in the proportion of CD45RA+CCR7+ T cells;

8) an increase in the number of CD45RA+CCR7+ T cells;

9) a decrease in the proportion of PD-1 positive immune effector cells;

10) an increase in the ratio of PD-1 negative immune effector cells / PD-1 positive immune effector cells;

11) a decrease in the proportion of PD-1+/Lag3+/Tim3+ immune effector cells;

12) an increase in the ratio of PD-1-/Lag3-/Tim3- immune effector cells to PD-1+/Lag3+/Tim3+ immune effector cells;

13) an increase in the proliferation of the immune effector cells;

14 an increase in the production of cytokines (e.g., IFNg and/or IL-2) from said population of immune effector cells; or

15) a combination of two, three, four, five, six, seven, eight, nine, ten, eleven, twelve, thirteen, or more (e.g., all) of the above.

In embodiments, the effect of 1)-15) is transient. In embodiments, the effect of 1)-15) is permanent. In embodiments, the effect of 1)-15) is as compared to cells not contacted with the LSD1 inhibitor.

In embodiments, the method further includes the step of b) inserting nucleic acid that encodes a CAR molecule, e.g., as described herein, into cells of the population of immune effector cells. In embodiments, the contacting of step a) occurs

1) prior to;

2) concurrently with;

3) after; or

4) both before and after;

said inserting of step b). In embodiments, the contacting of step a), and optionally the inserting of step b), is *ex vivo*.

In another aspect, the invention features cells, e.g., immune effector cells, e.g., a population of immune effector cells, e.g., engineered to express a CAR molecule, e.g., as described herein, made by any of the methods described in the foregoing aspects.

In another aspect, the invention features a population of immune effector cells engineered to express a CAR molecule, e.g., as described herein, wherein the CAR includes an antigen-binding domain, a transmembrane domain, and an intracellular signaling domain, and wherein expression and/or function of LSD1 in said cell has been reduced or eliminated. In an embodiment, the reduction or elimination is at least transient. In embodiments, the population of immune effector cells has been contacted with an LSD1 inhibitor. In embodiments the invention features a composition comprising the population of immune effector cells described above and an LSD1 inhibitor.

In any of the foregoing aspects and embodiments, the cells and/or population of cells are (or include) immune effector cells, e.g., the population of immune effector cells includes, e.g., consists of, T cells or NK cells. In embodiments, the cells are T cells, e.g., CD8+ T cells, CD4+ T cells, or a combination thereof. In embodiments, the cells are NK cells.

In embodiments, the cells are human cells. In embodiments, the cells are autologous, e.g., to the subject to be administered the cells. In embodiments, the cells are allogeneic, e.g., to the subject to be administered the cells.

In embodiments, the cells are, or include, cells engineered to express a CAR molecule, e.g., as described herein.

In any of the foregoing aspects and embodiments involving a CAR, the CAR includes an antigen binding domain (which is optionally an antibody or antibody fragment, TCR or TCR fragment), a transmembrane domain, and an intracellular signaling domain (which is optionally an intracellular signaling domain including a costimulatory domain and/or a primary signaling domain). In embodiments, the antigen-binding domain binds to a tumor antigen is selected from a group consisting of: TSHR, CD19, CD123, CD22, CD30, CD171, CS-1, CLL-1, CD33, EGFRvIII , GD2, GD3, BCMA, Tn Ag, PSMA, ROR1, FLT3, FAP, TAG72, CD38, CD44v6, CEA, EPCAM, B7H3, KIT, IL-13Ra2, Mesothelin, IL-11Ra, PSCA, PRSS21, VEGFR2, LewisY, CD24, PDGFR-beta, SSEA-4, CD20, Folate receptor alpha, ERBB2 (Her2/neu), MUC1, EGFR, NCAM, Prostase, PAP, ELF2M, Ephrin B2, IGF-I receptor, CAIX, LMP2, gp100, bcr-abl, tyrosinase, EphA2, Fucosyl GM1, sLe, GM3, TGS5, HMWMAA, o-acetyl-GD2, Folate receptor beta, TEM1/CD248, TEM7R, CLDN6, GPRC5D, CXORF61, CD97, CD179a, ALK, Polysialic acid, PLAC1, GloboH, NY-BR-1, UPK2, HAVCR1, ADRB3, PANX3, GPR20, LY6K, OR51E2, TARP, WT1, NY-ESO-1, LAGE-1a, MAGE-A1, legumain, HPV E6,E7, MAGE A1, ETV6-AML, sperm protein 17, XAGE1, Tie 2, MAD-CT-1, MAD-CT-2, Fos-related antigen 1, p53, p53 mutant, prostein, survivin and telomerase, PCTA-1/Galectin 8, MelanA/MART1, Ras mutant, hTERT, sarcoma translocation breakpoints, ML-IAP, ERG (TMPRSS2 ETS fusion gene), NA17, PAX3, Androgen receptor, Cyclin B1, MYCN, RhoC, TRP-2, CYP1B1, BORIS, SART3, PAX5, OY-TES1, LCK, AKAP-4, SSX2, RAGE-1, human telomerase reverse transcriptase, RU1, RU2, intestinal carboxyl esterase, mut hsp70-2, CD79a, CD79b, CD72, LAIR1, FCAR, LILRA2, CD300LF, CLEC12A, BST2, EMR2, LY75, GPC3, FCRL5, and IGLL1. In one embodiment, the antigen binding domain binds CD19, e.g., is an antigen binding domain as described in, e.g., WO2012/079000 or WO2014/153270. In one embodiment, the antigen binding domain binds BCMA, e.g., is an antigen binding domain as described in, e.g., WO2016/014565, e.g., is the antigen binding domain of CAR BCMA-10 (139109) from WO2016/014565.

In embodiments, the antigen-binding domain is an antibody or antibody fragment comprising:

(i) the amino acid sequence of a CD19 binding domain according to Tables 6-9, e.g., the amino acid sequence of CTL019 scFv domain according to Table 9 or an amino acid sequence according to SEQ ID NO: 957, or an amino acid sequence at least 95% identical thereto;

(ii) the amino acid sequence of a humanized CD19 binding domain according to Tables 6-9, e.g., the amino acid sequence of CAR2 scFv domain according to Table 9 or an amino acid sequence according to SEQ ID NO: 898, or an amino acid sequence at least 95% identical thereto; or

(iii) the amino acid sequence of a BCMA binding domain according to Tables 11A-11B, e.g., the amino acid sequence of 139109 scFv domain according to Table 11A or an amino acid sequence according to SEQ ID NO: 967, or an amino acid sequence at least 95% identical thereto.

In embodiments, the transmembrane domain includes:

(i) an amino acid sequence having at least one, two or three modifications but not more than 20, 10 or 5 modifications of an amino acid sequence of SEQ ID NO: 12, or a sequence with 95-99% identity to an amino acid sequence of SEQ ID NO: 12; or

(ii) the sequence of SEQ ID NO: 12.

In embodiments, the antigen binding domain is connected to the transmembrane domain by a hinge region, wherein said hinge region includes SEQ ID NO: 2 or SEQ ID NO: 6, or a sequence with 95-99% identity thereof.

In embodiments, the intracellular signaling domain includes a primary signaling domain and/or a costimulatory signaling domain, wherein the primary signaling domain includes a functional signaling domain of a protein chosen from CD3 zeta, CD3 gamma, CD3 delta, CD3 epsilon, common FcR gamma (FCER1G), FcR beta (Fc Epsilon R1b), CD79a, CD79b, Fcgamma RIIa, DAP10, or DAP12.

In embodiments, the primary signaling domain includes:

(i) an amino acid sequence having at least one, two or three modifications but not more than 20, 10 or 5 modifications of an amino acid sequence of SEQ ID NO: 18 or SEQ ID NO: 20, or a sequence with 95-99% identity to an amino acid sequence of SEQ ID NO: 18 or SEQ ID NO: 20; or

(ii) the amino acid sequence of SEQ ID NO:18 or SEQ ID NO: 20.

In embodiments, the intracellular signaling domain includes a costimulatory signaling domain, or a primary signaling domain and a costimulatory signaling domain, wherein the costimulatory signaling domain includes a functional signaling domain of a protein selected from the group consisting of CD27, CD28, 4-1BB (CD137), OX40, CD30, CD40, PD-1, ICOS, lymphocyte function-associated antigen-1 (LFA-1), CD2, CD7, LIGHT, NKG2C, B7-H3, a ligand that specifically binds with CD83, CDS, ICAM-1, GITR, BAFFR, HVEM (LIGHTR), SLAMF7, NKp80 (KLRF1), CD160, CD19, CD4, CD8alpha, CD8beta, IL2R beta, IL2R gamma, IL7R alpha, ITGA4, VLA1, CD49a, ITGA4, IA4, CD49D, ITGA6, VLA-6, CD49f, ITGAD, CD11d, ITGAE, CD103, ITGAL, CD11a, LFA-1, ITGAM, CD11b, ITGAX, CD11c, ITGB1, CD29, ITGB2, CD18, LFA-1, ITGB7, TNFR2, TRANCE/RANKL, DNAM1 (CD226), SLAMF4 (CD244, 2B4), CD84, CD96 (Tactile), CEACAM1, CRTAM, Ly9 (CD229), CD160 (BY55), PSGL1, CD100 (SEMA4D), CD69, SLAMF6 (NTB-A, Ly108), SLAM (SLAMF1, CD150, IPO-3), BLAME (SLAMF8), SELPLG (CD162), LTBR, LAT, GADS, SLP-76, PAG/Cbp, NKp44, NKp30, NKp46, and NKG2D, e.g., the costimulatory signaling domain includes an amino acid sequence having at least one, two or three modifications but not more than 20, 10 or 5 modifications of an amino acid sequence of SEQ ID NO:14 or SEQ ID NO: 16, or a sequence with 95-99% identity to an amino acid sequence of SEQ ID NO:14 or SEQ ID NO: 16, e.g., the costimulatory signaling domain includes a sequence of SEQ ID NO: 14 or SEQ ID NO: 16, e.g., the intracellular domain includes the sequence of SEQ ID NO: 14 or SEQ ID NO: 16, and the sequence of SEQ ID NO: 18 or SEQ ID NO: 20, wherein the sequences including the intracellular signaling domain are expressed in the same frame and as a single polypeptide chain.

In embodiments, the CAR includes a leader sequence including, e.g., consisting of, SEQ ID NO: 2.

In embodiments, the CAR comprises:

(i) the amino acid sequence of a CD19 CAR according to Tables 6-9, e.g., the amino acid sequence of CTL019 according to Table 9 or an amino acid sequence according to SEQ ID NO: 956 or an amino acid sequence at least 95% identical thereto;

(ii) the amino acid sequence of a humanized CD19 CAR according to Tables 6-9, e.g., the amino acid sequence of CAR2 according to Table 9 or an amino acid sequence according to SEQ ID NO: 902, or an amino acid sequence at least 95% identical thereto; or

(iii) the amino acid sequence of a BCMA CAR according to Tables 11A-11B, e.g., the amino acid sequence of 139109 CAR according to Table 11A or an amino acid sequence according to SEQ ID NO: 971, or an amino acid sequence at least 95% identical thereto.

In any of the foregoing aspects and embodiments, the LSD1 inhibitor may be: (1) a gene editing system targeted to one or more sites of the LSD1 gene, or its corresponding regulatory elements; (2) a nucleic acid (e.g., an siRNA or shRNA, or antisense oligonucleotide) including sequence complementary to a target sequence of the LSD1 gene; (3) a protein (e.g., a dominant negative LSD1, e.g., catalytically inactive LSD1, or a dominant negative binding partner of LSD1); (4) a small molecule; (5) a nucleic acid encoding any of (1)-(3); or (6) any combination of (1)–(5).

In one aspect, the LSD1 inhibitor is an shRNA or siRNA. In embodiments, the LSD1 inhibitor is a shRNA. In embodiments, the LSD1 inhibitor is as siRNA. In embodiments, the shRNA or siRNA includes sequence complementary to a target sequence of the LSD1 gene (KDM1A), e.g., listed in Table 1, e.g., selected from SEQ ID NO: [43] to SEQ ID NO: [82].

In another aspect, the LSD1 inhibitor is an shRNA encoded by nucleic acid including any sequence encoding an anti-LSD1 shRNA of Table 1, e.g., encoded by nucleic acid including a sequence selected from SEQ ID NO: [83] to SEQ ID NO: [122].

In another aspect, the LSD1 inhibitor is nucleic acid including any sequence encoding an anti-LSD1 shRNA of Table 1, e.g., a sequence selected from SEQ ID NO: [83] to SEQ ID NO: [122].

In another aspect, the LSD1 inhibitor is an antisense oligonucleotide. In embodiments, the antisense oligonucleotide includes sequence that is complementary to a sequence of an LSD1 mRNA. In embodiments, the antisense oligonucleotide includes sequence that is complementary to a sequence of an LSD1 pre-mRNA.

In embodiments, the nucleic acid encoding the LSD1 inhibitor is disposed on a vector, e.g., a vector further including a U6 or H1 promoter operably linked to said nucleic acid, e.g., a retroviral vector, a lentiviral vector, an adenoviral vector, an adeno-associated viral (AAV) vector, a herpes simplex virus (HSV) vector, a plasmid, a minicircle, a nanoplasmid, or an RNA vector. In embodiments the vector further includes sequence encoding a CAR molecule.

In another aspect, the LSD1 inhibitor is a genome editing system specific for a sequence of the LSD1 gene (KDM1A) or its regulatory elements selected from a CRISPR genome editing system, a zinc finger nuclease genome editing system, a TALEN genome editing system and a meganuclease genome editing system.

In another aspect, the LSD1 inhibitor is a CRISPR genome editing system including a gRNA molecule including a targeting domain complementary to a sequence of the LSD1 gene (KDM1A) or its regulatory elements, e.g., including any one of SEQ ID NO: [132] to [862].

In other aspect, the LSD1 inhibitor is a small molecule. In embodiments, the small molecule is a reversible LSD1 inhibitor. In embodiments, the small molecule is an irreversible LSD1 inhibitor. In embodiments, the small molecule LSD1 inhibitor is:

a) GSK2699537;

b) rel- 2-[[(1R,2S)-2-[4-[(4-chlorophenyl)methoxy]phenyl]cyclopropyl]amino]-1-(4-methyl-1-piperazinyl)-ethanone (described in PCT Publication No. WO 2010043721);

c) (R)-4-(5-(pyrrolidin-3-ylmethoxy)-2-(p-tolyl)pyridin-3-yl)benzonitrile;

d) (1S,2R)-N-((2-methoxypyridin-3-yl)methyl)-2-phenylcyclopropan-1-amine;

e) N,N-dimethyl-1-((4-(4-(4-(piperidin-4-yl)phenyl)-1H-indazol-1-yl)phenyl)sulfonyl)piperidin-4-amine;

f) 5-(6-chloro-4'-(methylsulfonyl)-[1,1'-biphenyl]-3-yl)-2-(piperazin-1-yl)-1H-pyrrole-3-carbonitrile;

g) rel- N-[(1R,2S)-2-Phenylcyclopropyl]-4-Piperidinamine; or

h) 2-(1R,2S)-2-(4-(Benzyloxy)phenyl)cyclopropylamino)-1-(4-methylpiperazin-1-yl)ethanone;

i) Trans-3-(3-amino-2-methylphenyl)-1-(4-hydroxycyclohexyl)-6-methyl-1H-indole-5-carbonitrile; or

j) a pharmaceutically acceptable salt of any of the foregoing. In embodiments, the LSD1 inhibitor is a small molecule and said LSD1 inhibitor is conjugated to an antibody or antigen-binding fragment thereof, e.g., an antibody or antigen-binding fragment thereof that recognizes an antigen on the surface of a T cell, e.g., CD3.

In another aspect, the LSD1 inhibitor is a protein, e.g., is a dominant negative binding partner of LSD1 (e.g., a histone deacetylase (HDAC) that interacts with LSD1 or other member of the Co-REST or AR co-activator complex), or nucleic acid encoding said dominant negative binding partner of LSD1.

In another aspect, the inhibitor of LSD1 is a protein, e.g., is a dominant negative (e.g., catalytically inactive) LSD1, or nucleic acid encoding said dominant negative LSD1.

In another aspect, the invention provides a method of treating a subject in need thereof, including administering to said subject an effective amount of the population of immune effector cells of any of the previous aspects and embodiments. In embodiments, the method further includes administering to said subject an LSD1 inhibitor. In embodiments, the subject receives a pre-treatment of an LSD1 inhibitor, prior to the administration of the population of immune effector cells; In embodiments, the subject receives concurrent treatment with an LSD1 inhibitor and the population of immune effector cells; In embodiments, the subject receives treatment with an LSD1 inhibitor after administration of the population of immune effector cells; In embodiments, the subject receives a combination of any of the foregoing.

In an aspect, including in the previous aspects relating to methods of treatment, the invention relates to methods of treating a subject, wherein the subject has a disease associated with expression of a tumor antigen, e.g., a proliferative disease, a precancerous condition, a cancer, or a non-cancer related indication associated with expression of the tumor antigen. In embodiments, the cancer is a hematologic cancer chosen from one or more of chronic lymphocytic leukemia (CLL), acute leukemias, acute lymphoid leukemia (ALL), acute myeloid leukemia (AML), B-cell acute lymphoid leukemia (B-ALL), T-cell acute lymphoid leukemia (T-ALL), chronic myelogenous leukemia (CML), B cell prolymphocytic leukemia, blastic plasmacytoid dendritic cell neoplasm, Burkitt's lymphoma, diffuse large B cell lymphoma, follicular lymphoma, hairy cell leukemia, small cell- or a large cell-follicular lymphoma, malignant lymphoproliferative conditions, MALT lymphoma, mantle cell lymphoma, marginal zone lymphoma, multiple myeloma, myelodysplasia and myelodysplastic syndrome, non-Hodgkin’s lymphoma, Hodgkin’s lymphoma, plasmablastic lymphoma, plasmacytoid dendritic cell neoplasm, Waldenstrom macroglobulinemia, or pre-leukemia. In embodiments, the cancer is selected from the group consisting of colon cancer, rectal cancer, renal-cell carcinoma, liver cancer, non-small cell carcinoma of the lung, cancer of the small intestine, cancer of the esophagus, melanoma, bone cancer, pancreatic cancer, skin cancer, cancer of the head or neck, cutaneous or intraocular malignant melanoma, uterine cancer, ovarian cancer, rectal cancer, cancer of the anal region, stomach cancer, testicular cancer, uterine cancer, carcinoma of the fallopian tubes, carcinoma of the endometrium, carcinoma of the cervix, carcinoma of the vagina, carcinoma of the vulva, Hodgkin's Disease, non-Hodgkin's lymphoma, cancer of the endocrine system, cancer of the thyroid gland, cancer of the

parathyroid gland, cancer of the adrenal gland, sarcoma of soft tissue, cancer of the urethra, cancer of the penis, solid tumors of childhood, cancer of the bladder, cancer of the kidney or ureter, carcinoma of the renal pelvis, neoplasm of the central nervous system (CNS), primary CNS lymphoma, tumor angiogenesis, spinal axis tumor, brain stem glioma, pituitary adenoma, Kaposi's sarcoma, epidermoid cancer, squamous cell cancer, T-cell lymphoma, environmentally induced cancers, combinations of said cancers, and metastatic lesions of said cancers.

In another aspect, the invention provides novel compounds. Such compounds are useful, for example, in the methods and compositions described herein, but such uses and compositions are not intended to be limiting. In an embodiment, the invention provides a compound selected from N,N-dimethyl-1-((4-(4-(4-(piperidin-4-yl)phenyl)-1H-indazol-1-yl)phenyl)sulfonyl)piperidin-4-amine and 5-(6-chloro-4'-(methylsulfonyl)biphenyl-3-yl)-2-(piperazin-1-yl)-1H-pyrrole-3-carbonitrile. In an embodiment, the invention provides N,N-dimethyl-1-((4-(4-(4-(piperidin-4-yl)phenyl)-1H-indazol-1-yl)phenyl)sulfonyl)piperidin-4-amine. In an embodiment, the invention provides 5-(6-chloro-4'-(methylsulfonyl)biphenyl-3-yl)-2-(piperazin-1-yl)-1H-pyrrole-3-carbonitrile. The invention further provides a pharmaceutically acceptable salt of any of the foregoing. The invention further provides a compound described above, for use in the manufacture of a medicament. The invention further provides a compound described above, for use as a medicament. The invention further provides a compound described above, for use in the manufacture of a medicament for use as an LSD1 inhibitor, e.g., for use as an LSD1 inhibitor in any of the methods described herein. In an embodiment, the invention provides a compound described above, for use in therapy, alone, or optionally in combination with at least another agent.

In another aspect, the invention provides a composition for use in *ex vivo* manufacturing a population of immune effector cells, that includes an LSD1 inhibitor, e.g., a small molecule LSD1 inhibitor. In embodiments, the composition includes the small molecule LSD1 inhibitor at a concentration of ranges from about 0.001 nM to about 10 mM, e.g., from about 0.001 nM to about 100 nM, or from, e.g., about 0.1 uM to about 10 uM.

In an aspect, the invention provides an LSD1 inhibitor, for use in treating a subject, wherein said subject has received, is receiving, or is about to receive therapy including an immune effector cell, e.g., an immune effector cell engineered to express a CAR molecule, e.g., as described herein.

In an aspect, the invention provides an LSD1 inhibitor, for use in the manufacture of an immune effector cell, e.g., an immune effector cell engineered to express a CAR molecule, e.g., as described herein.

In an aspect, the invention provides a method of manufacturing an immune effector cell, e.g., a population of immune effector cells, that includes introducing into said cells nucleic acid encoding a CAR molecule, e.g., as described herein, wherein the nucleic acid integrates into the genome of said cell within the LSD1 gene, such that LSD1 expression and/or function is reduced.

BRIEF DESCRIPTION OF THE DRAWINGS

Figure 1 depicts the percentage of CD8+ T cells from human donors that are CD45RA+CD62L+ after activation using CD3/CD28 in the presence of an LSD1 inhibitor shRNA (molecules 1A, 1B, 2, 3A, 3B, 4 or 6A) compared to control.

Figure 2 depicts the percentage of CD4+ T cells from human donors that are CD45RA+CD62L+ after activation using CD3/CD28 in the presence of an LSD1 inhibitor shRNA (molecules 1A, 1B, 2, 3A, 3B, 4 or 6A) compared to control.

Figure 3A shows the ability of the indicated compounds to produce T cells of a given phenotype was assessed. Naive human T cells (peripheral pan CD3+ T cells, pooled population) were isolated by negative selection and expanded with anti-CD3/CD28 beads at a 3:1 ratio for 10 days in the presence of the indicated compounds. Compounds were refreshed every 2 days. Following expansion, T cell phenotypes were determined by FACS staining. LSD1 inhibition significantly enhanced the percentage of Tscm cells while reducing the percentage of Tem cells in CD4+ T cells relative to controls and relative to other known conditions.

Figure 3B shows the ability of the indicated compounds to produce T cells of a given phenotype was assessed. Naive human T cells (peripheral pan CD3+ T cells, pooled population) were isolated by negative selection and expanded with anti-CD3/CD28 beads at a 3:1 ratio for 10 days in the presence of the indicated compounds. Compounds were refreshed every 2 days. Following expansion, T cell phenotypes were determined by FACS staining. LSD1 inhibition significantly enhanced the percentage of T_{SCM} cells while reducing the percentage of Tem cells in CD8+ T cells relative to controls and relative to other known conditions.

Figure 4A shows the effect of LSD1 inhibition in comparison to other compounds believed to affect T cell phenotype on T_{SCM} to T_{EM} ratio in CD4+ cells.

Figure 4B shows the effect of LSD1 inhibition in comparison to other compounds believed to affect T cell phenotype on T_{SCM} to T_{EM} ratio in CD8+ cells.

Figure 5 shows the expansion of T cells using CD3/CD28 stimulation in the presence of LSD1 inhibitors.

Figure 6 shows the expression of checkpoint proteins PD1, Tim3 and Lag3 on T cells expanded in the presence of LSD1 inhibitors.

Figure 7 shows the level of CAR expression on T cells expanded in the presence of LSD1 inhibitors.

Figure 8 shows the proportion of CD4+ and CD8+ CART and untransduced T cells in the presence of LSD1 inhibitors.

Figure 9 shows the expansion of CART cells and untransduced T cells in the presence of LSD1 inhibitors.

Figure 10 shows the cytokine production from CART cells expanded in the presence of LSD1 inhibitors, and then exposed to CD19+ or CD19- tumor cells.

Figure 11 shows the effect of LSD1 inhibition in significantly increased proliferative capacity following CAR stimulation. Naive human T cells (peripheral pan CD3+ T cells, pooled population) were isolated from PBMCs by negative selection and expanded with anti-CD3/CD28 beads at a 3:1 ratio for 10 days in the presence of the indicated compounds. T cells were transduced with an anti-CD19 scFV on day 1. Compounds were refreshed every 2 days until day 10 was washed out prior to functional assays. Following expansion, T cells were mixed with CD19+ tumor cells lines NALM6 and Raji, as well as CD19- tumor cell line K562. Tumor cells were irradiated and T cells and tumor cells were mixed at a 1:1 ratio. On day 4 following incubation, T cells were stained for CAR using Protein L and CAR+ T cell numbers were determined by FACS using countbright beads. Proliferation was measured as the number of FACS positive cells detected in the period of time used to count 2500 beads. Data expressed as fold no target (CD19-) control (K562).

Figure 12 shows the effects of the indicated compounds on naive human T cells (peripheral pan CD3+ T cells, pooled population) isolated by negative selection from PBMCs and expanded with anti-CD3/CD28 beads at a 3:1 ratio for 10 days in the presence of the indicated compounds. T cells were transduced with an anti-CD19 scFV on day 1. Compounds were refreshed every 2 days until day 10 was washed out prior to functional assays. Following expansion, T cell killing of the luciferized CD19+ NALM6 tumor cell line was assessed. After 20 hours luciferase signal was measured using the Bright-Glo™ Luciferase Assay on the EnVision instrument.

Figure 13 shows the in vivo anti-tumor efficacy of CART cells expanded ex vivo in the presence of LSD1 inhibitors.

Figure 14A shows the level of expansion of CD4+ T cells (e.g., T_{SCM}) cells in the presence of LSD1 inhibitors.

Figure 14B shows the level of expansion of CD8+ T cells (e.g., T_{SCM}) cells in the presence of LSD1 inhibitors.

Figure 15A shows the level of expression of checkpoint proteins PD1, Tim3 and Lag3 on CD4+ T cells expanded in the presence of LSD1 inhibitors.

Figure 15B shows the level of expression of checkpoint proteins PD1, Tim3 and Lag3 on CD8+ T cells expanded in the presence of LSD1 inhibitors.

Figure 16 depicts the percentage of total T cells expressing PD1, Tim3 or Lag3 (left panel) or co-expressing PD1/Lag3 or PD1/Lag3/Tim3 after expansion in the presence or absence of LSD1 inhibitor.

Figure 17 depicts the percentage of CD4+ T cells (left panel) and percentage of CD8+ T cells (right panel) which are Tscm after expansion in the presence or absence of LSD1 inhibitor.

Figure 18 depicts the percentage of CD4+ T cells (left panel) and percentage of CD8+ T cells (right panel) which are positive for co-expression of Tim3/Lag3/PD-1 after expansion in the presence or absence of LSD1 inhibitor.

Figure 19 depicts an ilustration of gating strategy for Tscm, Tcm and Tem by flow cytometry analysis.

Figure 20 depicts the effects of different concentrations of LSD1 inhibitors on the T cell phenotypic changes after 10 day activation, treatment and expansion in culture. Percentage of Tscm, Tem and Tcm of CD3+ T cells, as well as the ratio of subset CD8+/CD4+, Tscm/Tem, and Tscm/Tcm are shown.

Figure 21 depicts a dose response curve of the compound 93 (NVS Compound 1) on the subset of CD8+ and CD4+ T cells on induction of Tscm, Tem and Tcm.

Figure 22 depicts a dose response curve of the LSD1i-GSK on the subset of CD8+ and CD4+ T cells on induction of Tscm, Tem and Tcm.

Figure 23 depicts a dose response curve of the compound 93 and LSD1i-GSK on the CD3+ and CD8+ T cells on induction of Tscm (EC50 shown).

Figure 24 shows that Compound A and Compound B showed similar effects on inducing Tscm and reducing Tem at 100nM started dosing 24h after activation, when compared with Compound 93.

Figure 25 depicts an ilustration of gating strategy for Tscm, Tcm and Tem in total T cell and CAR+ T cells by flow cytometry analysis.

Figure 26 shows FACS analyses for total CD3+ T cells, CAR expression, and CD8+ CAR+ vs CD4+ CAR+ ratio in the final CART product in response to different concentrations of LSD1 inhibition by Compound 93.

Figure 27 shows FACS analyses for Tscm, Tem, Tcm in the total CD8+ T cells of the final CART products in response to different concentrations of LSD1 inhibition by Compound 93.

Figure 28 shows FACS analyses for the percentage of Tscm, Tem, Tcm in the CAR+CD8+ T cells in response to different concentrations of LSD1 inhibition by Compound 93.

Figure 29 shows results from in vitro cytokine assay for IFNg secretion by CART cells with or without LSD1 inhibitor treatment in response to their specific tumor target cells line at an effector to target cells ratio 1.25:1 incubation for 20 hours.

Figure 30 shows in vitro CD3+ (total) and CD3+CAR+ cell proliferation levels in response to their specific irradiated tumor cells lines at an effector to target cells ratio 1:1 for 4 days.

Figure 31 shows in vitro CD8+ and CD8+CAR+ T cell proliferation levels in response to their specific irradiated tumor cells lines at an effector to target cells ratio 1:1 for 4 days.

Figure 32 shows in vitro CD4+ and CD4+CAR+ T cell proliferation levels in response to their specific irradiated tumor cells lines at an effector to target cells ratio 1:1 for 4 days.

Figure 33 shows in vivo anti-tumor efficacy of BCMA CART cells against BCMA+ tumor line. UTD = T cells that were not transduced with the CAR gene; LSDi = indicates those populations that were expanded ex vivo in the presence of Compound 93; BCMA 0.167 million = indicates those populations which were transduced with the CAR gene, and the number of CAR+ cells in the population; PBS = no cell control (phosphate buffered saline injection only).

DETAILED DESCRIPTION

Definitions

Unless defined otherwise, all technical and scientific terms used herein have the same meaning as commonly understood by one of ordinary skill in the art to which the invention pertains.

The term “a” and “an” refers to one or to more than one (i.e., to at least one) of the grammatical object of the article. By way of example, “an element” means one element or more than one element.

The term “about” when referring to a measurable value such as an amount, a temporal duration, and the like, is meant to encompass variations of ±20% or in some instances ±10%, or in some instances ±5%, or in some instances ±1%, or in some instances ±0.1% from the specified value, as such variations are appropriate to perform the disclosed methods.

The term “Chimeric Antigen Receptor” or alternatively a “CAR” refers to a set of polypeptides, typically two in the simplest embodiments, which when in an immune effector cell, provides the cell with specificity for a target cell, typically a cancer cell, and with intracellular signal generation. In some embodiments, a CAR comprises at least an extracellular antigen binding domain, a transmembrane domain and a cytoplasmic signaling domain (also referred to herein as “an intracellular signaling domain”) comprising a functional signaling domain derived from a stimulatory molecule and/or costimulatory molecule as defined below. In some aspects, the set of polypeptides are contiguous with each other. In some embodiments, the set of polypeptides includes a dimerization switch that, upon

the presence of a dimerization molecule, can couple the polypeptides to one another, e.g., can couple an antigen binding domain to an intracellular signaling domain. In one aspect, the stimulatory molecule is the zeta chain associated with the T cell receptor complex. In one aspect, the cytoplasmic signaling domain further comprises one or more functional signaling domains derived from at least one costimulatory molecule as defined below. In one aspect, the costimulatory molecule is chosen from the costimulatory molecules described herein, e.g., 4-1BB (i.e., CD137), CD27 and/or CD28. In one aspect, the CAR comprises a chimeric fusion protein comprising an extracellular antigen binding domain, a transmembrane domain and an intracellular signaling domain comprising a functional signaling domain derived from a stimulatory molecule. In one aspect, the CAR comprises a chimeric fusion protein comprising an extracellular antigen binding domain, a transmembrane domain and an intracellular signaling domain comprising a functional signaling domain derived from a costimulatory molecule and a functional signaling domain derived from a stimulatory molecule. In one aspect, the CAR comprises a chimeric fusion protein comprising an extracellular antigen binding domain, a transmembrane domain and an intracellular signaling domain comprising two functional signaling domains derived from one or more costimulatory molecule(s) and a functional signaling domain derived from a stimulatory molecule. In one aspect, the CAR comprises a chimeric fusion protein comprising an extracellular antigen binding domain, a transmembrane domain and an intracellular signaling domain comprising at least two functional signaling domains derived from one or more costimulatory molecule(s) and a functional signaling domain derived from a stimulatory molecule. In one aspect the CAR comprises an optional leader sequence at the amino-terminus (N-ter) of the CAR fusion protein. In one aspect, the CAR further comprises a leader sequence at the N-terminus of the extracellular antigen binding domain, wherein the leader sequence is optionally cleaved from the antigen binding domain (e.g., a scFv) during cellular processing and localization of the CAR to the cellular membrane.

A CAR that comprises an antigen binding domain (e.g., a scFv, or TCR) that targets a specific tumor marker X, such as those described herein, is also referred to as XCAR. For example, a CAR that comprises an antigen binding domain that targets CD19 is referred to as CD19CAR.

The term “signaling domain” refers to the functional portion of a protein which acts by transmitting information within the cell to regulate cellular activity via defined signaling pathways by generating second messengers or functioning as effectors by responding to such messengers.

The term “antibody,” as used herein, refers to a protein, or polypeptide sequence derived from an immunoglobulin molecule which specifically binds with an antigen. Antibodies can be polyclonal or

monoclonal, multiple or single chain, or intact immunoglobulins, and may be derived from natural sources or from recombinant sources. Antibodies can be tetramers of immunoglobulin molecules.

The term “antibody fragment” refers to at least one portion of an antibody, that retains the ability to specifically interact with (e.g., by binding, steric hinderance, stabilizing/destabilizing, spatial distribution) an epitope of an antigen. Examples of antibody fragments include, but are not limited to, Fab, Fab', $F(ab')_2$, Fv fragments, scFv antibody fragments, disulfide-linked Fvs (sdFv), a Fd fragment consisting of the VH and CH1 domains, linear antibodies, single domain antibodies such as sdAb (either VL or VH), camelid VHH domains, multi-specific antibodies formed from antibody fragments such as a bivalent fragment comprising two Fab fragments linked by a disulfide bridge at the hinge region, and an isolated CDR or other epitope binding fragments of an antibody. An antigen binding fragment can also be incorporated into single domain antibodies, maxibodies, minibodies, nanobodies, intrabodies, diabodies, triabodies, tetrabodies, v-NAR and bis-scFv (see, e.g., Hollinger and Hudson, Nature Biotechnology 23:1126-1136, 2005). Antigen binding fragments can also be grafted into scaffolds based on polypeptides such as a fibronectin type III (Fn3) (see U.S. Patent No.: 6,703,199, which describes fibronectin polypeptide minibodies).

The term “scFv” refers to a fusion protein comprising at least one antibody fragment comprising a variable region of a light chain and at least one antibody fragment comprising a variable region of a heavy chain, wherein the light and heavy chain variable regions are contiguously linked, e.g., via a synthetic linker, e.g., a short flexible polypeptide linker, and capable of being expressed as a single chain polypeptide, and wherein the scFv retains the specificity of the intact antibody from which it is derived. Unless specified, as used herein an scFv may have the VL and VH variable regions in either order, e.g., with respect to the N-terminal and C-terminal ends of the polypeptide, the scFv may comprise VL-linker-VH or may comprise VH-linker-VL.

The portion of the CAR of the invention comprising an antibody or antibody fragment thereof may exist in a variety of forms where the antigen binding domain is expressed as part of a contiguous polypeptide chain including, for example, a single domain antibody fragment (sdAb), a single chain antibody (scFv), a humanized antibody or bispecific antibody (Harlow et al., 1999, In: Using Antibodies: A Laboratory Manual, Cold Spring Harbor Laboratory Press, NY; Harlow et al., 1989, In: Antibodies: A Laboratory Manual, Cold Spring Harbor, New York; Houston et al., 1988, Proc. Natl. Acad. Sci. USA 85:5879-5883; Bird et al., 1988, Science 242:423-426). In one aspect, the antigen binding domain of a CAR composition of the invention comprises an antibody fragment. In a further aspect, the CAR

comprises an antibody fragment that comprises a scFv. The precise amino acid sequence boundaries of a given CDR can be determined using any of a number of well-known schemes, including those described by Kabat et al. (1991), “Sequences of Proteins of Immunological Interest,” 5th Ed. Public Health Service, National Institutes of Health, Bethesda, MD (“Kabat” numbering scheme), Al-Lazikani et al., (1997) JMB 273,927-948 (“Chothia” numbering scheme), or a combination thereof.

As used herein, the term “binding domain” or "antibody molecule" refers to a protein, e.g., an immunoglobulin chain or fragment thereof, comprising at least one immunoglobulin variable domain sequence. The term “binding domain” or “antibody molecule” encompasses antibodies and antibody fragments. In an embodiment, an antibody molecule is a multispecific antibody molecule, e.g., it comprises a plurality of immunoglobulin variable domain sequences, wherein a first immunoglobulin variable domain sequence of the plurality has binding specificity for a first epitope and a second immunoglobulin variable domain sequence of the plurality has binding specificity for a second epitope. In an embodiment, a multispecific antibody molecule is a bispecific antibody molecule. A bispecific antibody has specificity for no more than two antigens. A bispecific antibody molecule is characterized by a first immunoglobulin variable domain sequence which has binding specificity for a first epitope and a second immunoglobulin variable domain sequence that has binding specificity for a second epitope.

The term “antibody heavy chain,” refers to the larger of the two types of polypeptide chains present in antibody molecules in their naturally occurring conformations, and which normally determines the class to which the antibody belongs.

The term “antibody light chain,” refers to the smaller of the two types of polypeptide chains present in antibody molecules in their naturally occurring conformations. Kappa (κ) and lambda (λ) light chains refer to the two major antibody light chain isotypes.

The term “recombinant antibody” refers to an antibody which is generated using recombinant DNA technology, such as, for example, an antibody expressed by a bacteriophage or yeast expression system. The term should also be construed to mean an antibody which has been generated by the synthesis of a DNA molecule encoding the antibody and which DNA molecule expresses an antibody protein, or an amino acid sequence specifying the antibody, wherein the DNA or amino acid sequence has been obtained using recombinant DNA or amino acid sequence technology which is available and well known in the art.

The term “antigen” or “Ag” refers to a molecule that provokes an immune response. This immune response may involve either antibody production, or the activation of specific immunologically-competent cells, or both. The skilled artisan will understand that any macromolecule, including virtually all proteins or peptides, can serve as an antigen. Furthermore, antigens can be derived from recombinant or genomic DNA. A skilled artisan will understand that any DNA, which comprises a nucleotide sequences or a partial nucleotide sequence encoding a protein that elicits an immune response, therefore encodes an “antigen” as that term is used herein. Furthermore, one skilled in the art will understand that an antigen need not be encoded solely by a full length nucleotide sequence of a gene. It is readily apparent that the present invention includes, but is not limited to, the use of partial nucleotide sequences of more than one gene and that these nucleotide sequences are arranged in various combinations to encode polypeptides that elicit the desired immune response. Moreover, a skilled artisan will understand that an antigen need not be encoded by a “gene” at all. It is readily apparent that an antigen can be synthesized or can be derived from a biological sample, or might be a macromolecule besides a polypeptide. Such a biological sample can include, but is not limited to a tissue sample, a tumor sample, a cell or a fluid with other biological components.

The term “anti-cancer effect” refers to a biological effect which can be manifested by various means, including but not limited to, e.g., a decrease in tumor volume, a decrease in the number of cancer cells, a decrease in the number of metastases, an increase in life expectancy, decrease in cancer cell proliferation, decrease in cancer cell survival, or amelioration of various physiological symptoms associated with the cancerous condition. An “anti-cancer effect” can also be manifested by the ability of the peptides, polynucleotides, cells and antibodies in prevention of the occurrence of cancer in the first place. The term “anti-tumor effect” refers to a biological effect which can be manifested by various means, including but not limited to, e.g., a decrease in tumor volume, a decrease in the number of tumor cells, a decrease in tumor cell proliferation, an increase in tumor cell death, an increase in tumor cell apoptosis, or a decrease in tumor cell survival.

The term “autologous” refers to any material derived from the same individual to whom it is later to be re-introduced into the individual.

The term “allogeneic” refers to any material derived from a different animal of the same species as the individual to whom the material is introduced. Two or more individuals are said to be allogeneic to one another when the genes at one or more loci are not identical. In some aspects, allogeneic

material from individuals of the same species may be sufficiently unlike genetically to interact antigenically.

The term “xenogeneic” refers to a graft derived from an animal of a different species.

The term “cancer” refers to a disease characterized by the uncontrolled growth of aberrant cells. Cancer cells can spread locally or through the bloodstream and lymphatic system to other parts of the body. Examples of various cancers are described herein and include but are not limited to, breast cancer, prostate cancer, ovarian cancer, cervical cancer, skin cancer, pancreatic cancer, colorectal cancer, renal cancer, liver cancer, brain cancer, lymphoma, leukemia, lung cancer and the like. The terms “tumor” and “cancer” are used interchangeably herein, e.g., both terms encompass solid and liquid, e.g., diffuse or circulating, tumors. As used herein, the term “cancer” or “tumor” includes premalignant, as well as malignant cancers and tumors.

“Derived from” as that term is used herein, indicates a relationship between a first and a second molecule. It generally refers to structural similarity between the first molecule and a second molecule and does not connotate or include a process or source limitation on a first molecule that is derived from a second molecule. For example, in the case of an intracellular signaling domain that is derived from a CD3zeta molecule, the intracellular signaling domain retains sufficient CD3zeta structure such that it has the required function, namely, the ability to generate a signal under the appropriate conditions. It does not connotate or include a limitation to a particular process of producing the intracellular signaling domain, e.g., it does not mean that, to provide the intracellular signaling domain, one must start with a CD3zeta sequence and delete unwanted sequence, or impose mutations, to arrive at the intracellular signaling domain.

The phrase “disease associated with expression of a tumor antigen as described herein” includes, but is not limited to, a disease associated with expression of a tumor antigen as described herein or condition associated with cells which express a tumor antigen as described herein including, e.g., proliferative diseases such as a cancer or malignancy or a precancerous condition such as a myelodysplasia, a myelodysplastic syndrome or a preleukemia; or a noncancer related indication associated with cells which express a tumor antigen as described herein. In one aspect, a cancer associated with expression of a tumor antigen as described herein is a hematological cancer. In one aspect, a cancer associated with expression of a tumor antigen as described herein is a solid cancer. Further diseases associated with expression of a tumor antigen described herein include, but not limited to, e.g., atypical and/or non-classical cancers, malignancies, precancerous conditions or proliferative

diseases associated with expression of a tumor antigen as described herein. Non-cancer related indications associated with expression of a tumor antigen as described herein include, but are not limited to, e.g., autoimmune disease, (e.g., lupus), inflammatory disorders (allergy and asthma) and transplantation. In some embodiments, the tumor antigen-expressing cells express, or at any time expressed, mRNA encoding the tumor antigen. In an embodiment, the tumor antigen -expressing cells produce the tumor antigen protein (e.g., wild-type or mutant), and the tumor antigen protein may be present at normal levels or reduced levels. In an embodiment, the tumor antigen -expressing cells produced detectable levels of a tumor antigen protein at one point, and subsequently produced substantially no detectable tumor antigen protein.

The term "conservative sequence modifications" refers to amino acid modifications that do not significantly affect or alter the binding characteristics of the antibody or antibody fragment containing the amino acid sequence. Such conservative modifications include amino acid substitutions, additions and deletions. Modifications can be introduced into an antibody or antibody fragment of the invention 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 a CAR of the invention can be replaced with other amino acid residues from the same side chain family and the altered CAR can be tested using the functional assays described herein.

The term "stimulation," refers to a primary response induced by binding of a stimulatory molecule (e.g., a TCR/CD3 complex or CAR) with its cognate ligand (or tumor antigen in the case of a CAR) thereby mediating a signal transduction event, such as, but not limited to, signal transduction via the TCR/CD3 complex or signal transduction via the appropriate NK receptor or signaling domains of the CAR. Stimulation can mediate altered expression of certain molecules.

The term "stimulatory molecule," refers to a molecule expressed by an immune cell (e.g., T cell, NK cell, B cell) that provides the cytoplasmic signaling sequence(s) that regulate activation of the

immune cell in a stimulatory way for at least some aspect of the immune cell signaling pathway. In one aspect, the-signal is a primary signal that is initiated by, for instance, binding of a TCR/CD3 complex with an MHC molecule loaded with peptide, and which leads to mediation of a T cell response, including, but not limited to, proliferation, activation, differentiation, and the like. A primary cytoplasmic signaling sequence (also referred to as a "primary signaling domain") that acts in a stimulatory manner may contain a signaling motif which is known as immunoreceptor tyrosine-based activation motif or ITAM. Examples of an ITAM containing-cytoplasmic signaling sequence that is of particular use in the invention include, but are not limited to, those derived from CD3 zeta, common FcR gamma (FCER1G), Fc gamma RIIa, FcR beta (Fc Epsilon R1b), CD3 gamma, CD3 delta, CD3 epsilon, CD79a, CD79b, DAP10, and DAP12. In a specific CAR of the invention, the intracellular signaling domain in any one or more CARS of the invention comprises an intracellular signaling sequence, e.g., a primary signaling sequence of CD3-zeta. In a specific CAR of the invention, the primary signaling sequence of CD3-zeta is the sequence provided as SEQ ID NO: 18, or the equivalent residues from a non-human species, e.g., mouse, rodent, monkey, ape and the like. In a specific CAR of the invention, the primary signaling sequence of CD3-zeta is the sequence as provided in SEQ ID NO:20, or the equivalent residues from a non-human species, e.g., mouse, rodent, monkey, ape and the like.

The term "antigen presenting cell" or "APC" refers to an immune system cell such as an accessory cell (e.g., a B-cell, a dendritic cell, and the like) that displays a foreign antigen complexed with major histocompatibility complexes (MHC's) on its surface. T-cells may recognize these complexes using their T-cell receptors (TCRs). APCs process antigens and present them to T-cells.

An "intracellular signaling domain," as the term is used herein, refers to an intracellular portion of a molecule. The intracellular signaling domain generates a signal that promotes an immune effector function of the CAR containing cell, e.g., a CART cell. Examples of immune effector function, e.g., in a CART cell, include cytolytic activity and helper activity, including the secretion of cytokines.

In an embodiment, the intracellular signaling domain can comprise a primary intracellular signaling domain. Exemplary primary intracellular signaling domains include those derived from the molecules responsible for primary stimulation, or antigen dependent simulation. In an embodiment, the intracellular signaling domain can comprise a costimulatory intracellular domain. Exemplary costimulatory intracellular signaling domains include those derived from molecules responsible for costimulatory signals, or antigen independent stimulation. For example, in the case of a CART, a primary intracellular signaling domain can comprise a cytoplasmic sequence of a T cell receptor, and a

costimulatory intracellular signaling domain can comprise cytoplasmic sequence from co-receptor or costimulatory molecule.

A primary intracellular signaling domain can comprise a signaling motif which is known as an immunoreceptor tyrosine-based activation motif or ITAM. Examples of ITAM containing primary cytoplasmic signaling sequences include, but are not limited to, those derived from CD3 zeta, common FcR gamma (FCER1G), Fc gamma RIIa, FcR beta (Fc Epsilon R1b), CD3 gamma, CD3 delta, CD3 epsilon, CD79a, CD79b, DAP10, and DAP12.

The term "zeta" or alternatively "zeta chain", "CD3-zeta" (or "CD3zeta , CD3 zeta or CD3z) or "TCR-zeta" is defined as the protein provided as GenBan Acc. No. BAG36664.1, or the equivalent residues from a non-human species, e.g., mouse, rodent, monkey, ape and the like, and a "zeta stimulatory domain" or alternatively a "CD3-zeta stimulatory domain" or a "TCR-zeta stimulatory domain" is defined as the amino acid residues from the cytoplasmic domain of the zeta chain, or functional derivatives thereof, that are sufficient to functionally transmit an initial signal necessary for T cell activation. In one aspect the cytoplasmic domain of zeta comprises residues 52 through 164 of GenBank Acc. No. BAG36664.1 or the equivalent residues from a non-human species, e.g., mouse, rodent, monkey, ape and the like, that are functional orthologs thereof. In one aspect, the "zeta stimulatory domain" or a "CD3-zeta stimulatory domain" is the sequence provided as SEQ ID NO:18. In one aspect, the "zeta stimulatory domain" or a "CD3-zeta stimulatory domain" is the sequence provided as SEQ ID NO:20.

The term a "costimulatory molecule" refers to a cognate binding partner on a T cell that specifically binds with a costimulatory ligand, thereby mediating a costimulatory response by the T cell, such as, but not limited to, proliferation. Costimulatory molecules are cell surface molecules other than antigen receptors or their ligands that are contribute to an efficient immune response. Costimulatory molecules include, but are not limited to an MHC class I molecule, BTLA and a Toll ligand receptor, as well as OX40, CD27, CD28, CDS, ICAM-1, LFA-1 (CD11a/CD18), ICOS (CD278), and 4-1BB (CD137). Further examples of such costimulatory molecules include CDS, ICAM-1, GITR, BAFFR, HVEM (LIGHTR), SLAMF7, NKp80 (KLRF1), NKp44, NKp30, NKp46, CD160, CD19, CD4, CD8alpha, CD8beta, IL2R beta, IL2R gamma, IL7R alpha, ITGA4, VLA1, CD49a, ITGA4, IA4, CD49D, ITGA6, VLA-6, CD49f, ITGAD, CD11d, ITGAE, CD103, ITGAL, CD11a, LFA-1, ITGAM, CD11b, ITGAX, CD11c, ITGB1, CD29, ITGB2, CD18, LFA-1, ITGB7, NKG2D, NKG2C, TNFR2, TRANCE/RANKL, DNAM1 (CD226), SLAMF4 (CD244, 2B4), CD84, CD96 (Tactile), CEACAM1, CRTAM, Ly9 (CD229), CD160 (BY55), PSGL1, CD100 (SEMA4D), CD69, SLAMF6 (NTB-A, Ly108),

SLAM (SLAMF1, CD150, IPO-3), BLAME (SLAMF8), SELPLG (CD162), LTBR, LAT, GADS, SLP-76, PAG/Cbp, CD19a, and a ligand that specifically binds with CD83.

A costimulatory intracellular signaling domain can be the intracellular portion of a costimulatory molecule. A costimulatory molecule can be represented in the following protein families: TNF receptor proteins, Immunoglobulin-like proteins, cytokine receptors, integrins, signaling lymphocytic activation molecules (SLAM proteins), and activating NK cell receptors. Examples of such molecules include CD27, CD28, 4-1BB (CD137), OX40, GITR, CD30, CD40, ICOS, BAFFR, HVEM, ICAM-1, lymphocyte function-associated antigen-1 (LFA-1), CD2, CDS, CD7, CD287, LIGHT, NKG2C, NKG2D, SLAMF7, NKp80, NKp30, NKp44, NKp46, CD160, B7-H3, and a ligand that specifically binds with CD83, and the like.

The intracellular signaling domain can comprise the entire intracellular portion, or the entire native intracellular signaling domain, of the molecule from which it is derived, or a functional fragment or derivative thereof.

The term "4-1BB" refers to a member of the TNFR superfamily with an amino acid sequence provided as GenBank Acc. No. AAA62478.2, or the equivalent residues from a non-human species, e.g., mouse, rodent, monkey, ape and the like; and a "4-1BB costimulatory domain" is defined as amino acid residues 214-255 of GenBank Acc. No. AAA62478.2, or the equivalent residues from a non-human species, e.g., mouse, rodent, monkey, ape and the like. In one aspect, the "4-1BB costimulatory domain" is the sequence provided as SEQ ID NO:14 or the equivalent residues from a non-human species, e.g., mouse, rodent, monkey, ape and the like.

"Immune effector cell," as that term is used herein, refers to a cell that is involved in an immune response, e.g., in the promotion of an immune effector response. Examples of immune effector cells include T cells, e.g., alpha/beta T cells and gamma/delta T cells, B cells, natural killer (NK) cells, natural killer T (NKT) cells, mast cells, and myeloic-derived phagocytes.

"Immune effector function or immune effector response," as that term is used herein, refers to function or response, e.g., of an immune effector cell, that enhances or promotes an immune attack of a target cell. E.g., an immune effector function or response refers a property of a T or NK cell that promotes killing or inhibition of growth or proliferation, of a target cell. In the case of a T cell, primary stimulation and co-stimulation are examples of immune effector function or response.

The term "encoding" refers to the inherent property of specific sequences of nucleotides in a polynucleotide, such as a gene, a cDNA, or an mRNA, to serve as templates for synthesis of other

polymers and macromolecules in biological processes having either a defined sequence of nucleotides (e.g., rRNA, tRNA and mRNA) or a defined sequence of amino acids and the biological properties resulting therefrom. Thus, a gene, cDNA, or RNA, encodes a protein if transcription and translation of mRNA corresponding to that gene produces the protein in a cell or other biological system. Both the coding strand, the nucleotide sequence of which is identical to the mRNA sequence and is usually provided in sequence listings, and the non-coding strand, used as the template for transcription of a gene or cDNA, can be referred to as encoding the protein or other product of that gene or cDNA.

Unless otherwise specified, a “nucleotide sequence encoding an amino acid sequence” includes all nucleotide sequences that are degenerate versions of each other and that encode the same amino acid sequence. The phrase nucleotide sequence that encodes a protein or a RNA may also include introns to the extent that the nucleotide sequence encoding the protein may in some version contain an intron(s).

The term “effective amount” or “therapeutically effective amount” are used interchangeably herein, and refer to an amount of a compound, formulation, material, or composition, as described herein effective to achieve a particular biological result.

The term “endogenous” refers to any material from or produced inside an organism, cell, tissue or system.

The term “exogenous” refers to any material introduced from or produced outside an organism, cell, tissue or system.

The term “expression” refers to the transcription and/or translation of a particular nucleotide sequence driven by a promoter.

The term “transfer vector” refers to a composition of matter which comprises an isolated nucleic acid and which can be used to deliver the isolated nucleic acid to the interior of a cell. Numerous vectors are known in the art including, but not limited to, linear polynucleotides, polynucleotides associated with ionic or amphiphilic compounds, plasmids, and viruses. Thus, the term “transfer vector” includes an autonomously replicating plasmid or a virus. The term should also be construed to further include non-plasmid and non-viral compounds which facilitate transfer of nucleic acid into cells, such as, for example, a polylysine compound, liposome, and the like. Examples of viral transfer vectors include, but are not limited to, adenoviral vectors, adeno-associated virus vectors, retroviral vectors, lentiviral vectors, and the like.

The term “expression vector” refers to a vector comprising a recombinant polynucleotide comprising expression control sequences operatively linked to a nucleotide sequence to be expressed. An expression vector comprises sufficient cis-acting elements for expression; other elements for expression can be supplied by the host cell or in an in vitro expression system. Expression vectors include all those known in the art, including cosmids, plasmids (e.g., naked or contained in liposomes) and viruses (e.g., lentiviruses, retroviruses, adenoviruses, and adeno-associated viruses) that incorporate the recombinant polynucleotide.

The term “lentivirus” refers to a genus of the Retroviridae family. Lentiviruses are unique among the retroviruses in being able to infect non-dividing cells; they can deliver a significant amount of genetic information into the DNA of the host cell, so they are one of the most efficient methods of a gene delivery vector. HIV, SIV, and FIV are all examples of lentiviruses.

The term “lentiviral vector” refers to a vector derived from at least a portion of a lentivirus genome, including especially a self-inactivating lentiviral vector as provided in Milone et al., Mol. Ther. 17(8): 1453–1464 (2009). Other examples of lentivirus vectors that may be used in the clinic, include but are not limited to, e.g., the LENTIVECTOR® gene delivery technology from Oxford BioMedica, the LENTIMAX™ vector system from Lentigen and the like. Nonclinical types of lentiviral vectors are also available and would be known to one skilled in the art.

The term “homologous” or “identity” refers to the subunit sequence identity between two polymeric molecules, e.g., between two nucleic acid molecules, such as, two DNA molecules or two RNA molecules, or between two polypeptide molecules. When a subunit position in both of the two molecules is occupied by the same monomeric subunit; e.g., if a position in each of two DNA molecules is occupied by adenine, then they are homologous or identical at that position. The homology between two sequences is a direct function of the number of matching or homologous positions; e.g., if half (e.g., five positions in a polymer ten subunits in length) of the positions in two sequences are homologous, the two sequences are 50% homologous; if 90% of the positions (e.g., 9 of 10), are matched or homologous, the two sequences are 90% homologous.

“Humanized” forms of non-human (e.g., murine) antibodies are chimeric immunoglobulins, immunoglobulin chains or fragments thereof (such as Fv, Fab, Fab', F(ab')2 or other antigen-binding subsequences of antibodies) which contain minimal sequence derived from non-human immunoglobulin. For the most part, humanized antibodies and antibody fragments thereof are human immunoglobulins (recipient antibody or antibody fragment) in which residues from a complementary-

determining region (CDR) of the recipient are replaced by residues from a CDR of a non-human species (donor antibody) such as mouse, rat or rabbit having the desired specificity, affinity, and capacity. In some instances, Fv framework region (FR) residues of the human immunoglobulin are replaced by corresponding non-human residues. Furthermore, a humanized antibody/antibody fragment can comprise residues which are found neither in the recipient antibody nor in the imported CDR or framework sequences. These modifications can further refine and optimize antibody or antibody fragment performance. In general, the humanized antibody or antibody fragment thereof will comprise substantially all of at least one, and typically two, variable domains, in which all or substantially all of the CDR regions correspond to those of a non-human immunoglobulin and all or a significant portion of the FR regions are those of a human immunoglobulin sequence. The humanized antibody or antibody fragment can also comprise at least a portion of an immunoglobulin constant region (Fc), typically that of a human immunoglobulin. For further details, see Jones et al., Nature, 321: 522-525, 1986; Reichmann et al., Nature, 332: 323-329, 1988; Presta, Curr. Op. Struct. Biol., 2: 593-596, 1992.

"Fully human" refers to an immunoglobulin, such as an antibody or antibody fragment, where the whole molecule is of human origin or consists of an amino acid sequence identical to a human form of the antibody or immunoglobulin.

The term "isolated" means altered or removed from the natural state. For example, a nucleic acid or a peptide naturally present in a living animal is not "isolated," but the same nucleic acid or peptide partially or completely separated from the coexisting materials of its natural state is "isolated." An isolated nucleic acid or protein can exist in substantially purified form, or can exist in a non-native environment such as, for example, a host cell.

The term "operably linked" or "transcriptional control" refers to functional linkage between a regulatory sequence and a heterologous nucleic acid sequence resulting in expression of the latter. For example, a first nucleic acid sequence is operably linked with a second nucleic acid sequence when the first nucleic acid sequence is placed in a functional relationship with the second nucleic acid sequence. For instance, a promoter is operably linked to a coding sequence if the promoter affects the transcription or expression of the coding sequence. Operably linked DNA sequences can be contiguous with each other and, e.g., where necessary to join two protein coding regions, are in the same reading frame.

The term “parenteral” administration of an immunogenic composition includes, e.g., subcutaneous (s.c.), intravenous (i.v.), intramuscular (i.m.), or intrasternal injection, intratumoral, or infusion techniques.

The term “nucleic acid” or “polynucleotide” refers to deoxyribonucleic acids (DNA) or ribonucleic acids (RNA) and polymers thereof in either single- or double-stranded form. Unless specifically limited, the term encompasses nucleic acids containing known analogues of natural nucleotides that have similar binding properties as the reference nucleic acid and are metabolized in a manner similar to naturally occurring nucleotides. Unless otherwise indicated, a particular nucleic acid sequence also implicitly encompasses conservatively modified variants thereof (e.g., degenerate codon substitutions), alleles, orthologs, SNPs, and complementary sequences as well as the sequence explicitly indicated. Specifically, degenerate codon substitutions may be achieved by generating sequences in which the third position of one or more selected (or all) codons is substituted with mixed-base and/or deoxyinosine residues (Batzer et al., Nucleic Acid Res. 19:5081 (1991); Ohtsuka et al., J. Biol. Chem. 260:2605-2608 (1985); and Rossolini et al., Mol. Cell. Probes 8:91-98 (1994)).

The terms “peptide,” “polypeptide,” and “protein” are used interchangeably, and refer to a compound comprised of amino acid residues covalently linked by peptide bonds. A protein or peptide must contain at least two amino acids, and no limitation is placed on the maximum number of amino acids that can comprise a protein’s or peptide’s sequence. Polypeptides include any peptide or protein comprising two or more amino acids joined to each other by peptide bonds. As used herein, the term refers to both short chains, which also commonly are referred to in the art as peptides, oligopeptides and oligomers, for example, and to longer chains, which generally are referred to in the art as proteins, of which there are many types. “Polypeptides” include, for example, biologically active fragments, substantially homologous polypeptides, oligopeptides, homodimers, heterodimers, variants of polypeptides, modified polypeptides, derivatives, analogs, fusion proteins, among others. A polypeptide includes a natural peptide, a recombinant peptide, or a combination thereof.

The term “promoter” refers to a DNA sequence recognized by the synthetic machinery of the cell, or introduced synthetic machinery, required to initiate the specific transcription of a polynucleotide sequence.

The term “promoter/regulatory sequence” refers to a nucleic acid sequence which is required for expression of a gene product operably linked to the promoter/regulatory sequence. In some instances, this sequence may be the core promoter sequence and in other instances, this sequence may

also include an enhancer sequence and other regulatory elements which are required for expression of the gene product. The promoter/regulatory sequence may, for example, be one which expresses the gene product in a tissue specific manner.

The term “constitutive” promoter refers to a nucleotide sequence which, when operably linked with a polynucleotide which encodes or specifies a gene product, causes the gene product to be produced in a cell under most or all physiological conditions of the cell.

The term “inducible” promoter refers to a nucleotide sequence which, when operably linked with a polynucleotide which encodes or specifies a gene product, causes the gene product to be produced in a cell substantially only when an inducer which corresponds to the promoter is present in the cell.

The term “tissue-specific” promoter refers to a nucleotide sequence which, when operably linked with a polynucleotide encodes or specified by a gene, causes the gene product to be produced in a cell substantially only if the cell is a cell of the tissue type corresponding to the promoter.

The terms “cancer associated antigen” or “tumor antigen” interchangeably refers to a molecule (typically a protein, carbohydrate or lipid) that is expressed on the surface of a cancer cell, either entirely or as a fragment (e.g., MHC/peptide), and which is useful for the preferential targeting of a pharmacological agent to the cancer cell. In some embodiments, a tumor antigen is a marker expressed by both normal cells and cancer cells, e.g., a lineage marker, e.g., CD19 on B cells. In some embodiments, a tumor antigen is a cell surface molecule that is overexpressed in a cancer cell in comparison to a normal cell, for instance, 1-fold over expression, 2-fold overexpression, 3-fold overexpression or more in comparison to a normal cell. In some embodiments, a tumor antigen is a cell surface molecule that is inappropriately synthesized in the cancer cell, for instance, a molecule that contains deletions, additions or mutations in comparison to the molecule expressed on a normal cell. In some embodiments, a tumor antigen will be expressed exclusively on the cell surface of a cancer cell, entirely or as a fragment (e.g., MHC/peptide), and not synthesized or expressed on the surface of a normal cell. In some embodiments, the CARs of the present invention includes CARs comprising an antigen binding domain (e.g., antibody or antibody fragment) that binds to a MHC presented peptide. Normally, peptides derived from endogenous proteins fill the pockets of Major histocompatibility complex (MHC) class I molecules, and are recognized by T cell receptors (TCRs) on CD8 + T lymphocytes. The MHC class I complexes are constitutively expressed by all nucleated cells. In cancer, virus-specific and/or tumor-specific peptide/MHC complexes represent a unique class of cell surface targets for

immunotherapy. TCR-like antibodies targeting peptides derived from viral or tumor antigens in the context of human leukocyte antigen (HLA)-A1 or HLA-A2 have been described (see, e.g., Sastry et al., J Virol. 2011 85(5):1935-1942; Sergeeva et al., Blood, 2011 117(16):4262-4272; Verma et al., J Immunol 2010 184(4):2156-2165; Willemsen et al., Gene Ther 2001 8(21) :1601-1608; Dao et al., Sci Transl Med 2013 5(176) :176ra33; Tassev et al., Cancer Gene Ther 2012 19(2):84-100). For example, TCR-like antibody can be identified from screening a library, such as a human scFv phage displayed library.

The term "tumor-supporting antigen" or "cancer-supporting antigen" interchangeably refer to a molecule (typically a protein, carbohydrate or lipid) that is expressed on the surface of a cell that is, itself, not cancerous, but supports the cancer cells, e.g., by promoting their growth or survival e.g., resistance to immune cells. Exemplary cells of this type include stromal cells and myeloid-derived suppressor cells (MDSCs). The tumor-supporting antigen itself need not play a role in supporting the tumor cells so long as the antigen is present on a cell that supports cancer cells.

The term "flexible polypeptide linker" or "linker" as used in the context of a scFv refers to a peptide linker that consists of amino acids such as glycine and/or serine residues used alone or in combination, to link variable heavy and variable light chain regions together. In one embodiment, the flexible polypeptide linker is a Gly/Ser linker and comprises the amino acid sequence (Gly-Gly-Gly-Ser)n, where n is a positive integer equal to or greater than 1. For example, n=1, n=2, n=3. n=4, n=5 and n=6, n=7, n=8, n=9 and n=10 (SEQ ID NO:28). In one embodiment, the flexible polypeptide linkers include, but are not limited to, (Gly4 Ser)4 (SEQ ID NO:29) or (Gly4 Ser)3 (SEQ ID NO:30). In another embodiment, the linkers include multiple repeats of (Gly2Ser), (GlySer) or (Gly3Ser) (SEQ ID NO:31). Also included within the scope of the invention are linkers described in WO2012/138475, incorporated herein by reference.

As used herein, a 5' cap (also termed an RNA cap, an RNA 7-methylguanosine cap or an RNA m^7G cap) is a modified guanine nucleotide that has been added to the "front" or 5' end of a eukaryotic messenger RNA shortly after the start of transcription. The 5' cap consists of a terminal group which is linked to the first transcribed nucleotide. Its presence is critical for recognition by the ribosome and protection from RNases. Cap addition is coupled to transcription, and occurs co-transcriptionally, such that each influences the other. Shortly after the start of transcription, the 5' end of the mRNA being synthesized is bound by a cap-synthesizing complex associated with RNA polymerase. This enzymatic complex catalyzes the chemical reactions that are required for mRNA capping. Synthesis proceeds as a

multi-step biochemical reaction. The capping moiety can be modified to modulate functionality of mRNA such as its stability or efficiency of translation.

As used herein, “in vitro transcribed RNA” refers to RNA, preferably mRNA, that has been synthesized in vitro. Generally, the in vitro transcribed RNA is generated from an in vitro transcription vector. The in vitro transcription vector comprises a template that is used to generate the in vitro transcribed RNA.

As used herein, a “poly(A)” is a series of adenosines attached by polyadenylation to the mRNA. In a preferred embodiment of a construct for transient expression, the polyA is between 50 and 5000 (SEQ ID NO: 34), preferably greater than 64, more preferably greater than 100, most preferably greater than 300 or 400 poly(A) sequences can be modified chemically or enzymatically to modulate mRNA functionality such as localization, stability or efficiency of translation.

As used herein, “polyadenylation” refers to the covalent linkage of a polyadenylyl moiety, or its modified variant, to a messenger RNA molecule. In eukaryotic organisms, most messenger RNA (mRNA) molecules are polyadenylated at the 3' end. The 3' poly(A) tail is a long sequence of adenine nucleotides (often several hundred) added to the pre-mRNA through the action of an enzyme, polyadenylate polymerase. In higher eukaryotes, the poly(A) tail is added onto transcripts that contain a specific sequence, the polyadenylation signal. The poly(A) tail and the protein bound to it aid in protecting mRNA from degradation by exonucleases. Polyadenylation is also important for transcription termination, export of the mRNA from the nucleus, and translation. Polyadenylation occurs in the nucleus immediately after transcription of DNA into RNA, but additionally can also occur later in the cytoplasm. After transcription has been terminated, the mRNA chain is cleaved through the action of an endonuclease complex associated with RNA polymerase. The cleavage site is usually characterized by the presence of the base sequence AAUAAA near the cleavage site. After the mRNA has been cleaved, adenosine residues are added to the free 3' end at the cleavage site.

As used herein in connection with expression, e.g., expression of a CAR molecule, “transient” refers to expression of a non-integrated transgene for a period of hours, days or weeks, wherein the period of time of expression is less than the period of time for expression of the gene if integrated into the genome or contained within a stable plasmid replicon in the host cell.

As used herein in connection with an effect, e.g., an effect of an LSD1 inhibitor, “transient” means the effect is present for a period of, for example, hours, days, weeks or months, but diminishes

(e.g., until the effect is no longer measurable) over a period of time. In embodiments the effect is as measured according to the assays described herein, e.g., in the examples.

As used herein, the terms “treat”, “treatment” and “treating” refer to the reduction or amelioration of the progression, severity and/or duration of a proliferative disorder, or the amelioration of one or more symptoms (preferably, one or more discernible symptoms) of a proliferative disorder resulting from the administration of one or more therapies (e.g., one or more therapeutic agents such as a CAR of the invention). In specific embodiments, the terms “treat”, “treatment” and “treating” refer to the amelioration of at least one measurable physical parameter of a proliferative disorder, such as growth of a tumor, not necessarily discernible by the patient. In other embodiments the terms “treat”, “treatment” and “treating” -refer to the inhibition of the progression of a proliferative disorder, either physically by, e.g., stabilization of a discernible symptom, physiologically by, e.g., stabilization of a physical parameter, or both. In other embodiments the terms “treat”, “treatment” and “treating” refer to the reduction or stabilization of tumor size or cancerous cell count.

The term “signal transduction pathway” refers to the biochemical relationship between a variety of signal transduction molecules that play a role in the transmission of a signal from one portion of a cell to another portion of a cell. The phrase “cell surface receptor” includes molecules and complexes of molecules capable of receiving a signal and transmitting signal across the membrane of a cell.

The term “subject” is intended to include living organisms in which an immune response can be elicited (e.g., mammals, human).

The term, a “substantially purified” cell refers to a cell that is essentially free of other cell types. A substantially purified cell also refers to a cell which has been separated from other cell types with which it is normally associated in its naturally occurring state. In some instances, a population of substantially purified cells refers to a homogenous population of cells. In other instances, this term refers simply to cell that have been separated from the cells with which they are naturally associated in their natural state. In some aspects, the cells are cultured in vitro. In other aspects, the cells are not cultured in vitro.

The term “therapeutic” as used herein means a treatment. A therapeutic effect is obtained by reduction, suppression, remission, or eradication of a disease state.

The term “prophylaxis” as used herein means the prevention of or protective treatment for a disease or disease state.

The term “transfected” or “transformed” or “transduced” refers to a process by which exogenous nucleic acid is transferred or introduced into the host cell. A “transfected” or “transformed” or “transduced” cell is one which has been transfected, transformed or transduced with exogenous nucleic acid. The cell includes the primary subject cell and its progeny.

The term “specifically binds,” refers to an antibody, or a ligand, which recognizes and binds with a binding partner (e.g., a tumor antigen) protein present in a sample, but which antibody or ligand does not substantially recognize or bind other molecules in the sample.

“Regulatable chimeric antigen receptor (RCAR),”as that term is used herein, refers to a set of polypeptides, typically two in the simplest embodiments, which when in a RCARX cell, provides the RCARX cell with specificity for a target cell, typically a cancer cell, and with regulatable intracellular signal generation or proliferation, which can optimize an immune effector property of the RCARX cell. An RCARX cell relies at least in part, on an antigen binding domain to provide specificity to a target cell that comprises the antigen bound by the antigen binding domain. In an embodiment, an RCAR includes a dimerization switch that, upon the presence of a dimerization molecule, can couple an intracellular signaling domain to the antigen binding domain.

“Membrane anchor” or “membrane tethering domain”, as that term is used herein, refers to a polypeptide or moiety, e.g., a myristoyl group, sufficient to anchor an extracellular or intracellular domain to the plasma membrane.

“Switch domain,” as that term is used herein, e.g., when referring to an RCAR, refers to an entity, typically a polypeptide-based entity, that, in the presence of a dimerization molecule, associates with another switch domain. The association results in a functional coupling of a first entity linked to, e.g., fused to, a first switch domain, and a second entity linked to, e.g., fused to, a second switch domain. A first and second switch domain are collectively referred to as a dimerization switch. In embodiments, the first and second switch domains are the same as one another, e.g., they are polypeptides having the same primary amino acid sequence, and are referred to collectively as a homodimerization switch. In embodiments, the first and second switch domains are different from one another, e.g., they are polypeptides having different primary amino acid sequences, and are referred to collectively as a heterodimerization switch. In embodiments, the switch is intracellular. In embodiments, the switch is extracellular. In embodiments, the switch domain is a polypeptide-based entity, e.g., FKBP or FRB-based, and the dimerization molecule is small molecule, e.g., a rapalogue. In embodiments, the switch domain is a polypeptide-based entity, e.g., an scFv that binds a myc peptide,

and the dimerization molecule is a polypeptide, a fragment thereof, or a multimer of a polypeptide, e.g., a myc ligand or multimers of a myc ligand that bind to one or more myc scFvs. In embodiments, the switch domain is a polypeptide-based entity, e.g., myc receptor, and the dimerization molecule is an antibody or fragments thereof, e.g., myc antibody.

"Dimerization molecule," as that term is used herein, e.g., when referring to an RCAR, refers to a molecule that promotes the association of a first switch domain with a second switch domain. In embodiments, the dimerization molecule does not naturally occur in the subject, or does not occur in concentrations that would result in significant dimerization. In embodiments, the dimerization molecule is a small molecule, e.g., rapamycin or a rapalogue, e.g, RAD001.

The term "bioequivalent" refers to an amount of an agent other than the reference compound, required to produce an effect equivalent to the effect produced by the reference dose or reference amount of the reference compound. In an embodiment the effect is the level of LSD1 inhibition, e.g., as measured by LSD1 protein levels, e.g., as evaluated in an in vivo or in vitro assay, e.g., as measured by an assay described herein, e.g., flow cytometry. In an embodiment, the effect is enhanced proliferation of T_{SCM} cells, e.g., CD45RA+CD62L+ cells, e.g., e.g., enhanced proliferation relative to other T cell phenotypes, e.g., T_{CM} (e.g., CD45RA-CD62L+), T_{EM} (e.g., CD45RA-CD62L-), T_{EFF} or T_{REG} cells, e.g., as measured by cell sorting. In an embodiment, the effect is enhanced proliferation of T_{SCM} cells, e.g., CD45RA+CCR7+ cells, e.g., e.g., enhanced proliferation relative to other T cell phenotypes, e.g., CD45RA-CCR7+, CD45RA-CCR7-, CD45RA+CCR7-, T_{EFF} or T_{REG} cells, e.g., as measured by cell sorting.

"Refractory" as used herein refers to a disease, e.g., cancer, that does not respond to a treatment. In embodiments, a refractory cancer can be resistant to a treatment before or at the beginning of the treatment. In other embodiments, the refractory cancer can become resistant during a treatment. A refractory cancer is also called a resistant cancer.

"Relapsed" as used herein refers to the return of a disease (e.g., cancer) or the signs and symptoms of a disease such as cancer after a period of improvement, e.g., after prior treatment of a therapy, e.g., cancer therapy

"LSD1," "lysine-specific demethylase 1A," "Lysine-specific histone demethylase 1A," "KDM1A," "AOF2," "KIAA0601" and "BHC110" are used interchangeably herein, and refer to the gene KDM1A (lysine-specific demethylase 1A) and the protein encoded by said gene, lysine-specific demethylase 1A (LSD1). This gene encodes a nuclear protein containing a SWIRM domain, a FAD-binding motif, and an amine oxidase domain. This protein is a component of several histone deacetylase complexes, through it

silences genes by functioning as a histone demethylase. In the human genome, KDM1A is located on chromosome 1 Chr1:23030596 (on Assembly GRCh38). Currently two isoforms of LSD1 are known, and the isoforms are described in GeneBank number NM_001009999.2 and NM_015013.3.

Examples of the protein sequence of human LSD1 is provided as UniProt accession number O60341-1 having an amino acid sequence as follows:

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          10         20         30         40         50
MLSGKKAAAA AAAAAAAATG TEAGPGTAGG SENGSEVAAQ PAGLSGPAEV
          60         70         80         90        100
GPGAVGERTP RKKEPPRASP PGGLAEPPGS AGPQAGPTVV PGSATPMETG
         110        120        130        140        150
IAETPEGRRT SRRKRAKVEY REMDESLANL SEDEYYSEEE RNAKAEKEKK
         160        170        180        190        200
LPPPPPQAPP EEENESEPEE PSGVEGAAFQ SRLPHDRMTS QEAACFPDII
         210        220        230        240        250
SGPQQTQKVF LFIRNRTLQL WLDNPKIQLT FEATLQQLEA PYNSDTVLVH
         260        270        280        290        300
RVHSYLERHG LINFGIYKRI KPLPTKKTGK VIIIGSGVSG LAAARQLQSF
         310        320        330        340        350
GMDVTLLEAR DRVGGRVATF RKGNYVADLG AMVVTGLGGN PMAVVSKQVN
         360        370        380        390        400
MELAKIKQKC PLYEANGQAV PKEKDEMVEQ EFNRLLEATS YLSHQLDFNV
         410        420        430        440        450
LNNKPVSLGQ ALEVVIQLQE KHVKDEQIEH WKKIVKTQEE LKELLNKMVN
         460        470        480        490        500
LKEKIKELHQ QYKEASEVKP PRDITAEFLV KSKHRDLTAL CKEYDELAET
         510        520        530        540        550
QGKLEEKLQE LEANPPSDVY LSSRDRQILD WHFANLEFAN ATPLSTLSLK
         560        570        580        590        600
HWDQDDDFEF TGSHLTVRNG YSCVPVALAE GLDIKLNTAV RQVRYTASGC
         610        620        630        640        650
EVIAVNTRST SQTFIYKCDA VLCTLPLGVL KQQPPAVQFV PPLPEWKTSA
         660        670        680        690        700
VQRMGFGNLN KVVLCFDRVF WDPSVNLFGH VGSTTASRGE LFLFWNLYKA
```

710 720 730 740 750

PILLALVAGE AAGIMENISD DVIVGRCLAI LKGIFGSSAV PQPKETVVSR

760 770 780 790 800

WRADPWARGS YSYVAAGSSG NDYDLMAQPI TPGPSIPGAP QPIPRLFFAG

810 820 830 840 850

EHTIRNYPAT VHGALLSGLR EAGRIADQFL GAMYTLPRQA TPGVPAQQSP

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SM

SEQ ID NO: 40.

The invention also includes other isoforms of LSD1, including Isoform 2, provided as UniProt accession number O60341-2.

Examples of nucleic acid sequences encoding LSD1 are provided below. There are 2 identified isoforms of human LSD1. The mRNA sequences are provided below (In embodiments, in each sequence, T may be replaced with U). In embodiments, LSD1 includes the proteins encoded by each of the sequences below:

Gene ID	Variant	Sequence
KDM1A/LSD1/AOF2; Gene ID: 23028 SEQ ID NO: 41	Isoform A; NCBI Reference Sequence: NM_001009999.2	GGCGCGGCGGGAGCGCGCTTGGCGCGTGCGTACGCG ACGGCGGTTGGCGGCGCGCGGGCAGCGTGAAGCG AGGCGAGGCAAGGCTTTTCGGACCCACGGAGCGACA GAGCGAGCGGCCCCTACGGCCGTCGGCGGCCCGG CGGCCCGAGATGTTATCTGGGAAGAAGGCGGCAGCC GCGGCGGCGGCGGCTGCAGCGGCAGCAACCGGGA CGGAGGCTGGCCCTGGGACAGCAGGCGGCTCCGAGA ACGGGTCTGAGGTGGCCGCGCAGCCCGCGGGCCT GTCGGGCCCAGCCGAGGTCGGGCCGGGGGCGGTGG GGGAGCGCACACCCCGCAAGAAAGAGCCTCCGCGG GCCTCGCCCCCCGGGGGCCTGGCGGAACCGCCGGGG TCCGCAGGGCCTCAGGCCGGCCCTACTGTCGTGC CTGGGTCTGCGACCCCCATGGAAACTGGAATAGCAGA GACTCCGGAGGGGCGTCGGACCAGCCGGCGCAA GCGGGCGAAGGTAGAGTACAGAGAGATGGATGAAA GCTTGGCCAACCTCTCAGAAGATGAGTATTATTCA GAAGAAGAGAGAAATGCCAAAGCAGAGAAGGAAAA GAAGCTTCCCCCACCACCCCCTCAAGCCCCACCTG AGGAAGAAAATGAAAGTGAGCCTGAAGAACCATCGG

		GGCAAGCAGGAGGACTTCAAGACGACAGTTCTGG AGGGTATGGAGACGGCCAAGCATCAGGTGTGGAGGG CGCAGCTTTCCAGAGCCGACTTCCTCATGACCGG ATGACTTCTCAAGAAGCAGCCTGTTTTCCAGATATTAT CAGTGGACCACAACAGACCCAGAAGGTTTTTC TTTTCATTAGAAACCGCACACTGCAGTTGTGGTTGGAT AATCCAAAGATTCAGCTGACATTTGAGGCTAC TCTCCAACAATTAGAAGCACCTTATAACAGTGATACTG TGCTTGTCCACCGAGTTCACAGTTATTTAGAG CGTCATGGTCTTATCAACTTCGGCATCTATAAGAGGAT AAAACCCCTACCAACTAAAAAGACAGGAAAGG TAATTATTATAGGCTCTGGGGTCTCAGGCTTGGCAGC AGCTCGACAGTTACAAAGTTTTGGAATGGATGT CACACTTTTGGAAGCCAGGGATCGTGTGGGTGGACGA GTTGCCACATTTCGCAAAGGAAACTATGTAGCT GATCTTGGAGCCATGGTGGTAACAGGTCTTGGAGGG AATCCTATGGCTGTGGTCAGCAAACAAGTAAATA TGGAACTGGCCAAGATCAAGCAAAAATGCCCACTTTA TGAAGCCAACGGACAAGCTGACACTGTCAAGGT TCCTAAAGAGAAAGATGAAATGGTAGAGCAAGAGTTT AACCGGTTGCTAGAAGCTACATCTTACCTTAGT CATCAACTAGACTTCAATGTCCTCAATAATAAGCCTGT GTCCCTTGGCCAGGCATTGGAAGTTGTCATTC AGTTACAAGAGAAGCATGTCAAAGATGAGCAGATTGA ACATTGGAAGAAGATAGTGAAAACTCAGGAAGA ATTGAAAGAACTTCTTAATAAGATGGTAAATTTGAAA GAGAAAATTAAAGAACTCCATCAGCAATACAAA GAAGCATCTGAAGTAAAGCCACCCAGAGATATTACTG CCGAGTTCTTAGTGAAAAGCAAACACAGGGATC TGACCGCCCTATGCAAGGAATATGATGAATTAGCTGA AACACAAGGAAAGCTAGAAGAAAAACTTCAGGA GTTGGAAGCGAATCCCCCAAGTGATGTATATCTCTCAT CAAGAGACAGACAAATACTTGATTGGCATTTT GCAAATCTTGAATTTGCTAATGCCACACCTCTCTCAACT CTCTCCCTTAAGCACTGGGATCAGGATGATG ACTTTGAGTTCACTGGCAGCCACCTGACAGTAAGGAA TGGCTACTCGTGTGTGCCTGTGGCTTTAGCAGA AGGCCTAGACATTAAACTGAATACAGCAGTGCGACAG GTTCGCTACACGGCTTCAGGATGTGAAGTGATA GCTGTGAATACCCGCTCCACGAGTCAAACCTTTATTTA TAAATGCGACGCAGTTCTCTGTACCCTTCCCC

		TGGGTGTGCTGAAGCAGCAGCCACCAGCCGTTCAGTT TGTGCCACCTCTCCCTGAGTGGAAAACATCTGC AGTCCAAAGGATGGGATTTGGCAACCTTAACAAGGTG GTGTTGTGTTTTGATCGGGTGTTCTGGGATCCA AGTGTCAATTTGTTCGGGCATGTTGGCAGTACGACTG CCAGCAGGGGTGAGCTCTTCCTCTTCTGGAACC TCTATAAAGCTCCAATACTGTTGGCACTAGTGGCAGG AGAAGCTGCTGGTATCATGGAAAACATAAGTGA CGATGTGATTGTTGGCCGATGCCTGGCCATTCTCAAA GGGATTTTTGGTAGCAGTGCAGTACCTCAGCCC AAAGAAACTGTGGTGTCTCGTTGGCGTGCTGATCCCT GGGCTCGGGGCTCTTATTCCTATGTTGCTGCAG GATCATCTGGAAATGACTATGATTTAATGGCTCAGCCA ATCACTCCTGGCCCCTCGATTCCAGGTGCCCC ACAGCCGATTCCACGACTCTTCTTTGCGGGAGAACATA CGATCCGTAACTACCCAGCCACAGTGCATGGT GCTCTGCTGAGTGGGCTGCGAGAAGCGGGAAGAATT GCAGACCAGTTTTTGGGGGCCATGTATACGCTGC CTCGCCAGGCCACACCAGGTGTTCCTGCACAGCAGTC CCCAAGCATGTGAGACAGATGCATTCTAAGGGA AGAGGCCCATGTGCCTGTTTCTGCCATGTAAGGAAGG CTCTTCTAGCAATACTAGATCCCACTGAGAAAA TCCACCCTGGCATCTGGGCTCCTGATCAGCTGATGGA GCTCCTGATTTGACAAAGGAGCTTGCCTCCTTT GAATGACCTAGAGCACAGGGAGGAACTTGTCCATTAG TTTGGAATTGTGTTCTTCGTAAAGACTGAGGCA AGCAAGTGCTGTGAAATAACATCATCTTAGTCCCTTGG TGTGTGGGGTTTTGTTTTTTTTTATATTTT GAGAATAAAACTTCATATAAAATTGGCAAAAAAAAAA AAAAAAAA
KDM1A/LSD1/AOF2; Gene ID: 23028 SEQ ID NO: 42	Isoform B; NCBI Reference Sequence: NM_015013.3	GGCGCGGCGGGAGCGCGCTTGGCGCGTGCGTACGCG ACGGCGGTTGGCGGCGCGCGGGCAGCGTGAAGCG AGGCGAGGCAAGGCTTTTCGGACCCACGGAGCGACA GAGCGAGCGGCCCCTACGGCCGTCGGCGGCCCGG CGGCCCGAGATGTTATCTGGGAAGAAGGCGGCAGCC GCGGCGGCGGCGGCTGCAGCGGCAGCAACCGGGA CGGAGGCTGGCCCTGGGACAGCAGGCGGCTCCGAGA ACGGGTCTGAGGTGGCCGCGCAGCCCGCGGGCCT GTCGGGCCCAGCCGAGGTCGGGCCGGGGGCGGTGG GGGAGCGCACACCCCGCAAGAAAGAGCCTCCGCGG GCCTCGCCCCCCGGGGGCCTGGCGGAACCGCCGGGG

	TCCGCAGGGCCTCAGGCCGGCCCTACTGTCGTGC CTGGGTCTGCGACCCCCATGGAAACTGGAATAGCAGA GACTCCGGAGGGGCGTCGGACCAGCCGGCGCAA GCGGGCGAAGGTAGAGTACAGAGAGATGGATGAAA GCTTGGCCAACCTCTCAGAAGATGAGTATTATTCA GAAGAAGAGAGAAATGCCAAAGCAGAGAAGGAAAA GAAGCTTCCCCCACCACCCCCTCAAGCCCCACCTG AGGAAGAAAATGAAAGTGAGCCTGAAGAACCATCGG GTGTGGAGGGCGCAGCTTTCCAGAGCCGACTTCC TCATGACCGGATGACTTCTCAAGAAGCAGCCTGTTTTC CAGATATTATCAGTGGACCACAACAGACCCAG AAGGTTTTTCTTTTCATTAGAAACCGCACACTGCAGTT GTGGTTGGATAATCCAAAGATTCAGCTGACAT TTGAGGCTACTCTCCAACAATTAGAAGCACCTTATAAC AGTGATACTGTGCTTGTCCACCGAGTTCACAG TTATTTAGAGCGTCATGGTCTTATCAACTTCGGCATCT ATAAGAGGATAAAACCCCTACCAACTAAAAAG ACAGGAAAGGTAATTATTATAGGCTCTGGGGTCTCAG GCTTGGCAGCAGCTCGACAGTTACAAAGTTTTG GAATGGATGTCACACTTTTGGAAGCCAGGGATCGTGT GGGTGGACGAGTTGCCACATTTCGCAAAGGAAA CTATGTAGCTGATCTTGGAGCCATGGTGGTAACAGGT CTTGGAGGGAATCCTATGGCTGTGGTCAGCAAA CAAGTAAATATGGAACTGGCCAAGATCAAGCAAAAAT GCCCACTTTATGAAGCCAACGGACAAGCTGTTC CTAAAGAGAAAGATGAAATGGTAGAGCAAGAGTTTA ACCGGTTGCTAGAAGCTACATCTTACCTTAGTCA TCAACTAGACTTCAATGTCCTCAATAATAAGCCTGTGT CCCTTGGCCAGGCATTGGAAGTTGTCATTCAG TTACAAGAGAAGCATGTCAAAGATGAGCAGATTGAAC ATTGGAAGAAGATAGTGAAAACTCAGGAAGAAT TGAAAGAACTTCTTAATAAGATGGTAAATTTGAAAGA GAAAATTAAAGAACTCCATCAGCAATACAAAGA AGCATCTGAAGTAAAGCCACCCAGAGATATTACTGCC GAGTTCTTAGTGAAAAGCAAACACAGGGATCTG ACCGCCCTATGCAAGGAATATGATGAATTAGCTGAAA CACAAGGAAAGCTAGAAGAAAAACTTCAGGAGT TGGAAGCGAATCCCCCAAGTGATGTATATCTCTCATCA AGAGACAGACAAATACTTGATTGGCATTTTGC AAATCTTGAATTTGCTAATGCCACACCTCTCTCAACTCT CTCCCTTAAGCACTGGGATCAGGATGATGAC

	TTTGAGTTCACTGGCAGCCACCTGACAGTAAGGAATG GCTACTCGTGTGTGCCTGTGGCTTTAGCAGAAG GCCTAGACATTAAACTGAATACAGCAGTGCGACAGGT TCGCTACACGGCTTCAGGATGTGAAGTGATAGC TGTGAATACCCGCTCCACGAGTCAAACCTTTATTTATA AATGCGACGCAGTTCTCTGTACCCTTCCCCTG GGTGTGCTGAAGCAGCAGCCACCAGCCGTTCAGTTTG TGCCACCTCTCCCTGAGTGGAAAACATCTGCAG TCCAAAGGATGGGATTTGGCAACCTTAACAAGGTGGT GTTGTGTTTTGATCGGGTGTTCTGGGATCCAAG TGTCAATTTGTTCGGGCATGTTGGCAGTACGACTGCCA GCAGGGGTGAGCTCTTCCTCTTCTGGAACCTC TATAAAGCTCCAATACTGTTGGCACTAGTGGCAGGAG AAGCTGCTGGTATCATGGAAAACATAAGTGACG ATGTGATTGTTGGCCGATGCCTGGCCATTCTCAAAGG GATTTTTGGTAGCAGTGCAGTACCTCAGCCCAA AGAAACTGTGGTGTCTCGTTGGCGTGCTGATCCCTGG GCTCGGGGCTCTTATTCCTATGTTGCTGCAGGA TCATCTGGAAATGACTATGATTTAATGGCTCAGCCAAT CACTCCTGGCCCCTCGATTCCAGGTGCCCCAC AGCCGATTCCACGACTCTTCTTTGCGGGAGAACATACG ATCCGTAACTACCCAGCCACAGTGCATGGTGC TCTGCTGAGTGGGCTGCGAGAAGCGGGAAGAATTGC AGACCAGTTTTTGGGGGCCATGTATACGCTGCCT CGCCAGGCCACACCAGGTGTTCCTGCACAGCAGTCCC CAAGCATGTGAGACAGATGCATTCTAAGGGAAG AGGCCCATGTGCCTGTTTCTGCCATGTAAGGAAGGCT CTTCTAGCAATACTAGATCCCACTGAGAAAATC CACCCTGGCATCTGGGCTCCTGATCAGCTGATGGAGC TCCTGATTTGACAAAGGAGCTTGCCTCCTTTGA ATGACCTAGAGCACAGGGAGGAACTTGTCCATTAGTT TGGAATTGTGTTCTTCGTAAAGACTGAGGCAAG CAAGTGCTGTGAAATAACATCATCTTAGTCCCTTGGTG TGTGGGGTTTTTGTTTTTTTTTTATATTTTGA GAATAAAACTTCATATAAAATTGGCAAAAAAAAAAAA AAAAAA

"LSD1 inhibitor" as the term is used herein, refers to a molecule, or a group of molecules (e.g., a system) that reduces or eliminates the function and/or expression of LSD1. In embodiments, an LSD1

inhibitor is a molecule that inhibits the expression of LSD1 e.g., reduces or eliminates expression of LSD1. In embodiments, the LSD1 inhibitor is a molecule that inhibits the function of LSD1. An example of an LSD1 inhibitor that inhibits the expression of LSD1 is a gene editing system, e.g., as described herein, that is targeted to nucleic acid within the LSD1 gene (e.g., within the KDM1A gene), or its regulatory elements, such that modification of the nucleic acid at or near the gene editing system binding site(s) is modified to reduce or eliminate expression of LSD1. Another example of an LSD1 inhibitor that inhibits the expression of LSD1 is a nucleic acid molecule, e.g., RNA molecule, e.g., a short hairpin RNA (shRNA) or short interfering RNA (siRNA), capable of hybridizing with LSD1 mRNA and causing a reduction or elimination of LSD1 translation. Another example of an LSD1 inhibitor that inhibits the expression of LSD1 is an antisense oligonucleotide. LSD1 inhibitors also include nucleic acids encoding molecules which inhibit LSD1 expression (e.g., nucleic acid encoding an anti-LSD1 shRNA or siRNA, or nucleic acid encoding one or more, e.g., all, components of an anti-LSD1 gene editing system). An example of a molecule that inhibits the function of LSD1 is a molecule, e.g., a protein or small molecule which inhibits one or more activities of LSD1. An example is a small molecule inhibitor of LSD1, e.g., as described herein. In an exemplary embodiment, a small molecule LSD1 inhibitor is a reversible LSD1 inhibitor. In another exemplary embodiment, a small molecule LSD1 inhibitor is an irreversible LSD1 inhibitor. A small molecule LSD1 inhibitor may bind LSD1 at the catalytic site or at a site other than the catalytic site. Another example is a dominant negative LSD1 protein. Another example is an anti-LSD1 antibody or antigen-binding fragment thereof. Another example is a molecule, e.g., a small molecule, which inhibits an LSD1 binding partner. LSD1 inhibitors also include nucleic acids encoding inhibitors of LSD1 function. Further description of LSD1 inhibitors is provided below in the section titled “LSD1 inhibitors.”

A “binding partner” as the term is used herein in the context of an LSD1 binding partner, refers to a molecule, e.g., a protein, which interacts, e.g., binds to, LSD1 protein. Without being bound by theory, it is believed that LSD1 binds to one or more HDAC proteins, e.g., HDAC1. Such HDAC proteins are considered examples of LSD1 binding partners. Other LSD1 binding partners include, for example, proteins of the Co-REST/REST complex, e.g., HDAC1, HDAC2, p40, p80, Co-REST and ZNF217 (Lee, M. G., Wynder, C., Cooch, N. & Shiekhattar, R. An essential role for CoREST in nucleosomal histone 3 lysine 4 demethylation. Nature 437, 432–435 (2005)); proteins of the Blimp1 complex (Mol Cell Biol. 2009 Mar;29(6):1421-31. doi: 10.1128/MCB.01158-08. Epub 2009 Jan 5); proteins of the NuRD complex (Cell. 2009 Aug. 21;138(4):660-72. doi: 10.1016/j.cell.2009.05.050); and the androgen receptor (Nature. 2005 Sep 15;437(7057):436-9. Epub 2005 Aug 3).

A “system” as the term is used herein in connection with, for example, gene editing, refers to a group of molecules, e.g., one or more molecules, which together act to produce a desired function.

A “gene editing system” as the term is used herein, refers to a system, e.g., one or more molecules, that direct and effect an alteration, e.g., a deletion, of one or more nucleic acids at or near a site of genomic DNA targeted by said system. Gene editing systems are known in the art, and are described more fully below.

A “dominant negative” gene product or protein is one that interferes with the function of a gene product or protein. The gene product affected can be the same or different from the dominant negative protein. Dominant negative gene products can be of many forms, including truncations, full length proteins with point mutations or fragments thereof, or fusions of full length wild type or mutant proteins or fragments thereof with other proteins. The level of inhibition observed can be very low. For example, it may require a large excess of the dominant negative protein compared to the functional protein or proteins involved in a process in order to see an effect. It may be difficult to see effects under normal biological assay conditions. In one embodiment, a dominant negative LSD1 is a catalytically inactive LSD1.

The term “proportion” refers to the ratio of the specified molecule to the total number of molecules in a population. In an exemplary embodiment, a proportion of T cells having a specific phenotype (e.g., T_{SCM} cells) refers to the ratio of the number of T cells having that phenotype relative to the total number of T cells in a population. In an exemplary embodiment, a proportion of T cells having a specific phenotype (e.g., CD45RA+CD62L+ cells) refers to the ratio of the number of T cells having that phenotype relative to the total number of T cells in a population. It will be understood that such proportions may be measured against certain subsets of cells, where indicted. For example, the proportion of CD4+ T_{SCM} cells may be measured against the total number of CD4+ T cells.

The term “population of immune effector cells” as used herein refers to a composition comprising at least two, e.g., two or more, e.g., more than one, immune effector cell, and does not denote any level of purity or the presence or absence of other cell types. In an exemplary embodiment, the population is substantially free of other cell types. In another exemplary embodiment, the population comprises at least two cells of the specified cell type, or having the specified function or property.

The terms "T_{SCM}-like cell," "naive T Cell' and "naïve T cell" are used interchangeably and refer to a less differentiated T cell state, that is characterized by surface expression of CD45RA and CD62L (e.g., is CD45RA positive and CD62L positive (sometimes written as CD45RA+CD62L+)). In general, T cell differentiation proceeds, from most "naive" to most "exhausted," T_{SCM}-like (e.g., a CD45RA+CD62L+ cell) >T_{CM} (e.g., a CD45RA-CD62L+ cell)>T_{EM} (e.g., a CD45RA-CD62L- cell)>T_{EFF}. Naive T cells may be characterized, for example, as having increased self-renewal, anti-tumor efficacy, proliferation and/or survival, relative to a more exhausted T cell phenotype. In an exemplary embodiment, a naive T cell refers to a CD45RA+CD62L+ T cell. In another exemplary embodiment, a naive T cell refers to a T_{SCM} cell, e.g., a CD45RA+CD62L+CCR7+CD27+CD95+ T cell.

The term "T_{SCM}" refers to a T cell having a stem cell memory phenotype, characterized in that it expresses CD45RA, CD62L, CCR7, CD27 and CD95 on its cell surface (e.g., is CD45RA positive, CD62L positive, CCR7 positive, CD27 positive and CD95 positive (sometimes written as CD45RA+CD62L+CCR7+CD27+CD95+)). A T_{SCM} cell is an example of a naive T cell. The T cell may be CD4+ and/or CD8+ T cell.

Ranges: throughout this disclosure, various aspects of the invention can be presented in a range format. It should be understood that the description in range format is merely for convenience and brevity and should not be construed as an inflexible limitation on the scope of the invention. Accordingly, the description of a range should be considered to have specifically disclosed all the possible subranges as well as individual numerical values within that range. For example, description of a range such as from 1 to 6 should be considered to have specifically disclosed subranges such as from 1 to 3, from 1 to 4, from 1 to 5, from 2 to 4, from 2 to 6, from 3 to 6 etc., as well as individual numbers within that range, for example, 1, 2, 2.7, 3, 4, 5, 5.3, and 6. As another example, a range such as 95-99% identity, includes something with 95%, 96%, 97%, 98% or 99% identity, and includes subranges such as 96-99%, 96-98%, 96-97%, 97-99%, 97-98% and 98-99% identity. This applies regardless of the breadth of the range.

Headings, sub-headings or numbered or lettered elements, e.g., (a), (b), (i) etc, are presented merely for ease of reading. The use of headings or numbered or lettered elements in this document does not require the steps or elements be performed in alphabetical order or that the steps or elements are necessarily discrete from one another.

All publications, patent applications, patents, and other references mentioned herein are incorporated by reference in their entirety.

Other features, objects, and advantages of the invention will be apparent from the description and drawings, and from the claims.

LSD1 Inhibitors

Any molecule that inhibits LSD1 may be useful in the aspects of the invention described herein, e.g., in connection with the cells, compositions and methods disclosed herein. The following sections provide exemplary LSD1 inhibitors, and are not intended to be limiting.

In embodiments, the LSD1 inhibitor is a molecule or system that results in increased or prolonged proliferation or persistence of CAR-expressing cells with a naive phenotype (e.g., T_{SCM} cells), e.g., in culture or in a subject, e.g., as compared to non-treated CAR-expressing cells or a non-treated subject. In embodiments, increased proliferation or persistence is associated with in an increase in the number of CAR-expressing cells. Methods for measuring increased or prolonged proliferation are described in Example 4. In another embodiment, administration or contacting with an LSD1 inhibitor results in increased cytokine release or increased killing of cancer cells by CAR-expressing cells, e.g., in culture or in a subject, e.g., as compared to non-treated CAR-expressing cells or a non-treated subject. Methods for measuring increased cytokine release are described in, e.g., Example 4. In embodiments, increased killing of cancer cells is associated with in a decrease in tumor volume. Methods for measuring increased killing of cancer cells are described in Example 4 and, e.g., in International Application WO2014/153270, which is herein incorporated by reference in its entirety.

Nucleic acid inhibitors

In one aspect the LSD1 inhibitor is a nucleic acid molecule. In embodiments, the nucleic acid is a DNA molecule, e.g., an antisense oligonucleotide (e.g., Watts et al., *J. Pathol.*, 2012, 226(2), pp. 365-379). In an embodiment, the antisense oligonucleotide is complementary to an LSD1 mRNA or pre-mRNA molecule. In another aspect, the LSD1 inhibitor includes nucleic acid encoding said antisense oligonucleotide.

In embodiments, the nucleic acid LSD1 inhibitor is an interfering RNA molecule, e.g., a shRNA or siRNA, that inhibits expression, e.g., translation, of LSD1. In another aspect, the LSD1 inhibitor includes nucleic acid encoding said interfering RNA molecule. In embodiments, the interfering RNA molecule, e.g., a shRNA or siRNA, that inhibits expression, e.g., translation, of LSD1 comprises a domain complementary to a sequence of an LSD1 mRNA (such sequence referred to herein in relation to an

interfering RNA molecule, e.g., a shRNA or siRNA, as a “target sequence”). Examples of such target sequences are provided in Table 1.

Exemplary Target Sequences for shRNA and siRNA LSD1 inhibitors, and exemplary nucleic acids encoding shRNA LSD1 inhibitors are provided in Table 1 below,

Table 1

TARGET	SHRNA_ NAME	Target Sequence	SEQ ID NO:	Sequence encoding anti-LSD1 shRNA	SEQ ID NO:
AOF2	KDM1A-917_58263	CCGAGTTCACAGTTATTTAGA	43	CCGAGTTCACAGTTATTTAGACTCGAGTCTAAATAACTGTGAACTCGGTTTTTTGAATTCGCACCAGCACGCTACGCAACACGTTGACGTTGACCACATGTTCGCCGTCTTC	83
AOF2	KDM1A-1265_58264	CGGACAAGCTGTTCCTAAAGA	44	CGGACAAGCTGTTCCTAAAGACTCGAGTCTTTAGGAACAGCTTGTCCGTTTTTTGAATTCGCACCAGCACGCTACGCAGTCAGTTGACGTTGACCAACGTTTCGCCGTCTTC	84
AOF2	KDM1A-2928_58265	GAATTGTGTTCTTCGTAAAGA	45	GAATTGTGTTCTTCGTAAAGACTCGAGTCTTTACGAAGAACACAATTCTTTTTTGAATTCGCACCAGCACGCTACGCATGACGTTGACGTTGACCAACCATTCGCCGTCTTC	85
AOF2	KDM1A-914_58266	CCACCGAGTTCACAGTTATTT	46	CCACCGAGTTCACAGTTATTTCTCGAGAAATAACTGTGAACTCGGTGGTTTTTTGAATTCGCACCAGCACGCTACGCAACGTGTTGACGTTGACCAACACTTCGCCGTCTTC	86
AOF2	KDM1A-1000_58267	CAGGAAAGGTAATTATTATAG	47	CAGGAAAGGTAATTATTATAGCTCGAGCTATAATAATTACCTTTCCTGTTTTTTGAATTCGCACCAGCACGCTACGCACATGGTTGACGTTGACCAACTGTTCGCCGTCTTC	87
AOF2	KDM1A-2759_58268	GCCTGTTTCTGCCATGTAAGG	48	GCCTGTTTCTGCCATGTAAGGCTCGAGCCTTACATGGCAGAAACAGGCTTTTTTGAATTCGCACCAGCACGCTACGCATGACGTTGACGTTGACCCATGGTTTCGCCGTCTTC	88
AOF2	KDM1A-998_58269	GACAGGAAAGGTAATTATTAT	49	GACAGGAAAGGTAATTATTATCTCGAGATAATAATTACCTTTCCTGTCTTTTTTGAATTCGCACCAGCACGCTACGCAACTGGTTGACGTTGACCATGCATTCGCCGTCTTC	89
AOF2	KDM1A-1271_58	AGCTGTTCCTAAAGAGA	50	AGCTGTTCCTAAAGAGAAAGACTCGAGTCTTTCTCTTTAGGAACAGCTTTTTTTGAATTCGCA	90

	270	AAGA		CCAGCACGCTACGCACACAGTTGACGTTGACCATGACTTCGCCGTCTTC	
AOF2	KDM1A-1996_58271	CGAGTCAAACCTTTATTTATA	51	CGAGTCAAACCTTTATTTATACTCGAGTATAAATAAAGGTTTGACTCGTTTTTTGAATTCGCACCAGCACGCTACGCAGTGTGTTGACGTTGACCATGTGTTCGCCGTCTTC	91
AOF2	KDM1A-863_58272	GGCTACTCTCCAACAATTAGA	52	GGCTACTCTCCAACAATTAGACTCGAGTCTAATTGTTGGAGAGTAGCCTTTTTTGAATTCGCACCAGCACGCTACGCAGTACGTTGACGTTGACACGTGTTTCGCCGTCTTC	92
AOF2	KDM1A-1044_58273	GCAGCTCGACAGTTACAAAGT	53	GCAGCTCGACAGTTACAAAGTCTCGAGACTTTGTAACTGTCGAGCTGCTTTTTTGAATTCGCACCAGCACGCTACGCATGTGGTTGACGTTGACACGTCATTCGCCGTCTTC	93
AOF2	KDM1A-1903_58274	CAGAAGGCCTAGACATTAAAC	54	CAGAAGGCCTAGACATTAAACCTCGAGGTTTAATGTCTAGGCCTTCTGTTTTTTGAATTCGCACCAGCACGCTACGCAACCAGTTGACGTTGACACGTACTTCGCCGTCTTC	94
AOF2	KDM1A-563_58275	AGATGAGTATTATTCAGAAGA	55	AGATGAGTATTATTCAGAAGACTCGAGTCTTCTGAATAATACTCATCTTTTTTTGAATTCGCACCAGCACGCTACGCACAGTGTTGACGTTGACACGTTGTTCGCCGTCTTC	95
AOF2	KDM1A-876_58276	CAATTAGAAGCACCTTATAAC	56	CAATTAGAAGCACCTTATAACCTCGAGGTTATAAGGTGCTTCTAATTGTTTTTTGAATTCGCACCAGCACGCTACGCATGTGGTTGACGTTGACACCAGTTTCGCCGTCTTC	96
AOF2	KDM1A-2908_58277	GGAACTTGTCCATTAGTTTGG	57	GGAACTTGTCCATTAGTTTGGCTCGAGCCAAACTAATGGACAAGTTCCTTTTTTGAATTCGCACCAGCACGCTACGCAACGTGTTGACGTTGACACCACATTCGCCGTCTTC	97
AOF2	KDM1A-885_58278	GCACCTTATAACAGTGATACT	58	GCACCTTATAACAGTGATACTCTCGAGAGTATCACTGTTATAAGGTGCTTTTTTGAATTCGCACCAGCACGCTACGCACAACGTTGACGTTGACACCAACTTCGCCGTCTTC	98
AOF2	KDM1A-960_58279	GGCATCTATAAGAGGATAAAA	59	GGCATCTATAAGAGGATAAAACTCGAGTTTTATCCTCTTATAGATGCCTTTTTTGAATTCGCACCAGCACGCTACGCAGTCAGTTGACGTTGACACCATGTTCGCCGTCTTC	99
AOF2	KDM1A-1399_58	CATTGGAAGTTGTCATTC	60	CATTGGAAGTTGTCATTCAGTCTCGAGACTGAATGACAACTTCCAATGTTTTTTGAATTCGCA	100

	280	AGT		CCAGCACGCTACGCAACGTGTTGACGTTGACACACGTTTCGCCGTCTTC	
AOF2	KDM1A-2474_58281	TGGAAATGACTATGATTTAAT	61	TGGAAATGACTATGATTTAATCTCGAGATTAAATCATAGTCATTTCCATTTTTTGAATTCGCACCAGCACGCTACGCACACAGTTGACGTTGACACACCATTCGCCGTCTTC	101
AOF2	KDM1A_NM_015013.3_21174	CGGTTGCTAGAAGCTACATCT	62	GAAGACGCACCGGCGGTTGCTAGAAGTTACATTTGTTAATATTCATAGCAGATGTAGCTTCTAGCAACCGTTTTTTGAATTCGCACCAGCACGCTACGCAACGTGTCACAGTACACTGCAACTTCGCCGT	102
AOF2	KDM1A_NM_015013.3_21173	GCCACATTTCGCAAAGGAAAC	63	GAAGACGCACCGGGCCACATTTCGTAAAGGAAATGTTAATATTCATAGCGTTTCCTTTGCGAAATGTGGCTTTTTTGAATTCGCACCAGCACGCTACGCATGACGTCACAGTACACTGCACATTCGCCGT	103
AOF2	KDM1A_NM_015013.3_21172	CCTCATGACCGGATGACTTCT	64	GAAGACGCACCGGCCTCATGACTGGATGATTTCTGTTAATATTCATAGCAGAAGTCATCCGGTCATGAGGTTTTTTGAATTCGCACCAGCACGCTACGCAGTCAGTCACAGTACACTGCAGTTTCGCCGT	104
AOF2	KDM1A_NM_015013.3_21171	CGAGTTGCCACATTTCGCAAA	65	GAAGACGCACCGGCGAGTTGCCATATTTCGTAAAGTTAATATTCATAGCTTTGCGAAATGTGGCAACTCGTTTTTTGAATTCGCACCAGCACGCTACGCAACACGTCACAGTACACTGGTTGTTCGCCGT	105
AOF2	KDM1A_NM_015013.3_21170	CGTCATGGTCTTATCAACTTC	66	GAAGACGCACCGGCGTCATGGTCTTATCAATTTCGTTAATATTCATAGCGAAGTTGATAAGACCATGACGTTTTTTGAATTCGCACCAGCACGCTACGCATGTGGTCACAGTACACTGGTACTTCGCCGT	106
AOF2	KDM1A_NM_015013.3_21169	GAGCGTCATGGTCTTATCAAC	67	GAAGACGCACCGGGAGCGTTATGGTCTTATTAACGTTAATATTCATAGCGTTGATAAGACCATGACGCTCTTTTTTGAATTCGCACCAGCACGCTACGCAGTCAGTCACAGTACACTGGTCATTCGCCGT	107
AOF2	KDM1A_NM_015013.3_21168	CCGGATGACTTCTCAAGAAGC	68	GAAGACGCACCGGCCGGATGACTTCTTAAGAAGTGTTAATATTCATAGCGCTTCTTGAGAAGTCATCCGGTTTTTTGAATTCGCACCAGCACGCTACGCACAGTGTCACAGTACACTGGTGTTTCGCCGT	108

AOF2	KDM1A_NM_015013.3_21167	CACAGGGAGGAACTTGTCCAT	69	GAAGACGCACCGGCACAGGGAGGAATTTGTCTATGTTAATATTCATAGCATGGACAAGTTCCTCCCTGTGTTTTTGAATTCGCACCAGCACGCTACGCAGTACGTCACAGTACACACTGTGTTCGCCGT	109
AOF2	KDM1A_NM_015013.3_21166	GCCTCCTTTGAATGACCTAGA	70	GAAGACGCACCGGGCCTTCTTTGAATGACTTAGAGTTAATATTCATAGCTCTAGGTCATTCAAAGGAGGCTTTTTTGAATTCGCACCAGCACGCTACGCACATGGTCACAGTACACACTGACTTCGCCGT	110
AOF2	KDM1A_NM_015013.3_21165	CCTATGGCTGTGGTCAGCAAA	71	GAAGACGCACCGGCCTATGGTTGTGGTCAGTAAAGTTAATATTCATAGCTTTGCTGACCACAGCCATAGGTTTTTTGAATTCGCACCAGCACGCTACGCAACCAGTCACAGTACACACTGCATTCGCCGT	111
AOF2	KDM1A_NM_015013.3_21164	GAGCTTGCCTCCTTTGAATGA	72	GAAGACGCACCGGGAGCTTGTGTTCTTTGAATGAGTTAATATTCATAGCTCATTCAAAGGAGGCAAGCTCTTTTTTGAATTCGCACCAGCACGCTACGCATGGTGTCACAGTACACACTGGTTTCGCCGT	112
AOF2	KDM1A_NM_015013.3_21163	CAGGTCTTGGAGGGAATCCTA	73	GAAGACGCACCGGCAGGTCTTGGAGGGGATCTTAGTTAATATTCATAGCTAGGATTCCCTCCAAGACCTGTTTTTTGAATTCGCACCAGCACGCTACGCACACAGTCACAGTACACACACTGTTCGCCGT	113
AOF2	KDM1A_NM_015013.3_21162	CAGTACCTCAGCCCAAAGAAA	74	GAAGACGCACCGGCAGTACTTCAGCCTAAAGAAAGTTAATATTCATAGCTTTCTTTGGGCTGAGGTACTGTTTTTTGAATTCGCACCAGCACGCTACGCAACACGTCACAGTACACACACACTTCGCCGT	114
AOF2	KDM1A_NM_015013.3_21161	CTGGCCATTCTCAAAGGGATT	75	GAAGACGCACCGGCTGGCTATTCTTAAAGGGATTGTTAATATTCATAGCAATCCCTTTGAGAATGGCCAGTTTTTTGAATTCGCACCAGCACGCTACGCATGGTGTCACAGTACACACACCATTCGCCGT	115
AOF2	KDM1A_NM_015013.3_21160	GCCATGGTGGTAACAGGTCTT	76	GAAGACGCACCGGGCCATGGTGGTAATAGGTCTTGTTAATATTCATAGCAAGACCTGTTACCACCATGGCTTTTTTGAATTCGCACCAGCACGCTACGCAGTTGGTCACAGTACACACACGTTTCGCCGT	116
AOF2	KDM1A_	CAGGAGAAGCTGCTGGT	77	GAAGACGCACCGGCAGGAGAAGCTGTTGGT	117

	NM_015013.3_21159	ATCA		ATTAGTTAATATTCATAGCTGATACCAGCAGCTTCTCCTGTTTTTTGAATTCGCACCAGCACGCTACGCAACGTGTCACAGTACACACCATGTTCGCCGT	
AOF2	KDM1A_NM_015013.3_15448	CCACGAGTCAAACCTTTATTT	78	GAAGACGCACCGGCCACGAGTCAAATCTTTATTTGTTAATATTCATAGCAAATAAAGGTTTGACTCGTGGTTTTTTGAATTCGCACCAGCACGCTACGCACACAGTGTTGTGGTCACAACGTTTCGCCGT	118
AOF2	KDM1A_NM_015013.3_15447	GCTACATCTTACCTTAGTCAT	79	GAAGACGCACCGGGCTACATCTTACCTTAGTTATGTTAATATTCATAGCATGACTAAGGTAAGATGTAGCTTTTTTGAATTCGCACCAGCACGCTACGCATGACGTGTTGTGGTCACACATGTTCGCCGT	119
AOF2	KDM1A-725_58282	CCGGATGACTTCTCAAGAAGC	80	CCGGATGACTTCTCAAGAAGCCTCGAGGCTTCTTGAGAAGTCATCCGGTTTTTTGAATTCGCACCAGCACGCTACGCAGTTGGTTGACGTTGACACACACTTCGCCGTCTTC	120
AOF2	KDM1A_NM_015013.3_15445	GCTCCAATACTGTTGGCACTA	81	GAAGACGCACCGGGCTCCAATATTGTTGGCATTAGTTAATATTCATAGCTAGTGCCAACAGTATTGGAGCTTTTTTGAATTCGCACCAGCACGCTACGCACACAGTGTTGTGGTCACACACATTCGCCGT	121
AOF2	KDM1A_NM_015013.3_15446	CCAACAATTAGAAGCACCTTA	82	GAAGACGCACCGGCCAACAATTAGAAGCATCTTAGTTAATATTCATAGCTAAGGTGCTTCTAATTGTTGGTTTTTTGAATTCGCACCAGCACGCTACGCAGTTGGTGTTGTGGTCACACAACTTCGCCGT	122

Nucleic acid LSD1 inhibitor molecules include nucleic acid molecules comprising chemical modifications, e.g., modifications to the nucleic acid base, the sugar and/or the phosphate backbone, including, for example, peptide nucleic acids, phospho morpholino backbones, phosphorothioate backbones, 5' and 3' end caps, 2'-Omethyl modification, 2'-F modifications, and other modifications known in the art.

Genome Editing System LSD1 inhibitors

Genome editing systems are known in the art, and include zinc finger nuclease gene editing systems, TALEN gene editing systems, meganuclease gene editing systems, and CRISPR gene editing systems. As used herein, the term "genome editing system" (used herein synonymously with "gene editing system") refers to a molecule or set of molecules necessary and sufficient to direct modification, e.g., insertion or deletion, of nucleic acids, at or near a site targeted by said system. As the term is used herein, the term "genome editing system" also refers to nucleic acid encoding one or more components (e.g., molecules) of the genome editing system. Exemplary gene editing systems are known in the art, and are described more fully below.

CRISPR Gene Editing Systems

As used herein, the terms "CRISPR System" "CRISPR/Cas System", "CRISPR/Cas gene editing system", "CRISPR/Cas genome editing system", "CRISPR genome editing system" and "CRISPR gene editing system" are used synonymously herein. Naturally-occurring CRISPR systems are found in approximately 40% of sequenced eubacteria genomes and 90% of sequenced archaea. Grissa *et al.* (2007) *BMC Bioinformatics* 8: 172. This system is a type of prokaryotic immune system that confers resistance to foreign genetic elements such as plasmids and phages and provides a form of acquired immunity. Barrangou *et al.* (2007) *Science* 315: 1709-1712; Marragini *et al.* (2008) *Science* 322: 1843-1845.

The CRISPR system has been modified for use in gene editing (silencing, enhancing or changing specific genes) in eukaryotes such as mice, primates and humans. Wiedenheft *et al.* (2012) *Nature* 482: 331-8. This is accomplished by, for example, introducing into the eukaryotic cell one or more vectors encoding a specifically engineered guide RNA (gRNA) (e.g., a gRNA comprising sequence complementary to sequence of a eukaryotic genome) and one or more appropriate RNA-guided nucleases, e.g., Cas proteins. The RNA guided nuclease forms a complex with the gRNA, which is then directed to the target DNA site by hybridization of the gRNA's sequence to complementary sequence of a eukaryotic genome, where the RNA-guided nuclease then induces a double or single-strand break in the DNA. Insertion or deletion of nucleotides at or near the strand break creates the modified genome.

As these naturally occur in many different types of bacteria, the exact arrangements of the CRISPR and structure, function and number of Cas genes and their product differ somewhat from species to species. Haft *et al.* (2005) *PLoS Comput. Biol.* 1: e60; Kunin *et al.* (2007) *Genome Biol.* 8: R61; Mojica *et al.* (2005) *J. Mol. Evol.* 60: 174-182; Bolotin *et al.* (2005) *Microbiol.* 151: 2551-2561; Pourcel et

al. (2005) *Microbiol.* 151: 653-663; and Stern *et al.* (2010) *Trends. Genet*. 28: 335-340. For example, the Cse (Cas subtype, E. coli) proteins (e.g., CasA) form a functional complex, Cascade, that processes CRISPR RNA transcripts into spacer-repeat units that Cascade retains. Brouns et al. (2008) *Science* 321: 960-964. In other prokaryotes, Cas6 processes the CRISPR transcript. The CRISPR-based phage inactivation in E. coli requires Cascade and Cas3, but not Cas1 or Cas2. The Cmr (Cas RAMP module) proteins in Pyrococcus furiosus and other prokaryotes form a functional complex with small CRISPR RNAs that recognizes and cleaves complementary target RNAs. A simpler CRISPR system relies on the protein Cas9, which is a nuclease with two active cutting sites, one for each strand of the double helix. Combining Cas9 and modified CRISPR locus RNA can be used in a system for gene editing. Pennisi (2013) *Science* 341: 833-836.

With respect to general information on CRISPR-Cas Systems, components thereof, and delivery of such components, including methods, materials, delivery vehicles, vectors, particles, AAV, and making and using thereof, including as to amounts and formulations, all useful in the practice of the instant invention, reference is made to: US Patents Nos. 8,697,359, 8,771,945, 8,795,965, 8,865,406, 8,871,445, 8,889,356, 8,889,418 and 8,895,308; US Patent Publications US 2014-0310830 (US APP. Ser. No. 14/105,031), US 2014-0287938 Al (U.S. App. Ser. No. 14/213,991), US 2014-0273234 Al (U.S. App. Ser. No. 14/293,674), US2014-0273232 Al (U.S. App. Ser. No. 14/290,575), US 2014-0273231 (U.S. App. Ser. No. 14/259,420), US 2014-0256046 Al (U.S. App. Ser. No. 14/226,274), US 2014-0248702 Al (U.S. App. Ser. No. 14/258,458), US 2014-0242700 Al (U.S. App. Ser. No. 14/222,930), US 2014-0242699 Al (U.S. App. Ser. No. 14/183,512), US 2014-0242664 Al (U.S. App. Ser. No. 14/104,990), US 2014-0234972 Al (U.S. App. Ser. No. 14/183,471), US 2014-0227787 Al (U.S. App. Ser. No. 14/256,912), US 2014-0189896 Al (U.S. App. Ser. No. 14/105,035), US 2014-0186958 (U.S. App. Ser. No. 14/105,017), US 2014-0186919 Al (U.S. App. Ser. No. 14/104,977), US 2014-0186843 Al (U.S. App. Ser. No. 14/104,900), US 2014-0179770 Al (U.S. App. Ser. No. 14/104,837) and US 2014-0179006 Al (U.S. App. Ser. No. 14/183,486), US 2014-0170753 (US App Ser No 14/183,429); European Patent Applications EP 2 771 468 (EP13818570.7), EP 2 764 103 (EP13824232.6), and EP 2 784 162 (EP14170383.5); and PCT Patent Publications WO 2014/093661 (PCT/US2013/074743), WO 2014/093694 (PCT/US2013/074790), WO 2014/093595 (PCT/US2013/074611), WO 2014/093718 (PCT/US2013/074825), WO 2014/093709 (PCT/US2013/074812), WO 2014/093622 (PCT/US2013/074667), WO 2014/093635 (PCT/US2013/074691), WO 2014/093655 (PCT/US2013/074736), WO 2014/093712 (PCT/US2013/074819), WO 2014/093701 (PCT7US2013/074800), WO 2014/018423 (PCT/US2013/051418) ,WO 2014/204723 (PCT/US2014/041790), WO 2014/204724

(PCT/US2014/041800), WO 2014/204725 (PCT/US2014/041803), WO 2014/204726 (PCT US2014/041804), WO 2014/204727 (PCT US2014/041806), WO 2014/204728 (PCT/US2014/041808), and WO 2014/204729 (PCT US2014/041809). Reference is also made to US provisional patent applications 61/758,468; 61/802,174; 61/806,375; 61/814,263; 61/819,803 and 61/828,130, filed on January 30, 2013; March 15, 2013; March 28, 2013; April 20, 2013; May 6, 2013 and May 28, 2013 respectively. Reference is also made to US provisional patent application 61/836,123, filed on June 17, 2013. Reference is additionally made to US provisional patent applications 61/835,931, 61/835,936, 61/836,127, 61/836, 101, 61/836,080 and 61/835,973, each filed June 17, 2013. Further reference is made to US provisional patent applications 61/862,468 and 61/862,355 filed on August 5, 2013; 61/871,301 filed on August 28, 2013; 61/960,777 filed on September 25, 2013 and 61/961 ,980 filed on October 28, 2013. Reference is yet further made to: PCT Patent applications Nos: PCT/US2014/041803, PCT/US2014/041800, PCT/US2014/041809, PCT/US2014/041804 and PCT US2014/041806, each filed June 10, 2014 6/10/14; PCT US2014/041808 filed June 1 1, 2014; and PCT/US2014/62558 filed October 28, 2014, and US Provisional Patent Applications Serial Nos.: 61/915,150, 61/915,301, 61/915,267 and 61/915,260, each filed December 12, 2013; 61/757,972 and 61/768,959, filed on January 29, 2013 and February 25, 2013; 61/835,936, 61/836,127, 61/836,101, 61/836,080, 61/835,973, and 61/835,931, filed June 17, 2013; 62/010,888 and 62/010,879, both filed June 1 1, 2014; 62/010,329 and 62/010,441, each filed June 10, 2014; 61/939,228 and 61/939,242, each filed February 12, 2014; 61/980,012, filed April 15,2014; 62/038,358, filed August 17, 2014; 62/054,490, 62/055,484, 62/055,460 and 62/055,487, each filed September 25, 2014; and 62/069,243, filed October 27, 2014. Reference is also made to US provisional patent applications Nos. 62/055,484, 62/055,460, and 62/055,487, filed September 25, 2014; US provisional patent application 61/980,012, filed April 15, 2014; and US provisional patent application 61/939,242 filed February 12, 2014. Reference is made to PCT application designating, inter alia, the United States, application No. PCT/US 14/41806, filed June 10, 2014. Reference is made to US provisional patent application 61/930,214 filed on January 22, 2014. Reference is made to US provisional patent applications 61/915,251, 61/915,260 and 61/915,267, each filed on December 12, 2013. Reference is made to US provisional patent application USSN 61/980,012 filed April 15, 2014. Reference is made to PCT application designating, inter alia, the United States, application No. PCT/US 14/41806, filed June 10, 2014. Reference is made to US provisional patent application 61/930,214 filed on January 22, 2014. Reference is made to US provisional patent applications 61/915,251; 61/915,260 and 61/915,267, each filed on December 12, 2013. [0054] Mention is also made of US application 62/091,455, filed, 12-Dec-14, PROTECTED GUIDE RNAS (PGRNAS); US application 62/096,708, 24-Dec-

14, PROTECTED GUIDE RNAS (PGRNAS); US application 62/091,462, 12-Dec-14, DEAD GUIDES FOR CRISPR TRANSCRIPTION FACTORS; US application 62/096,324, 23-Dec-14, DEAD GUIDES FOR CRISPR TRANSCRIPTION FACTORS; US application 62/091,456, 12-Dec-14, ESCORTED AND FUNCTI ON AL IZED GUIDES FOR CRISPR-CAS SYSTEMS; US application 62/091,461, 12-Dec-14, DELIVERY, USE AND THERAPEUTIC APPLICATIONS OF THE CRISPR-CAS SYSTEMS AND COMPOSITIONS FOR GENOME EDITING AS TO HEMATOPOETIC STEM CELLS (HSCs); US application 62/094,903, 19-Dec-14, UNBIASED IDENTIFICATION OF DOUBLE-STRAND BREAKS AND GENOMIC REARRANGEMENT BY GENOME- WISE INSERT CAPTURE SEQUENCING; US application 62/096,761 , 24-Dec-14, ENGINEERING OF SYSTEMS, METHODS AND OPTIMIZED ENZYME AND GUIDE SCAFFOLDS FOR SEQUENCE MANIPULATION; US application 62/098,059, 30-Dec-14, RNA-TARGETING SYSTEM; US application 62/096,656, 24-Dec-14, CRISPR HAVING OR ASSOCIATED WITH DESTABILIZATION DOMAINS; US application 62/096,697, 24-Dec-14, CRISPR HAVING OR ASSOCIATED WITH AAV; US application 62/098,158, 30-Dec-14, ENGINEERED CRISPR COMPLEX INSERTIONAL TARGETING SYSTEMS; US application 62/151,052, 22-Apr-15, CELLULAR TARGETING FOR EXTRACELLULAR EXOSOMAL REPORTING; US application 62/054,490, 24-Sep-14, DELIVERY, USE AND THERAPEUTIC APPLICATIONS OF THE CRISPR-CAS SYSTEMS AND COMPOSITIONS FOR TARGETING DISORDERS AND DISEASES USING PARTICLE DELIVERY COMPONENTS; US application 62/055,484, 25-Sep-14, SYSTEMS, METHODS AND COMPOSITIONS FOR SEQUENCE MANIPULATION WITH OPTIMIZED FUNCTIONAL CRISPR-CAS SYSTEMS; US application 62/087,537, 4-Dec-14, SYSTEMS, METHODS AND COMPOSITIONS FOR SEQUENCE MANIPULATION WITH OPTIMIZED FUNCTIONAL CRISPR-CAS SYSTEMS; US application 62/054,651, 24-Sep-14, DELIVERY, USE AND THERAPEUTIC APPLICATIONS OF THE CRISPR-CAS SYSTEMS AND COMPOSITIONS FOR MODELING COMPETITION OF MULTIPLE CANCER MUTATIONS IN VIVO; US application 62/067,886, 23-Oct-14, DELIVERY, USE AND THERAPEUTIC APPLICATIONS OF THE CRISPR-CAS SYSTEMS AND COMPOSITIONS FOR MODELING COMPETITION OF MULTIPLE CANCER MUTATIONS IN VIVO; US application 62/054,675, 24-Sep-14, DELIVERY, USE AND THERAPEUTIC APPLICATIONS OF THE CRISPR-CAS SYSTEMS AND COMPOSITIONS IN NEURONAL CELLS/TISSUES; US application 62/054,528, 24-Sep-14, DELIVERY, USE AND THERAPEUTIC APPLICATIONS OF THE CRISPR-CAS SYSTEMS AND COMPOSITIONS IN IMMUNE DISEASES OR DISORDERS; US application 62/055,454, 25-Sep-14, DELIVERY, USE AND THERAPEUTIC APPLICATIONS OF THE CRISPR-CAS SYSTEMS AND COMPOSITIONS FOR TARGETING DISORDERS AND DISEASES USING CELL PENETRATION PEPTIDES (CPP); US application 62/055,460, 25-Sep-14, MULTIFUNCTIONAL-CRISPR COMPLEXES AND/OR OPTIMIZED ENZYME LINKED FUNCTION AL-CRISPR COMPLEXES; US application 62/087,475, 4-Dec-14, FUNCTIONAL SCREENING WITH OPTIMIZED FUNCTIONAL CRISPR-CAS SYSTEMS; US application

62/055,487, 25-Sep-14, FUNCTIONAL SCREENING WITH OPTIMIZED FUNCTIONAL CRISPR-CAS SYSTEMS; US application 62/087,546, 4-Dec-14, MULTIFUNCTIONAL CRISPR COMPLEXES AND/OR OPTIMIZED ENZYME LINKED FUNCTIONAL-CRISPR COMPLEXES; and US application 62/098,285, 30-Dec-14, CRISPR MEDIATED IN VIVO MODELING AND GENETIC SCREENING OF TUMOR GROWTH AND METASTASIS.

Each of these patents, patent publications, and applications, and all documents cited therein or during their prosecution ("appln cited documents") and all documents cited or referenced in the appln cited documents, together with any instructions, descriptions, product specifications, and product sheets for any products mentioned therein or in any document therein and incorporated by reference herein, are hereby incorporated herein by reference, and may be employed in the practice of the invention. All documents (e.g., these patents, patent publications and applications and the appln cited documents) are incorporated herein by reference to the same extent as if each individual document was specifically and individually indicated to be incorporated by reference.

Also with respect to general information on CRISPR-Cas Systems, mention is made of the following (also hereby incorporated herein by reference):

Multiplex genome engineering using CRISPR/Cas systems, Cong, L., Ran, F.A., Cox, D., Lin, S., Barretto, R., Habib, N., Hsu, P.D., Wu, X., Jiang, W., Marraffini, L.A., & Zhang, F. Science Feb 15;339(6121):819-23 (2013);RNA-gided editing of bacterial genomes using CRISPR-Cas systems. Jiang W., Bikard D., Cox D., Zhang F, Marraffini LA. Nat Biotechnol Mar;31(3):233-9 (2013); One-Step Generation of Mice Carrying Mutations in Multiple Genes by CRISPR/Cas-Mediated Genome Engineering. Wang H., Yang H., Shivalila CS., Dawlaty MM., Cheng AW., Zhang F., Jaenisch R. Cell May 9;153(4):910-8 (2013); Optical control of mammalian endogenous transcription and epigenetic states. Konermann S, Brigham MD, Trevino AE, Hsu PD, Heidenreich M, Cong L, Piatt RJ, Scott DA, Church GM, Zhang F. Nature. 2013 Aug 22;500(7463):472-6. doi: 10.1038/Nature 12466. Epub 2013 Aug 23;

Double Nicking by RNA-Guided CRISPR Cas9 for Enhanced Genome Editing Specificity. Ran, FA., Hsu, PD., Lin, CY., Gootenberg, JS., Konermann, S., Trevino, AE,, Scott, DA., Inoue, A., Matoba, S., Zhang, Y,, & Zhang, F. Cell Aug 28. pii: S0092-8674(13)01015-5. (2013

DNA targeting specificity of RNA-guided Cas9 nucleases. Hsu, P., Scott, D., Weinstein, J., Ran, FA., Konermann, S., Agarwala, V., Li, Y., Fine, E., Wu, X., Shalem, O., Cradick, TJ., Marraffini, LA., Bao, G., & Zhang, F. Nat Biotechnol doi:10.1038/nbt.2647 (2013);

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Genome-wide CRISPR Screen in a Mouse Model of Tumor Growth and Metastasis, Chen S, Sanjana NE, Zheng , Shalem O, Lee , Shi X, Scott DA, Song J, Pan JQ, Weissleder R, Lee H, Zhang F, Sharp PA. Cell 160, 1246-1260, March 12, 2015 (multiplex screen in mouse), and

In vivo genome editing using Staphylococcus aureus Cas9, Ran FA, Cong L, Yan WX, Scott DA, Gootenberg JS, Kriz AJ, Zetsche B, Shalem O, Wu X, Makarova KS, oonin EV, Sharp PA, Zhang F., (published online 01 April 2015), Nature. Apr 9;520(7546): 186-91 (2015).

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Sequence determinants of improved CRISPR sgRNA design, Xu et al., Genome Research 25, 1 147-1 157 (August 2015).

A Genome-wide CRISPR Screen in Primary Immune Cells to Dissect Regulatory Networks, Parnas et al., Cell 162, 675-686 (July 30, 2015).

CRISPR/Cas9 cleavage of viral DNA efficiently suppresses hepatitis B virus, Ramanan et al., Scientific Reports 5:10833. doi: 10.1038/srepl0833 (June 2, 2015).

Crystal Structure of Staphylococcus aureus Cas9, Nishimasu et al., Cell 162, 1113-1126 (Aug. 27, 2015).

BCL11 A enhancer dissection by Cas9-mediated in situ saturating mutagenesis, Canver et al., Nature 527(7577): 192-7 (Nov. 12, 2015) doi: 10.1038/naturel 5521. Epub 2015 Sep 16. each of which is incorporated herein by reference, and discussed briefly below:

Cong et al. engineered type II CRISPR/Cas systems for use in eukaryotic cells based on both Streptococcus thermophilus Cas9 and also Streptoccocus pyogenes Cas9 and demonstrated that Cas9 nucleases can be directed by short RNAs to induce precise cleavage of DNA in human and mouse cells. Their study further showed that Cas9 as converted into a nicking enzyme can be used to facilitate homology-directed repair in eukaryotic cells with minimal mutagenic activity. Additionally, their study demonstrated that multiple guide sequences can be encoded into a single CRISPR array to enable simultaneous editing of several at endogenous genomic loci sites within the mammalian genome, demonstrating easy programmability and wide applicability of the RNA-guided nuclease technology. This ability to use RNA to program sequence specific DNA cleavage in cells defined a new class of genome engineering tools. These studies further showed that other CRISPR loci are likely to be transplantable into mammalian cells and can also mediate mammalian genome cleavage. Importantly, it can be envisaged that several aspects of the CRISPR/Cas system can be further improved to increase its efficiency and versatility.

Jiang et al. used the clustered, regularly interspaced, short palindromic repeats (CRISPR)-associated Cas9 endonuclease complexed with dual-RNAs to introduce precise mutations in the genomes of Streptococcus pneumoniae and Escherichia coli. The approach relied on dual-RNA:Cas9-directed cleavage at the targeted genomic site to kill unmutated cells and circumvents the need for selectable markers or counter-selection systems, The study reported reprogramming dual-RNA:Cas9 specificity by changing the sequence of short CRISPR RNA (crRNA) to make single- and multinucleotide changes carried on editing templates. The study showed that simultaneous use of two crRNAs enabled multiplex mutagenesis. Furthermore, when the approach was used in combination with recombineering, in S. pneumoniae, nearly 100% of cells that were recovered using the described approach contained the desired mutation, and in E. coli, 65% that were recovered contained the mutation.

Wang et al. (2013) used the CRISPR/Cas system for the one-step generation of mice carrying mutations in multiple genes which were traditionally generated in multiple steps by sequential recombination in embryonic stem cells and/or time-consuming intercrossing of mice with a single mutation. The CRISPR/Cas system will greatly accelerate the in vivo study of functionally redundant genes and of epistatic gene interactions.

Konermann et al. addressed the need in the art for versatile and robust technologies that enable optical and chemical modulation of DNA-binding domains based CRISPR Cas9 enzyme and also Transcriptional Activator Like Effectors.

Ran et al. (2013-A) described an approach that combined a Cas9 nickase mutant with paired guide RNAs to introduce targeted double-strand breaks. This addresses the issue of the Cas9 nuclease from the microbial CRISPR-Cas system being targeted to specific genomic loci by a guide sequence, which can tolerate certain mismatches to the DNA target and thereby promote undesired off-target mutagenesis. Because individual nicks in the genome are repaired with high fidelity, simultaneous nicking via appropriately offset guide RNAs is required for double-stranded breaks and extends the number of specifically recognized bases for target cleavage. The authors demonstrated that using paired nicking can reduce off-target activity by 50- to 1,500-fold in cell lines and to facilitate gene knockout in mouse zygotes without sacrificing on-target cleavage efficiency. This versatile strategy enables a wide variety of genome editing applications that require high specificity. Hsu et al. (2013) characterized SpCas9 targeting specificity in human cells to inform the selection of target sites and avoid off-target effects. The study evaluated >700 guide RNA variants and SpCas9-induced indel mutation levels at > 100 predicted genomic off-target loci in 293T and 293FT cells. The authors that SpCas9 tolerates mismatches

between guide RNA and target DNA at different positions in a sequence-dependent manner, sensitive to the number, position and distribution of mismatches. The authors further showed that SpCas9-mediated cleavage is unaffected by DNA methylation and that the dosage of SpCas9 and sgRNA can be titrated to minimize off-target modification. Additionally, to facilitate mammalian genome engineering applications, the authors reported providing a web-based software tool to guide the selection and validation of target sequences as well as off-target analyses.

Ran et al. (2013-B) described a set of tools for Cas9-mediated genome editing via non-homologous end joining (NHEJ) or homology-directed repair (HDR) in mammalian cells, as well as generation of modified cell lines for downstream functional studies. To minimize off-target cleavage, the authors further described a double-nicking strategy using the Cas9 nickase mutant with paired guide RNAs. The protocol provided by the authors experimentally derived guidelines for the selection of target sites, evaluation of cleavage efficiency and analysis of off-target activity. The studies showed that beginning with target design, gene modifications can be achieved within as little as 1-2 weeks, and modified clonal cell lines can be derived within 2-3 weeks.

Shaiem et al. described a new way to interrogate gene function on a genome-wide scale. Their studies showed that delivery of a genome-scale CRISPR-Cas9 knockout (GeC O) library targeted 18,080 genes with 64,751 unique guide sequences enabled both negative and positive selection screening in human cells. First, the authors showed use of the GeCKO library to identify genes essential for cell viability in cancer and pluripotent stem cells. Next, in a melanoma model, the authors screened for genes whose loss is involved in resistance to vemurafenib, a therapeutic that inhibits mutant protein kinase BRAF. Their studies showed that the highest-ranking candidates included previously validated genes NF1 and MED 12 as well as novel hits NF2, CUL3, TADA2B, and TADAL The authors observed a high level of consistency between independent guide RNAs targeting the same gene and a high rate of hit confirmation, and thus demonstrated the promise of genome-scale screening with Cas9.

Nishimasu et al. reported the crystal structure of Streptococcus pyogenes Cas9 in complex with sgRNA and its target DNA at 2.5 A° resolution. The structure revealed a bilobed architecture composed of target recognition and nuclease lobes, accommodating the sgRNA:DNA heteroduplex in a positively charged groove at their interface. Whereas the recognition lobe is essential for binding sgRNA and DNA, the nuclease lobe contains the HNH and RuvC nuclease domains, which are properly positioned for cleavage of the complementary and non-complementary strands of the target DNA, respectively. The nuclease lobe also contains a carboxyl-terminal domain responsible for the interaction with the

protospacer adjacent motif (PAM). This high-resolution structure and accompanying functional analyses have revealed the molecular mechanism of RNA-guided DNA targeting by Cas9, thus paving the way for the rational design of new, versatile genome-editing technologies.

Wu et al. mapped genome-wide binding sites of a catalytically inactive Cas9 (dCas9) from Streptococcus pyogenes loaded with single guide RNAs (sgRNAs) in mouse embryonic stem cells (mESCs). The authors showed that each of the four sgRNAs tested targets dCas9 to between tens and thousands of genomic sites, frequently characterized by a 5-nucleotide seed region in the sgRNA and an NGG protospacer adjacent motif (PAM). Chromatin inaccessibility decreases dCas9 binding to other sites with matching seed sequences; thus 70% of off-target sites are associated with genes. The authors showed that targeted sequencing of 295 dCas9 binding sites in mESCs transfected with catalytically active Cas9 identified only one site mutated above background levels. The authors proposed a two-state model for Cas9 binding and cleavage, in which a seed match triggers binding but extensive pairing with target DNA is required for cleavage.

Piatt et al. established a Cre-dependent Cas9 knockin mouse. The authors demonstrated in vivo as well as ex vivo genome editing using adeno-associated virus (AAV)-, lentivirus-, or particle-mediated delivery of guide RNA in neurons, immune cells, and endothelial cells.

Hsu et al. (2014) is a review article that discusses generally CRISPR-Cas9 history from yogurt to genome editing, including genetic screening of cells.

Wang et al, (2014) relates to a pooled, loss-of-function genetic screening approach suitable for both positive and negative selection that uses a genome-scale lentiviral single guide RNA (sgRNA) library.

Doench et al. created a pool of sgRNAs, tiling across all possible target sites of a panel of six endogenous mouse and three endogenous human genes and quantitatively assessed their ability to produce null alleles of their target gene by antibody staining and flow cytometry. The authors showed that optimization of the PAM improved activity and also provided an on-line tool for designing sgRNAs.

Swiech et al. demonstrate that AAV-mediated SpCas9 genome editing can enable reverse genetic studies of gene function in the brain.

Konermann et al. (2015) discusses the ability to attach multiple effector domains, e.g., transcriptional activator, functional and epigenomic regulators at appropriate positions on the guide such as stem or tetraloop with and without linkers.

Zetsche et al. demonstrates that the Cas9 enzyme can be split into two and hence the assembly of Cas9 for activation can be controlled.

Chen et al relates to multiplex screening by demonstrating that a genome- wide in vivo CRISPR-Cas9 screen in mice reveals genes regulating lung metastasis. > Ran et al. (2015) relates to SaCas9 and its ability to edit genomes and demonstrates that one cannot extrapolate from biochemical assays. Shalem et al. (2015) described ways in which catalytically inactive Cas9 (dCas9) fusions are used to synthetically repress (CRISPRi) or activate (CRISPRa) expression, showing, advances using Cas9 for genome-scale screens, including arrayed and pooled screens, knockout approaches that inactivate genomic loci and strategies that modulate transcriptional activity.

Shalem et al. (2015) described ways in which catalytically inactive Cas9 (dCas9) fusions are used to synthetically repress (CRISPRi) or activate (CRISPRa) expression, showing, advances using Cas9 for genome-scale screens, including arrayed and pooled screens, knockout approaches that inactivate genomic loci and strategies that modulate transcriptional activity.

Xu et al. (2015) assessed the DNA sequence features that contribute to single guide RNA (sgRNA) efficiency in CRISPR-based screens. The authors explored efficiency of CRISPR/Cas9 knockout and nucleotide preference at the cleavage site. The authors also found that the sequence preference for CRISPRi/a is substantially different from that for CRISPR Cas9 knockout.

Parnas et al. (2015) introduced genome- wide pooled CRISPR-Cas9 libraries into dendritic cells (DCs) to identify genes that control the induction of tumor necrosis factor (Tnf) by bacterial lipopolysaccharide (LPS). Known regulators of TIr4 signaling and previously unknown candidates were identified and classified into three functional modules with distinct effects on the canonical responses to LPS.

Ramanan et al (2015) demonstrated cleavage of viral episomal DNA (cccDNA) in infected cells. The HBV genome exists in the nuclei of infected hepatocytes as a 3.2kb double-stranded episomal DNA species called covalently closed circular DNA (cccDNA), which is a key component in the HBV life cycle whose replication is not inhibited by current therapies. The authors showed that sgRNAs specifically

targeting highly conserved regions of HBV robustly suppresses viral replication and depleted cccDNA. Nishimasu et al. (2015) reported the crystal structures of SaCas9 in complex with a single guide RNA (sgRNA) and its double-stranded DNA targets, containing the 5'-TTGAAT-3' PAM and the 5'-TTGGGT-3' PAM. A structural comparison of SaCas9 with SpCas9 highlighted both structural conservation and divergence, explaining their distinct PAM specificities and orthologous sgRNA recognition.

Slaymaker et al (2015) reported the use of structure-guided protein engineering to improve the specificity of Streptococcus pyogenes Cas9 (SpCas9). The authors developed "enhanced specificity" SpCas9 (eSpCas9) variants which maintained robust on-target cleavage with reduced off-target effects.

Tsai et al, "Dimeric CRISPR A-guided Fokl nucleases for highly specific genome editing," Nature Biotechnology 32(6): 569-77 (2014) which is not believed to be prior art to the instant invention or application, but which may be considered in the practice of the instant invention. Mention is also made of Konermann et al., "Genome-scale transcription activation by an engineered CRISPR-Cas9 complex," doi:10.1038/naturel4136, incorporated herein by reference.

In general, the CRISPR-Cas or CRISPR system is as used in the foregoing documents, such as WO 2014/093622 (PCT/US2013/074667) and refers collectively to transcripts and other elements involved in the expression of or directing the activity of CRISPR-associated (Cas) genes, including sequences encoding a Cas gene, a tracr (trans-activating CRISPR) sequence (e.g. tracrRNA or an active partial tracrRNA), a tracr-mate sequence (encompassing a direct repeat and a tracrRNA-processed partial direct repeat in the context of an endogenous CRISPR system), a guide sequence (also referred to as a spacer in the context of an endogenous CRISPR system), or gRNA(s) as that term is herein used (including, e.g., single guide RNA (sgRNA) (chimeric RNA) and dual guide RNAs (dgRNAs)). In the context CRISPR systems a, "target sequence" refers to a sequence to which the targeting domain sequence of a gRNA molecule is designed to have complementarity, where hybridization between a target sequence and a targeting domain sequence of a gRNA directs the CRISPR system to the locus comprising the target sequence. A target sequence may comprise any polynucleotide, such as DNA or RNA polynucleotides. In some embodiments, a target sequence is located in the nucleus or cytoplasm of a cell. In some embodiments, direct repeats may be identified in silico by searching for repetitive motifs that fulfill any or all of the following criteria: 1. found in a 2Kb window of genomic sequence flanking the type II CRISPR locus; 2. span from 20 to 50 bp; and 3. interspaced by 20 to 50 bp. In some embodiments, 2 of these criteria may be used, for instance 1 and 2, 2 and 3, or 1 and 3. In some embodiments, all 3 criteria may be used. In some embodiments it may be preferred in a CRISPR system that the tracr sequence has one or more

hairpins and is 30 or more nucleotides in length, 40 or more nucleotides in length, or 50 or more nucleotides in length; the guide sequence is between 10 to 30 nucleotides in length, the CRISPR/Cas enzyme is a Type II Cas9 enzyme. In embodiments of the invention the terms guide sequence and guide RNA are used interchangeably as in foregoing cited documents such as WO 2014/093622 (PCT US2013/074667). In general, a guide sequence is any polynucleotide sequence having sufficient complementarity with a target sequence to hybridize with the target sequence and direct sequence-specific binding of a CRISPR system to the target sequence. In some embodiments, the degree of complementarity between a guide sequence and its corresponding target sequence, when optimally aligned using a suitable alignment algorithm, is about or more than about 50%, 60%, 75%, 80%, 85%, 90%, 95%, 97.5%, 99%, or more. Optimal alignment may be determined with the use of any suitable algorithm for aligning sequences, non-limiting example of which include the Smith- Waterman algorithm, the Needleman-Wunsch algorithm, algorithms based on the Burrows-Wheeler Transform (e.g. the Burrows Wheeler Aligner), ClustalW, Clustal X, BLAT, Novoalign (Novocraft Technologies; available at www.novocraft.com), ELAND (Illumina, San Diego, CA), SOAP (available at soap.genomics.org.cn), and aq (available at maq.sourceforge.net). In some embodiments, a guide sequence is about or more than about 5, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, 30, 35, 40, 45, 50, 75, or more nucleotides in length. In some embodiments, a guide sequence is less than about 75, 50, 45, 40, 35, 30, 25, 20, 15, 12, or fewer nucleotides in length. Preferably the guide sequence is 10 - 30 nucleotides long. The ability of a guide sequence to direct sequence- specific binding of a CRISPR system to a target sequence may be assessed by any suitable assay. For example, the components of a CRISPR system sufficient to form a CRISPR system, including the guide sequence to be tested, may be provided to a host cell having the corresponding target sequence, such as by transfection with vectors encoding the components of the CRISPR sequence, followed by an assessment of preferential cleavage within the target sequence, such as by Surveyor assay as described in the literature and known to those skilled in the art. Similarly, cleavage of a target polynucleotide sequence may be evaluated in a test tube by providing the target sequence, components of a CRISPR system, including the guide sequence to be tested and a control guide sequence different from the test guide sequence, and comparing binding or rate of cleavage at the target sequence between the test and control guide sequence reactions. Other assays are possible, and will occur to those skilled in the art. A guide sequence may be selected to target any target sequence. In some embodiments, the target sequence is a sequence within a genome of a cell. Exemplary target sequences include those that are unique in the target genome. For example, for the S. pyogenes Cas9, a unique sequence in a genome

may include a Cas9 target sequence of the form NNNNNNNNNNNNNNNNNNNNXGG (where N is A, G, T, or C; and X can be anything), wherein the sequence has a single occurrence in the genome. For the S. thermophilus CRISPR/Cas9 system, a unique sequence in a genome may include a Cas9 target site of the form NNNNNNNNNNNNNNNNNNNNXXAGAAW (SEQ ID NO: 863) (where N is A, G, T, or C; X can be anything; and W is A or T), wherein the sequence has a single occurrence in the genome. For the S. pyogenes Cas9 or the S. Thermophilius Cas9, a unique sequence in a genome may include a Cas9 target site of the form NNNNNNNNNNNNNNNNNNNNXGGXG (where N is A, G, T, or C; and X can be anything), wherein the sequence has a single occurrence in the genome. In some embodiments, a guide RNA sequence is selected to reduce the degree secondary structure within the guide RNA sequence. In some embodiments, about or less than about 75%, 50%, 40%, 30%, 25%, 20%, 15%, 10%, 5%, 1%, or fewer of the nucleotides of the guide sequence participate in self-complementary base pairing when optimally folded. Optimal folding may be determined by any suitable polynucleotide folding algorithm. Some programs are based on calculating the minimal Gibbs free energy. An example of one such algorithm is mFold, as described by Zuker and Stiegler (Nucleic Acids Res. 9 (1981), 133-148). Another example folding algorithm is the online webserver RNAfold, developed at Institute for Theoretical Chemistry at the University of Vienna, using the centroid structure prediction algorithm (see e.g. A.R. Gruber et al., 2008, Cell 106(1): 23-24; and PA Carr and GM Church, 2009, Nature Biotechnology 27(12): 1 151-62).

In some embodiments, the RNA-guided nuclease is a Cas molecule, e.g., a Cas9 molecule. Cas9 molecules of a variety of species can be used in the methods and compositions described herein. In preferred embodiments, the Cas9 molecule is a S. pyogenes Cas9 molecule. In embodiments, the Cas9 molecule is derived from a S. pyogenes Cas9 molecule (e.g., UniProt Q99ZW2). While the S. pyogenes Cas9 molecule are the subject of much of the disclosure herein, Cas9 molecules of, derived from, or based on the Cas9 proteins of other species listed herein can be used as well. In other words, other Cas9 molecules, e.g., S. thermophilus, Staphylococcus aureus and/or Neisseria meningitidis Cas9 molecules, may be used in the systems, methods and compositions described herein. Additional Cas9 species include: Acidovorax avenae, Actinobacillus pleuropneumoniae, Actinobacillus succinogenes, Actinobacillus suis, Actinomyces sp., cycliphilus denitrificans, Aminomonas paucivorans, Bacillus cereus, Bacillus smithii, Bacillus thuringiensis, Bacteroides sp., Blastopirellula marina, Bradyrhiz' obium sp., Brevibacillus latemsporus, Campylobacter coli, Campylobacter jejuni, Campylobacter lad, Candidatus Puniceispirillum, Clostridiu cellulolyticum, Clostridium perfringens, Corynebacterium accolens, Corynebacterium diphtheria, Corynebacterium matruchotii, Dinoroseobacter sliibae, Eubacterium dolichum, gamma proteobacterium, Gluconacetobacler diazotrophicus, Haemophilus parainfluenzae,

Haemophilus sputorum, Helicobacter canadensis, Helicobacter cinaedi, Helicobacter mustelae, Ilyobacler polytropus, Kingella kingae, Lactobacillus crispatus, Listeria ivanovii, Listeria monocytogenes, Listeriaceae bacterium, Methylocystis sp., Methylosinus trichosporium, Mobiluncus mulieris, Neisseria bacilliformis, Neisseria cinerea, Neisseria flavescens, Neisseria lactamica. Neisseria sp., Neisseria wadsworthii, Nitrosomonas sp., Parvibaculum lavamentivorans, Pasteurella multocida, Phascolarctobacterium succinatutens, Ralstonia syzygii, Rhodopseudomonas palustris, Rhodovulum sp., Simonsiella muelleri, Sphingomonas sp., Sporolactobacillus vineae, Staphylococcus lugdunensis, Streptococcus sp., Subdoligranulum sp., Tislrella mobilis, Treponema sp., or Verminephrobacter eiseniae.

A Cas9 molecule, as that term is used herein, refers to a molecule that can interact with a gRNA molecule (e.g., sequence of a domain of a tracr) and, in concert with the gRNA molecule, localize (e.g., target or home) to a site which comprises a target sequence and PAM sequence.

In embodiments, the ability of an active Cas9 molecule to interact with and cleave a target nucleic acid is PAM sequence dependent. A PAM sequence is a sequence in the target nucleic acid. In an embodiment, cleavage of the target nucleic acid occurs upstream from the PAM sequence. Active Cas9 molecules from different bacterial species can recognize different sequence motifs (e.g., PAM sequences). In an embodiment, an active Cas9 molecule of S. pyogenes recognizes the sequence motif NGG and directs cleavage of a target nucleic acid sequence 1 to 10, e.g., 3 to 5, base pairs upstream from that sequence. See, e.g., Mali el al, SCIENCE 2013; 339(6121): 823- 826. In an embodiment, an active Cas9 molecule of S. thermophilus recognizes the sequence motif NGGNG and NNAGAAW (W = A or T) and directs cleavage of a core target nucleic acid sequence 1 to 10, e.g., 3 to 5, base pairs upstream from these sequences. See, e.g., Horvath et al., SCIENCE 2010; 327(5962): 167- 170, and Deveau et al, J BACTERIOL 2008; 190(4): 1390- 1400. In an embodiment, an active Cas9 molecule of S. mutans recognizes the sequence motif NGG or NAAR (R - A or G) and directs cleavage of a core target nucleic acid sequence 1 to 10, e.g., 3 to 5 base pairs, upstream from this sequence. See, e.g., Deveau et al., J BACTERIOL 2008; 190(4): 1390- 1400.

In an embodiment, an active Cas9 molecule of S. aureus recognizes the sequence motif NNGRR (R = A or G) and directs cleavage of a target nucleic acid sequence 1 to 10, e.g., 3 to 5, base pairs upstream from that sequence. See, e.g., Ran F. et al., NATURE, vol. 520, 2015, pp. 186-191. In an embodiment, an active Cas9 molecule of N. meningitidis recognizes the sequence motif NNNNGATT and directs cleavage of a target nucleic acid sequence 1 to 10, e.g., 3 to 5, base pairs upstream from that

sequence. See, e.g., Hou et al., PNAS EARLY EDITION 2013, 1 -6. The ability of a Cas9 molecule to recognize a PAM sequence can be determined, e.g., using a transformation assay described in Jinek et al, SCIENCE 2012, 337:816.

Exemplary naturally occurring Cas9 molecules are described in Chylinski et al , RNA Biology 2013; 10:5, 727-737. Such Cas9 molecules include Cas9 molecules of a cluster 1 bacterial family, cluster 2 bacterial family, cluster 3 bacterial family, cluster 4 bacterial family, cluster 5 bacterial family, cluster 6 bacterial family, a cluster 7 bacterial family, a cluster 8 bacterial family, a cluster 9 bacterial family, a cluster 10 bacterial family, a cluster 1 1 bacterial family, a cluster 12 bacterial family, a cluster 13 bacterial family, a cluster 14 bacterial family, a cluster 1 bacterial family, a cluster 16 bacterial family, a cluster 17 bacterial family, a cluster 1 8 bacterial family, a cluster 19 bacterial family, a cluster 20 bacterial family, a cluster 21 bacterial family, a cluster 22 bacterial family, a cluster 23 bacterial family, a cluster 24 bacterial family, a cluster 25 bacterial family, a cluster 26 bacterial family, a cluster 27 bacterial family, a cluster 28 bacterial family, a cluster 29 bacterial family, a cluster 30 bacterial family, a cluster 31 bacterial family, a cluster 32 bacterial family, a cluster 33 bacterial family, a cluster 34 bacterial family, a cluster 35 bacterial family, a cluster 36 bacterial family, a cluster 37 bacterial family, a cluster 38 bacterial family, a cluster 39 bacterial family, a cluster 40 bacterial family, a cluster 41 bacterial family, a cluster 42 bacterial family, a cluster 43 bacterial family, a cluster 44 bacterial family, a cluster 45 bacterial family, a cluster 46 bacterial family, a cluster 47 bacterial family, a cluster 48 bacterial family,. a cluster 49 bacterial family, a cluster 50 bacterial family, a cluster 5 1 bacterial family, a cluster 52 bacterial family, a cluster 53 bacterial family, a cluster 54 bacterial family, a cluster 55 bacterial family, a cluster 56 bacterial family, a cluster 57 bacterial family, a cluster 58 bacterial family, a cluster 59 bacterial family, a cluster 60 bacterial family, a cluster 61 bacterial family, a cluster 62 bacterial family, a cluster 63 bacterial family, a cluster 64 bacterial family, a cluster 65 bacterial family, a cluster 66 bacterial family, a cluster 67 bacterial family, a cluster 68 bacterial family, a cluster 69 bacterial family, a cluster 70 bacterial family, a cluster 71 bacterial family, a cluster 72 bacterial family, a cluster 73 bacterial family, a cluster 74 bacterial family, a cluster 75 bacterial family, a cluster 76 bacterial family, a cluster 77 bacterial family, or a cluster 78 bacterial family.

Exemplary naturally occurring Cas9 molecules include a Cas9 molecule of a cluster 1 bacterial family. Examples include a Cas9 molecule of: S. pyogenes (e.g., strain SF370, MGAS 10270, MGAS 10750, MGAS2096, MGAS315, MGAS5005, MGAS6180, MGAS9429, NZ131 and SSI- 1), S. thermophilus (e.g., strain LMD-9), S. pseudoporcinus (e.g., strain SPIN 20026), S. mutans (e.g., strain UA 159, NN2025), S.

macacae (e.g., strain NCTC1 1558), S. gallolylicus (e.g., strain UCN34, ATCC BAA-2069), S. equines (e.g., strain ATCC 9812, MGCS 124), S. dysdalactiae (e.g., strain GGS 124), S. bovis (e.g., strain ATCC 700338), S. cmginosus (e.g.; strain F021 1), S. agalactia* (e.g., strain NEM316, A909), Listeria monocytogenes (e.g., strain F6854), Listeria innocua (L. innocua, e.g., strain Clip l1262), EtUerococcus italicus (e.g., strain DSM 15952), or Enterococcus faecium (e.g., strain 1,23 ,408). Additional exemplary Cas9 molecules are a Cas9 molecule of Neisseria meningitidis (Hou et al. PNAS Early Edition 2013, 1-6) and a S. aureus Cas9 molecule.

In an embodiment, a Cas9 molecule, e.g., an active Cas9 molecule or inactive Cas9 molecule, comprises an amino acid sequence: having 60%, 65%, 70%, 75%, 80%, 85%, 90%, 95%, 96%, 97%, 98%, or 99% homology with; differs at no more than 1%, 2%, 5%, 10%, 15%, 20%, 30%, or 40% of the amino acid residues when compared with; differs by at least 1, 2, 5, 10 or 20 amino acids but by no more than 100, 80, 70, 60, 50, 40 or 30 amino acids from; or is identical to; any Cas9 molecule sequence described herein or a naturally occurring Cas9 molecule sequence, e.g., a Cas9 molecule from a species listed herein or described in Chylinski et al. , RNA Biology 2013, 10:5, 12l-T,1 Hou et al. PNAS Early Edition 2013, 1-6.

In an embodiment, a Cas9 molecule comprises an amino acid sequence having 60%, 65%, 70%, 75%, 80%, 85%, 90%, 95%, 96%, 97%, 98%, or 99% homology with; differs at no more than 1%, 2%, 5%, 10%, 15%, 20%, 30%, or 40% of the amino acid residues when compared with; differs by at least 1, 2, 5, 10 or 20 amino acids but by no more than 100, 80, 70, 60, 50, 40 or 30 amino acids from; or is identical to; *S. pyogenes* Cas9 (UniProt Q99ZW2). In embodiments, the Cas9 molecule is a *S. pyogenes* Cas9 variant, such as a variant described in Slaymaker et al., *Science Express*, available online December 1, 2015 at Science DOI: 10.1126/science.aad5227; Kleinstiver et al., *Nature*, 529, 2016, pp. 490-495, available online January 6, 2016 at doi:10.1038/nature16526; or US2016/0102324, the contents of which are incorporated herein in their entirety. In an embodiment, the Cas9 molecule is catalytically inactive, e.g., dCas9. Tsai et al. (2014), Nat. Biotech. 32:569-577; U.S. Patent No.: 8,871,445; 8,865,406; 8,795,965; 8,771,945; and 8,697,359, the contents of which are hereby incorporated by reference in their entirety. A catalytically inactive Cas9, e.g., dCas9, molecule may be fused with a transcription modulator, e.g., a transcription repressor or transcription activator.

In embodiments, the Cas9 molecule, e.g, a Cas9 of *S. pyogenes*, may additionally comprise one or more amino acid sequences that confer additional activity. In some aspects, the Cas9 molecule may comprise one or more nuclear localization sequences (NLSs), such as at least 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, or

more NLSs. Typically, an NLS consists of one or more short sequences of positively charged lysines or arginines exposed on the protein surface, but other types of NLS are known. Non-limiting examples of NLSs include an NLS sequence comprising or derived from: the NLS of the SV40 virus large T-antigen, having the amino acid sequence PKKKRKV (SEQ ID NO: 864). Other suitable NLS sequences are known in the art (e.g., Sorokin, Biochemistry (Moscow) (2007) 72:13, 1439-1457; Lange J Biol Chem. (2007) 282:8, 5101-5). In any of the aforementioned embodiments, the Cas9 molecule may additionally (or alternatively) comprise a tag, e.g., a His tag, e.g., a His(6) tag (SEQ ID NO: 865) or His(8) tag (SEQ ID NO: 866), e.g., at the N terminus and/or the C terminus.

Thus, engineered CRISPR gene editing systems, e.g., for gene editing in eukaryotic cells, typically involve (1) a guide RNA molecule (gRNA) comprising a targeting domain (which is capable of hybridizing to the genomic DNA target sequence), and sequence which is capable of binding to a Cas, e.g., Cas9 enzyme, and (2) a Cas, e.g., Cas9, protein. This second domain may comprise a domain referred to as a tracr domain. The targeting domain and the sequence which is capable of binding to a Cas, e.g., Cas9 enzyme, may be disposed on the same (sometimes referred to as a single gRNA, chimeric gRNA or sgRNA) or different molecules (sometimes referred to as a dual guide RNA, dual gRNA or dgRNA). If disposed on different molecules, each includes a hybridization domain which allows the molecules to associate, e.g., through hybridization.

gRNA molecule formats are known in the art. An exemplary gRNA molecule, e.g., dgRNA molecule, of the present invention comprises, e.g., consists of, a first nucleic acid having the sequence:
nnnnnnnnnnnnnnnnnnnnGUUUUAGAGCUAUGCUGUUUUG (SEQ ID NO: 867),
where the “n”’s refer to the residues of the targeting domain, e.g., a targeting domain to KDM1A, e.g., as described herein, and may consist of 15-25 nucleotides, e.g., consists of 20 nucleotides;
and a second nucleic acid sequence having the exemplary sequence:
AACUUACCAAGGAACAGCAUAGCAAGUUAAAAUAAGGCUAGUCCGUUAUCAACUUGAAAAAGUGGCACCGA GUCGGUGC, optionally with 1, 2, 3, 4, 5, 6, or 7 (e.g., 4 or 7, e.g., 7) additional U nucleotides at the 3’ end (SEQ ID NO: 868).

The second nucleic acid molecule may alternatively consist of a fragment of the sequence above, wherein such fragment is capable of hybridizing to the first nucleic acid. An example of such second nucleic acid molecule is:

AACAGCAUAGCAAGUUAAAAUAAGGCUAGUCCGUUAUCAACUUGAAAAAGUGGCACCGAGUCGGU GC, optionally with 1, 2, 3, 4, 5, 6, or 7 (e.g., 4 or 7, e.g., 7) additional U nucleotides at the 3’ end (SEQ ID NO: 869).

Another exemplary gRNA molecule, e.g., a sgRNA molecule, of the present invention comprises, e.g., consists of a first nucleic acid having the sequence:

nnnnnnnnnnnnnnnnnnnnGUUUUAGAGCUAGAAAUAGCAAGUUAAAAUAAGGCUAGUCCGUUAUCA ACUUGAAAAAGUGGCACCGAGUCGGUGC (SEQ ID NO: 870), where the "n"'s refer to the residues of the targeting domain, e.g., a targeting domain to KDM1A, e.g., as described herein, and may consist of 15-25 nucleotides, e.g., consist of 20 nucleotides, optionally with 1, 2, 3, 4, 5, 6, or 7 (e.g., 4 or 7, e.g., 4) additional U nucleotides at the 3' end.

Exemplary sequences of gRNA molecule targeting domains useful in the present invention (e.g., which target an LSD1 gene, e.g., KDM1A) are provided in Table 2.

Additional components and/or elements of CRISPR gene editing systems known in the art, e.g., are described in U.S. Publication No.2014/0068797, WO2015/048577, and Cong (2013) Science 339: 819-823, the contents of which are hereby incorporated by reference in their entirety. Such systems can be generated which inhibit a target gene, by, for example, engineering a CRISPR gene editing system to include a gRNA molecule comprising a targeting domain that hybridizes to a sequence of the target gene. In embodiments, the gRNA comprises a targeting domain which is fully complementarity to 15-25 nucleotides, e.g., 20 nucleotides, of a target gene, e.g., KDM1A or its regulatory elements. In embodiments, the 15-25 nucleotides, e.g., 20 nucleotides, of the target gene, are disposed immediately 5' to a protospacer adjacent motif (PAM) sequence recognized by the RNA-guided nuclease, e.g., Cas protein, of the CRISPR gene editing system (e.g., where the system comprises a *S. pyogenes* Cas9 protein, the PAM sequence comprises NGG, where N can be any of A, T, G or C).

In embodiments, the gRNA molecule and RNA-guided nuclease, e.g., Cas protein, of the CRISPR gene editing system can be complexed to form a RNP complex. In other embodiments, nucleic acid encoding one or more components of the CRISPR gene editing system may be used.

In embodiments, foreign DNA can be introduced into the cell along with the CRISPR gene editing system, e.g., DNA encoding a desired transgene, with or without a promoter active in the target cell type. Depending on the sequences of the foreign DNA and target sequence of the genome, this process can be used to integrate the foreign DNA into the genome, at or near the site targeted by the CRISPR gene editing system. For example, 3' and 5' sequences flanking the transgene may be included in the foreign DNA which are homologous to the gene sequence 3' and 5' (respectively) of the site in the genome cut by the gene editing system. Such foreign DNA molecule can be referred to "template DNA."

In an embodiment, the CRISPR gene editing system of the present invention comprises Cas9, e.g., S. pyogenes Cas9, and a gRNA comprising a targeting domain which hybridizes to a sequence of a gene of interest, e.g., KDM1A. In an embodiment, the gRNA and Cas9 are complexed to form a RNP. In an embodiment, the CRISPR gene editing system comprises nucleic acid encoding a gRNA and nucleic acid encoding a Cas protein, e.g., Cas9, e.g., S. pyogenes Cas9. In an embodiment, the CRISPR gene editing system comprises a gRNA and nucleic acid encoding a Cas protein, e.g., Cas9, e.g., S. pyogenes Cas9.

In an exemplary embodiment, the genome editing system LSD1 inhibitor is a CRISPR system. CRISPR genome editing systems useful in the practice of this invention are described in, for example, Artificial CRISPR/Cas systems can be generated which inhibit LSD1, using technology known in the art, e.g., that are described in U.S. Publication No. 20140068797, and Cong (2013) Science 339: 819-823. Other artificial CRISPR/Cas systems that are known in the art may also be generated which inhibit TCR and/or HLA, e.g., that described in Tsai (2014) Nature Biotechnol., 32:6 569-576, U.S. Patent No.: 8,871,445; 8,865,406; 8,795,965; 8,771,945; and 8,697,359.

Table 2: Exemplary LSD1 gRNA targeting domain sequences.

Target Gene	gRNA Targeting Domain Sequence	SEQ ID NO:
KDM1A	AGUGCGACAGGUUCGCUACA	132
KDM1A	UGGAAUAGCAGAGACUCCGG	133
KDM1A	CUAAAUAACUGUGAACUCGG	134
KDM1A	UGCUAUUCCAGUUUCCAUGG	135
KDM1A	CAGACCCAGGCACGACAGUA	136
KDM1A	UUUCUGAAACAGGAUCGUGU	137
KDM1A	UGAGAAGUCAUCCGGUCAUG	138
KDM1A	GAAUUGCAGACCAGUUUUUG	139
KDM1A	CGAGUUGCCACAUUUCGCAA	140
KDM1A	CUGAAACAGGAUCGUGUGGG	141
KDM1A	GGCGGCCCGAGAUGUUAUCU	142
KDM1A	UGUAUACCACACCUUGCAUA	143
KDM1A	AUGUAUACCACACCUUGCAU	144
KDM1A	UCUUAGUGAAAAGCAAACAC	145
KDM1A	GGGAUUUGGCAACCUUAACA	146
KDM1A	UUGUGCCACCUCUCCCUGAG	147
KDM1A	ACCAAGACCUGUUACCACCA	148
KDM1A	UCGGCAGUAAUAUCUCUGGG	149
KDM1A	GGAUCUGACCGCCCUAUGCA	150

KDM1A	CGUCAUGGUCUUAUCAACUU	151
KDM1A	UUCAAGACGACAGUUCUGGA	152
KDM1A	UGAAGUCUAGUUGAUGACUA	153
KDM1A	CACAGUUAUUUAGAGCGUCA	154
KDM1A	UGCCAACAGUAUUGGAGCUG	155
KDM1A	AGUGCCAACAGUAUUGGAGC	156
KDM1A	GCAUCUGUCUCACAUGCUUG	157
KDM1A	ACAUCUGCAGUCCAAAGGAU	158
KDM1A	GGCACAAACUGAACGGCUGG	159
KDM1A	GGUAAUUAUUAUAGGCUCUG	160
KDM1A	AAGAUGAGCAGAUUGAACAU	161
KDM1A	UGGCAGUACGACUGCCAGCA	162
KDM1A	CGCGGAGGCUCUUUCUUGCG	163
KDM1A	AUAAGUGACGAUGUGAUUGU	164
KDM1A	GGCAGUACGACUGCCAGCAG	165
KDM1A	GGAAAACAUCUGCAGUCCAA	166
KDM1A	UGCCCGAACAAAUUGACACU	167
KDM1A	ACUGAAUACAGCAGUGCGAC	168
KDM1A	CUGGCAGCCACCUGACAGUA	169
KDM1A	GCAGACCCAGGCACGACAGU	170
KDM1A	UUUGCUUUUCACUAAGAACU	171
KDM1A	GGGACACAGGCUUAUUAUUG	172
KDM1A	UGGCUACUCGUGUGUGCCUG	173
KDM1A	UCUCCCGCAAAGAAGAGUCG	174
KDM1A	AACAUCUGCAGUCCAAAGGA	175
KDM1A	CUUAGUGAAAAGCAAACACA	176
KDM1A	CCCAGGCACGACAGUAGGGC	177
KDM1A	GUCAAUUUGUUCGGGCAUGU	178
KDM1A	GAAUAAGAGCCCCGAGCCCA	179
KDM1A	CUGCUAUUCCAGUUUCCAUG	180
KDM1A	GAGAGGUGGCACAAACUGAA	181
KDM1A	AAGUGAGCCUGAAGAACCAU	182
KDM1A	UUUGCUUUCUUCGUUAGGUG	183
KDM1A	AUUUCUGAAACAGGAUCGUG	184
KDM1A	ACGACAGUAGGGCCGGCCUG	185
KDM1A	GUCCGUUGGCUUCAUAAAGU	186
KDM1A	AUACCACACCUUGCAUAGGG	187
KDM1A	GCUCUAAAUAACUGUGAACU	188
KDM1A	AGGAGGGAAUCCUAUGGCUG	189
KDM1A	AUAAAUAAAGGUUUGACUCG	190
KDM1A	AGCUGAUCUUGGAGCCAUGG	191

KDM1A	UAAAGGUUUGACUCGUGGAG	192
KDM1A	UGGACCACAACAGACCCAGA	193
KDM1A	GAUGAAUUAGCUGAAACACA	194
KDM1A	AGUAGCCAUUCCUUACUGUC	195
KDM1A	CUGGCGAGGCAGCGUAUACA	196
KDM1A	GUUUUCCACUCAGGGAGAGG	197
KDM1A	CGGAACCGCCGGGGUCCGCA	198
KDM1A	AGUGAGCCUGAAGAACCAUC	199
KDM1A	UCAACUUCGGCAUCUAUAAG	200
KDM1A	GUGGGGCACCUGGAAUCGAG	201
KDM1A	GGAAUAGCAGAGACUCCGGA	202
KDM1A	AGCCACCUGACAGUAAGGAA	203
KDM1A	AUAGUUUCCUUUGCGAAAUG	204
KDM1A	AGUUGGAAGCGAAUCCCCCA	205
KDM1A	AUGGCUCAGCCAAUCACUCC	206
KDM1A	AGCGGCAGCAACCGGGACGG	207
KDM1A	CUUUUAGCCCAAAGAAACUG	208
KDM1A	AAGAUGUAGCUUCUAGCAAC	209
KDM1A	AAUAAUACUCAUCUUCUGAG	210
KDM1A	CCGGCCCUACUGUCGUGCCU	211
KDM1A	CUGCUUCUUGAGAAGUCAUC	212
KDM1A	AGAGCCGACUUCCUCAUGAC	213
KDM1A	AGGCCUUCUGCUAAAGCCAC	214
KDM1A	GAAUGACAACUUCCAAUGCC	215
KDM1A	UGGUUAAAAGGACACUGUCA	216
KDM1A	UGCUGCUUCAGCACACCCAG	217
KDM1A	CCGCAAGAAAGAGCCUCCGC	218
KDM1A	AACUUCCAAUGCCUGGCCAA	219
KDM1A	AAAGGUUUGACUCGUGGAGC	220
KDM1A	GUGCCAACAGUAUUGGAGCU	221
KDM1A	UCAUCCGGUCAUGAGGAAGU	222
KDM1A	GCCACUAGUGCCAACAGUAU	223
KDM1A	UAGGGCAAGCUACCUUGUUA	224
KDM1A	GAUGCCUGGCCAUUCUCAAA	225
KDM1A	CAACUCUCUCCCUUAAGCAC	226
KDM1A	UGUUUUCCAGAUAUUAUCAG	227
KDM1A	CUUCAAGACGACAGUUCUGG	228
KDM1A	GAUUCCACGACUCUUCUUUG	229
KDM1A	AUGCAUCUGUCUCACAUGCU	230
KDM1A	AACUCUCUCCCUUAAGCACU	231
KDM1A	UCGACUUACAGCUUGUCCGU	232

KDM1A	UGGUAGCAGUGCAGUACCUC	233
KDM1A	AGGAGGUCCUUACUUGGUAG	234
KDM1A	AUAGGUAGAGUACAGAGAGA	235
KDM1A	GCACUGUGGCUGGGUAGUUA	236
KDM1A	UUUUCUAGGAGGGAAUCCUA	237
KDM1A	CGGACCCCGGCGGUUCCGCC	238
KDM1A	UGCCCACUUUAUGAAGCCAA	239
KDM1A	GAGCACCAUGCACUGUGGCU	240
KDM1A	GCAAAGAAGAGUCGUGGAAU	241
KDM1A	CUCCCUUAAGCACUGGGAUC	242
KDM1A	AGAUGAUCCUGCAGCAACAU	243
KDM1A	CCCGCGGAGGCUCUUUCUUG	244
KDM1A	UCUGCUAUUCCAGUUUCCAU	245
KDM1A	UGUGGUCCACUGAUAAUAUC	246
KDM1A	AAUUGCAGACCAGUUUUUGG	247
KDM1A	UUUAGCUAAAAAGACAGGAA	248
KDM1A	ACAGCAGGCGGCUCCGAGAA	249
KDM1A	AUGUCACACUUUUGGAAGCC	250
KDM1A	UUCUUCCCAGAUAACAUCUC	251
KDM1A	UCGUUGGCGUGCUGAUCCCU	252
KDM1A	GCUGAGUGGGCUGCGAGAAG	253
KDM1A	UCGGACCAGCCGGCGCAAGC	254
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KDM1A	GGUUCCGCCAGGCCCCCGGG	256
KDM1A	UGUGAUUGUUGGCCGAUGCC	257
KDM1A	AGCCAUUCCUUACUGUCAGG	258
KDM1A	AAUAAUAAGCCUGUGUCCCU	259
KDM1A	UCCAAUACUGUUGGCACUAG	260
KDM1A	UCCAUGGGGGUCGCAGACCC	261
KDM1A	GGUGGCACAAACUGAACGGC	262
KDM1A	AAAUCCCUUUGAGAAUGGCC	263
KDM1A	UGUAGCUGAUCUUGGAGCCA	264
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KDM1A	UACCUGAUCCCAGUGCUUAA	372
KDM1A	UGGGAAGAAGGCGGCAGCCG	373
KDM1A	GCUGGGCCCGACAGGCCCGC	374
KDM1A	CUCUUCUGGAACCUCUAUAA	375
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KDM1A	CUGUGCAGGAACACCUGGUG	385
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KDM1A	CGGAGCCGCCUGCUGUCCCA	388
KDM1A	CAGAGACUCCGGAGGGGCGU	389
KDM1A	CUGCUGCUUCAGCACACCCA	390
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KDM1A	GAGGUGGCCGCGCAGCCCGC	446
KDM1A	UUUCAUUUUCUUCCUCAGGU	447
KDM1A	AUCGAGGGGCCAGGAGUGAU	448
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KDM1A	GGCGGUUCCGCCAGGCCCCC	475
KDM1A	ACCGGGACGGAGGCUGGCCC	476
KDM1A	CACCGCCCCCGGCCCGACCU	477
KDM1A	CGCCAGGCCCCCGGGGGGCG	478

KDM1A	GGGGUGAGCUCUUCCUCUUC	479
KDM1A	ACCAUUUCAUCUUUCUCUUU	480
KDM1A	GGCCUCGCCCCCCGGGGGCC	481
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KDM1A	GGUCGGGCCGGGGGCGGUGG	488
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KDM1A	CUUCCUCAGGUGGGGCUUGA	490
KDM1A	GUGGGGCUUGAGGGGGUGGU	491
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KDM1A	GAAGGCGGCAGCCGCGGCGG	862

TALEN gene editing systems

TALENs are produced artificially by fusing a TAL effector DNA binding domain to a DNA cleavage domain. Transcription activator-like effects (TALEs) can be engineered to bind any desired DNA sequence, e.g., a target gene. By combining an engineered TALE with a DNA cleavage domain, a restriction enzyme can be produced which is specific to any desired DNA sequence. These can then be used, for example, as components of gene editing systems, e.g., TALEN gene editing systems, by for example, being introduced into a cell, wherein they can be used for genome editing. Boch (2011) *Nature Biotech*. 29: 135-6; and Boch et al. (2009) *Science* 326: 1509-12; Moscou et al. (2009) *Science* 326: 3501.

TALEs are proteins secreted by Xanthomonas bacteria. The DNA binding domain contains a repeated, highly conserved 33-34 amino acid sequence, with the exception of the 12th and 13th amino acids. These two positions are highly variable, showing a strong correlation with specific nucleotide recognition. They can thus be engineered to bind to a desired DNA sequence.

To produce a TALEN, a TALE protein is fused to a nuclease (N), which is, for example, a wild-type or mutated FokI endonuclease. Several mutations to FokI have been made for its use in TALENs; these, for example, improve cleavage specificity or activity. Cermak et al. (2011) *Nucl. Acids Res.* 39: e82; Miller et al. (2011) *Nature Biotech*. 29: 143-8; Hockemeyer et al. (2011) *Nature Biotech*. 29: 731-734; Wood et al. (2011) *Science* 333: 307; Doyon et al. (2010) *Nature Methods* 8: 74-79; Szczepek et al. (2007) *Nature Biotech.* 25: 786-793; and Guo et al. (2010) *J. Mol. Biol.* 200: 96.

The FokI domain functions as a dimer, requiring two constructs with unique DNA binding domains for sites in the target genome with proper orientation and spacing. Both the number of amino acid residues between the TALE DNA binding domain and the FokI cleavage domain and the number of bases between the two individual TALEN binding sites appear to be important parameters for achieving high levels of activity. Miller et al. (2011) *Nature Biotech.* 29: 143-8.

A TALEN (or pair of TALENs) can be used inside a cell to produce a double-stranded break (DSB). A mutation can be introduced at the break site if the repair mechanisms improperly repair the break via non-homologous end joining. For example, improper repair may introduce a frame shift mutation. Alternatively, foreign DNA can be introduced into the cell along with the TALEN, e.g., DNA encoding a transgene, and depending on the sequences of the foreign DNA and chromosomal sequence, this process can be used to integrate the transgene at or near the site targeted by the TALEN. TALENs specific to a target gene, e.g., LSD1, can be constructed using any method known in the art, including various schemes using modular components. Zhang et al. (2011) *Nature Biotech.* 29: 149-53; Geibler et al. (2011) *PLoS ONE* 6: e19509; US 8,420,782 ; US 8,470,973, the contents of which are hereby incorporated by reference in their entirety.

Thus, in exemplary embodiments, the genome editing system LSD1 inhibitor is a TALEN gene editing system directed to a sequence of an LSD1 gene, e.g., KDM1A. Such systems are known generally in the art and TALEN genome editing systems specific for LSD1 can be generated using known methods. See, e.g., Boch (2011) *Nature Biotech*. 29: 135-6; and Boch et al. (2009) *Science* 326: 1509-12; Moscou et al. (2009) *Science* 326: 3501; Zhang et al. (2011) *Nature Biotech.* 29: 149-53; Geibler et al. (2011) *PLoS ONE* 6: e19509; US patent number 8,420,782; US Patent Number 8,470,973.

Zinc finger nuclease (ZFN) gene editing systems

"ZFN" or "Zinc Finger Nuclease" refer to a zinc finger nuclease, an artificial nuclease or pair of nucleases which can be used, e.g., as part of a ZFN gene editing system to modify, e.g., insert or delete, one or more nucleic acids at or near a desired nucleic acid sequence, e.g., desired sequence of an LSD1 gene.

Like a TALEN, a ZFN comprises a FokI nuclease domain (or derivative thereof) fused to a DNA-binding domain. In the case of a ZFN, the DNA-binding domain comprises one or more zinc fingers. Carroll et al. (2011) *Genetics Society of America* 188: 773-782; and Kim et al. (1996) *Proc. Natl. Acad. Sci. USA* 93: 1156-1160.

A zinc finger is a small protein structural motif stabilized by one or more zinc ions. A zinc finger can comprise, for example, Cys2His2, and can recognize an approximately 3-bp sequence. Various zinc

fingers of known specificity can be combined to produce multi-finger polypeptides which recognize about 6, 9, 12, 15 or 18-bp sequences. Various selection and modular assembly techniques are available to generate zinc fingers (and combinations thereof) recognizing specific sequences, including phage display, yeast one-hybrid systems, bacterial one-hybrid and two-hybrid systems, and mammalian cells.

Like a TALEN, a ZFN must dimerize to cleave DNA. Thus, a pair of ZFNs are required to target non-palindromic DNA sites. The two individual ZFNs must bind opposite strands of the DNA with their nucleases properly spaced apart. Bitinaite et al. (1998) *Proc. Natl. Acad. Sci. USA* 95: 10570-5.

Also like a TALEN, a ZFN can create a double-stranded break in the DNA, which can create a frame-shift mutation if improperly repaired, leading to a decrease in the expression and amount of the target gene in a cell. ZFNs can also be used with homologous recombination to mutate the target gene or locus, or to introduce nucleic acid encoding a desired transgene at a site at or near the targeted sequence.

ZFNs specific to sequences in a target gene can be constructed using any method known in the art. See, e.g., Provasi (2011) Nature Med. 18: 807-815; Torikai (2013) Blood 122: 1341-1349; Cathomen et al. (2008) *Mol. Ther.* 16: 1200-7; and Guo et al. (2010) *J. Mol. Biol.* 400: 96; U.S. Patent Publication 2011/0158957; and U.S. Patent Publication 2012/0060230, the contents of which are hereby incorporated by reference in their entirety. In embodiments, The ZFN gene editing system may also comprise nucleic acid encoding one or more components of the ZFN gene editing system.

Thus, in exemplary embodiments, the genome editing system LSD1 inhibitor is a zinc finger nuclease gene editing system specific for a LSD1 gene, e.g., KDM1A. Such systems are known generally in the art and zinc finger nuclease genome editing systems specific for LSD1 can be generated using known methods. See, e.g., Provasi (2011) Nature Med. 18: 807-815; Torikai (2013) Blood 122: 1341-1349; Cathomen et al. (2008) *Mol. Ther.* 16: 1200-7; Guo et al. (2010) *J. Mol. Biol.* 400: 96; U.S. Patent Publication 2011/0158957; and U.S. Patent Publication 2012/0060230.

In an exemplary embodiment, the genome editing system LSD1 inhibitor is a meganuclease system. Such systems are known generally in the art and meganuclease genome editing systems specific for LSD1 can be generated using known methods.

Small Molecule LSD1 Inhibitors

In one aspect the LSD1 inhibitor is a small molecule. Exemplary small molecule LSD1 inhibitors are provided below, and additional candidate molecules may be identified by known assays, such as LSD1 binding assays and the assays described herein.

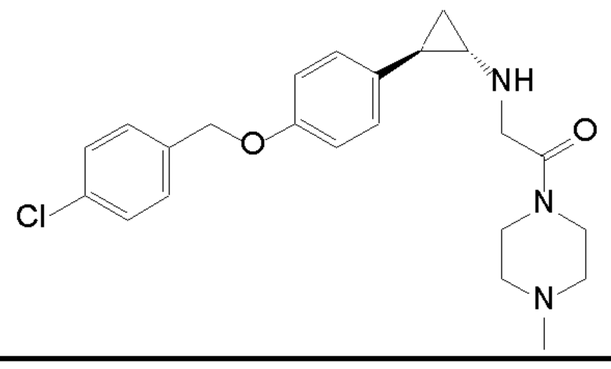
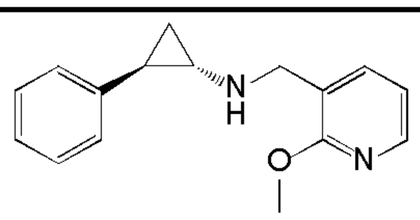
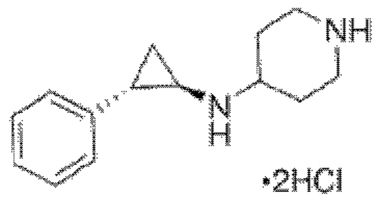
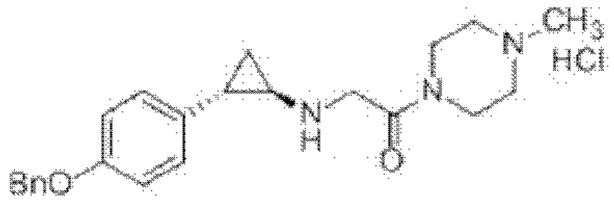
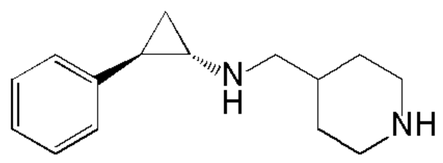
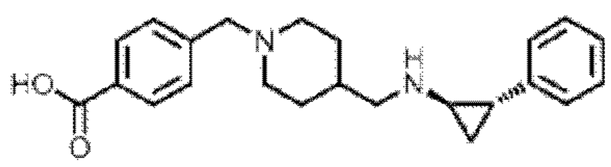
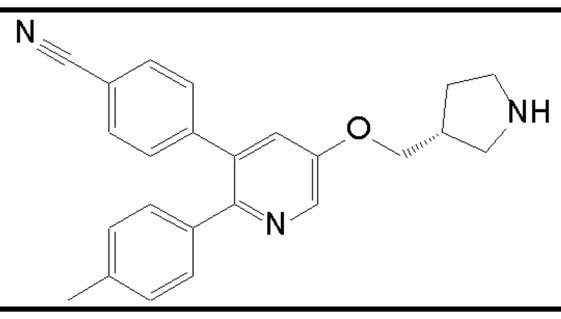
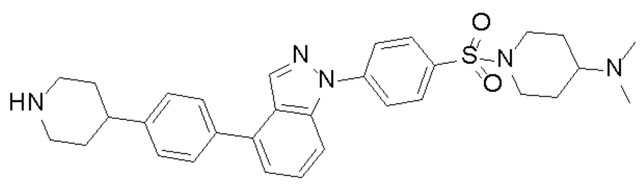
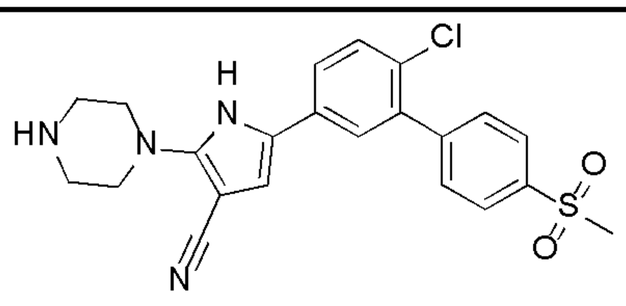
Useful lysine specific demethylase 1 (LSD1) inhibitors include both irreversible and reversible inhibitors. Reviews describing a variety of reversible and irreversible LSD1 inhibitors were published by Mould, Daniel P.,et al., "Reversible Inhibitors of LSD1 as Therapeutic Agents in Acute Myeloid Leukemia: Clinical Significance and Progress to Date," Med. Res. Rev., **35**, No. 3, 586–618, (2015); and Xheng, Yi-Choa, et. al., "A Systematic Review of Histone Lysine-Specific Demethylase 1 and Its Inhibitors" Med. Res. Rev., **35**, No. 5, 1032–1071, (2015), incorporated herein by reference. Suitable LSD1 inhibitors are also disclosed in PCT Patent Publication Nos. WO07/021839; WO2010/043721; WO2010/084160; WO2011/035941; WO2011/042217; WO2012/013727; WO2012/034116; WO2012/071469; WO2012/135113; WO2013/057320; WO2013/057322; WO2014/205213; WO2015/031564; WO2015/123408; WO2015/123437; WO2015/123465; and WO2015/156417. Representative examples of irreversible and reversible LSD1 inhibitors are described herein below.

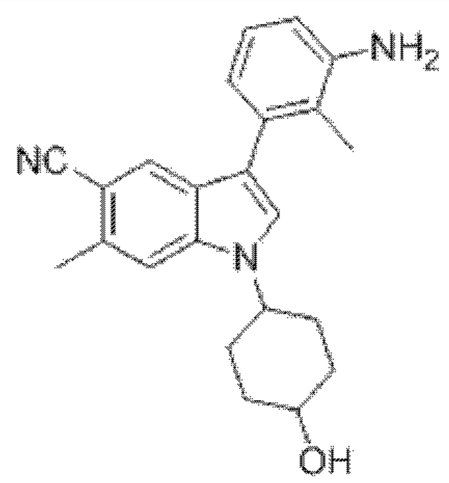
Exemplary irreversible LSD1 inhibitors include: GSK-LSD1 (trans-racemic) dihydrochloride, rel-N-[(1R,2S)-2-Phenylcyclopropyl]-4-piperidinamine hydrochloride (1:2) (available from Sigma-Aldrich); Tranylcypromine; N-[(1S,2R)-2-phenylcyclopropyl]-4-piperidinemethanamine (GSK2699537, described in PCT publication Nos. WO 2013057320 and WO 2012135113); 4-[[4-[[[(1R,2S)-2-phenylcyclopropyl]amino]methyl]-1-piperidinyl]methyl]-benzoic acid or a pharmaceutically acceptable salt thereof (GSK2879552, described in PCT publication No. WO 2012135113); *trans*-N1-[(1R,2S)-2-phenylcyclopropyl]-1,4-cyclohexanediamine or a pharmaceutically acceptable salt thereof (ORY-1001, described in PCT publication No. WO 2013057322); *rel*-1-(4-methyl-1-piperazinyl)-2-[[(1R*,2S*)-2-[4-phenylmethoxy)phenyl]cyclopropyl]amino]ethanone or a pharmaceutically acceptable salt thereof (RN-1, described in PCT Publication No. WO 2010043721); *rel*- 2-[[(1R,2S)-2-[4-[(4-chlorophenyl)methoxy] phenyl]cyclopropyl]amino]-1-(4-methyl-1-piperazinyl)-ethanone or a pharmaceutically acceptable salt thereof (described in PCT Publication No. WO 2010043721); 4'-((1R,2S)-2-Aminocyclopropyl)biphenyl-3-ol or a pharmaceutically acceptable salt thereof (OG-L002, described in PCT Publication No. WO 2012013727); (1S,2R)-N-((2-methoxypyridin-3-yl)methyl)-2-phenylcyclopropan-1-amine (described in PCT Publication No. WO2010/084160) or a pharmaceutically acceptable salt thereof.

Examplary reversible LSD1 inhibitors include: Namoline (available from ChemBridge, San Diego, CA); 3-(4-morpholinylsulfonyl)-benzoic acid, (2E)-2-[1-(5-chloro-2-hydroxyphenyl)ethylidene]hydrazide (SP-2509, described in PCT Publication No. WO 2014205213); 3-[[4-[4-(Aminoiminomethyl)benzoyl]-1-piperazinyl]carbonyl]-5-[[4-(aminoiminomethyl)-1-piperazinyl]methyl]-benzoic acid, methyl ester (CBB-1007, available from DSK Biopharma, Inc., and described in PCT Publication No. WO2012/071469); (R)-4-(5-(pyrrolidin-3-ylmethoxy)-2-(p-tolyl)pyridin-3-yl)benzonitrile (GSK354); N,N-dimethyl-1-((4-(4-(4-(piperidin-4-yl)phenyl)-1H-indazol-1-yl)phenyl)sulfonyl)piperidin-4-amine; 5-(6-chloro-4'-(methylsulfonyl)-[1,1'-biphenyl]-3-yl)-2-(piperazin-1-yl)-1H-pyrrole-3-carbonitrile; and trans-3-(3-amino-2-methylphenyl)-1-(4-hydroxycyclohexyl)-6-methyl-1H-indole-5-carbonitrile; or a pharmaceutically acceptable salt of any of the foregoing.

Additional exemplary LSD1 inhibitors are provided in Table 3, below.

Table 3.

Chemical Name	Structure	LCMS IC50 (uM)*	Also Referred to in this Application As
rel- 2-[[(1R,2S)-2-[4-[(4-chlorophenyl)methoxy]phenyl]cyclopropyl]amino]-1-(4-methyl-1-piperazinyl)-ethanone		0.01	
(1S,2R)-N-((2-methoxypyridin-3-yl)methyl)-2-phenylcyclopropan-1-amine		0.02	Compound A
rel- N-[(1R,2S)-2-Phenylcyclopropyl]-4-Piperidinamine hydrochloride (1:2)		--	GSK-LSD1; LSD1i-GSK
2-(1R,2S)-2-(4-(Benzyloxy)phenyl)cyclopropylamino)-1-(4-methylpiperazin-1-yl)ethanone, HCl		--	LSD1i-IV; LSD1i-EMD
GSK2699537		0.0007	Compound B
GSK2879552		--	
(R)-4-(5-(pyrrolidin-3-ylmethoxy)-2-(p-tolyl)pyridin-3-yl)benzonitrile		0.03	GSK354; Compound C
N,N-dimethyl-1-((4-(4-(4-(piperidin-4-yl)phenyl)-1H-indazol-1-yl)phenyl)sulfonyl)piperidin-4-amine		0.009	described in Example 3
5-(6-chloro-4'-(methylsulfonyl)-[1,1'-biphenyl]-3-yl)-2-(piperazin-1-yl)-1H-pyrrole-3-carbonitrile		0.012	described in Example 2

Trans-3-(3-amino-2-methylphenyl)-1-(4-hydroxycyclohexyl)-6-methyl-1H-indole-5-carbonitrile (Described in Example 6)		0.003	NVS Compound 1; Compound 93

* LSD1 IC50 as measured by LCMS.

Small molecule LSD1 inhibitors useful according to the present invention also include prodrugs, derivatives, pharmaceutically acceptable salts, or analogs thereof of any of the foregoing.

Small molecule LSD1 inhibitors may be formulated for delivery based on well-established methods in the art based on the particular dosages described herein.

In embodiments, the LSD1 small molecule inhibitor may be conjugated to an antibody or antigen binding fragment thereof. In an embodiment, the antibody or antigen-binding fragment thereof has specificity for an antigen expressed on the surface of a T cell.

Protein LSD1 Inhibitors

In embodiments, the LSD1 inhibitor may be a protein LSD1 inhibitor. In embodiments, the protein LSD1 inhibitor is a dominant negative binding partner of LSD1 (e.g., a histone deacetylase (HDAC) that interacts with LSD1 or other member of the Co-REST or AR co-activator complex), or nucleic acid encoding said dominant negative binding partner of LSD1. In embodiments, the protein LSD1 inhibitor is a dominant negative (e.g., catalytically inactive) LSD1, or nucleic acid encoding said molecule.

Methods of Preparing Populations of Immune Effector Cells Using LSD1 inhibitors

The invention features the use of LSD1 inhibitors in the manufacture of a population of immune effector cells, e.g., engineered to express a CAR molecule, e.g., as described herein. Without being bound by theory, the invention in part rests upon the surprising and unexpected discovery that inhibition of LSD1 in immune effector cells, e.g., T cells, results in a population of immune effector cells, e.g., T cells, with a higher number and/or higher proportion of naive immune effector cells, e.g., T cells, and with improved therapeutic properties. The inhibition of LSD1 in said immune effector cells may occur before and/or concurrently with therapy that includes said cells. Thus, one aspect of the

invention relates to compositions for and use of LSD1 inhibitors in the manufacture of immune effector cells, e.g., T cells.

In one aspect, the invention provides a method of making a population of immune effector cells, which is optionally a population of T cells, including the steps of:

a) contacting a population of immune effector cells with an LSD1 inhibitor;

thereby making a population of immune effector cells, which is optionally a population of T cells, wherein the contacting with the LSD1 inhibitor causes one or more of the following to occur:

1) an increase in the proportion of naive T cells, e.g., T_{SCM} cells;

2) an increase in the number of naive T cells, e.g., T_{SCM} cells;

3) a decrease in the number of T_{EM} cells;

4) a decrease in the proportion of T_{EM} cells;

5) an increase in the proportion of CD45RA+CD62L+ T cells;

6) an increase in the number of CD45RA+CD62L+ T cells;

7) an increase in the proportion of CD45RA+CCR7+ T cells;

8) an increase in the number of CD45RA+CCR7+ T cells;

9) a decrease in the proportion of PD-1 positive immune effector cells;

10) an increase in the ratio of PD-1 negative immune effector cells / PD-1 positive immune effector cells;

11) a decrease in the proportion of PD-1+/Lag3+/Tim3+ immune effector cells;

12) an increase in the ratio of PD-1-/Lag3-/Tim3- immune effector cells to PD-1+/Lag3+/Tim3+ immune effector cells;

13) an increase in the proliferation of the immune effector cells;

14) an increase in the production of cytokines (e.g., IFNg and/or IL-2) from said population of immune effector cells; or

15) a combination of two, three, four, five, six, seven, eight, nine, ten, eleven, twelve, thirteen or more (e.g., all) of the above;

optionally, as compared to a non-contacted population of immune effector cells.

In embodiments, the method further includes the step of b) inserting nucleic acid that encodes the CAR into cells of the population of immune effector cells. In embodiments, the contacting of step a) occurs 1) prior to; 2) concurrently with; 3) after; or 4) both before and after; said inserting of step b). In embodiments, the contacting of step a), and optionally the inserting of step b), is *ex vivo*.

In another aspect, the invention provides a method of making a population of immune effector cells, which is optionally a population of T cells, including the steps, optionally in the order listed, of:

a) providing a population of immune effector cells *ex vivo*;

b) contacting a population of immune effector cells *ex vivo* with an LSD1 inhibitor;

thereby making a population of immune effector cells, which is optionally a population of T cells, wherein the contacting with the LSD1 inhibitor causes one or more of the following to occur:

1) an increase in the proportion of naive T cells, e.g., T_{SCM} cells;

2) an increase in the number of naive T cells, e.g., T_{SCM} cells;

3) a decrease in the number of T_{EM} cells;

4) a decrease in the proportion of T_{EM} cells;

5) an increase in the proportion of CD45RA+CD62L+ T cells;

6) an increase in the number of CD45RA+CD62L+ T cells;

7) an increase in the proportion of CD45RA+CCR7+ T cells;

8) an increase in the number of CD45RA+CCR7+ T cells;

9) a decrease in the proportion of PD-1 positive immune effector cells;

10) an increase in the ratio of PD-1 negative immune effector cells / PD-1 positive immune effector cells;

11) a decrease in the proportion of PD-1+/Lag3+/Tim3+ immune effector cells;

12) an increase in the ratio of PD-1-/Lag3-/Tim3- immune effector cells to PD-1+/Lag3+/Tim3+ immune effector cells;

13) an increase in the proliferation of the immune effector cells;

14 an increase in the production of cytokines (e.g., IFNg and/or IL-2) from said population of immune effector cells; or

15) a combination of two, three, four, five, six, seven, eight, nine, ten, eleven, twelve, thirteen or more (e.g., all) of the above;

optionally, as compared to a non-contacted population of immune effector cells.

In embodiments, the method further includes the step of c) inserting nucleic acid that encodes the CAR into cells of the population of immune effector cells. In embodiments, the contacting of step b) occurs 1) prior to; 2) concurrently with; 3) after; or 4) both before and after; said inserting of step c). In embodiments, the contacting of step b), and optionally the inserting of step c), is *ex vivo*.

In another aspect, the invention provides a method of making a population of immune effector cells, which is optionally a population of T cells, including the steps, optionally in the order listed, of:

a) administering to a subject an LSD1 inhibitor;

wherein the administering the LSD1 inhibitor causes one or more of the following to occur in said subject:

1) an increase in the proportion of naive T cells, e.g., T_{SCM} cells;

2) an increase in the number of naive T cells, e.g., T_{SCM} cells;

3) a decrease in the number of T_{EM} cells;

4) a decrease in the proportion of T_{EM} cells;

5) an increase in the proportion of CD45RA+CD62L+ T cells;

6) an increase in the number of CD45RA+CD62L+ T cells;

7) an increase in the proportion of CD45RA+CCR7+ T cells;

8) an increase in the number of CD45RA+CCR7+ T cells;

9) a decrease in the proportion of PD-1 positive immune effector cells;

10) an increase in the ratio of PD-1 negative immune effector cells / PD-1 positive immune effector cells;

11) a decrease in the proportion of PD-1+/Lag3+/Tim3+ immune effector cells;

12) an increase in the ratio of PD-1-/Lag3-/Tim3- immune effector cells to PD-1+/Lag3+/Tim3+ immune effector cells;

13) an increase in the proliferation of the immune effector cells;

14) an increase in the production of cytokines (e.g., IFNg and/or IL-2) from said population of immune effector cells; or

15) a combination of two, three, four, five, six, seven, eight, nine, ten, eleven, twelve, thirteen or more (e.g., all) of the above;

optionally, as compared to a population of immune effector cells from a non-administered subject;

b) providing a population of immune effector cells from said subject *ex vivo*;

thereby making a population of immune effector cells, which is optionally a population of T cells.

In embodiments, the method further includes the step of c) inserting nucleic acid that encodes the CAR into cells of the population of immune effector cells.

In another aspect, the invention provides a method of making a population of immune effector cells, which is optionally a population of T cells, including the steps, optionally in the order listed, of:

a) administering to a subject an LSD1 inhibitor;

b) providing a population of immune effector cells from said subject *ex vivo;*

c) contacting a population of immune effector cells *ex vivo* with an LSD1 inhibitor;

thereby making a population of immune effector cells, which is optionally a population of T cells, wherein one or more of the following occurs:

1) an increase in the proportion of naive T cells, e.g., T_{SCM} cells;

2) an increase in the number of naive T cells, e.g., T_{SCM} cells;

3) a decrease in the number of T_{EM} cells;

4) a decrease in the proportion of T_{EM} cells;

5) an increase in the proportion of CD45RA+CD62L+ T cells;

6) an increase in the number of CD45RA+CD62L+ T cells;

7) an increase in the proportion of CD45RA+CCR7+ T cells;

8) an increase in the number of CD45RA+CCR7+ T cells;

9) a decrease in the proportion of PD-1 positive immune effector cells;

10) an increase in the ratio of PD-1 negative immune effector cells / PD-1 positive immune effector cells;

11) a decrease in the proportion of PD-1+/Lag3+/Tim3+ immune effector cells;

12) an increase in the ratio of PD-1-/Lag3-/Tim3- immune effector cells to PD-1+/Lag3+/Tim3+ immune effector cells;

13) an increase in the proliferation of the immune effector cells;

14 an increase in the production of cytokines (e.g., IFNg and/or IL-2) from said population of immune effector cells; or

15) a combination of two, three, four, five, six, seven, eight, nine, ten, eleven, twelve, thirteen or more (e.g., all) of the above;

optionally, as compared to a non-contacted and non-administered population of immune effector cells.

In embodiments, the method further includes the step of d) inserting nucleic acid that encodes the CAR into cells of the population of immune effector cells. In embodiments, the contacting of step c) occurs 1) prior to; 2) concurrently with; 3) after; or 4) both before and after; said inserting of step d).

In aspects the administration of the LSD1 inhibitor to the subject prior to collection of the population of immune effector cells from said subject may be of sufficient time and/or at a sufficient dose so that one or more of the following occurs:

1) an increase in the proportion of naive T cells, e.g., T_{SCM} cells;

2) an increase in the number of naive T cells, e.g., T_{SCM} cells;

3) a decrease in the number of T_{EM} cells;

4) a decrease in the proportion of T_{EM} cells;

5) an increase in the proportion of CD45RA+CD62L+ T cells;

6) an increase in the number of CD45RA+CD62L+ T cells;

7) an increase in the proportion of CD45RA+CCR7+ T cells;

8) an increase in the number of CD45RA+CCR7+ T cells;

9) a decrease in the proportion of PD-1 positive immune effector cells;

10) an increase in the ratio of PD-1 negative immune effector cells / PD-1 positive immune effector cells;

11) a decrease in the proportion of PD-1+/Lag3+/Tim3+ immune effector cells;

12) an increase in the ratio of PD-1-/Lag3-/Tim3- immune effector cells to PD-1+/Lag3+/Tim3+ immune effector cells;

13) an increase in the proliferation of the immune effector cells;

14) an increase in the production of cytokines (e.g., IFNg and/or IL-2) from said population of immune effector cells; or

15) a combination of two, three, four, five, six, seven, eight, nine, ten, eleven, twelve, thirteen or more (e.g., all) of the above;

optionally, as compared to non-administered population of immune effector cells. The assays described herein may be utilized in order to determine the proper dose and or time of administration. In embodiments, the LSD1 inhibitor is administered for a period of at least 1 day prior to collection of the population of immune effector cells from said subject. In embodiments, the LSD1 inhibitor is administered for a period of at least 2 days prior to collection of the population of immune effector cells from said subject. In embodiments, the LSD1 inhibitor is administered for a period of at least 3 days prior to collection of the population of immune effector cells from said subject. In embodiments, the LSD1 inhibitor is administered for a period of at least 4 days prior to collection of the population of immune effector cells from said subject. In embodiments, the LSD1 inhibitor is administered for a period of at least 5 days prior to collection of the population of immune effector cells from said subject. In embodiments, the LSD1 inhibitor is administered for a period of at least 6 days prior to collection of the population of immune effector cells from said subject. In embodiments, the LSD1 inhibitor is administered for a period of at least 7 days prior to collection of the population of immune effector cells

from said subject. In embodiments, the LSD1 inhibitor is administered for a period of at least a week or weeks prior to collection of the population of immune effector cells from said subject.

In embodiments, the administration of the LSD1 inhibitor to the subject continues after collection of the immune effector cells from said subject, e.g., continues for a period of 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20 or more days after collection of the immune effector cells, e.g., continues at least until the immune effector cells (modified *ex vivo*) are administered back to the subject, e.g., continues past the time when the immune effector cells (modified *ex vivo*) are administered back to the subject.

In aspects the contacting (e.g., ex vivo) of the LSD1 inhibitor to the population of immune effector cells may be of sufficient time and/or at a sufficient dose so that one or more of the following occurs:

1) an increase in the proportion of naive T cells, e.g., T_{SCM} cells;

2) an increase in the number of naive T cells, e.g., T_{SCM} cells;

3) a decrease in the number of T_{EM} cells;

4) a decrease in the proportion of T_{EM} cells;

5) an increase in the proportion of CD45RA+CD62L+ T cells;

6) an increase in the number of CD45RA+CD62L+ T cells;

7) an increase in the proportion of CD45RA+CCR7+ T cells;

8) an increase in the number of CD45RA+CCR7+ T cells;

9) a decrease in the proportion of PD-1 positive immune effector cells;

10) an increase in the ratio of PD-1 negative immune effector cells / PD-1 positive immune effector cells;

11) a decrease in the proportion of PD-1+/Lag3+/Tim3+ immune effector cells;

12) an increase in the ratio of PD-1-/Lag3-/Tim3- immune effector cells to PD-1+/Lag3+/Tim3+ immune effector cells;

13) an increase in the proliferation of the immune effector cells;

14 an increase in the production of cytokines (e.g., IFNg and/or IL-2) from said population of immune effector cells; or

15) a combination of two, three, four, five, six, seven, eight, nine, ten, eleven, twelve, thirteen or more (e.g., all) of the above;

optionally, as compared to non-contacted population of immune effector cells. The assays described herein may be utilized in order to determine the proper dose and or time of administration. In embodiments, the population of immune effector cells is contacted with an LSD1 inhibitor for a period of at least 1 day. In embodiments, the population of immune effector cells is contacted with an LSD1 inhibitor for a period of at least 2 days. In embodiments, the population of immune effector cells is contacted with an LSD1 inhibitor for a period of at least 3 days. In embodiments, the population of immune effector cells is contacted with an LSD1 inhibitor for a period of at least 4 days. In embodiments, the population of immune effector cells is contacted with an LSD1 inhibitor for a period of at least 5 days. In embodiments, the population of immune effector cells is contacted with an LSD1 inhibitor for a period of at least 6 days. In embodiments, the population of immune effector cells is contacted with an LSD1 inhibitor for a period of at least 7 days. In embodiments, the population of immune effector cells is contacted with an LSD1 inhibitor for a period of at least a week or weeks. In embodiments, media containing the LSD1 inhibitor is replaced with fresh media containing the LSD1 inhibitor, e.g., once, twice, three times, 4 times, 5 times, 6 times, 7 times, or more than 7 times (e.g., every day or every other day) during the time the immune effector cells are ex vivo. The concentration of LSD1 inhibitor can be adjusted in order that the desired effect occurs, and may be, for example, about 0.001 nM to about 10 mM, e.g., about 0.01 nM to about 1 mM, e.g., about 0.1 nM to about 100 uM, e.g., from about 1 nM to about 100 uM, e.g., from about 10 nM to about 100 uM, e.g., from about 100 nM to about 10 uM, e.g., from about 0.001 nM to about 100 nM, or e.g., from about 0.1 uM to about 10 uM. In embodiments, the concentration of LSD1 inhibitor is 100 nM. In embodiments, the concentration of LSD1 inhibitor is about 100 uM. In embodiments, the concentration of LSD1 inhibitor is 200 nM. In embodiments, the concentration of LSD1 inhibitor is about 200 uM.

In another aspect the invention provides a composition for use in *ex vivo* manufacturing a population of immune effector cells, that includes an LSD1 inhibitor, e.g., a small molecule LSD1 inhibitor. In embodiments, the composition includes the small molecule LSD1 inhibitor at a concentration of from about 0.001 nM to about 10 mM, e.g., from about 0.001 nM to about 100 nM, or, e.g., from about 0.1 uM to about 10 uM.

In embodiments involving immune effector cells engineered to express a CAR molecule, e.g., as described herein, it is understood that the method may further include any of the aspects, steps or features described below in the section relating to Chimeric Antigen Receptors.

Methods of Treatment with Immune Effector Cells and LSD1 inhibitors

The invention features the use of LSD1 inhibitors in the treatment of a disease, e.g., cancer, in a patient wherein such treatment is in combination with administration of a population of immune effector cells, e.g., immune effector cells engineered to express a CAR molecule, e.g., as described herein. Without being bound by theory, the invention in part rests upon the surprising and unexpected discovery that inhibition of LSD1 in immune effector cells, e.g., T cells, results in a population of immune effector cells, e.g., T cells, with a higher number and/or higher proportion of naive immune effector cells, e.g., T cells, and with improved therapeutic properties. Thus, one aspect of the invention provides treatment of a disease, e.g., a cancer, with a combination of a population of immune effector cells, e.g., engineered to express a CAR molecule, e.g., as described herein, and an LSD1 inhibitor.

In one aspect, the invention features a method of treating a subject that includes administering an LSD1 inhibitor to the subject, wherein said subject has received, is receiving or is about to receive a population of immune effector cells engineered to express a chimeric antigen receptor (CAR). In embodiments, the method includes administering to said subject an LSD1 inhibitor and a population of immune effector cells engineered to express a CAR molecule, e.g., as described herein. In embodiments, the LSD1 inhibitor is administered before the population of immune effector cells engineered to express a CAR molecule, e.g., as described herein, and wherein said administration of the LSD1 inhibitor is continued for a period of time after the administration of the population of immune effector cells engineered to express a CAR molecule, e.g., as described herein. In other embodiments, the administration of the LSD1 inhibitor after the administration of the population of immune effector cells engineered to express a CAR molecule, e.g., as described herein is in an amount sufficient to increase an anti-tumor effect of the population of immune effector cells engineered to express a CAR molecule, e.g., as described herein relative to an equivalent population of immune effector cells engineered to express a CAR molecule, e.g., as described herein administered in the absence of said LSD1 inhibitor.

In another aspect, the invention features a method of increasing the therapeutic efficacy in a subject of a population of immune effector cells engineered to express a CAR molecule, e.g., as described herein, e.g., a CAR19 (e.g., CTL019), including a step of decreasing the activity or expression of LSD1 in said cell, at least transiently. In embodiments, the step of decreasing the activity or expression of LSD1 in said cell includes contacting the cell with an LSD1 inhibitor. In embodiments, the contacting is

done ex vivo. In embodiments, the contacting is done in vivo (e.g., the population of immune effector cells and the LSD1 inhibitor are coadministered to the subject).

In embodiments of any of the forgoing aspect, the administration or the contacting of the LSD1 inhibitor results in:

1) an increase in the proportion of naive T cells, e.g., T_{SCM} cells;

2) an increase in the number of naive T cells, e.g., T_{SCM} cells;

3) a decrease in the number of T_{EM} cells;

4) a decrease in the proportion of T_{EM} cells;

5) an increase in the proportion of CD45RA+CD62L+ T cells;

6) an increase in the number of CD45RA+CD62L+ T cells;

7) an increase in the proportion of CD45RA+CCR7+ T cells;

8) an increase in the number of CD45RA+CCR7+ T cells;

9) a decrease in the proportion of PD-1 positive immune effector cells;

10) an increase in the ratio of PD-1 negative immune effector cells / PD-1 positive immune effector cells;

11) a decrease in the proportion of PD-1+/Lag3+/Tim3+ immune effector cells;

12) an increase in the ratio of PD-1-/Lag3-/Tim3- immune effector cells to PD-1+/Lag3+/Tim3+ immune effector cells;

13) an increase in the proliferation of the immune effector cells;

14) an increase in the production of cytokines (e.g., IFNg and/or IL-2) from said population of immune effector cells; or

15) a combination of two, three, four, five, six, seven, eight, nine, ten, eleven, twelve, thirteen or more (e.g., all) of the above.

In embodiments, the effect is as compared to cells not contacted with the LSD1 inhibitor. In embodiments, the effect is as compared to cells of the same subject not contacted with the LSD1 inhibitor.

In another aspect, the invention provides a method of treating a subject, that includes:

a) administering an LSD1 inhibitor to said subject;

b) collecting a population of immune effector cells from said subject after said administration of the LSD1 inhibitor;

c) providing said population of immune effector cells *ex vivo*;

d) contacting said *ex vivo* population of immune effector cells with the LSD1 inhibitor, wherein the contacting with the LSD1 inhibitor causes one or more of the following to occur:

1) an increase in the proportion of naive T cells, e.g., T_{SCM} cells;

2) an increase in the number of naive T cells, e.g., T_{SCM} cells;

3) a decrease in the number of T_{EM} cells;

4) a decrease in the proportion of T_{EM} cells;

5) an increase in the proportion of CD45RA+CD62L+ T cells;

6) an increase in the number of CD45RA+CD62L+ T cells;

7) an increase in the proportion of CD45RA+CCR7+ T cells;

8) an increase in the number of CD45RA+CCR7+ T cells;

9) a decrease in the proportion of PD-1 positive immune effector cells;

10) an increase in the ratio of PD-1 negative immune effector cells / PD-1 positive immune effector cells;

11) a decrease in the proportion of PD-1+/Lag3+/Tim3+ immune effector cells;

12) an increase in the ratio of PD-1-/Lag3-/Tim3- immune effector cells to PD-1+/Lag3+/Tim3+ immune effector cells;

13) an increase in the proliferation of the immune effector cells;

14) an increase in the production of cytokines (e.g., IFNg and/or IL-2) from said population of immune effector cells; or

15) a combination of two, three, four, five, six, seven, eight, nine, ten, eleven, twelve, thirteen or more (e.g., all) of the above;

optionally, as compared to a non-contacted *ex vivo* population of immune effector cells; and

e) administering the population of immune effector cells to the subject.

In embodiments, step of e) further includes administering the LSD1 inhibitor to the subject. In embodiments, the method further includes the step of inserting nucleic acid that encodes a CAR into cells of the *ex vivo* population of immune effector cells.

In another aspect the invention provides a method of treating a subject in need thereof, including administering to said subject an effective amount of the population of immune effector cells of any of the previous aspects and embodiments. In embodiments, the method further includes

administering to said subject an LSD1 inhibitor. In embodiments, the subject receives a pre-treatment of the LSD1 inhibitor, prior to the administration of the population of immune effector cells. In embodiments, the subject receives concurrent treatment with an LSD1 inhibitor and the population of immune effector cells. In embodiments, the subject receives treatment with an LSD1 inhibitor after administration of the population of immune effector cells; In embodiments, the subject receives a combination of any of the foregoing.

In an aspect, including in the previous aspects relating to methods of treatment, the invention relates to methods of treating a subject, wherein the subject has a disease associated with expression of a tumor antigen, e.g., a proliferative disease, a precancerous condition, a cancer, and a non-cancer related indication associated with expression of the tumor antigen. In embodiments, the cancer is a hematologic cancer chosen from one or more of chronic lymphocytic leukemia (CLL), acute leukemias, acute lymphoid leukemia (ALL), acute myeloid leukemia (AML), B-cell acute lymphoid leukemia (B-ALL), T-cell acute lymphoid leukemia (T-ALL), chronic myelogenous leukemia (CML), B cell prolymphocytic leukemia, blastic plasmacytoid dendritic cell neoplasm, Burkitt's lymphoma, diffuse large B cell lymphoma, follicular lymphoma, hairy cell leukemia, small cell- or a large cell-follicular lymphoma, malignant lymphoproliferative conditions, MALT lymphoma, mantle cell lymphoma, marginal zone lymphoma, multiple myeloma, myelodysplasia and myelodysplastic syndrome, non-Hodgkin’s lymphoma, Hodgkin’s lymphoma, plasmablastic lymphoma, plasmacytoid dendritic cell neoplasm, Waldenstrom macroglobulinemia, or pre-leukemia. In embodiments, the cancer is selected from the group consisting of colon cancer, rectal cancer, renal-cell carcinoma, liver cancer, non-small cell carcinoma of the lung, cancer of the small intestine, cancer of the esophagus, melanoma, bone cancer, pancreatic cancer, skin cancer, cancer of the head or neck, cutaneous or intraocular malignant melanoma, uterine cancer, ovarian cancer, rectal cancer, cancer of the anal region, stomach cancer, testicular cancer, uterine cancer, carcinoma of the fallopian tubes, carcinoma of the endometrium, carcinoma of the cervix, carcinoma of the vagina, carcinoma of the vulva, Hodgkin's Disease, non-Hodgkin's lymphoma, cancer of the endocrine system, cancer of the thyroid gland, cancer of the parathyroid gland, cancer of the adrenal gland, sarcoma of soft tissue, cancer of the urethra, cancer of the penis, solid tumors of childhood, cancer of the bladder, cancer of the kidney or ureter, carcinoma of the renal pelvis, neoplasm of the central nervous system (CNS), primary CNS lymphoma, tumor angiogenesis, spinal axis tumor, brain stem glioma, pituitary adenoma, Kaposi's sarcoma, epidermoid cancer, squamous cell cancer, T-cell lymphoma, environmentally induced cancers, combinations of said cancers, and metastatic lesions of said cancers.

In embodiments involving immune effector cells engineered to express a CAR molecule, e.g., as described herein, it is understood that the treatment method may further include any of the steps, aspects or features described below in the section relating to Chimeric Antigen Receptors.

Cells

As will be readily apparent to the skilled artisan from this disclosure, the invention relates to cells comprising LSD1 inhibitors. The invention further includes cells that have been contacted with an LSD1 inhibitor, e.g., for a period of time and/or at a dose sufficient for one or more of the following to occur:

1) an increase in the proportion of naive T cells, e.g., T_{SCM} cells;
2) an increase in the number of naive T cells, e.g., T_{SCM} cells;
3) a decrease in the number of T_{EM} cells;
4) a decrease in the proportion of T_{EM} cells;
5) an increase in the proportion of CD45RA+CD62L+ T cells;
6) an increase in the number of CD45RA+CD62L+ T cells;
7) an increase in the proportion of CD45RA+CCR7+ T cells;
8) an increase in the number of CD45RA+CCR7+ T cells;
9) a decrease in the proportion of PD-1 positive immune effector cells;
10) an increase in the ratio of PD-1 negative immune effector cells / PD-1 positive immune effector cells;
11) a decrease in the proportion of PD-1+/Lag3+/Tim3+ immune effector cells;
12) an increase in the ratio of PD-1-/Lag3-/Tim3- immune effector cells to PD-1+/Lag3+/Tim3+ immune effector cells;
13) an increase in the proliferation of the immune effector cells;
14 an increase in the production of cytokines (e.g., IFNg and/or IL-2) from said population of immune effector cells; or
15) a combination of two, three, four, five, six, seven, eight, nine, ten, eleven, twelve, thirteen or more (e.g., all) of the above;

optionally, relative to un-contacted cells.

The invention further relates to cells made by any of the methods described herein.

The cells are preferably immune effector cells. In an embodiment, the cells are T cells. In an embodiment, the cells are NK cells. In embodiments, the invention relates to a population of cells of the invention, e.g., a population of immune effector cells of the invention. In embodiments, the population of cells of the invention comprises cells of the type indicated, and may comprise other types (e.g., a population of immune effector cells, e.g., T cells, engineered to express a CAR molecule, e.g., as described herein, may include T cells engineered to express a CAR molecule as well as T cells (or other cell types) that have not been engineered to express a CAR molecule). In embodiments, the population of cells of the invention consists essentially of cells of the type indicated. In embodiments, the population of cells of the invention is substantially free of other cell types. In embodiments, the population of cells of the invention consists of the indicated cell type.

In any of the foregoing aspects and embodiments, the cells and/or population of cells are or include immune effector cells, e.g., the population of immune effector cells includes, e.g., consists of, T cells or NK cells. In embodiments the cells are T cells, e.g., CD8+ T cells, CD4+ T cells, or a combination thereof. In embodiments the cells are NK cells.

In embodiments the cells are human cells. In embodiments, the cells are autologous, e.g., to the subject to be administered the cells. In embodiments, the cells are allogeneic, e.g., to the subject to be administered the cells.

In embodiments, the cells are, or include, cells engineered to express a CAR molecule, e.g., as described herein. Additional features and/or aspects of the cells useful in the invention are described below in the section entitled Chimeric Antigen Receptors.

In one embodiment, the immune effector cells expressing a CAR molecule, e.g., a CAR molecule described herein, are obtained from a subject that has received an LSD1 inhibitor. In an embodiment, the population of immune effector cells, e.g., T cells, to be engineered to express a CAR molecule, are harvested after a sufficient time, or after sufficient dosing of the LSD1 inhibitor, such that the level of PD1 negative immune effector cells, e.g., T cells, or the ratio of PD1 negative immune effector cells, e.g., T cells/ PD1 positive immune effector cells, e.g., T cells, in the subject or harvested from the subject has been, at least transiently, increased.

In other embodiments, a population of immune effector cells, e.g., T cells, which have, or will be engineered to express a CAR molecule, e.g., as described herein, can be treated ex vivo by contact with an amount of an LSD1 inhibitor that increases the number of PD1 negative immune effector cells, e.g., T

cells or increases the ratio of PD1 negative immune effector cells, e.g., T cells/ PD1 positive immune effector cells, e.g., T cells.

In an embodiment, the NK cells are obtained from the subject. In another embodiment, the NK cells are an NK cell line, e.g., NK-92 cell line (Conkwest).

In an embodiment, immune effector cells, e.g., T cells, are obtained or harvested from a subject after administration to the subject of an LSD1 inhibitor.

In an embodiment, the immune effector cells, e.g., T cells, are collected after an increase in the number of PD1 negative immune effector, e.g., T cells, or after an increase in the ratio of PD1 negative immune effector, e.g., T cells/ PD1 positive immune effector, e.g., T cells, has occurred.

In an embodiment, the immune effector cells, e.g., T cells, are collected after an increase in the number of naive T cells has occurred.

In an embodiment, the immune effector cells, e.g., T cells, are collected after one or more of the following:

1) an increase in the proportion of naive T cells, e.g., T_{SCM} cells;

2) an increase in the number of naive T cells, e.g., T_{SCM} cells;

3) a decrease in the number of T_{EM} cells;

4) a decrease in the proportion of T_{EM} cells;

5) an increase in the proportion of CD45RA+CD62L+ T cells;

6) an increase in the number of CD45RA+CD62L+ T cells;

7) an increase in the proportion of CD45RA+CCR7+ T cells;

8) an increase in the number of CD45RA+CCR7+ T cells;

9) a decrease in the proportion of PD-1 positive immune effector cells;

10) an increase in the ratio of PD-1 negative immune effector cells / PD-1 positive immune effector cells;

11) a decrease in the proportion of PD-1+/Lag3+/Tim3+ immune effector cells;

12) an increase in the ratio of PD-1-/Lag3-/Tim3- immune effector cells to PD-1+/Lag3+/Tim3+ immune effector cells;

13) an increase in the proliferation of the immune effector cells;

14) an increase in the production of cytokines (e.g., IFNg and/or IL-2) from said population of immune effector cells; or

15) a combination of two, three, four, five, six, seven, eight, nine, ten, eleven, twelve, thirteen or more (e.g., all) of the above;

The increase or decrease can be transient. The increase or decrease can be permanent. The increase or decrease can be as compared with a standard, e.g., cells from an untreated subject.

In embodiment, immune effector cells, e.g., T cells, are contacted, ex vivo (after removal from the subject or a donor and before introduction into the subject), with an LSD1 inhibitor.

In an embodiment, the contact is at a level which results in an increase in the number of PD1 negative immune effector, e.g., T cells, or an increase in the ratio of PD1 negative immune effector cells, e.g., T cells/ PD1 positive immune effector, e.g., T cells.

In an embodiment, immune effector cells, e.g., T cells, are contacted, ex vivo (after removal from the subject or a donor and before introduction into the subject), with an LSD1 inhibitor, at a level which results in an increase in the number of naive T cells.

In an embodiment, immune effector cells, e.g., T cells, are contacted, ex vivo (after removal from the subject or a donor and before introduction into the subject), with an LSD1 inhibitor, at a level which results in one or more of the following:

1) an increase in the proportion of naive T cells, e.g., T_{SCM} cells;

2) an increase in the number of naive T cells, e.g., T_{SCM} cells;

3) a decrease in the number of T_{EM} cells;

4) a decrease in the proportion of T_{EM} cells;

5) an increase in the proportion of CD45RA+CD62L+ T cells;

6) an increase in the number of CD45RA+CD62L+ T cells;

7) an increase in the proportion of CD45RA+CCR7+ T cells;

8) an increase in the number of CD45RA+CCR7+ T cells;

9) a decrease in the proportion of PD-1 positive immune effector cells;

10) an increase in the ratio of PD-1 negative immune effector cells / PD-1 positive immune effector cells;

11) a decrease in the proportion of PD-1+/Lag3+/Tim3+ immune effector cells;

12) an increase in the ratio of PD-1-/Lag3-/Tim3- immune effector cells to PD-1+/Lag3+/Tim3+ immune effector cells;

13) an increase in the proliferation of the immune effector cells;

14) an increase in the production of cytokines (e.g., IFNg and/or IL-2) from said population of immune effector cells; or

15) a combination of two, three, four, five, six, seven, eight, nine, ten, eleven, twelve, thirteen or more (e.g., all) of the above;

The increase or decrease can be transient. The increase or decrease can be permanent. The increase or decrease can be as compared with a standard, e.g., cells from an untreated subject.

In an embodiment a preparation of T cells is evaluated for the level of increase in the number of PD1 negative immune effector, e.g., T cells, or an increase in the ratio of PD1 negative immune effector cells, e.g., T cells/ PD1 positive immune effector, e.g., T cells.

In an embodiment, a preparation of T cells is evaluated for the level of increase in the number of naive T cells. In an embodiment, a preparation of T cells is evaluated for one or more of the following:

1) an increase in the proportion of naive T cells, e.g., T_{SCM} cells;

2) an increase in the number of naive T cells, e.g., T_{SCM} cells;

3) a decrease in the number of T_{EM} cells;

4) a decrease in the proportion of T_{EM} cells;

5) an increase in the proportion of CD45RA+CD62L+ T cells;

6) an increase in the number of CD45RA+CD62L+ T cells;

7) an increase in the proportion of CD45RA+CCR7+ T cells;

8) an increase in the number of CD45RA+CCR7+ T cells;

9) a decrease in the proportion of PD-1 positive immune effector cells;

10) an increase in the ratio of PD-1 negative immune effector cells / PD-1 positive immune effector cells;

11) a decrease in the proportion of PD-1+/Lag3+/Tim3+ immune effector cells;

12) an increase in the ratio of PD-1-/Lag3-/Tim3- immune effector cells to PD-1+/Lag3+/Tim3+ immune effector cells;

13) an increase in the proliferation of the immune effector cells;

14) an increase in the production of cytokines (e.g., IFNg and/or IL-2) from said population of immune effector cells; or

15) a combination of two, three, four, five, six, seven, eight, nine, ten, eleven, twelve, thirteen or more (e.g., all) of the above;

The increase or decrease can be transient. The increase or decrease can be permanent. The increase or decrease can be as compared with a standard, e.g., cells of an untreated subject.

Pharmaceutical Compositions: LSD1 Inhibitors

In one aspect, the present invention relates to pharmaceutical compositions comprising an LSD1 inhibitor, e.g., an LSD1 inhibitor as described herein, formulated for use as a medicament.

In one aspect, the present invention relates to pharmaceutical compositions comprising an LSD1 inhibitor, e.g., an LSD1 inhibitor as described herein, formulated for use in the manufacture of a population of immune effector cells.

In one aspect, the present invention relates to pharmaceutical compositions comprising an LSD1 inhibitor, e.g., an LSD1 inhibitor as described herein, formulated for use in combination with CAR cells described herein.

In some embodiments, the LSD1 inhibitor is formulated for administration in combination with another agent, in addition to a CAR cell, e.g., as described herein.

In general, compounds of the invention will be administered in therapeutically effective amounts as described above via any of the usual and acceptable modes known in the art, either singly or in combination with one or more therapeutic agents.

The pharmaceutical formulations may be prepared using conventional dissolution and mixing procedures. For example, the bulk drug substance (e.g., an LSD1 inhibitor or stabilized form of the compound (e.g., complex with a cyclodextrin derivative or other known complexation agent) is dissolved in a suitable solvent in the presence of one or more of the excipients described herein. The LSD1 inhibitor is typically formulated into pharmaceutical dosage forms to provide an easily controllable dosage of the drug and to give the patient an elegant and easily handleable product.

Compounds of the invention can be administered as pharmaceutical compositions by any conventional route, in particular enterally, e.g., orally, e.g., in the form of tablets or capsules, or parenterally, e.g., in the form of injectable solutions or suspensions, topically, e.g., in the form of lotions, gels, ointments or creams, or in a nasal or suppository form. Where an LSD1 inhibitor is administered in combination with (either simultaneously with or separately from) another agent as described herein, in one aspect, both components can be administered by the same route (e.g., parenterally). Alternatively, another agent may be administered by a different route relative to the LSD1 inhibitor. For example, an LSD1 inhibitor may be administered orally and the other agent may be administered parenterally. Pharmaceutical compositions comprising an LSD1 inhibitor in free form or in a pharmaceutically acceptable salt form in association with at least one pharmaceutically acceptable carrier or diluent can

be manufactured in a conventional manner by mixing, granulating or coating methods. For example, oral compositions can be tablets or gelatin capsules comprising the active ingredient together with a) diluents, e.g., lactose, dextrose, sucrose, mannitol, sorbitol, cellulose and/or glycine; b) lubricants, e.g., silica, talcum, stearic acid, its magnesium or calcium salt and/or polyethyleneglycol; for tablets also c) binders, e.g., magnesium aluminum silicate, starch paste, gelatin, tragacanth, methylcellulose, sodium carboxymethylcellulose and or polyvinylpyrrolidone; if desired d) disintegrants, e.g., starches, agar, alginic acid or its sodium salt, or effervescent mixtures; and/or e) absorbents, colorants, flavors and sweeteners. Oral formulations can also comprise the active ingredient along with 20-60% Eudragit EPO, Hydroxypropyl cellulose EF, Hydroxypropyl methylcellulose, or Kollidon VA64, and up to 5% of pluronic F68, Cremophor EL, or Gelucire 44/14. Injectable compositions can be aqueous isotonic solutions or suspensions, and suppositories can be prepared from fatty emulsions or suspensions. The compositions may be sterilized and/or contain adjuvants, such as preserving, stabilizing, wetting or emulsifying agents, solution promoters, salts for regulating the osmotic pressure and/or buffers. In addition, they may also contain other therapeutically valuable substances. Suitable formulations for transdermal applications include an effective amount of a compound of the present invention with a carrier. A carrier can include absorbable pharmacologically acceptable solvents to assist passage through the skin of the host. For example, transdermal devices are in the form of a bandage comprising a backing member, a reservoir containing the compound optionally with carriers, optionally a rate controlling barrier to deliver the compound to the skin of the host at a controlled and predetermined rate over a prolonged period of time, and means to secure the device to the skin. Matrix transdermal formulations may also be used. In a further aspect, the LSD1 inhibitors described herein may be administered via a microneedle patch. Microneedle based drug delivery is well known in the art (See, e.g., U.S. Pat. 8,162,901) and these technologies and methods may be adapted by one of skill in the art for administration of an LSD1 inhibitor as described herein. Suitable formulations for topical application, e.g., to the skin and eyes, are preferably aqueous solutions, ointments, creams or gels well-known in the art. Such formulations may contain solubilizers, stabilizers, tonicity enhancing agents, buffers and preservatives.

The pharmaceutical composition (or formulation) for application may be packaged in a variety of ways depending upon the method used for administering the drug. Generally, an article for distribution includes a container having deposited therein the pharmaceutical formulation in an appropriate form. Suitable containers are well-known to those skilled in the art and include materials such as bottles (plastic and glass), sachets, ampoules, plastic bags, metal cylinders, and the like. The container may also include a tamper-proof assemblage to prevent indiscreet access to the contents of the package. In

addition, the container has deposited thereon a label that describes the contents of the container. The label may also include appropriate warnings. The invention also provides for a pharmaceutical combinations, e.g. a kit, comprising a) a first agent which is an LSD1 inhibitor as disclosed herein, in free form or in pharmaceutically acceptable salt form, and b) at least one additional agent. The kit can comprise instructions for its administration.

The term "pharmaceutical combination" as used herein means a product that results from the mixing or combining of more than one active ingredient and includes both fixed and non-fixed combinations of the active ingredients. The term "fixed combination" means that the active ingredients, e.g. an LSD1 inhibitor and other agent, are both administered to a patient simultaneously in the form of a single entity or dosage. The term "non-fixed combination" means that the active ingredients, e.g. an LSD1 inhibitor and other agent, are both administered to a patient as separate entities either simultaneously, concurrently or sequentially with no specific time limits, wherein such administration provides therapeutically effective levels of the 2 compounds in the body of the patient. The latter also applies to cocktail therapy, e.g. the administration of 3 or more active ingredients.

Chimeric Antigen Receptors

General Description of Chimeric Antigen Receptor Technology Relevant to the Invention

Described herein are methods for combining the administration of LSD1 inhibitors with administration of a population of immune effector cells, e.g., T cells or NK cells, engineered to express a CAR molecule, e.g., as described herein (the cell is engineered to express a CAR, and in embodiments, expresses the CAR by the time at which it is administered to the subject. In other embodiments, expression initiates after administration.) In some embodiments, the cell is a T cell engineered to express a CAR molecule, e.g., as described herein, wherein the CAR T cell ("CART") exhibits an anticancer property. Also described herein are methods for using LSD1 inhibitors for the manufacture, e.g., the activation and/or expansion, a population of immune effector cells, e.g., T cells or NK cells, engineered to express a CAR molecule, e.g., as described herein, wherein the cells have enhanced activity (e.g., proliferation, cytokine release, and/or tumor targeting efficacy) and/or a more naive phenotype, relative to cells manufactured without the use of LSD1 inhibitors. In general, the molecules, cells, methods or other aspects discussed in this section may be useful in the methods, compositions, cells and other aspects of the invention, e.g., in combination with LSD1 inhibitors.

In general, the invention pertains to an isolated nucleic acid molecule encoding a chimeric antigen receptor (CAR), wherein the CAR comprises an antigen binding domain (e.g., antibody or

antibody fragment, TCR or TCR fragment) that binds to a tumor antigen as described herein, a transmembrane domain (e.g., a transmembrane domain described herein), and an intracellular signaling domain (e.g., an intracellular signaling domain described herein) (e.g., an intracellular signaling domain comprising a costimulatory domain (e.g., a costimulatory domain described herein) and/or a primary signaling domain (e.g., a primary signaling domain described herein). In other aspects, the invention includes: host cells containing the above nucleic acids and isolated proteins encoded by such nucleic acid molecules. CAR nucleic acid constructs, encoded proteins, containing vectors, host cells, pharmaceutical compositions, and methods of administration and treatment related to the present invention are disclosed in detail in International Patent Application Publication No. WO2015/142675, which is incorporated by reference in its entirety.

In one aspect, the invention pertains to an isolated nucleic acid molecule encoding a chimeric antigen receptor (CAR), wherein the CAR comprises an antigen binding domain (e.g., antibody or antibody fragment, TCR or TCR fragment) that binds to a tumor-supporting antigen (e.g., a tumor-supporting antigen as described herein), a transmembrane domain (e.g., a transmembrane domain described herein), and an intracellular signaling domain (e.g., an intracellular signaling domain described herein) (e.g., an intracellular signaling domain comprising a costimulatory domain (e.g., a costimulatory domain described herein) and/or a primary signaling domain (e.g., a primary signaling domain described herein). In some embodiments, the tumor-supporting antigen is an antigen present on a stromal cell or a myeloid-derived suppressor cell (MDSC). In other aspects, the invention features polypeptides encoded by such nucleic acids and host cells containing such nucleic acids and/or polypeptides. In other aspects, the invention features cells (e.g., a population of cells), e.g., immune effector cells, e.g., T cells or NK cells, engineered to express a CAR molecule, e.g., as described herein.

Targets

The present invention provides immune effector cells (e.g., T cells, NK cells) that are engineered to contain one or more CARs that direct the immune effector cells to undesired cells (e.g., cancer cells). This is achieved through an antigen binding domain on the CAR that is specific for a cancer associated antigen. There are two classes of cancer associated antigens (tumor antigens) that can be targeted by the CARs of the instant invention: (1) cancer associated antigens that are expressed on the surface of cancer cells; and (2) cancer associated antigens that itself is intracellar, however, a fragment of such antigen (peptide) is presented on the surface of the cancer cells by MHC (major histocompatibility complex).

In some embodiments, the tumor antigen is chosen from one or more of: CD19; CD123; CD22; CD30; CD171; CS-1 (also referred to as CD2 subset 1, CRACC, SLAMF7, CD319, and 19A24); C-type lectin-like molecule-1 (CLL-1 or CLECL1); CD33; epidermal growth factor receptor variant III (EGFRvIII); ganglioside G2 (GD2); ganglioside GD3 (aNeu5Ac(2-8)aNeu5Ac(2-3)bDGalp(1-4)bDGlcp(1-1)Cer); TNF receptor family member B cell maturation (BCMA); Tn antigen ((Tn Ag) or (GalNAcα-Ser/Thr)); prostate-specific membrane antigen (PSMA); Receptor tyrosine kinase-like orphan receptor 1 (ROR1); Fms-Like Tyrosine Kinase 3 (FLT3); Tumor-associated glycoprotein 72 (TAG72); CD38; CD44v6; Carcinoembryonic antigen (CEA); Epithelial cell adhesion molecule (EPCAM); B7H3 (CD276); KIT (CD117); Interleukin-13 receptor subunit alpha-2 (IL-13Ra2 or CD213A2); Mesothelin; Interleukin 11 receptor alpha (IL-11Ra); prostate stem cell antigen (PSCA); Protease Serine 21 (Testisin or PRSS21); vascular endothelial growth factor receptor 2 (VEGFR2); Lewis(Y) antigen; CD24; Platelet-derived growth factor receptor beta (PDGFR-beta); Stage-specific embryonic antigen-4 (SSEA-4); CD20; Folate receptor alpha; Receptor tyrosine-protein kinase ERBB2 (Her2/neu); Mucin 1, cell surface associated (MUC1); epidermal growth factor receptor (EGFR); neural cell adhesion molecule (NCAM); Prostase; prostatic acid phosphatase (PAP); elongation factor 2 mutated (ELF2M); Ephrin B2; fibroblast activation protein alpha (FAP); insulin-like growth factor 1 receptor (IGF-I receptor), carbonic anhydrase IX (CAIX); Proteasome (Prosome, Macropain) Subunit, Beta Type, 9 (LMP2); glycoprotein 100 (gp100); oncogene fusion protein consisting of breakpoint cluster region (BCR) and Abelson murine leukemia viral oncogene homolog 1 (Abl) (bcr-abl); tyrosinase; ephrin type-A receptor 2 (EphA2); Fucosyl GM1; sialyl Lewis adhesion molecule (sLe); ganglioside GM3 (aNeu5Ac(2-3)bDGalp(1-4)bDGlcp(1-1)Cer); transglutaminase 5 (TGS5); high molecular weight-melanoma-associated antigen (HMWMAA); o-acetyl-GD2 ganglioside (OAcGD2); Folate receptor beta; tumor endothelial marker 1 (TEM1/CD248); tumor endothelial marker 7-related (TEM7R); claudin 6 (CLDN6); thyroid stimulating hormone receptor (TSHR); G protein-coupled receptor class C group 5, member D (GPRC5D); chromosome X open reading frame 61 (CXORF61); CD97; CD179a; anaplastic lymphoma kinase (ALK); Polysialic acid; placenta-specific 1 (PLAC1); hexasaccharide portion of globoH glycoceramide (GloboH); mammary gland differentiation antigen (NY-BR-1); uroplakin 2 (UPK2); Hepatitis A virus cellular receptor 1 (HAVCR1); adrenoceptor beta 3 (ADRB3); pannexin 3 (PANX3); G protein-coupled receptor 20 (GPR20); lymphocyte antigen 6 complex, locus K 9 (LY6K); Olfactory receptor 51E2 (OR51E2); TCR Gamma Alternate Reading Frame Protein (TARP); Wilms tumor protein (WT1); Cancer/testis antigen 1 (NY-ESO-1); Cancer/testis antigen 2 (LAGE-1a); Melanoma-associated antigen 1 (MAGE-A1); ETS translocation-variant gene 6, located on chromosome 12p (ETV6-AML); sperm protein 17 (SPA17); X Antigen Family, Member 1A (XAGE1); angiopoietin-binding cell surface receptor 2

(Tie 2); melanoma cancer testis antigen-1 (MAD-CT-1); melanoma cancer testis antigen-2 (MAD-CT-2); Fos-related antigen 1; tumor protein p53 (p53); p53 mutant; prostein; surviving; telomerase; prostate carcinoma tumor antigen-1 (PCTA-1 or Galectin 8), melanoma antigen recognized by T cells 1 (MelanA or MART1); Rat sarcoma (Ras) mutant; human Telomerase reverse transcriptase (hTERT); sarcoma translocation breakpoints; melanoma inhibitor of apoptosis (ML-IAP); ERG (transmembrane protease, serine 2 (TMPRSS2) ETS fusion gene); N-Acetyl glucosaminyl-transferase V (NA17); paired box protein Pax-3 (PAX3); Androgen receptor; Cyclin B1; v-myc avian myelocytomatosis viral oncogene neuroblastoma derived homolog (MYCN); Ras Homolog Family Member C (RhoC); Tyrosinase-related protein 2 (TRP-2); Cytochrome P450 1B1 (CYP1B1); CCCTC-Binding Factor (Zinc Finger Protein)-Like (BORIS or Brother of the Regulator of Imprinted Sites), Squamous Cell Carcinoma Antigen Recognized By T Cells 3 (SART3); Paired box protein Pax-5 (PAX5); proacrosin binding protein sp32 (OY-TES1); lymphocyte-specific protein tyrosine kinase (LCK); A kinase anchor protein 4 (AKAP-4); synovial sarcoma, X breakpoint 2 (SSX2); Receptor for Advanced Glycation Endproducts (RAGE-1); renal ubiquitous 1 (RU1); renal ubiquitous 2 (RU2); legumain; human papilloma virus E6 (HPV E6); human papilloma virus E7 (HPV E7); intestinal carboxyl esterase; heat shock protein 70-2 mutated (mut hsp70-2); CD79a; CD79b; CD72; Leukocyte-associated immunoglobulin-like receptor 1 (LAIR1); Fc fragment of IgA receptor (FCAR or CD89); Leukocyte immunoglobulin-like receptor subfamily A member 2 (LILRA2); CD300 molecule-like family member f (CD300LF); C-type lectin domain family 12 member A (CLEC12A); bone marrow stromal cell antigen 2 (BST2); EGF-like module-containing mucin-like hormone receptor-like 2 (EMR2); lymphocyte antigen 75 (LY75); Glypican-3 (GPC3); Fc receptor-like 5 (FCRL5); and immunoglobulin lambda-like polypeptide 1 (IGLL1).

A CAR described herein can comprise an antigen binding domain (e.g., antibody or antibody fragment, TCR or TCR fragment) that binds to a tumor-supporting antigen (e.g., a tumor-supporting antigen as described herein). In some embodiments, the tumor-supporting antigen is an antigen present on a stromal cell or a myeloid-derived suppressor cell (MDSC). Stromal cells can secrete growth factors to promote cell division in the microenvironment. MDSC cells can inhibit T cell proliferation and activation. Without wishing to be bound by theory, in some embodiments, the CAR-expressing cells destroy the tumor-supporting cells, thereby indirectly inhibiting tumor growth or survival.

In embodiments, the stromal cell antigen is chosen from one or more of: bone marrow stromal cell antigen 2 (BST2), fibroblast activation protein (FAP) and tenascin. In an embodiment, the FAP-specific antibody is, competes for binding with, or has the same CDRs as, sibrotuzumab. In

embodiments, the MDSC antigen is chosen from one or more of: CD33, CD11b, C14, CD15, and CD66b. Accordingly, in some embodiments, the tumor-supporting antigen is chosen from one or more of: bone marrow stromal cell antigen 2 (BST2), fibroblast activation protein (FAP) or tenascin, CD33, CD11b, C14, CD15, and CD66b.

Antigen Binding Domain Structures

In some embodiments, the antigen binding domain of the encoded CAR molecule comprises an antibody, an antibody fragment, an scFv, a Fv, a Fab, a (Fab')2, a single domain antibody (SDAB), a VH or VL domain, a camelid VHH domain or a bi-functional (e.g. bi-specific) hybrid antibody (e.g., Lanzavecchia et al., Eur. J. Immunol. 17, 105 (1987)).

In some instances, scFvs can be prepared according to method known in the art (see, for example, Bird et al., (1988) Science 242:423-426 and Huston et al., (1988) Proc. Natl. Acad. Sci. USA 85:5879-5883). ScFv molecules can be produced by linking VH and VL regions together using flexible polypeptide linkers. The scFv molecules comprise a linker (e.g., a Ser-Gly linker) with an optimized length and/or amino acid composition. The linker length can greatly affect how the variable regions of a scFv fold and interact. In fact, if a short polypeptide linker is employed (e.g., between 5-10 amino acids) intrachain folding is prevented. Interchain folding is also required to bring the two variable regions together to form a functional epitope binding site. For examples of linker orientation and size see, e.g., Hollinger et al. 1993 Proc Natl Acad. Sci. U.S.A. 90:6444-6448, U.S. Patent Application Publication Nos. 2005/0100543, 2005/0175606, 2007/0014794, and PCT publication Nos. WO2006/020258 and WO2007/024715, is incorporated herein by reference.

An scFv can comprise a linker of at least 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 25, 30, 35, 40, 45, 50, or more amino acid residues between its VL and VH regions. The linker sequence may comprise any naturally occurring amino acid. In some embodiments, the linker sequence comprises amino acids glycine and serine. In another embodiment, the linker sequence comprises sets of glycine and serine repeats such as $(Gly_4Ser)n$, where n is a positive integer equal to or greater than 1 (SEQ ID NO:22). In one embodiment, the linker can be $(Gly_4Ser)_4$ (SEQ ID NO:29) or $(Gly_4Ser)_3$(SEQ ID NO:30). Variation in the linker length may retain or enhance activity, giving rise to superior efficacy in activity studies.

In another aspect, the antigen binding domain is a T cell receptor ("TCR"), or a fragment thereof, for example, a single chain TCR (scTCR). Methods to make such TCRs are known in the art. See, e.g., Willemsen RA et al, Gene Therapy 7: 1369–1377 (2000); Zhang T et al, Cancer Gene Ther 11: 487–496

(2004); Aggen et al, Gene Ther. 19(4):365-74 (2012) (references are incorporated herein by its entirety). For example, scTCR can be engineered that contains the Vα and Vβ genes from a T cell clone linked by a linker (e.g., a flexible peptide). This approach is very useful to cancer associated target that itself is intracellar, however, a fragment of such antigen (peptide) is presented on the surface of the cancer cells by MHC.

In certain embodiments, the encoded antigen binding domain has a binding affinity KD of 10^{-4} M to 10^{-8} M.

In one embodiment, the encoded CAR molecule comprises an antigen binding domain that has a binding affinity KD of 10-4 M to 10-8 M, e.g., 10-5 M to 10-7 M, e.g., 10-6 M or 10-7 M, for the target antigen. In one embodiment, the antigen binding domain has a binding affinity that is at least five-fold, 10-fold, 20-fold, 30-fold, 50-fold, 100-fold or 1,000-fold less than a reference antibody, e.g., an antibody described herein. In one embodiment, the encoded antigen binding domain has a binding affinity at least 5-fold less than a reference antibody (e.g., an antibody from which the antigen binding domain is derived). In one aspect such antibody fragments are functional in that they provide a biological response that can include, but is not limited to, activation of an immune response, inhibition of signal-transduction origination from its target antigen, inhibition of kinase activity, and the like, as will be understood by a skilled artisan.

In one aspect, the antigen binding domain of the CAR is a scFv antibody fragment that is humanized compared to the murine sequence of the scFv from which it is derived.

In one aspect, the antigen binding domain of a CAR of the invention (e.g., a scFv) is encoded by a nucleic acid molecule whose sequence has been codon optimized for expression in a mammalian cell. In one aspect, entire CAR construct of the invention is encoded by a nucleic acid molecule whose entire sequence has been codon optimized for expression in a mammalian cell. Codon optimization refers to the discovery that the frequency of occurrence of synonymous codons (i.e., codons that code for the same amino acid) in coding DNA is biased in different species. Such codon degeneracy allows an identical polypeptide to be encoded by a variety of nucleotide sequences. A variety of codon optimization methods is known in the art, and include, e.g., methods disclosed in at least US Patent Numbers 5,786,464 and 6,114,148.

Specific Antigen Binding Domains

In some embodiments, the portion of the CAR comprising the antigen binding domain comprises an antigen binding domain that targets a tumor antigen, e.g., a tumor antigen described herein (e.g., in

the section entitled “Targets.” In some embodiments, the tumor antigen is a tumor antigen described in International Application WO2015/142675, filed March 13, 2015, which is herein incorporated by reference in its entirety. Exemplary target antigens that can be targeted using the CAR-expressing cells, include, but are not limited to, CD19, CD123, EGFRvIII, CD33, mesothelin, BCMA, and GFR ALPHA-4, among others, as described in, for example, WO2014/153270, WO 2014/130635, WO2016/028896, WO 2014/130657, WO2016/014576, WO 2015/090230, WO2016/014565, WO2016/014535, and WO2016/025880, each of which is herein incorporated by reference in its entirety.

In embodiments, the antigen binding domain comprises one, two, three (e.g., all three) heavy chain CDRs, HC CDR1, HC CDR2 and HC CDR3, from an antibody described herein or in any of the publications incorporated by reference herein, and/or one, two, three (e.g., all three) light chain CDRs, LC CDR1, LC CDR2 and LC CDR3, from an antibody described herein or in any of the publications incorporated by reference herein. In one embodiment, the antigen binding domain comprises a heavy chain variable region and/or a variable light chain region of an antibody described herein or in any of the publications incorporated by reference herein. In one embodiment, the antigen binding domain of any of the CAR molecules described herein (e.g., any of CD19, CD123, EGFRvIII, CD33, mesothelin, BCMA, and GFR ALPHA-4) comprises one, two, three (e.g., all three) heavy chain CDRs, HC CDR1, HC CDR2 and HC CDR3, from an antibody described herein or in any of the publications incorporated by reference herein, and/or one, two, three (e.g., all three) light chain CDRs, LC CDR1, LC CDR2 and LC CDR3, from an antigen binding domain described herein or in any of the publications incorporated by reference herein. In one embodiment, the antigen binding domain comprises a heavy chain variable region and/or a variable light chain region of an antibody described herein or in any of the publications incorporated by reference herein.

In one embodiment, the antigen binding domain comprises one, two three (e.g., all three) heavy chain CDRs, HC CDR1, HC CDR2 and HC CDR3, from an antibody described herein (e.g., an antibody described in WO2015/142675, US-2015-0283178-A1, US-2016-0046724-A1, US2014/0322212A1, US2016/0068601A1, US2016/0051651A1, US2016/0096892A1, US2014/0322275A1, or WO2015/090230, incorporated herein by reference), and/or one, two, three (e.g., all three) light chain CDRs, LC CDR1, LC CDR2 and LC CDR3, from an antibody described herein (e.g., an antibody described in WO2015/142675, US-2015-0283178-A1, US-2016-0046724-A1, US2014/0322212A1, US2016/0068601A1, US2016/0051651A1, US2016/0096892A1, US2014/0322275A1, or WO2015/090230, incorporated herein by reference). In one embodiment, the antigen binding domain comprises a heavy chain variable region and/or a variable light chain region of an antibody described

herein. In embodiments, the antigen binding domain is an antigen binding domain described in WO2015/142675, US-2015-0283178-A1, US-2016-0046724-A1, US2014/0322212A1, US2016/0068601A1, US2016/0051651A1, US2016/0096892A1, US2014/0322275A1, or WO2015/090230, incorporated herein by reference.

In one embodiment, an antigen binding domain against CD19 is an antigen binding portion, e.g., CDRs, of a CAR (e.g., CD19 CAR), antibody or antigen-binding fragment thereof described in, e.g., PCT publication WO2012/079000; PCT publication WO2014/153270; Kochenderfer, J.N. et al., J. Immunother. 32 (7), 689-702 (2009); Kochenderfer, J.N., et al., Blood, 116 (20), 4099-4102 (2010); PCT publication WO2014/031687; Bejcek, Cancer Research, 55, 2346-2351, 1995; or U.S. Patent No. 7,446,190.

In one embodiment, the CD19 CAR includes a CAR molecule, or an antigen binding domain (e.g., a humanized antigen binding domain) according to Table 3 of WO2014/153270, incorporated herein by reference. The amino acid and nucleotide sequences encoding the CD19 CAR molecules and antigen binding domains (e.g., including one, two, three VH CDRs; and one, two, three VL CDRs according to Kabat or Chothia), are specified in WO2014/153270. In embodiments, the CD19 CAR, or antigen binding domain, comprises an amino acid, or has a nucleotide sequence shown in WO2014/153270 incorporated herein by reference, or a sequence substantially identical to any of the aforesaid sequences (e.g., at least 85%, 90%, 95% or more identical to any of the aforesaid sequences).

In another embodiment, the antigen binding domain comprises an anti-CD19 antibody, or fragment thereof, e.g., an scFv. For example, the antigen binding domain comprises a variable heavy chain and a variable light chain listed in Tables 6-9, or a sequence substantially identical to any of the aforesaid sequences (e.g., at least 85%, 90%, 95% or more identical to any of the aforesaid sequences). The linker sequence joining the variable heavy and variable light chains can be, e.g., any of the linker sequences described herein, or alternatively, can be GSTSGSGKPGSGEGSTKG (SEQ ID NO: 871).

Table 6: Anti-CD19 antibody binding domains

CD19	huscFv1 (SEQ ID NO: 872)	EIVMTQSPATLSLSPGERATLSCRASQDISKYLNWYQQKPGQAPRLLIYHTSRL HSGIPARFSGSGSGTDYTLTISSLQPEDFAVYFCQQGNTLPYTFGQGTKLEIKG GGGSGGGGSGGGGSQVQLQESGPGLVKPSETLSLTCTVSGVSLPDYGVSWIRQP PGKGLEWIGVIWGSETTYYSSSLKSRVTISKDNSKNQVSLKLSSVTAADTAVYY CAKHYYYGGSYAMDYWGQGTLVTVSS
CD19	huscFv2 (SEQ ID	Eivmtqspatlslspgeratlscrasqdiskylnwyqqkpgqaprlliyhtsrlhsgip arfsgsgsgtdytltisslqpedfavyfcqqgntlpytfgqgtkleikggggsggggsg gggsqvqlqesgpglvkpsetlsltctvsgvslpdygvswirqppgkglewigviwgse

	NO: 873)	ttyyqsslksrvtiskdnsknqvslklssvtaadtavyycakhyyyggsyamdywgqgtlvtvss
CD19	huscFv3 (SEQ ID NO: 874)	Qvqlqesgpglvkpsetlsltctvsgvslpdygvswirqppgkglewigviwgsettyyssslksrvtiskdnsknqvslklssvtaadtavyycakhyyyggsyamdywgqgtlvtvssggggsggggsggggseivmtqspatlslspgeratlscrasqdiskylnwyqqkpgqaprlliyhtsrlhsgiparfsgsgsgtdytltisslqpedfavyfcqqgntlpytfgqgtkleik
CD19	huscFv4 (SEQ ID NO: 875)	Qvqlqesgpglvkpsetlsltctvsgvslpdygvswirqppgkglewigviwgsettyyqsslksrvtiskdnsknqvslklssvtaadtavyycakhyyyggsyamdywgqgtlvtvssggggsggggsggggseivmtqspatlslspgeratlscrasqdiskylnwyqqkpgqaprlliyhtsrlhsgiparfsgsgsgtdytltisslqpedfavyfcqqgntlpytfgqgtkleik
CD19	huscFv5 (SEQ ID NO: 876)	Eivmtqspatlslspgeratlscrasqdiskylnwyqqkpgqaprlliyhtsrlhsgiparfsgsgsgtdytltisslqpedfavyfcqqgntlpytfgqgtkleikggggsggggsggggsggggsqvqlqesgpglvkpsetlsltctvsgvslpdygvswirqppgkglewigviwgsettyyssslksrvtiskdnsknqvslklssvtaadtavyycakhyyyggsyamdywgqgtlvtvss
CD19	huscFv6 (SEQ ID NO: 877)	Eivmtqspatlslspgeratlscrasqdiskylnwyqqkpgqaprlliyhtsrlhsgiparfsgsgsgtdytltisslqpedfavyfcqqgntlpytfgqgtkleikggggsggggsggggsggggsqvqlqesgpglvkpsetlsltctvsgvslpdygvswirqppgkglewigviwgsettyyqsslksrvtiskdnsknqvslklssvtaadtavyycakhyyyggsyamdywgqgtlvtvss
CD19	huscFv7 (SEQ ID NO: 878)	Qvqlqesgpglvkpsetlsltctvsgvslpdygvswirqppgkglewigviwgsettyyssslksrvtiskdnsknqvslklssvtaadtavyycakhyyyggsyamdywgqgtlvtvssggggsggggsggggsggggseivmtqspatlslspgeratlscrasqdiskylnwyqqkpgqaprlliyhtsrlhsgiparfsgsgsgtdytltisslqpedfavyfcqqgntlpytfgqgtkleik
CD19	huscFv8 (SEQ ID NO: 879)	Qvqlqesgpglvkpsetlsltctvsgvslpdygvswirqppgkglewigviwgsettyyqsslksrvtiskdnsknqvslklssvtaadtavyycakhyyyggsyamdywgqgtlvtvssggggsggggsggggsggggseivmtqspatlslspgeratlscrasqdiskylnwyqqkpgqaprlliyhtsrlhsgiparfsgsgsgtdytltisslqpedfavyfcqqgntlpytfgqgtkleik
CD19	huscFv9 (SEQ ID NO: 880)	Eivmtqspatlslspgeratlscrasqdiskylnwyqqkpgqaprlliyhtsrlhsgiparfsgsgsgtdytltisslqpedfavyfcqqgntlpytfgqgtkleikggggsggggsggggsggggsqvqlqesgpglvkpsetlsltctvsgvslpdygvswirqppgkglewigviwgsettyynsslksrvtiskdnsknqvslklssvtaadtavyycakhyyyggsyamdywgqgtlvtvss
CD19	Hu scFv10 (SEQ ID NO: 881)	Qvqlqesgpglvkpsetlsltctvsgvslpdygvswirqppgkglewigviwgsettyynsslksrvtiskdnsknqvslklssvtaadtavyycakhyyyggsyamdywgqgtlvtvssggggsggggsggggsggggseivmtqspatlslspgeratlscrasqdiskylnwyqqkpgqaprlliyhtsrlhsgiparfsgsgsgtdytltisslqpedfavyfcqqgntlpytfgqgtkleik
CD19	Hu scFv11 (SEQ ID NO: 882)	Eivmtqspatlslspgeratlscrasqdiskylnwyqqkpgqaprlliyhtsrlhsgiparfsgsgsgtdytltisslqpedfavyfcqqgntlpytfgqgtkleikggggsggggsggggsqvqlqesgpglvkpsetlsltctvsgvslpdygvswirqppgkglewigviwgsettyynsslksrvtiskdnsknqvslklssvtaadtavyycakhyyyggsyamdywgqgtlvtvss
CD19	Hu scFv12 (SEQ ID NO: 883)	Qvqlqesgpglvkpsetlsltctvsgvslpdygvswirqppgkglewigviwgsettyynsslksrvtiskdnsknqvslklssvtaadtavyycakhyyyggsyamdywgqgtlvtvssggggsggggsggggseivmtqspatlslspgeratlscrasqdiskylnwyqqkpgqaprlliyhtsrlhsgiparfsgsgsgtdytltisslqpedfavyfcqqgntlpytfgqgtkleik
CD19	muCTL019 (SEQ ID NO:	Diqmtqttsslsaslgdrvtiscrasqdiskylnwyqqkpdgtvkllliyhtsrlhsgvpsrfsgsgsgtdysltisnleqediatyfcqqgntlpytfgggtkleitggggsggggsggggsevklqesgpglvapsqslsvtctvsgvslpdygvswirqpprkglewlgviwgse

	884)	ttyynsalksrltiikdnsksqvflkmnslqtddtaiyycakhyyyyggsyamdywgqgt svtvss

Table 7: Additional anti-CD19 antibody binding domains

Antibody	VH Sequence	VL Sequence
SSJ25-C1	QVQLLESGAELVRPGSSVKISCKASGYAFSS YWMNWVKQRPGQGLEWIGQIYPGDGDTNYNG KFKGQATLTADKSSSTAYMQLSGLTSEDSAV YSCARKTISSVVDFYFDYWGQGTTVT (SEQ ID NO: 885)	ELVLTQSPKFMSTSVGDRVSVTCKASQNVGTNVA WYQQKPGQSPKPLIYSATYRNSGVPDRFTGSGSG TDFTLTITNVQSKDLADYFYFCQYNRYPYTSGGG TKLEIKRRS (SEQ ID NO: 886)

Table 8: Additional murine anti-CD19 antibody binding domains

mCAR1 scFv	SEQ ID NO: 887	QVQLLESGAELVRPGSSVKISCKASGYAFSSYWMNWVKQRPGQGLEWIGQIYPGDG DTNYNGKFKGQATLTADKSSSTAYMQLSGLTSEDSAVYSCARKTISSVVDFYFDYW GQGTTVTGGGSGGGSGGGSGGGSELVLTQSPKFMSTSVGDRVSVTCKASQNVGTNV AWYQQKPGQSPKPLIYSATYRNSGVPDRFTGSGSGTDFTLTITNVQSKDLADYFCQ YNRYPYTSFFFTKLEIKRRS
mCAR2 scFv	SEQ ID NO: 888	DIQMTQTTSSLSASLGDRVTISCRASQDISKYLNWYQQKPDGTVKLLIYHTSRLHS GVPSRFSGSGSGTDYSLTISNLEQEDIATYFCQQGNTLPYTFGGGTKLEITGSTSG SGKPGSGEGSTKGEVKLQESGPGLVAPSQSLSVTCTVSGVSLPDYGVSWIRQPPRK GLEWLGVIWGSETTYYNSALKSRLTIIKDNSKSQVFLKMNSLQTDDTAIYYCAKHY YYGGSYAMDYWGQGTSVTVSSE
mCAR3 scFv	SEQ ID NO: 889	DIQMTQTTSSLSASLGDRVTISCRASQDISKYLNWYQQKPDGTVKLLIYHTSRLHS GVPSRFSGSGSGTDYSLTISNLEQEDIATYFCQQGNTLPYTFGGGTKLEITGSTSG SGKPGSGEGSTKGEVKLQESGPGLVAPSQSLSVTCTVSGVSLPDYGVSWIRQPPRK GLEWLGVIWGSETTYYNSALKSRLTIIKDNSKSQVFLKMNSLQTDDTAIYYCAKHY YYGGSYAMDYWGQGTSVTVSS

Any CD19 CAR, e.g., the CD19 antigen binding domain of any known CD19 CAR, can be used in accordance with the present disclosure. For example, LG-740; CD19 CAR described in the US Pat. No. 8,399,645; US Pat. No. 7,446,190; Xu et al., Leuk Lymphoma. 2013 54(2):255-260(2012); Cruz et al., Blood 122(17):2965-2973 (2013); Brentjens et al., Blood, 118(18):4817-4828 (2011); Kochenderfer et al., Blood 116(20):4099-102 (2010); Kochenderfer et al., Blood 122 (25):4129-39(2013); and 16th Annu Meet Am Soc Gen Cell Ther (ASGCT) (May 15-18, Salt Lake City) 2013, Abst 10.

In one embodiment, an antigen binding domain against EGFRvIII is an antigen binding portion, e.g., CDRs, of a CAR, antibody or antigen-binding fragment thereof described in, e.g., PCT publication

WO2014/130657 or US2014/0322275A1. In one embodiment, the CAR molecule comprises an EGFRvIII CAR, or an antigen binding domain according to Table 2 or SEQ ID NO:11 of WO 2014/130657, incorporated herein by reference, or a sequence substantially identical thereto (e.g., at least 85%, 90%, 95% or more identical thereto). The amino acid and nucleotide sequences encoding the EGFRvIII CAR molecules and antigen binding domains (*e.g.*, including one, two, three VH CDRs; and one, two, three VL CDRs according to Kabat or Chothia), are specified in WO 2014/130657.

In one embodiment, an antigen binding domain against mesothelin is an antigen binding portion, e.g., CDRs, of an antibody, antigen-binding fragment or CAR described in, e.g., PCT publication WO2015/090230. In one embodiment, an antigen binding domain against mesothelin is an antigen binding portion, e.g., CDRs, of an antibody, antigen-binding fragment, or CAR described in, e.g., PCT publication WO1997/025068, WO1999/028471, WO2005/014652, WO2006/099141, WO2009/045957, WO2009/068204, WO2013/142034, WO2013/040557, or WO2013/063419.

In an embodiment, the CAR molecule comprises a mesothelin CAR described herein, e.g., a mesothelin CAR described in WO 2015/090230, incorporated herein by reference. In embodiments, the mesothelin CAR comprises an amino acid, or has a nucleotide sequence shown in WO 2015/090230 incorporated herein by reference, or a sequence substantially identical to any of the aforesaid sequences (e.g., at least 85%, 90%, 95% or more identical to any of the aforesaid mesothelin CAR sequences). In one embodiment, the CAR molecule comprises a mesothelin CAR, or an antigen binding domain according to Tables 2-3 of WO 2015/090230, incorporated herein by reference, or a sequence substantially identical thereto (e.g., at least 85%, 90%, 95% or more identical thereto). The amino acid and nucleotide sequences encoding the mesothelin CAR molecules and antigen binding domains (e.g., including one, two, three VH CDRs; and one, two, three VL CDRs according to Kabat or Chothia), are specified in WO 2015/090230.

In one embodiment, an antigen binding domain against CD123 is an antigen binding portion, e.g., CDRs, of an antibody, antigen-binding fragment or CAR described in, e.g., PCT publication WO2016/028896. In one embodiment, an antigen binding domain against CD123 is an antigen binding portion, e.g., CDRs, of an antibody, antigen-binding fragment, or CAR described in, e.g., PCT publication WO2014/138805, WO2014/138819, WO2013/173820, WO2014/144622, WO2001/66139, WO2010/126066, WO2014/144622, or US2009/0252742.

In one embodiment, an antigen binding domain against CD123 is an antigen binding portion, e.g., CDRs, of an antibody, antigen-binding fragment or CAR described in, e.g.,US2014/0322212A1 or US2016/0068601A1, both incorporated herein by reference. In embodiments, the CD123 CAR comprises an amino acid, or has a nucleotide sequence shown in US2014/0322212A1 or US2016/0068601A1, both incorporated herein by reference, or a sequence substantially identical to any of the aforesaid sequences (e.g., at least 85%, 90%, 95% or more identical to any of the aforesaid CD123 CAR sequences). In one embodiment, the CAR molecule comprises a CD123 CAR (*e.g.,* any of the CAR1-CAR8), or an antigen binding domain according to Tables 1-2 of WO 2014/130635, incorporated herein by reference, or a sequence substantially identical thereto (e.g., at least 85%, 90%, 95% or more identical to any of the aforesaid CD123 CAR sequences). The amino acid and nucleotide sequences encoding the CD123 CAR molecules and antigen binding domains (e.g., including one, two, three VH CDRs; and one, two, three VL CDRs according to Kabat or Chothia), are specified in WO 2014/130635.

In other embodiments, the CAR molecule comprises a CD123 CAR comprises a CAR molecule (e.g., any of the CAR123-1 to CAR123-4 and hzCAR123-1 to hzCAR123-32), or an antigen binding domain according to Tables 2, 6, and 9 of WO2016/028896, incorporated herein by reference, or a sequence substantially identical thereto (e.g., at least 85%, 90%, 95% or more identical to any of the aforesaid CD123 CAR sequences). The amino acid and nucleotide sequences encoding the CD123 CAR molecules and antigen binding domains (e.g., including one, two, three VH CDRs; and one, two, three VL CDRs according to Kabat or Chothia), are specified in WO2016/028896.

In one embodiment, an antigen binding domain against CD22 is an antigen binding portion, e.g., CDRs, of an antibody described in, e.g., Haso et al., Blood, 121(7): 1165-1174 (2013); Wayne et al., Clin Cancer Res 16(6): 1894-1903 (2010); Kato et al., Leuk Res 37(1):83-88 (2013); Creative BioMart (creativebiomart.net): MOM-18047-S(P).

In one embodiment, an antigen binding domain against CS-1 is an antigen binding portion, e.g., CDRs, of Elotuzumab (BMS), see e.g., Tai et al., 2008, Blood 112(4):1329-37; Tai et al., 2007, Blood. 110(5):1656-63.

In one embodiment, an antigen binding domain against CLL-1 is an antigen binding portion, e.g., CDRs, of an antibody available from R&D, ebiosciences, Abcam, for example, PE-CLL1-hu Cat# 353604 (BioLegend); and PE-CLL1 (CLEC12A) Cat# 562566 (BD).

In other embodiments, the CLL1 CAR includes a CAR molecule, or an antigen binding domain according to Table 2 of WO2016/014535, incorporated herein by reference. The amino acid and

nucleotide sequences encoding the CLL-1 CAR molecules and antigen binding domains (e.g., including one, two, three VH CDRs; and one, two, three VL CDRs according to Kabat or Chothia), are specified in WO2016/014535.

In one embodiment, an antigen binding domain against CD33 is an antigen binding portion, e.g., CDRs, of an antibody described in, e.g., Bross et al., Clin Cancer Res 7(6):1490-1496 (2001) (Gemtuzumab Ozogamicin, hP67.6),Caron et al., Cancer Res 52(24):6761-6767 (1992) (Lintuzumab, HuM195), Lapusan et al., Invest New Drugs 30(3):1121-1131 (2012) (AVE9633), Aigner et al., Leukemia 27(5): 1107-1115 (2013) (AMG330, CD33 BiTE), Dutour et al., Adv hematol 2012:683065 (2012), and Pizzitola et al., Leukemia doi:10.1038/Lue.2014.62 (2014).

In one embodiment, an antigen binding domain against CD33 is an antigen binding portion, e.g., CDRs, of an antibody described in, US2016/0096892A1, incorporated herein by reference. In embodiments, the CD33 CAR comprises an amino acid, or has a nucleotide sequence shown in US2016/0096892A1, incorporated herein by reference, or a sequence substantially identical to any of the aforesaid sequences (e.g., at least 85%, 90%, 95% or more identical to any of the aforesaid CD33 CAR sequences). In other embodiments, the CD33 CAR CAR or antigen binding domain thereof can include a CAR molecule (e.g., any of CAR33-1 to CAR-33-9), or an antigen binding domain according to Table 2 or 9 of WO2016/014576, incorporated herein by reference, or a sequence substantially identical to any of the aforesaid sequences (e.g., at least 85%, 90%, 95% or more identical to any of the aforesaid CD33 CAR sequences). The amino acid and nucleotide sequences encoding the CD33 CAR molecules and antigen binding domains (e.g., including one, two, three VH CDRs; and one, two, three VL CDRs according to Kabat or Chothia), are specified in WO2016/014576.

In one embodiment, an antigen binding domain against GD2 is an antigen binding portion, e.g., CDRs, of an antibody described in, e.g., Mujoo et al., Cancer Res. 47(4):1098-1104 (1987); Cheung et al., Cancer Res 45(6):2642-2649 (1985), Cheung et al., J Clin Oncol 5(9):1430-1440 (1987), Cheung et al., J Clin Oncol 16(9):3053-3060 (1998), Handgretinger et al., Cancer Immunol Immunother 35(3):199-204 (1992). In some embodiments, an antigen binding domain against GD2 is an antigen binding portion of an antibody selected from mAb 14.18, 14G2a, ch14.18, hu14.18, 3F8, hu3F8, 3G6, 8B6, 60C3, 10B8, ME36.1, and 8H9, see e.g., WO2012033885, WO2013040371, WO2013192294, WO2013061273, WO2013123061, WO2013074916, and WO201385552. In some embodiments, an antigen binding domain against GD2 is an antigen binding portion of an antibody described in US Publication No.: 20100150910 or PCT Publication No.: WO 2011160119.

In one embodiment, an antigen binding domain against BCMA is an antigen binding portion, e.g., CDRs, of an antibody, antigen-binding fragment or CAR described in, e.g., PCT publication WO2016/014565, e.g., the antigen binding portion of CAR BCMA-10 as described in WO2016/014565. In one embodiment, an antigen binding domain against BCMA is an antigen binding portion, e.g., CDRs, of an antibody, antigen-binding fragment or CAR described in, e.g., PCT publication WO2016/014789. In one embodiment, an antigen binding domain against BCMA is an antigen binding portion, e.g., CDRs, of an antibody described in, e.g., WO2012/163805, WO2001/12812, and WO2003/062401.

In other embodiment, the CAR molecule comprises a BCMA CAR molecule, or an antigen binding domain against BCMA described herein, e.g., a BCMA CAR described in US-2016-0046724-A1 or WO2016/014565. In embodiments, the BCMA CAR comprises an amino acid, or has a nucleotide sequence of a CAR molecule, or an antigen binding domain according to US-2016-0046724-A1, or Table 1 or 16, SEQ ID NO: 271 or SEQ ID NO: 273 of WO2016/014565, incorporated herein by reference, or a sequence substantially identical to any of the aforesaid sequences (e.g., at least 85%, 90%, 95% or more identical to any of the aforesaid BCMA CAR sequences). The amino acid and nucleotide sequences encoding the BCMA CAR molecules and antigen binding domains (*e.g.*, including one, two, three VH CDRs; and one, two, three VL CDRs according to Kabat or Chothia), are specified in WO2016/014565.

In one embodiment, an antigen binding domain against GFR ALPHA-4 CAR antigen is an antigen binding portion, e.g., CDRs, of an antibody described in, e.g., WO2016/025880, incorporated herein by reference. In one embodiment, the CAR molecule comprises an a GFR ALPHA-4 CAR, e.g., a CAR molecule, or an antigen binding domain according to Table 2 of WO2016/025880, incorporated herein by reference, or a sequence substantially identical to any of the aforesaid sequences (e.g., at least 85%, 90%, 95% or more identical to any of the aforesaid GFR ALPHA-4 sequences). The amino acid and nucleotide sequences encoding the GFR ALPHA-4 CAR molecules and antigen binding domains (e.g., including one, two, three VH CDRs; and one, two, three VL CDRs according to Kabat or Chothia), are specified in WO2016/025880.

In one embodiment, an antigen binding domain against Tn antigen is an antigen binding portion, e.g., CDRs, of an antibody described in, e.g., US8,440,798; Brooks et al., PNAS 107(22):10056-10061 (2010), and Stone et al., OncoImmunology 1(6):863-873(2012).

In one embodiment, an antigen binding domain against PSMA is an antigen binding portion, e.g., CDRs, of an antibody described in, e.g., Parker et al., Protein Expr Purif 89(2):136-145 (2013), US

20110268656 (J591 ScFv); Frigerio et al, European J Cancer 49(9):2223-2232 (2013) (scFvD2B); WO 2006125481 (mAbs 3/A12, 3/E7 and 3/F11) and single chain antibody fragments (scFv A5 and D7).

In one embodiment, an antigen binding domain against ROR1 is an antigen binding portion, e.g., CDRs, of an antibody described in, e.g., Hudecek et al., Clin Cancer Res 19(12):3153-3164 (2013); WO 2011159847; and US20130101607.

In one embodiment, an antigen binding domain against FLT3 is an antigen binding portion, e.g., CDRs, of an antibody described in, e.g., WO2011076922, US5777084, EP0754230, US20090297529, and several commercial catalog antibodies (R&D, ebiosciences, Abcam).

In one embodiment, an antigen binding domain against TAG72 is an antigen binding portion, e.g., CDRs, of an antibody described in, e.g., Hombach et al., Gastroenterology 113(4):1163-1170 (1997); and Abcam ab691.

In one embodiment, an antigen binding domain against FAP is an antigen binding portion, e.g., CDRs, of an antibody described in, e.g., Ostermann et al., Clinical Cancer Research 14:4584-4592 (2008) (FAP5), US Pat. Publication No. 2009/0304718; sibrotuzumab (see e.g., Hofheinz et al., Oncology Research and Treatment 26(1), 2003); and Tran et al., J Exp Med 210(6):1125-1135 (2013).

In one embodiment, an antigen binding domain against CD38 is an antigen binding portion, e.g., CDRs, of daratumumab (see, e.g., Groen et al., Blood 116(21):1261-1262 (2010); MOR202 (see, e.g., US8,263,746); or antibodies described in US8,362,211.

In one embodiment, an antigen binding domain against CD44v6 is an antigen binding portion, e.g., CDRs, of an antibody described in, e.g., Casucci et al., Blood 122(20):3461-3472 (2013).

In one embodiment, an antigen binding domain against CEA is an antigen binding portion, e.g., CDRs, of an antibody described in, e.g., Chmielewski et al., Gastoenterology 143(4):1095-1107 (2012).

In one embodiment, an antigen binding domain against EPCAM is an antigen binding portion, e.g., CDRS, of an antibody selected from MT110, EpCAM-CD3 bispecific Ab (see, e.g., clinicaltrials.gov/ct2/show/NCT00635596); Edrecolomab; 3622W94; ING-1; and adecatumumab (MT201).

In one embodiment, an antigen binding domain against PRSS21 is an antigen binding portion, e.g., CDRs, of an antibody described in US Patent No.: 8,080,650.

In one embodiment, an antigen binding domain against B7H3 is an antigen binding portion, e.g., CDRs, of an antibody MGA271 (Macrogenics).

In one embodiment, an antigen binding domain against KIT is an antigen binding portion, e.g., CDRs, of an antibody described in, e.g., US7915391, US20120288506 , and several commercial catalog antibodies.

In one embodiment, an antigen binding domain against IL-13Ra2 is an antigen binding portion, e.g., CDRs, of an antibody described in, e.g., WO2008/146911, WO2004087758, several commercial catalog antibodies, and WO2004087758.

In one embodiment, an antigen binding domain against CD30 is an antigen binding portion, e.g., CDRs, of an antibody described in, e.g., US7090843 B1, and EP0805871.

In one embodiment, an antigen binding domain against GD3 is an antigen binding portion, e.g., CDRs, of an antibody described in, e.g., US7253263; US 8,207,308; US 20120276046; EP1013761; WO2005035577; and US6437098.

In one embodiment, an antigen binding domain against CD171 is an antigen binding portion, e.g., CDRs, of an antibody described in, e.g., Hong et al., J Immunother 37(2):93-104 (2014).

In one embodiment, an antigen binding domain against IL-11Ra is an antigen binding portion, e.g., CDRs, of an antibody available from Abcam (cat# ab55262) or Novus Biologicals (cat# EPR5446). In another embodiment, an antigen binding domain again IL-11Ra is a peptide, see, e.g., Huang et al., Cancer Res 72(1):271-281 (2012).

In one embodiment, an antigen binding domain against PSCA is an antigen binding portion, e.g., CDRs, of an antibody described in, e.g., Morgenroth et al., Prostate 67(10):1121-1131 (2007) (scFv 7F5); Nejatollahi et al., J of Oncology 2013(2013), article ID 839831 (scFv C5-II); and US Pat Publication No. 20090311181.

In one embodiment, an antigen binding domain against VEGFR2 is an antigen binding portion, e.g., CDRs, of an antibody described in, e.g., Chinnasamy et al., J Clin Invest 120(11):3953-3968 (2010).

In one embodiment, an antigen binding domain against LewisY is an antigen binding portion, e.g., CDRs, of an antibody described in, e.g., Kelly et al., Cancer Biother Radiopharm 23(4):411-423 (2008) (hu3S193 Ab (scFvs)); Dolezal et al., Protein Engineering 16(1):47-56 (2003) (NC10 scFv).

In one embodiment, an antigen binding domain against CD24 is an antigen binding portion, e.g., CDRs, of an antibody described in, e.g., Maliar et al., Gastroenterology 143(5):1375-1384 (2012).

In one embodiment, an antigen binding domain against PDGFR-beta is an antigen binding portion, e.g., CDRs, of an antibody Abcam ab32570.

In one embodiment, an antigen binding domain against SSEA-4 is an antigen binding portion, e.g., CDRs, of antibody MC813 (Cell Signaling), or other commercially available antibodies.

In one embodiment, an antigen binding domain against CD20 is an antigen binding portion, e.g., CDRs, of the antibody Rituximab, Ofatumumab, Ocrelizumab, Veltuzumab, or GA101.

In one embodiment, an antigen binding domain against Folate receptor alpha is an antigen binding portion, e.g., CDRs, of the antibody IMGN853, or an antibody described in US20120009181; US4851332, LK26: US5952484.

In one embodiment, an antigen binding domain against ERBB2 (Her2/neu) is an antigen binding portion, e.g., CDRs, of the antibody trastuzumab, or pertuzumab.

In one embodiment, an antigen binding domain against MUC1 is an antigen binding portion, e.g., CDRs, of the antibody SAR566658.

In one embodiment, the antigen binding domain against EGFR is antigen binding portion, e.g., CDRs, of the antibody cetuximab, panitumumab, zalutumumab, nimotuzumab, or matuzumab.

In one embodiment, an antigen binding domain against NCAM is an antigen binding portion, e.g., CDRs, of the antibody clone 2-2B: MAB5324 (EMD Millipore).

In one embodiment, an antigen binding domain against Ephrin B2 is an antigen binding portion, e.g., CDRs, of an antibody described in, e.g., Abengozar et al., Blood 119(19):4565-4576 (2012).

In one embodiment, an antigen binding domain against IGF-I receptor is an antigen binding portion, e.g., CDRs, of an antibody described in, e.g., US8344112 B2; EP2322550 A1; WO 2006/138315, or PCT/US2006/022995.

In one embodiment, an antigen binding domain against CAIX is an antigen binding portion, e.g., CDRs, of the antibody clone 303123 (R&D Systems).

In one embodiment, an antigen binding domain against LMP2 is an antigen binding portion, e.g., CDRs, of an antibody described in, e.g., US7,410,640, or US20050129701.

In one embodiment, an antigen binding domain against gp100 is an antigen binding portion, e.g., CDRs, of the antibody HMB45, NKIbetaB, or an antibody described in WO2013165940, or US20130295007

In one embodiment, an antigen binding domain against tyrosinase is an antigen binding portion, e.g., CDRs, of an antibody described in, e.g., US5843674; or US19950504048.

In one embodiment, an antigen binding domain against EphA2 is an antigen binding portion, e.g., CDRs, of an antibody described in, e.g., Yu et al., Mol Ther 22(1):102-111 (2014).

In one embodiment, an antigen binding domain against GD3 is an antigen binding portion, e.g., CDRs, of an antibody described in, e.g., US7253263; US 8,207,308; US 20120276046; EP1013761 A3; 20120276046; WO2005035577; or US6437098.

In one embodiment, an antigen binding domain against fucosyl GM1 is an antigen binding portion, e.g., CDRs, of an antibody described in, e.g., US20100297138; or WO2007/067992.

In one embodiment, an antigen binding domain against sLe is an antigen binding portion, e.g., CDRs, of the antibody G193 (for lewis Y), see Scott AM et al, Cancer Res 60: 3254-61 (2000), also as described in Neeson et al, J Immunol May 2013 190 (Meeting Abstract Supplement) 177.10.

In one embodiment, an antigen binding domain against GM3 is an antigen binding portion, e.g., CDRs, of the antibody CA 2523449 (mAb 14F7).

In one embodiment, an antigen binding domain against HMWMAA is an antigen binding portion, e.g., CDRs, of an antibody described in, e.g., Kmiecik et al., Oncoimmunology 3(1):e27185 (2014) (PMID: 24575382) (mAb9.2.27); US6528481; WO2010033866; or US 20140004124.

In one embodiment, an antigen binding domain against o-acetyl-GD2 is an antigen binding portion, e.g., CDRs, of the antibody 8B6.

In one embodiment, an antigen binding domain against TEM1/CD248 is an antigen binding portion, e.g., CDRs, of an antibody described in, e.g., Marty et al., Cancer Lett 235(2):298-308 (2006); Zhao et al., J Immunol Methods 363(2):221-232 (2011).

In one embodiment, an antigen binding domain against CLDN6 is an antigen binding portion, e.g., CDRs, of the antibody IMAB027 (Ganymed Pharmaceuticals), see e.g., clinicaltrial.gov/show/NCT02054351.

In one embodiment, an antigen binding domain against TSHR is an antigen binding portion, e.g., CDRs, of an antibody described in, e.g., US8,603,466; US8,501,415; or US8,309,693.

In one embodiment, an antigen binding domain against GPRC5D is an antigen binding portion, e.g., CDRs, of the antibody FAB6300A (R&D Systems); or LS-A4180 (Lifespan Biosciences).

In one embodiment, an antigen binding domain against CD97 is an antigen binding portion, e.g., CDRs, of an antibody described in, e.g., US6,846,911;de Groot et al., J Immunol 183(6):4127-4134 (2009); or an antibody from R&D:MAB3734.

In one embodiment, an antigen binding domain against ALK is an antigen binding portion, e.g., CDRs, of an antibody described in, e.g., Mino-Kenudson et al., Clin Cancer Res 16(5):1561-1571 (2010).

In one embodiment, an antigen binding domain against polysialic acid is an antigen binding portion, e.g., CDRs, of an antibody described in, e.g., Nagae et al., J Biol Chem 288(47):33784-33796 (2013).

In one embodiment, an antigen binding domain against PLAC1 is an antigen binding portion, e.g., CDRs, of an antibody described in, e.g., Ghods et al., Biotechnol Appl Biochem 2013 doi:10.1002/bab.1177.

In one embodiment, an antigen binding domain against GloboH is an antigen binding portion of the antibody VK9; or an antibody described in, e.g., Kudryashov V et al, Glycoconj J.15(3):243-9 (1998), Lou et al., Proc Natl Acad Sci USA 111(7):2482-2487 (2014) ; MBr1: Bremer E-G et al. J Biol Chem 259:14773–14777 (1984).

In one embodiment, an antigen binding domain against NY-BR-1 is an antigen binding portion, e.g., CDRs of an antibody described in, e.g., Jager et al., Appl Immunohistochem Mol Morphol 15(1):77-83 (2007).

In one embodiment, an antigen binding domain against WT-1 is an antigen binding portion, e.g., CDRs, of an antibody described in, e.g., Dao et al., Sci Transl Med 5(176):176ra33 (2013); or WO2012/135854.

In one embodiment, an antigen binding domain against MAGE-A1 is an antigen binding portion, e.g., CDRs, of an antibody described in, e.g., Willemsen et al., J Immunol 174(12):7853-7858 (2005) (TCR-like scFv).

In one embodiment, an antigen binding domain against sperm protein 17 is an antigen binding portion, e.g., CDRs, of an antibody described in, e.g., Song et al., Target Oncol 2013 Aug 14 (PMID: 23943313); Song et al., Med Oncol 29(4):2923-2931 (2012).

In one embodiment, an antigen binding domain against Tie 2 is an antigen binding portion, e.g., CDRs, of the antibody AB33 (Cell Signaling Technology).

In one embodiment, an antigen binding domain against MAD-CT-2 is an antigen binding portion, e.g., CDRs, of an antibody described in, e.g., PMID: 2450952; US7635753.

In one embodiment, an antigen binding domain against Fos-related antigen 1 is an antigen binding portion, e.g., CDRs, of the antibody 12F9 (Novus Biologicals).

In one embodiment, an antigen binding domain against MelanA/MART1 is an antigen binding portion, e.g., CDRs, of an antibody described in, EP2514766 A2; or US 7,749,719.

In one embodiment, an antigen binding domain against sarcoma translocation breakpoints is an antigen binding portion, e.g., CDRs, of an antibody described in, e.g., Luo et al, EMBO Mol. Med. 4(6):453-461 (2012).

In one embodiment, an antigen binding domain against TRP-2 is an antigen binding portion, e.g., CDRs, of an antibody described in, e.g., Wang et al, J Exp Med. 184(6):2207-16 (1996).

In one embodiment, an antigen binding domain against CYP1B1 is an antigen binding portion, e.g., CDRs, of an antibody described in, e.g., Maecker et al, Blood 102 (9): 3287-3294 (2003).

In one embodiment, an antigen binding domain against RAGE-1 is an antigen binding portion, e.g., CDRs, of the antibody MAB5328 (EMD Millipore).

In one embodiment, an antigen binding domain against human telomerase reverse transcriptase is an antigen binding portion, e.g., CDRs, of the antibody cat no: LS-B95-100 (Lifespan Biosciences)

In one embodiment, an antigen binding domain against intestinal carboxyl esterase is an antigen binding portion, e.g., CDRs, of the antibody 4F12: cat no: LS-B6190-50 (Lifespan Biosciences).

In one embodiment, an antigen binding domain against mut hsp70-2 is an antigen binding portion, e.g., CDRs, of the antibody Lifespan Biosciences: monoclonal: cat no: LS-C133261-100 (Lifespan Biosciences).

In one embodiment, an antigen binding domain against CD79a is an antigen binding portion, e.g., CDRs, of the antibody Anti-CD79a antibody [HM47/A9] (ab3121), available from Abcam; antibody CD79A Antibody #3351 available from Cell Signalling Technology; or antibody HPA017748 - Anti-CD79A antibody produced in rabbit, available from Sigma Aldrich.

In one embodiment, an antigen binding domain against CD79b is an antigen binding portion, e.g., CDRs, of the antibody polatuzumab vedotin, anti-CD79b described in Dornan et al., "Therapeutic potential of an anti-CD79b antibody-drug conjugate, anti-CD79b-vc-MMAE, for the treatment of non-Hodgkin lymphoma" Blood. 2009 Sep 24;114(13):2721-9. doi: 10.1182/blood-2009-02-205500. Epub 2009 Jul 24, or the bispecific antibody Anti-CD79b/CD3 described in "4507 Pre-Clinical Characterization of T Cell-Dependent Bispecific Antibody Anti-CD79b/CD3 As a Potential Therapy for B Cell Malignancies" Abstracts of 56th ASH Annual Meeting and Exposition, San Francisco, CA December 6-9 2014.

In one embodiment, an antigen binding domain against CD72 is an antigen binding portion, e.g., CDRs, of the antibody J3-109 described in Myers, and Uckun, "An anti-CD72 immunotoxin against therapy-refractory B-lineage acute lymphoblastic leukemia." Leuk Lymphoma. 1995 Jun;18(1-2):119-22, or anti-CD72 (10D6.8.1, mIgG1) described in Polson et al., "Antibody-Drug Conjugates for the Treatment of Non–Hodgkin's Lymphoma: Target and Linker-Drug Selection" Cancer Res March 15, 2009 69; 2358.

In one embodiment, an antigen binding domain against LAIR1 is an antigen binding portion, e.g., CDRs, of the antibody ANT-301 LAIR1 antibody, available from ProSpec; or anti-human CD305 (LAIR1) Antibody, available from BioLegend.

In one embodiment, an antigen binding domain against FCAR is an antigen binding portion, e.g., CDRs, of the antibody CD89/FCARAntibody (Catalog#10414-H08H), available from Sino Biological Inc.

In one embodiment, an antigen binding domain against LILRA2 is an antigen binding portion, e.g., CDRs, of the antibody LILRA2 monoclonal antibody (M17), clone 3C7, available from Abnova, or Mouse Anti-LILRA2 antibody, Monoclonal (2D7), available from Lifespan Biosciences..

In one embodiment, an antigen binding domain against CD300LF is an antigen binding portion, e.g., CDRs, of the antibody Mouse Anti-CMRF35-like molecule 1 antibody, Monoclonal[UP-D2], available from BioLegend, or Rat Anti-CMRF35-like molecule 1 antibody, Monoclonal[234903], available from R&D Systems..

In one embodiment, an antigen binding domain against CLEC12A is an antigen binding portion, e.g., CDRs, of the antibody Bispecific T cell Engager (BiTE) scFv-antibody and ADC described in Noordhuis

et al., “Targeting of CLEC12A In Acute Myeloid Leukemia by Antibody-Drug-Conjugates and Bispecific CLL-1xCD3 BiTE Antibody” 53rd ASH Annual Meeting and Exposition, December 10-13, 2011, and MCLA-117 (Merus).

In one embodiment, an antigen binding domain against BST2 (also called CD317) is an antigen binding portion, e.g., CDRs, of the antibody Mouse Anti-CD317 antibody, Monoclonal[3H4], available from Antibodies-Online or Mouse Anti-CD317 antibody, Monoclonal[696739], available from R&D Systems.

In one embodiment, an antigen binding domain against EMR2 (also called CD312) is an antigen binding portion, e.g., CDRs, of the antibody Mouse Anti-CD312 antibody, Monoclonal[LS-B8033] available from Lifespan Biosciences, or Mouse Anti-CD312 antibody, Monoclonal[494025] available from R&D Systems.

In one embodiment, an antigen binding domain against LY75 is an antigen binding portion, e.g., CDRs, of the antibody Mouse Anti-Lymphocyte antigen 75 antibody, Monoclonal[HD30] available from EMD Millipore or Mouse Anti-Lymphocyte antigen 75 antibody, Monoclonal[A15797] available from Life Technologies.

In one embodiment, an antigen binding domain against GPC3 is an antigen binding portion, e.g., CDRs, of the antibody hGC33 described in Nakano K, Ishiguro T, Konishi H, et al. Generation of a humanized anti-glypican 3 antibody by CDR grafting and stability optimization. Anticancer Drugs. 2010 Nov;21(10):907–916, or MDX-1414, HN3, or YP7, all three of which are described in Feng et al., “Glypican-3 antibodies: a new therapeutic target for liver cancer.” FEBS Lett. 2014 Jan 21;588(2):377-82.

In one embodiment, an antigen binding domain against FCRL5 is an antigen binding portion, e.g., CDRs, of the anti-FcRL5 antibody described in Elkins et al., “FcRL5 as a target of antibody-drug conjugates for the treatment of multiple myeloma” Mol Cancer Ther. 2012 Oct;11(10):2222-32. .

In one embodiment, an antigen binding domain against IGLL1 is an antigen binding portion, e.g., CDRs, of the antibody Mouse Anti-Immunoglobulin lambda-like polypeptide 1 antibody, Monoclonal[AT1G4] available from Lifespan Biosciences, Mouse Anti-Immunoglobulin lambda-like polypeptide 1 antibody, Monoclonal[HSL11] available from BioLegend.

In one embodiment, the antigen binding domain comprises one, two three (e.g., all three) heavy chain CDRs, HC CDR1, HC CDR2 and HC CDR3, from an antibody listed above, and/or one, two, three (e.g., all three) light chain CDRs, LC CDR1, LC CDR2 and LC CDR3, from an antibody listed above. In one

embodiment, the antigen binding domain comprises a heavy chain variable region and/or a variable light chain region of an antibody listed above.

In another aspect, the antigen binding domain comprises a humanized antibody or an antibody fragment. In some aspects, a non-human antibody is humanized, where specific sequences or regions of the antibody are modified to increase similarity to an antibody naturally produced in a human or fragment thereof. In one aspect, the antigen binding domain is humanized.

Bispecific CARs

In certain embodiments, the antigen binding domain is a bi- or multi- specific molecule (e.g., a multispecific antibody molecule). In an embodiment a multispecific antibody molecule is a bispecific antibody molecule. A bispecific antibody has specificity for no more than two antigens. A bispecific antibody molecule is characterized by a first immunoglobulin variable domain sequence which has binding specificity for a first epitope and a second immunoglobulin variable domain sequence that has binding specificity for a second epitope. In an embodiment the first and second epitopes are on the same antigen, e.g., the same protein (or subunit of a multimeric protein). In an embodiment the first and second epitopes overlap. In an embodiment the first and second epitopes do not overlap. In an embodiment the first and second epitopes are on different antigens, e.g., different proteins (or different subunits of a multimeric protein). In an embodiment a bispecific antibody molecule comprises a heavy chain variable domain sequence and a light chain variable domain sequence which have binding specificity for a first epitope and a heavy chain variable domain sequence and a light chain variable domain sequence which have binding specificity for a second epitope. In an embodiment a bispecific antibody molecule comprises a half antibody having binding specificity for a first epitope and a half antibody having binding specificity for a second epitope. In an embodiment a bispecific antibody molecule comprises a half antibody, or fragment thereof, having binding specificity for a first epitope and a half antibody, or fragment thereof, having binding specificity for a second epitope. In an embodiment a bispecific antibody molecule comprises a scFv, or fragment thereof, have binding specificity for a first epitope and a scFv, or fragment thereof, have binding specificity for a second epitope.

In certain embodiments, the antibody molecule is a multi-specific (*e.g.*, a bispecific or a trispecific) antibody molecule. Such molecules include bispecific fusion proteins, *e.g.*, an expression construct containing two scFvs with a hydrophilic helical peptide linker between them and a full constant region, as described in, *e.g.*, US5637481; minibody constructs with linked VL and VH chains

further connected with peptide spacers to an antibody hinge region and CH3 region, which can be dimerized to form bispecific/multivalent molecules, as described in, *e.g.*, US5837821; String of VH domains (or VL domains in family members) connected by peptide linkages with crosslinkable groups at the C-terminus futher associated with VL domains to form a series of FVs (or scFvs), as described in, *e.g.*, US5864019; and single chain binding polypeptides with both a VH and a VL domain linked through a peptide linker are combined into multivalent structures through non-covalent or chemical crosslinking to form, *e.g.*, homobivalent, heterobivalent, trivalent, and tetravalent structures using both scFV or diabody type format, as described in, *e.g.*, US5869620. The contents of the above-referenced applications are incorporated herein by reference in their entireties.

Within each antibody or antibody fragment (e.g., scFv) of a bispecific antibody molecule, the VH can be upstream or downstream of the VL. In some embodiments, the upstream antibody or antibody fragment (e.g., scFv) is arranged with its VH (VH_1) upstream of its VL (VL_1) and the downstream antibody or antibody fragment (e.g., scFv) is arranged with its VL (VL_2) upstream of its VH (VH_2), such that the overall bispecific antibody molecule has the arrangement VH_1-VL_1-VL_2-VH_2. In other embodiments, the upstream antibody or antibody fragment (e.g., scFv) is arranged with its VL (VL_1) upstream of its VH (VH_1) and the downstream antibody or antibody fragment (e.g., scFv) is arranged with its VH (VH_2) upstream of its VL (VL_2), such that the overall bispecific antibody molecule has the arrangement VL_1-VH_1-VH_2-VL_2. Optionally, a linker is disposed between the two antibodies or antibody fragments (e.g., scFvs), e.g., between VL_1 and VL_2 if the construct is arranged as VH_1-VL_1-VL_2-VH_2, or between VH_1 and VH_2 if the construct is arranged as VL_1-VH_1-VH_2-VL_2. The linker may be a linker as described herein, e.g., a (Gly_4-Ser)n linker, wherein n is 1, 2, 3, 4, 5, or 6, preferably 4 (SEQ ID NO: 890). In general, the linker between the two scFvs should be long enough to avoid mispairing between the domains of the two scFvs. Optionally, a linker is disposed between the VL and VH of the first scFv. Optionally, a linker is disposed between the VL and VH of the second scFv. In constructs that have multiple linkers, any two or more of the linkers can be the same or different. Accordingly, in some embodiments, a bispecific CAR comprises VLs, VHs, and optionally one or more linkers in an arrangement as described herein.

Transmembrane domains

With respect to the transmembrane domain, in various embodiments, a chimeric molecule of the invention (e.g., a CAR) can be designed to comprise a transmembrane domain that is attached to the extracellular domain of the chimeric molecule. A transmembrane domain can include one or more additional amino acids adjacent to the transmembrane region, e.g., one or more amino acid associated

with the extracellular region of the protein from which the transmembrane was derived (e.g., 1, 2, 3, 4, 5, 6, 7, 8, 9, 10 up to 15 amino acids of the extracellular region) and/or one or more additional amino acids associated with the intracellular region of the protein from which the transmembrane protein is derived (e.g., 1, 2, 3, 4, 5, 6, 7, 8, 9, 10 up to 15 amino acids of the intracellular region). In one aspect, the transmembrane domain is one that is associated with one of the other domains of the chimeric protein (e.g., CAR) e.g., in one embodiment, the transmembrane domain may be from the same protein that the signaling domain, costimulatory domain or the hinge domain is derived from. In another aspect, the transmembrane domain is not derived from the same protein that any other domain of the chimeric protein (e.g., CAR) is derived from. In some instances, the transmembrane domain can be selected or modified by amino acid substitution to avoid binding of such domains to the transmembrane domains of the same or different surface membrane proteins, e.g., to minimize interactions with other members of the receptor complex. In one aspect, the transmembrane domain is capable of homodimerization with another CAR on the cell surface of a CAR-expressing cell. In a different aspect, the amino acid sequence of the transmembrane domain may be modified or substituted so as to minimize interactions with the binding domains of the native binding partner present in the same CAR-expressing cell.

The transmembrane domain may be derived either from a natural or from a recombinant source. Where the source is natural, the domain may be derived from any membrane-bound or transmembrane protein. In one aspect the transmembrane domain is capable of signaling to the intracellular domain(s) whenever the CAR has bound to a target. A transmembrane domain of particular use in this invention may include at least the transmembrane region(s) of e.g., the alpha, beta or zeta chain of the T-cell receptor, CD28, CD27, CD3 epsilon, CD45, CD4, CD5, CD8, CD9, CD16, CD22, CD33, CD37, CD64, CD80, CD86, CD134, CD137, CD154. In some embodiments, a transmembrane domain may include at least the transmembrane region(s) of, e.g., KIRDS2, OX40, CD2, CD27, LFA-1 (CD11a, CD18), ICOS (CD278), 4-1BB (CD137), GITR, CD40, BAFFR, HVEM (LIGHTR), SLAMF7, NKp80 (KLRF1), NKp44, NKp30, NKp46, CD160, CD19, IL2R beta, IL2R gamma, IL7R α, ITGA1, VLA1, CD49a, ITGA4, IA4, CD49D, ITGA6, VLA-6, CD49f, ITGAD, CD11d, ITGAE, CD103, ITGAL, CD11a, LFA-1, ITGAM, CD11b, ITGAX, CD11c, ITGB1, CD29, ITGB2, CD18, LFA-1, ITGB7, TNFR2, DNAM1 (CD226), SLAMF4 (CD244, 2B4), CD84, CD96 (Tactile), CEACAM1, CRTAM, Ly9 (CD229), CD160 (BY55), PSGL1, CD100 (SEMA4D), SLAMF6 (NTB-A, Ly108), SLAM (SLAMF1, CD150, IPO-3), BLAME (SLAMF8), SELPLG (CD162), LTBR, PAG/Cbp, NKG2D, or NKG2C.

In some instances, the transmembrane domain can be attached to the extracellular region of the CAR, e.g., the antigen binding domain of the CAR, via a hinge, e.g., a hinge from a human protein. For example, in one embodiment, the hinge can be a human Ig (immunoglobulin) hinge (e.g., an IgG4 hinge, an IgD hinge), a GS linker (e.g., a GS linker described herein), a KIR2DS2 hinge or a CD8a hinge. In one embodiment, the hinge or spacer comprises (e.g., consists of) the amino acid sequence of SEQ ID NO: 4. In one aspect, the transmembrane domain comprises (e.g., consists of) a transmembrane domain of SEQ ID NO: 12.

In certain embodiments, the encoded transmembrane domain comprises an amino acid sequence of a CD8 transmembrane domain having at least one, two or three modifications but not more than 20, 10 or 5 modifications of an amino acid sequence of SEQ ID NO: 12, or a sequence with 95-99% identity to an amino acid sequence of SEQ ID NO: 12. In one embodiment, the encoded transmembrane domain comprises the sequence of SEQ ID NO: 12.

In other embodiments, the nucleic acid molecule encoding the CAR comprises a nucleotide sequence of a CD8 transmembrane domain, e.g., comprising the sequence of SEQ ID NO: 13, or a sequence with 95-99% identity thereof.

In certain embodiments, the encoded antigen binding domain is connected to the transmembrane domain by a hinge region. In one embodiment, the encoded hinge region comprises the amino acid sequence of a CD8 hinge, e.g., SEQ ID NO: 4; or the amino acid sequence of an IgG4 hinge, e.g., SEQ ID NO: 6, or a sequence with 95-99% identity to SEQ ID NO:4 or 6. In other embodiments, the nucleic acid sequence encoding the hinge region comprises a sequence of SEQ ID NO: 5 or SEQ ID NO: 7, corresponding to a CD8 hinge or an IgG4 hinge, respectively, or a sequence with 95-99% identity to SEQ ID NO:5 or 7.

In one aspect, the hinge or spacer comprises an IgG4 hinge. For example, in one embodiment, the hinge or spacer comprises a hinge of the amino acid sequence:

ESKYGPPCPPCPAPEFLGGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSQEDPEVQFNWYVDGVEVHNAKTKPREEQF NSTYRVVSVLTVLHQDWLNGKEYKCKVSNKGLPSSIEKTISKAKGQPREPQVYTLPPSQEEMTKNQVSLTCLVKGFYPSD IAVEWESNGQPENNYKTTPPVLDSDGSFFLYSRLTVDKSRWQEGNVFSCSVMHEALHNHYTQKSLSLSLGKM (SEQ ID NO:6).

In some embodiments, the hinge or spacer comprises a hinge encoded by a nucleotide sequence of:

GAGAGCAAGTACGGCCCTCCCTGCCCCCCTTGCCCTGCCCCCGAGTTCCTGGGCGGACCCAGCGTGTTCCTGTTCC

CCCCCAAGCCCAAGGACACCCTGATGATCAGCCGGACCCCCGAGGTGACCTGTGTGGTGGTGGACGTGTCCCAGG
AGGACCCCGAGGTCCAGTTCAACTGGTACGTGGACGGCGTGGAGGTGCACAACGCCAAGACCAAGCCCCGGGAG
GAGCAGTTCAATAGCACCTACCGGGTGGTGTCCGTGCTGACCGTGCTGCACCAGGACTGGCTGAACGGCAAGGA
ATACAAGTGTAAGGTGTCCAACAAGGGCCTGCCCAGCAGCATCGAGAAAACCATCAGCAAGGCCAAGGGCCAGC
CTCGGGAGCCCCAGGTGTACACCCTGCCCCCTAGCCAAGAGGAGATGACCAAGAACCAGGTGTCCCTGACCTGCC
TGGTGAAGGGCTTCTACCCCAGCGACATCGCCGTGGAGTGGGAGAGCAACGGCCAGCCCGAGAACAACTACAAG
ACCACCCCCCCTGTGCTGGACAGCGACGGCAGCTTCTTCCTGTACAGCCGGCTGACCGTGGACAAGAGCCGGTGG
CAGGAGGGCAACGTCTTTAGCTGCTCCGTGATGCACGAGGCCCTGCACAACCACTACACCCAGAAGAGCCTGAGC
CTGTCCCTGGGCAAGATG (SEQ ID NO:7).

In one aspect, the hinge or spacer comprises an IgD hinge. For example, in one embodiment, the hinge or spacer comprises a hinge of the amino acid sequence RWPESPKAQASSVPTAQPQAEGSLAKATTAPATTRNTGRGGEEKKKEKEKEEQEERETKTPECPSHTQPLGVYLLTPAV QDLWLRDKATFTCFVVGSDLKDAHLTWEVAGKVPTGGVEEGLLERHSNGSQSQHSRLTLPRSLWNAGTSVTCTLNHP SLPPQRLMALREPAAQAPVKLSLNLLASSDPPEAASWLLCEVSGFSPPNILLMWLEDQREVNTSGFAPARPPPQPGSTT FWAWSVLRVPAPPSPQPATYTCVVSHEDSRTLLNASRSLEVSYVTDH (SEQ ID NO:8).

In some embodiments, the hinge or spacer comprises a hinge encoded by a nucleotide sequence of:

AGGTGGCCCGAAAGTCCCAAGGCCCAGGCATCTAGTGTTCCTACTGCACAGCCCCAGGCAGAAGGCAGCCTAGCC
AAAGCTACTACTGCACCTGCCACTACGCGCAATACTGGCCGTGGCGGGGAGGAGAAGAAAAAGGAGAAAGAGA
AAGAAGAACAGGAAGAGAGGGAGACCAAGACCCCTGAATGTCCATCCCATACCCAGCCGCTGGGCGTCTATCTCT
TGACTCCCGCAGTACAGGACTTGTGGCTTAGAGATAAGGCCACCTTTACATGTTTCGTCGTGGGCTCTGACCTGAA
GGATGCCCATTTGACTTGGGAGGTTGCCGGAAAGGTACCCACAGGGGGGGTTGAGGAAGGGTTGCTGGAGCGC
CATTCCAATGGCTCTCAGAGCCAGCACTCAAGACTCACCCTTCCGAGATCCCTGTGGAACGCCGGGACCTCTGTCA
CATGTACTCTAAATCATCCTAGCCTGCCCCCACAGCGTCTGATGGCCCTTAGAGAGCCAGCCGCCCAGGCACCAGT
TAAGCTTAGCCTGAATCTGCTCGCCAGTAGTGATCCCCCAGAGGCCGCCAGCTGGCTCTTATGCGAAGTGTCCGGC
TTTAGCCCGCCCAACATCTTGCTCATGTGGCTGGAGGACCAGCGAGAAGTGAACACCAGCGGCTTCGCTCCAGCC
CGGCCCCCACCCCAGCCGGGTTCTACCACATTCTGGGCCTGGAGTGTCTTAAGGGTCCCAGCACCACCTAGCCCCC
AGCCAGCCACATACACCTGTGTTGTGTCCCATGAAGATAGCAGGACCCTGCTAAATGCTTCTAGGAGTCTGGAGGT
TTCCTACGTGACTGACCATT (SEQ ID NO:9).

In one aspect, the transmembrane domain may be recombinant, in which case it will comprise predominantly hydrophobic residues such as leucine and valine. In one aspect a triplet of phenylalanine, tryptophan and valine can be found at each end of a recombinant transmembrane domain.

Optionally, a short oligo- or polypeptide linker, between 2 and 10 amino acids in length may form the linkage between the transmembrane domain and the cytoplasmic region of the CAR. A glycine-serine doublet provides a particularly suitable linker. For example, in one aspect, the linker comprises the amino acid sequence of GGGGSGGGGS (SEQ ID NO:10). In some embodiments, the linker is encoded by a nucleotide sequence of GGTGGCGGAGGTTCTGGAGGTGGAGGTTCC (SEQ ID NO:11).

In one aspect, the hinge or spacer comprises a KIR2DS2 hinge.

Signaling domains

In embodiments of the invention having an intracellular signaling domain, such a domain can contain, e.g., one or more of a primary signaling domain and/or a costimulatory signaling domain. In some embodiments, the intracellular signaling domain comprises a sequence encoding a primary signaling domain. In some embodiments, the intracellular signaling domain comprises a costimulatory signaling domain. In some embodiments, the intracellular signaling domain comprises a a primary signaling domain and a costimulatory signaling domain.

The intracellular signaling sequences within the cytoplasmic portion of the CAR of the invention may be linked to each other in a random or specified order. Optionally, a short oligo- or polypeptide linker, for example, between 2 and 10 amino acids (e.g., 2, 3, 4, 5, 6, 7, 8, 9, or 10 amino acids) in length may form the linkage between intracellular signaling sequences. In one embodiment, a glycine-serine doublet can be used as a suitable linker. In one embodiment, a single amino acid, e.g., an alanine, a glycine, can be used as a suitable linker.

In one aspect, the intracellular signaling domain is designed to comprise two or more, e.g., 2, 3, 4, 5, or more, costimulatory signaling domains. In an embodiment, the two or more, e.g., 2, 3, 4, 5, or more, costimulatory signaling domains, are separated by a linker molecule, e.g., a linker molecule described herein. In one embodiment, the intracellular signaling domain comprises two costimulatory signaling domains. In some embodiments, the linker molecule is a glycine residue. In some embodiments, the linker is an alanine residue.

Primary Signaling domains

A primary signaling domain regulates primary activation of the TCR complex either in a stimulatory way, or in an inhibitory way. Primary intracellular signaling domains that act in a stimulatory manner may contain signaling motifs which are known as immunoreceptor tyrosine-based activation motifs or ITAMs.

Examples of ITAM containing primary intracellular signaling domains that are of particular use in the invention include those of CD3 zeta, common FcR gamma (FCER1G), Fc gamma RIIa, FcR beta (Fc Epsilon R1b), CD3 gamma, CD3 delta, CD3 epsilon, CD79a, CD79b, DAP10, and DAP12. In one embodiment, a CAR of the invention comprises an intracellular signaling domain, e.g., a primary signaling domain of CD3-zeta.

In one embodiment, the encoded primary signaling domain comprises a functional signaling domain of CD3 zeta. The encoded CD3 zeta primary signaling domain can comprise an amino acid sequence having at least one, two or three modifications but not more than 20, 10 or 5 modifications of an amino acid sequence of SEQ ID NO: 18 or SEQ ID NO: 20, or a sequence with 95-99% identity to an amino acid sequence of SEQ ID NO:18 or SEQ ID NO: 20. In some embodiments, the encoded primary signaling domain comprises a sequence of SEQ ID NO:18 or SEQ ID NO: 20. In other embodiments, the nucleic acid sequence encoding the primary signaling domain comprises a sequence of SEQ ID NO:19 or SEQ ID NO: 21, or a sequence with 95-99% identity thereof.

Costimulatory Signaling Domains

In some embodiments, the encoded intracellular signaling domain comprises a a costimulatory signaling domain. For example, the intracellular signaling domain can comprise a primary signaling domain and a costimulatory signaling domain. In some embodiments, the encoded costimulatory signaling domain comprises a functional signaling domain of a protein chosen from one or more of CD27, CD28, 4-1BB (CD137), OX40, CD30, CD40, PD-1, ICOS, lymphocyte function-associated antigen-1 (LFA-1), CD2, CD7, LIGHT, NKG2C, B7-H3, a ligand that specifically binds with CD83, CDS, ICAM-1, GITR, BAFFR, HVEM (LIGHTR), SLAMF7, NKp80 (KLRF1), CD160, CD19, CD4, CD8alpha, CD8beta, IL2R beta, IL2R gamma, IL7R alpha, ITGA4, VLA1, CD49a, ITGA4, IA4, CD49D, ITGA6, VLA-6, CD49f, ITGAD, CD11d, ITGAE, CD103, ITGAL, CD11a, LFA-1, ITGAM, CD11b, ITGAX, CD11c, ITGB1, CD29, ITGB2, CD18, LFA-1, ITGB7, TNFR2, TRANCE/RANKL, DNAM1 (CD226), SLAMF4 (CD244, 2B4), CD84, CD96 (Tactile), CEACAM1, CRTAM, Ly9 (CD229), CD160 (BY55), PSGL1, CD100 (SEMA4D), CD69, SLAMF6 (NTB-A, Ly108), SLAM

(SLAMF1, CD150, IPO-3), BLAME (SLAMF8), SELPLG (CD162), LTBR, LAT, GADS, SLP-76, PAG/Cbp, NKp44, NKp30, NKp46, and NKG2D.

In certain embodiments, the encoded costimulatory signaling domain comprises an amino acid sequence having at least one, two or three modifications but not more than 20, 10 or 5 modifications of an amino acid sequence of SEQ ID NO:14 or SEQ ID NO: 16, or a sequence with 95-99% identity to an amino acid sequence of SEQ ID NO:14 or SEQ ID NO: 16. In one embodiment, the encoded costimulatory signaling domain comprises a sequence of SEQ ID NO: 14 or SEQ ID NO: 16. In other embodiments, the nucleic acid sequence encoding the costimulatory signaling domain comprises a sequence of SEQ ID NO:15 or SEQ ID NO: 17, or a sequence with 95-99% identity thereof.

In other embodiments, the encoded intracellular domain comprises the sequence of SEQ ID NO: 14 or SEQ ID NO: 16, and the sequence of SEQ ID NO: 18 or SEQ ID NO: 20, wherein the sequences comprising the intracellular signaling domain are expressed in the same frame and as a single polypeptide chain.

In certain embodiments, the nucleic acid sequence encoding the intracellular signaling domain comprises a sequence of SEQ ID NO:15 or SEQ ID NO: 17, or a sequence with 95-99% identity thereof, and a sequence of SEQ ID NO:19 or SEQ ID NO:21, or a sequence with 95-99% identity thereof.

In some embodiments, the nucleic acid molecule further encodes a leader sequence. In one embodiment, the leader sequence comprises the sequence of SEQ ID NO: 2.

In one aspect, the intracellular signaling domain is designed to comprise the signaling domain of CD3-zeta and the signaling domain of CD28. In one aspect, the intracellular signaling domain is designed to comprise the signaling domain of CD3-zeta and the signaling domain of 4-1BB. In one aspect, the signaling domain of 4-1BB is a signaling domain of SEQ ID NO: 14. In one aspect, the signaling domain of CD3-zeta is a signaling domain of SEQ ID NO: 18.

In one aspect, the intracellular signaling domain is designed to comprise the signaling domain of CD3-zeta and the signaling domain of CD27. In one aspect, the signaling domain of CD27 comprises an amino acid sequence of QRRKYRSNKGESPVEPAEPCRYSCPREEEGSTIPIQEDYRKPEPACSP (SEQ ID NO:16). In one aspect, the signaling domain of CD27 is encoded by a nucleic acid sequence of: AGGAGTAAGAGGAGCAGGCTCCTGCACAGTGACTACATGAACATGACTCCCCGCCGCCCCGGGCCCACCCGCAAG CATTACCAGCCCTATGCCCCACCACGCGACTTCGCAGCCTATCGCTCC (SEQ ID NO:17).

Exemplary CAR Molecules

The CAR molecules disclosed herein can comprise a binding domain that binds to a target, *e.g.*, a target as described herein; a transmembrane domain, e.g., a transmembrane domain as described herein; and an intracellular signaling domain, e.g., an intracellular domain as described herein. In embodiments, the binding domain comprises a heavy chain complementary determining region 1 (HC CDR1), a heavy chain complementary determining region 2 (HC CDR2), and a heavy chain complementary determining region 3 (HC CDR3) of a heavy chain binding domain described herein, and/or a light chain complementary determining region 1 (LC CDR1), a light chain complementary determining region 2 (LC CDR2), and a light chain complementary determining region 3 (LC CDR3) of a light chain binding domain described herein.

In other embodiments, the CAR molecule comprises a CD19 CAR molecule described herein, e.g., a CD19 CAR molecule described in US-2015-0283178-A1, e.g., CTL019. In embodiments, the CD19 CAR comprises an amino acid, or has a nucleotide sequence shown in US-2015-0283178-A1, incorporated herein by reference, or a sequence substantially identical thereto (e.g., at least 85%, 90%, 95% or more identical thereto).

In one embodiment, the CAR T cell that specifically binds to CD19 has the USAN designation TISAGENLECLEUCEL-T. CTL019 is made by a gene modification of T cells is mediated by stable insertion via transduction with a self-inactivating, replication deficient Lentiviral (LV) vector containing the CTL019 transgene under the control of the EF-1 alpha promoter. CTL019 can be a mixture of transgene positive and negative T cells that are delivered to the subject on the basis of percent transgene positive T cells.

In other embodiments, the CD19 CAR includes a CAR molecule, or an antigen binding domain (e.g., a humanized antigen binding domain) according to Table 3 of WO2014/153270, incorporated herein by reference. The amino acid and nucleotide sequences encoding the CD19 CAR molecules and antigen binding domains (e.g., including one, two, three VH CDRs; and one, two, three VL CDRs according to Kabat or Chothia), are specified in WO2014/153270. In embodiments, the CD19 CAR comprises an amino acid, or has a nucleotide sequence shown in WO2014/153270 incorporated herein by reference, or a sequence substantially identical to any of the aforesaid sequences (e.g., at least 85%, 90%, 95% or more identical to any of the aforesaid CD19 CAR sequences).

In one embodiment, the parental murine scFv sequence is the CAR19 construct provided in PCT publication WO2012/079000 (incorporated herein by reference) and provided herein in Table 9. In one embodiment, the anti-CD19 binding domain is a scFv described in WO2012/079000 and provided herein in Table 9.

In one embodiment, the CD19 CAR comprises an amino acid sequence provided as SEQ ID NO: 12 in PCT publication WO2012/079000. In embodiment, the amino acid sequence is:

MALPVTALLLPLALLLHAARPdiqmtqttsslsaslgdrvtiscrasqdiskylnwyqqkpdgtvklliyhtsrlhsgvpsrfsgsgsg tdysltisnleqediatyfcqqgntlpytfgggtkleitggggsggggsggggsevklqesgpglvapsqslsvtctvsgvslpdygvswirqpprkgle wlgviwgsettyynsalksrltiikdnsksqvflkmnslqtddtaiyycakhyyyggsyamdywgqgtsvtvsstttpaprpptpaptiasqplslrp eacrpaaggavhtrgldfacdiyiwaplagtcgvlllslvitlyckrgrkkllyifkqpfmrpvqttqeedgcscrfpeeeeggcelrvkfsrsadapayk qgqnqlynelnlgrreeydvldkrrgrdpemggkprrknpqeglynelqkdkmaeayseigmkgerrrgkghdglyqglstatkdtydalhmq alppr (SEQ ID NO: 891), or a sequence substantially identical thereto (e.g., at least 85%, 90% or 95% or higher identical thereto), with or without the signal peptide sequence indicated in capital letters.

In embodiment, the amino acid sequence is:

diqmtqttsslsaslgdrvtiscrasqdiskylnwyqqkpdgtvklliyhtsrlhsgvpsrfsgsgsgtdysltisnleqediatyfcqqgntl pytfgggtkleitggggsggggsggggsevklqesgpglvapsqslsvtctvsgvslpdygvswirqpprkglewlgviwgsettyynsalksrltiikd nsksqvflkmnslqtddtaiyycakhyyyggsyamdywgqgtsvtvsstttpaprpptpaptiasqplslrpeacrpaaggavhtrgldfacdiyi waplagtcgvlllslvitlyckrgrkkllyifkqpfmrpvqttqeedgcscrfpeeeeggcelrvkfsrsadapaykqgqnqlynelnlgrreeydvldk rrgrdpemggkprrknpqeglynelqkdkmaeayseigmkgerrrgkghdglyqglstatkdtydalhmqalppr (SEQ ID NO: 892), or a sequence substantially homologous thereto (e.g., at least 85%, 90% or 95% or higher identical thereto).

In embodiments, the CAR molecule is a CD19 CAR molecule described herein, e.g., a humanized CAR molecule described herein, e.g., a humanized CD19 CAR molecule of Table 6-9 or having CDRs as set out in Tables 10A and 10B.

In embodiments, the CAR molecule is a CD19 CAR molecule described herein, e.g., a murine CAR molecule described herein, e.g., a murine CD19 CAR molecule of Table 9 or having CDRs as set out in Tables 10A and 10B.

In some embodiments, the CAR molecule comprises one, two, and/or three CDRs from the heavy chain variable region and/or one, two, and/or three CDRs from the light chain variable region of the murine or humanized CD19 CAR of Table 10A and 10B.

In one embodiment, the antigen binding domain comprises one, two three (e.g., all three) heavy chain CDRs, HC CDR1, HC CDR2 and HC CDR3, from an antibody listed herein, and/or one, two, three

(e.g., all three) light chain CDRs, LC CDR1, LC CDR2 and LC CDR3, from an antibody listed herein. In one embodiment, the antigen binding domain comprises a heavy chain variable region and/or a variable light chain region of an antibody listed or described herein.

Exemplary CD19 CARs include any of the CD19 CARs or anti-CD19 binding domains described herein, e.g., in one or more tables (e.g., Tables 6-9) described herein (e.g., , or an anti-CD19 CAR described in Xu et al. Blood 123.24(2014):3750-9; Kochenderfer et al. Blood 122.25(2013):4129-39, Cruz et al. Blood 122.17(2013):2965-73, NCT00586391, NCT01087294, NCT02456350, NCT00840853, NCT02659943, NCT02650999, NCT02640209, NCT01747486, NCT02546739, NCT02656147, NCT02772198, NCT00709033, NCT02081937, NCT00924326, NCT02735083, NCT02794246, NCT02746952, NCT01593696, NCT02134262, NCT01853631, NCT02443831, NCT02277522, NCT02348216, NCT02614066, NCT02030834, NCT02624258, NCT02625480, NCT02030847, NCT02644655, NCT02349698, NCT02813837, NCT02050347, NCT01683279, NCT02529813, NCT02537977, NCT02799550, NCT02672501, NCT02819583, NCT02028455, NCT01840566, NCT01318317, NCT01864889, NCT02706405, NCT01475058, NCT01430390, NCT02146924, NCT02051257, NCT02431988, NCT01815749, NCT02153580, NCT01865617, NCT02208362, NCT02685670, NCT02535364, NCT02631044, NCT02728882, NCT02735291, NCT01860937, NCT02822326, NCT02737085, NCT02465983, NCT02132624, NCT02782351, NCT01493453, NCT02652910, NCT02247609, NCT01029366, NCT01626495, NCT02721407, NCT01044069, NCT00422383, NCT01680991, NCT02794961, or NCT02456207, each of which is incorporated herein by reference in its entirety.

Exemplary CD19 CAR and antigen binding domain constructs that can be used in the methods described herein are shown in Tables 6-9. The light and heavy chain CDR sequences according to Kabat are shown by the bold and underlined text, and are also summarized in Tables 9 and 10A-10B below. The location of the signal sequence and histidine tag are also underlined. In embodiments, the CD19 CAR sequences and antigen binding fragments thereof do not include the signal sequence and/or histidine tag sequences.

In embodiments, the CD19 CAR comprises an anti- CD19 binding domain (e.g., murine or humanized anti- CD19 binding domain), a transmembrane domain, and an intracellular signaling domain, and wherein said anti- CD19 binding domain comprises a heavy chain complementary determining region 1 (HC CDR1), a heavy chain complementary determining region 2 (HC CDR2), and a heavy chain complementary determining region 3 (HC CDR3) of any anti- CD19 heavy chain binding domain amino acid sequences listed in Tables 6-9 and 10A-10B, or a sequence at least 85%, 90%, 95% or

more identical thereto (e.g., having less than 5, 4, 3, 2 or 1 amino acid substitutions, e.g., conservative substitutions).

In one embodiment, the anti- CD19 binding domain comprises a light chain variable region described herein (e.g., in Tables 6-9) and/or a heavy chain variable region described herein (e.g., in Table 9), or a sequence at least 85%, 90%, 95% or more identical thereto.

In one embodiment, the encoded anti- CD19 binding domain is a scFv comprising a light chain and a heavy chain of an amino acid sequence of Tables 6-9, or a sequence at least 85%, 90%, 95% or more identical thereto.

In an embodiment, the human or humanized anti- CD19 binding domain (e.g., an scFv) comprises: a light chain variable region comprising an amino acid sequence having at least one, two or three modifications (e.g., substitutions, e.g., conservative substitutions) but not more than 30, 20 or 10 modifications (e.g., substitutions, e.g., conservative substitutions) of an amino acid sequence of a light chain variable region provided in Tables 6-9, or a sequence at least 85%, 90%, 95% or more identical thereto; and/or a heavy chain variable region comprising an amino acid sequence having at least one, two or three modifications (e.g., substitutions, e.g., conservative substitutions) but not more than 30, 20 or 10 modifications (e.g., substitutions, e.g., conservative substitutions) of an amino acid sequence of a heavy chain variable region provided in Tables 6-9, or a sequence at least 85%, 90%, 95% or more identical thereto.

Table 9: CD19 CAR Constructs

Name	SEQ ID NO:	Sequence
CAR 1		
CAR1 scFv domain	893	EIVMTQSPATLSLSPGERATLSCRASQDISKYLNWYQQKPGQAPRLLIYHT SRLHSGIPARFSGSGSGTDYTLTISSLQPEDFAVYFCQQGNTLPYTFGQGT KLEIKGGGGSGGGGSGGGGSQVQLQESGPGLVKPSETLSLTCTVSGVSLPD YGVSWIRQPPGKGLEWIGVIWGSETTYYSSSLKSRVTISKDNSKNQVSLKL SSVTAADTAVYYCAKHYYYGGSYAMDYWGQGTLVTVSS
103101 CAR1 Soluble scFv - nt	894	atggccctccctgtcaccgccctgctgcttccgctggctcttctgctccacgccgc tcggcccgaaattgtgatgacccagtcacccgccactcttagcctttcacccggtg agcgcgcaaccctgtcttgcagagcctcccaagacatctcaaaataccttaattgg tatcaacagaagcccggacaggctcctcgccttctgatctaccacaccagccggct ccattctggaatccctgccaggttcagcggtagcggatctgggaccgactacaccc tcactatcagctcactgcagccagaggacttcgctgtctatttctgtcagcaaggg aacaccctgccctacacctttggacagggcaccaagctcgagattaaaggtggagg

		tggcagcggaggaggtgggtccggcggtggaggaagccaggtccaactccaagaaa gcggaccgggtcttgtgaagccatcagaaactctttcactgacttgtactgtgagc ggagtgtctctccccgattacggggtgtcttggatcagacagccaccggggaaggg tctggaatggattggagtgatttggggctctgagactacttactactcttcatccc tcaagtcacgcgtcaccatctcaaaggacaactctaagaatcaggtgtcactgaaa ctgtcatctgtgaccgcagccgacaccgccgtgtactattgcgctaagcattacta ttatggcgggagctacgcaatggattactggggacagggtactctggtcaccgtgt ccagccaccaccatcatcaccatcaccat
103101 **CAR1** **Soluble scFv - aa**	895	**MALPVTALLLPLALLLHAARP**eivmtqspatlslspgeratlscrasqdiskylnw yqqkpgqaprlliyhtsrlhsgiparfsgsgsgtdytltisslqpedfavyfcqqg ntlpytfgqgtkleikggggsggggsggggsqvqlqesgpglvkpsetlsltctvs gvslpdygvswirqppgkglewigviwgsettyyssslksrvtiskdnsknqvslk lssvtaadtavyycakhyyyggsyamdywgqgtlvtvss**hhhhhhh**
104875 **CAR 1 – Full - nt**	896	atggccctccctgtcaccgccctgctgcttccgctggctcttctgctccacgccgc tcggcccgaaattgtgatgacccagtcacccgccactcttagcctttcacccggtg agcgcgcaaccctgtcttgcagagcctcccaagacatctcaaaataccttaattgg tatcaacagaagcccggacaggctcctcgccttctgatctaccacaccagccggct ccattctggaatccctgccaggttcagcggtagcggatctgggaccgactacaccc tcactatcagctcactgcagccagaggacttcgctgtctatttctgtcagcaaggg aacaccctgccctacacctttggacagggcaccaagctcgagattaaaggtggagg tggcagcggaggaggtgggtccggcggtggaggaagccaggtccaactccaagaaa gcggaccgggtcttgtgaagccatcagaaactctttcactgacttgtactgtgagc ggagtgtctctccccgattacggggtgtcttggatcagacagccaccggggaaggg tctggaatggattggagtgatttggggctctgagactacttactactcttcatccc tcaagtcacgcgtcaccatctcaaaggacaactctaagaatcaggtgtcactgaaa ctgtcatctgtgaccgcagccgacaccgccgtgtactattgcgctaagcattacta ttatggcgggagctacgcaatggattactggggacagggtactctggtcaccgtgt ccagcaccactaccccagcaccgaggccacccaccccggctcctaccatcgcctcc cagcctctgtccctgcgtccggaggcatgtagacccgcagctggtggggccgtgca tacccggggtcttgacttcgcctgcgatatctacatttgggcccctctggctggta cttgcggggtcctgctgctttcactcgtgatcactctttactgtaagcgcggtcgg aagaagctgctgtacatctttaagcaacccttcatgaggcctgtgcagactactca agaggaggacggctgttcatgccggttcccagaggaggaggaaggcggctgcgaac tgcgcgtgaaattcagccgcagcgcagatgctccagcctacaagcaggggcagaac cagctctacaacgaactcaatcttggtcggagagaggagtacgacgtgctggacaa gcggagaggacgggacccagaaatgggcgggaagccgcgcagaaagaatccccaag agggcctgtacaacgagctccaaaaggataagatggcagaagcctatagcgagatt ggtatgaaaggggaacgcagaagaggcaaaggccacgacggactgtaccagggact cagcaccgccaccaaggacacctatgacgctcttcacatgcaggccctgccgcctc gg
104875 **CAR 1 – Full - aa**	897	MALPVTALLLPLALLLHAARPeivmtqspatlslspgeratlsc**rasqdiskyln**w yqqkpgqaprlliy**htsrlhs**giparfsgsgsgtdytltisslqpedfavyfc**qqg ntlpyt**fgqgtkleikggggsggggsggggsqvqlqesgpglvkpsetlsltctvs gvslp**dygvs**wirqppgkglewig**viwgsettyyssslks**rvtiskdnsknqvslk lssvtaadtavyycak**hyyyggsyamdy**wgqgtlvtvssttpaprpptpaptias

		qplslrpeacrpaaggavhtrgldfacdiyiwaplagtcgvlllslvitlyckrgr kkllyifkqpfmrpvqttqeedgcscrfpeeeeggcelrvkfsrsadapaykqgqn qlynelnlgrreeydvldkrrgrdpemggkprrknpqeglynelqkdkmaeaysei gmkgerrrgkghdglyqglstatkdtydalhmqalppr
CAR 2		
CAR2 scFv domain	898	eivmtqspatlslspgeratlscrasqdiskylnwyqqkpgqaprlliyhtsrlhs giparfsgsgsgtdytltisslqpedfavyfcqqgntlpytfgqgtkleikggggs ggggsggggsqvqlqesgpglvkpsetlsltctvsgvslpdygvswirqppgkgle wigviwgsettyyqsslksrvtiskdnsknqvslklssvtaadtavyycakhyyyg gsyamdywgqgtlvtvss
103102 **CAR2 - Soluble scFv - nt**	899	atggccctccctgtcaccgccctgctgcttccgctggctcttctgctccacgccgc tcggcccgaaattgtgatgacccagtcacccgccactcttagcctttcacccggtg agcgcgcaaccctgtcttgcagagcctcccaagacatctcaaaataccttaattgg tatcaacagaagcccggacaggctcctcgccttctgatctaccacaccagccggct ccattctggaatccctgccaggttcagcggtagcggatctgggaccgactacaccc tcactatcagctcactgcagccagaggacttcgctgtctatttctgtcagcaaggg aacaccctgccctacacctttggacagggcaccaagctcgagattaaaggtggagg tggcagcggaggaggtgggtccggcggtggaggaagccaggtccaactccaagaaa gcggaccgggtcttgtgaagccatcagaaactctttcactgacttgtactgtgagc ggagtgtctctccccgattacggggtgtcttggatcagacagccaccggggaaggg tctggaatggattggagtgatttggggctctgagactacttactaccaatcatccc tcaagtcacgcgtcaccatctcaaaggacaactctaagaatcaggtgtcactgaaa ctgtcatctgtgaccgcagccgacaccgccgtgtactattgcgctaagcattacta ttatggcgggagctacgcaatggattactggggacagggtactctggtcaccgtgt ccagccaccaccatcatcaccatcaccat
103102 **CAR2 - Soluble scFv - aa**	900	**MALPVTALLLPLALLLHAARP**eivmtqspatlslspgeratlscrasqdiskylnw yqqkpgqaprlliyhtsrlhsgiparfsgsgsgtdytltisslqpedfavyfcqqg ntlpytfgqgtkleikggggsggggsggggsqvqlqesgpglvkpsetlsltctvs gvslpdygvswirqppgkglewigviwgsettyyqsslksrvtiskdnsknqvslk lssvtaadtavyycakhyyyggsyamdywgqgtlvtvss**hhhhhhhh**
104876 **CAR 2 - Full - nt**	901	atggccctccctgtcaccgccctgctgcttccgctggctcttctgctccacgccgc tcggcccgaaattgtgatgacccagtcacccgccactcttagcctttcacccggtg agcgcgcaaccctgtcttgcagagcctcccaagacatctcaaaataccttaattgg tatcaacagaagcccggacaggctcctcgccttctgatctaccacaccagccggct ccattctggaatccctgccaggttcagcggtagcggatctgggaccgactacaccc tcactatcagctcactgcagccagaggacttcgctgtctatttctgtcagcaaggg aacaccctgccctacacctttggacagggcaccaagctcgagattaaaggtggagg tggcagcggaggaggtgggtccggcggtggaggaagccaggtccaactccaagaaa gcggaccgggtcttgtgaagccatcagaaactctttcactgacttgtactgtgagc ggagtgtctctccccgattacggggtgtcttggatcagacagccaccggggaaggg tctggaatggattggagtgatttggggctctgagactacttactaccaatcatccc tcaagtcacgcgtcaccatctcaaaggacaactctaagaatcaggtgtcactgaaa ctgtcatctgtgaccgcagccgacaccgccgtgtactattgcgctaagcattacta ttatggcgggagctacgcaatggattactggggacagggtactctggtcaccgtgt

		ccagcaccactaccccagcaccgaggccacccaccccggctcctaccatcgcctcc cagcctctgtccctgcgtccggaggcatgtagacccgcagctggtggggccgtgca tacccggggtcttgacttcgcctgcgatatctacatttgggcccctctggctggta cttgcggggtcctgctgctttcactcgtgatcactctttactgtaagcgcggtcgg aagaagctgctgtacatctttaagcaacccttcatgaggcctgtgcagactactca agaggaggacggctgttcatgccggttcccagaggaggaggaaggcggctgcgaac tgcgcgtgaaattcagccgcagcgcagatgctccagcctacaagcaggggcagaac cagctctacaacgaactcaatcttggtcggagagaggagtacgacgtgctggacaa gcggagaggacgggacccagaaatgggcgggaagccgcgcagaaagaatccccaag agggcctgtacaacgagctccaaaaggataagatggcagaagcctatagcgagatt ggtatgaaaggggaacgcagaagaggcaaaggccacgacggactgtaccagggact cagcaccgccaccaaggacacctatgacgctcttcacatgcaggccctgccgcctc gg
104876 **CAR 2 - Full - aa**	902	MALPVTALLLPLALLLHAARPeivmtqspatlslspgeratlsc**rasqdiskyln**w yqqkpgqaprlliy**htsrlhs**giparfsgsgsgtdytltisslqpedfavyfc**qqg** **ntlpyt**fgqgtkleikggggsggggsggggsqvqlqesgpglvkpsetlsltctvs gvslp**dygvs**wirqppgkglewig**viwgsettyyqsslks**rvtiskdnsknqvslk lssvtaadtavyycak**hyyyggsyamdy**wgqgtlvtvsstttpaprpptpaptias qplslrpeacrpaaggavhtrgldfacdiyiwaplagtcgvlllslvitlyckrgr kkllyifkqpfmrpvqttqeedgcscrfpeeeeggcelrvkfsrsadapaykqgqn qlynelnlgrreeydvldkrrgrdpemggkprrknpqeglynelqkdkmaeaysei gmkgerrrgkghdglyqglstatkdtydalhmqalppr
CAR 3		
CAR3 scFv domain	903	qvqlqesgpglvkpsetlsltctvsgvslpdygvswirqppgkglewigviwgset tyyssslksrvtiskdnsknqvslklssvtaadtavyycakhyyyggsyamdywgq gtlvtvssggggsggggsggggseivmtqspatlslspgeratlscrasqdiskyl nwyqqkpgqaprlliyhtsrlhsgiparfsgsgsgtdytltisslqpedfavyfcq qgntlpytfgqgtkleik
103104 **CAR 3 - Soluble scFv - nt**	904	atggctctgcccgtgaccgcactcctcctgccactggctctgctgcttcacgccgc tcgcccacaagtccagcttcaagaatcagggcctggtctggtgaagccatctgaga ctctgtccctcacttgcaccgtgagcggagtgtccctcccagactacggagtgagc tggattagacagcctcccggaaagggactggagtggatcggagtgatttggggtag cgaaaccacttactattcatcttccctgaagtcacgggtcaccatttcaaaggata actcaaagaatcaagtgagcctcaagctctcatcagtcaccgccgctgacaccgcc gtgtattactgtgccaagcattactactatggagggtcctacgccatggactactg gggccagggaactctggtcactgtgtcatctggtggaggaggtagcggaggaggcg ggagcggtggaggtggctccgaaatcgtgatgacccagagccctgcaaccctgtcc ctttctcccggggaacgggctacccttttcttgtcgggcatcacaagatatctcaaa atacctcaattggtatcaacagaagccgggacaggcccctaggcttcttatctacc acacctctcgcctgcatagcgggattcccgcacgctttagcgggtctggaagcggg accgactacactctgaccatctcatctctccagcccgaggacttcgccgtctactt ctgccagcagggtaacaccctgccgtacaccttcggccagggcaccaagcttgaga tcaaacatcaccaccatcatcaccatcac
103104	905	**MALPVTALLLPLALLLHAARP**qvqlqesgpglvkpsetlsltctvsgvslpdygvs wirqppgkglewigviwgsettyyssslksrvtiskdnsknqvslklssvtaadta

CAR 3 - Soluble scFv - aa		vyycakhyyyggsyamdywgqgtlvtvssggggsggggsggggseivmtqspatls lspgeratlscrasqdiskylnwyqqkpgqaprlliyhtsrlhsgiparfsgsgsg tdytltisslqpedfavyfcqqgntlpytfgqgtkleik**hhhhhhhh**
104877 **CAR 3 – Full - nt**	906	atggctctgcccgtgaccgcactcctcctgccactggctctgctgcttcacgccgc tcgcccacaagtccagcttcaagaatcagggcctggtctggtgaagccatctgaga ctctgtccctcacttgcaccgtgagcggagtgtccctcccagactacggagtgagc tggattagacagcctcccggaaagggactggagtggatcggagtgatttggggtag cgaaaccacttactattcatcttccctgaagtcacgggtcaccatttcaaaggata actcaaagaatcaagtgagcctcaagctctcatcagtcaccgccgctgacaccgcc gtgtattactgtgccaagcattactactatggagggtcctacgccatggactactg gggccagggaactctggtcactgtgtcatctggtggaggaggtagcggaggaggcg ggagcggtggaggtggctccgaaatcgtgatgacccagagccctgcaaccctgtcc ctttctcccggggaacgggctacccttcttgtcgggcatcacaagatatctcaaa atacctcaattggtatcaacagaagccgggacaggcccctaggcttcttatctacc acacctctcgcctgcatagcgggattcccgcacgctttagcgggtctggaagcggg accgactacactctgaccatctcatctctccagcccgaggacttcgccgtctactt ctgccagcagggtaacaccctgccgtacaccttcggccagggcaccaagcttgaga tcaaaaccactactcccgctccaaggccacccacccctgccccgaccatcgcctct cagccgctttccctgcgtccggaggcatgtagacccgcagctggtggggccgtgca tacccggggtcttgacttcgcctgcgatatctacatttgggcccctctggctggta cttgcggggtcctgctgctttcactcgtgatcactctttactgtaagcgcggtcgg aagaagctgctgtacatctttaagcaacccttcatgaggcctgtgcagactactca agaggaggacggctgttcatgccggttcccagaggaggaggaaggcggctgcgaac tgcgcgtgaaattcagccgcagcgcagatgctccagcctacaagcaggggcagaac cagctctacaacgaactcaatcttggtcggagagaggagtacgacgtgctggacaa gcggagaggacgggacccagaaatgggcgggaagccgcgcagaaagaatccccaag agggcctgtacaacgagctccaaaaggataagatggcagaagcctatagcgagatt ggtatgaaaggggaacgcagaagaggcaaaggccacgacggactgtaccagggact cagcaccgccaccaaggacacctatgacgctcttcacatgcaggccctgccgcctc gg
104877 **CAR 3 – Full - aa**	907	MALPVTALLLPLALLLHAARPqvqlqesgpglvkpsetlsltctvsgvslp**dygvs** wirqppgkglewig**viwgsettyyssslks**rvtiskdnsknqvslklssvtaadta vyycak**hyyyggsyamdy**wgqgtlvtvssggggsggggsggggseivmtqspatls lspgeratlsc**rasqdiskyln**wyqqkpgqaprlliy**htsrlhs**giparfsgsgsg tdytltisslqpedfavyfc**qqgntlpyt**fgqgtkleikpttpaprppptpaptias qplslrpeacrpaaggavhtrgldfacdiyiwaplagtcgvlllslvitlyckrgr kkllyifkqpfmrpvqttqeedgcscrfpeeeeggcelrvkfsrsadapaykqgqn qlynelnlgrreeydvldkrrgrdpemggkprrknpqeglynelqkdkmaeaysei gmkgerrrgkghdglyqglstatkdtydalhmqalppr
CAR 4		
CAR4 scFv domain	908	qvqlqesgpglvkpsetlsltctvsgvslpdygvswirqppgkglewigviwgset tyyqsslksrvtiskdnsknqvslklssvtaadtavyycakhyyyggsyamdywgq gtlvtvssggggsggggsggggseivmtqspatlslspgeratlscrasqdiskyl nwyqqkpgqaprlliyhtsrlhsgiparfsgsgsgtdytltisslqpedfavyfcq

		qgntlpytfgqgtkleik
103106 CAR4 – Soluble scFv - nt	909	atggctctgcccgtgaccgcactcctcctgccactggctctgctgcttcacgccgc tcgcccacaagtccagcttcaagaatcagggcctggtctggtgaagccatctgaga ctctgtccctcacttgcaccgtgagcggagtgtccctcccagactacggagtgagc tggattagacagcctcccggaaagggactggagtggatcggagtgatttggggtag cgaaaccacttactatcaatcttccctgaagtcacgggtcaccatttcaaaggata actcaaagaatcaagtgagcctcaagctctcatcagtcaccgccgctgacaccgcc gtgtattactgtgccaagcattactactatggagggtcctacgccatggactactg gggccagggaactctggtcactgtgtcatctggtggaggaggtagcggaggaggcg ggagcggtggaggtggctccgaaatcgtgatgacccagagccctgcaaccctgtcc ctttctccccggggaacgggctacccttttcttgtcgggcatcacaagatatctcaaa atacctcaattggtatcaacagaagccgggacaggcccctaggcttcttatctacc acacctctcgcctgcatagcgggattcccgcacgctttagcgggtctggaagcggg accgactacactctgaccatctcatctctccagcccgaggacttcgccgtctactt ctgccagcagggtaacaccctgccgtacaccttcggccagggcaccaagcttgaga tcaaacatcaccaccatcatcaccatcac
103106 CAR4 – Soluble scFv -aa	910	**MALPVTALLLPLALLLHAARP**qvqlqesgpglvkpsetlsltctvsgvslpdygvs wirqppgkglewigviwgsettyyqsslksrvtiskdnsknqvslklssvtaadta vyycakhyyyggsyamdywgqgtlvtvssggggsggggsggggseivmtqspatls lspgeratlscrasqdiskylnwyqqkpgqaprlliyhtsrlhsgiparfsgsgsg tdytltisslqpedfavyfcqqgntlpytfgqgtkleik**hhhhhhhh**
104878 CAR 4 – Full - nt	911	atggctctgcccgtgaccgcactcctcctgccactggctctgctgcttcacgccgc tcgcccacaagtccagcttcaagaatcagggcctggtctggtgaagccatctgaga ctctgtccctcacttgcaccgtgagcggagtgtccctcccagactacggagtgagc tggattagacagcctcccggaaagggactggagtggatcggagtgatttggggtag cgaaaccacttactatcaatcttccctgaagtcacgggtcaccatttcaaaggata actcaaagaatcaagtgagcctcaagctctcatcagtcaccgccgctgacaccgcc gtgtattactgtgccaagcattactactatggagggtcctacgccatggactactg gggccagggaactctggtcactgtgtcatctggtggaggaggtagcggaggaggcg ggagcggtggaggtggctccgaaatcgtgatgacccagagccctgcaaccctgtcc ctttctccccggggaacgggctacccttttcttgtcgggcatcacaagatatctcaaa atacctcaattggtatcaacagaagccgggacaggcccctaggcttcttatctacc acacctctcgcctgcatagcgggattcccgcacgctttagcgggtctggaagcggg accgactacactctgaccatctcatctctccagcccgaggacttcgccgtctactt ctgccagcagggtaacaccctgccgtacaccttcggccagggcaccaagcttgaga tcaaaaccactactcccgctccaaggccacccacccctgccccgaccatcgcctct cagccgctttccctgcgtccggaggcatgtagacccgcagctggtggggccgtgca tacccggggtcttgacttcgcctgcgatatctacatttgggcccctctggctggta cttgcggggtcctgctgctttcactcgtgatcactctttactgtaagcgcggtcgg aagaagctgctgtacatctttaagcaacccttcatgaggcctgtgcagactactca agaggaggacggctgttcatgccggttcccagaggaggaggaaggcggctgcgaac tgcgcgtgaaattcagccgcagcgcagatgctccagcctacaagcaggggcagaac cagctctacaacgaactcaatcttggtcggagagaggagtacgacgtgctggacaa gcggagaggacgggacccagaaatgggcgggaagccgcgcagaaagaatccccaag agggcctgtacaacgagctccaaaaggataagatggcagaagcctatagcgagatt ggtatgaaaggggaacgcagaagaggcaaaggccacgacggactgtaccagggact cagcaccgccaccaaggacacctatgacgctcttcacatgcaggccctgccgcctc

		gg
104878 CAR 4 – Full - aa	912	MALPVTALLLPLALLLHAARPqvqlqesgpglvkpsetlsltctvsgvslp**dygvs** wirqppgkglewig**viwgsettyyqsslks**rvtiskdnsknqvslklssvtaadta vyycak**hyyyggsyamdy**wgqgtlvtvssggggsggggsggggsggggseivmtqspatls lspgeratlsc**rasqdiskyln**wyqqkpgqaprlliy**htsrlhs**giparfsgsgsg tdytltisslqpedfavyfc**qqgntlpyt**fgqgtkleikttpaprpptpaptias qplslrpeacrpaaggavhtrgldfacdiyiwaplagtcgvlllslvitlyckrgr kkllyifkqpfmrpvqttqeedgcscrfpeeeeggcelrvkfsrsadapaykqgqn qlynelnlgrreeydvldkrrgrdpemggkprrknpqeglynelqkdkmaeaysei gmkgerrrgkghdglyqglstatkdtydalhmqalppr
CAR 5		
CAR5 scFv domain	913	eivmtqspatlslspgeratlscrasqdiskylnwyqqkpgqaprlliyhtsrlhs giparfsgsgsgtdytltisslqpedfavyfcqqgntlpytfgqgtkleikggggs ggggsggggsggggsqvqlqesgpglvkpsetlsltctvsgvslpdygvswirqpp gkglewigviwgsettyyssslksrvtiskdnsknqvslklssvtaadtavyycak hyyyggsyamdywgqgtlvtvss
99789 **CAR5 - Soluble scFv - nt**	914	atggccctcccagtgaccgctctgctgctgcctctcgcacttcttctccatgccgc tcggcctgagatcgtcatgacccaaagccccgctaccctgtccctgtcacccggcg agagggcaacccctttcatgcagggccagccaggacatttctaagtacctcaactgg tatcagcagaagccagggcaggctcctcgcctgctgatctaccacaccagccgcct ccacagcggtatccccgccagattttccgggagcgggtctggaaccgactacaccc tcaccatctcttctctgcagcccgaggatttcgccgtctatttctgccagcagggg aatactctgccgtacaccttcggtcaaggtaccaagctggaaatcaagggaggcgg aggatcaggcggtggcggaagcggaggaggtggctccggaggaggaggttcccaag tgcagcttcaagaatcaggacccggacttgtgaagccatcagaaaccctctccctg acttgtaccgtgtccggtgtgagcctccccgactacggagtctcttggattcgcca gcctccggggaagggtcttgaatggattggggtgatttggggatcagagactactt actactcttcatcacttaagtcacgggtcaccatcagcaaagataatagcaagaac caagtgtcacttaagctgtcatctgtgaccgccgctgacaccgccgtgtactattg tgccaaacattactattacggagggtcttatgctatggactactggggacagggga ccctggtgactgtctctagccatcaccatcaccaccatcatcac
99789 **CAR5 - Soluble scFv -aa**	915	**MALPVTALLLPLALLLHAARP**eivmtqspatlslspgeratlscrasqdiskylnw yqqkpgqaprlliyhtsrlhsgiparfsgsgsgtdytltisslqpedfavyfcqqg ntlpytfgqgtkleikggggsggggsggggsggggsqvqlqesgpglvkpsetlsl tctvsgvslpdygvswirqppgkglewigviwgsettyyssslksrvtiskdnskn qvslklssvtaadtavyycakhyyyggsyamdywgqgtlvtvss**hhhhhhh**
104879 **CAR 5 – Full - nt**	916	atggccctccctgtcaccgccctgctgcttccgctggctcttctgctccacgccgc tcggcccgaaattgtgatgacccagtcacccgccactcttagcctttcacccggtg agcgcgcaaccctgtcttgcagagcctcccaagacatctcaaaataccttaattgg tatcaacagaagcccggacaggctcctcgccttctgatctaccacaccagccggct ccattctggaatccctgccaggttcagcggtagcggatctgggaccgactacaccc tcactatcagctcactgcagccagaggacttcgctgtctatttctgtcagcaaggg aacaccctgccctacacctttggacagggcaccaagctcgagattaaaggtggagg tggcagcggaggaggtgggtccggcggtggaggaagcggcggaggcgggagccagg

		tccaactccaagaaagcggaccgggtcttgtgaagccatcagaaactctttcactg acttgtactgtgagcggagtgtctctccccgattacggggtgtcttggatcagaca gccaccggggaagggtctggaatggattggagtgatttggggctctgagactactt actactcttcatccctcaagtcacgcgtcaccatctcaaaggacaactctaagaat caggtgtcactgaaactgtcatctgtgaccgcagccgacaccgccgtgtactattg cgctaagcattactattatggcgggagctacgcaatggattactggggacagggta ctctggtcaccgtgtccagcaccactaccccagcaccgaggccacccaccccggct cctaccatcgcctcccagcctctgtccctgcgtccggaggcatgtagacccgcagc tggtggggccgtgcatacccggggtcttgacttcgcctgcgatatctacatttggg cccctctggctggtacttgcggggtcctgctgctttcactcgtgatcactctttac tgtaagcgcggtcggaagaagctgctgtacatctttaagcaacccttcatgaggcc tgtgcagactactcaagaggaggacggctgttcatgccggttcccagaggaggagg aaggcggctgcgaactgcgcgtgaaattcagccgcagcgcagatgctccagcctac aagcaggggcagaaccagctctacaacgaactcaatcttggtcggagagaggagta cgacgtgctggacaagcggagaggacgggacccagaaatgggcgggaagccgcgca gaaagaatccccaagagggcctgtacaacgagctccaaaaggataagatggcagaa gcctatagcgagattggtatgaaaggggaacgcagaagaggcaaaggccacgacgg actgtaccagggactcagcaccgccaccaaggacacctatgacgctcttcacatgc aggccctgccgcctcgg
104879 **CAR 5 – Full - aa**	917	MALPVTALLLPLALLLHAARPeivmtqspatlslspgeratlsc**rasqdiskyln**w yqqkpgqaprlliy**htsrlhs**giparfsgsgsgtdytltisslqpedfavyfc**qqg** **ntlpyt**fgqgtkleikggggsggggsggggsggggsqvqlqesgpglvkpsetlsl tctvsgvslp**dygvs**wirqppgkglewig**viwgsettyyssslks**rvtiskdnskn qvslklssvtaadtavyycak**hyyyggsyamdy**wgqgtlvtvssttpaprpptpa ptiasqplslrpeacrpaaggavhtrgldfacdiyiwaplagtcgvlllslvitly ckrgrkkllyifkqpfmrpvqttqeedgcscrfpeeeeggcelrvkfsrsadapay kqgqnqlynelnlgrreeydvldkrrgrdpemggkprrknpqeglynelqkdkmae ayseigmkgerrrgkghdglyqglstatkdtydalhmqalppr
CAR 6		
CAR6 **scFv domain**	918	eivmtqspatlslspgeratlscrasqdiskylnwyqqkpgqaprlliyhtsrlhs giparfsgsgsgtdytltisslqpedfavyfcqqgntlpytfgqgtkleikggggs ggggsggggsggggsqvqlqesgpglvkpsetlsltctvsgvslpdygvswirqpp gkglewigviwgsettyyqsslksrvtiskdnsknqvslklssvtaadtavyycak hyyyggsyamdywgqgtlvtvss
99790 **CAR6 - Soluble scFv - nt**	919	atggccctcccagtgaccgctctgctgctgcctctcgcacttcttctccatgccgc tcggcctgagatcgtcatgacccaaagccccgctaccctgtccctgtcacccggcg agagggcaacccctttcatgcagggccagccaggacatttctaagtacctcaactgg tatcagcagaagccagggcaggctcctcgcctgctgatctaccacaccagccgcct ccacagcggtatccccgccagattttccgggagcgggtctggaaccgactacaccc tcaccatctcttctctgcagcccgaggatttcgccgtctatttctgccagcagggg aatactctgccgtacaccttcggtcaaggtaccaagctggaaatcaagggaggcgg aggatcaggcggtggcggaagcggaggaggtggctccggaggaggaggttcccaag tgcagcttcaagaatcaggacccggacttgtgaagccatcagaaaccctctccctg acttgtaccgtgtccggtgtgagcctccccgactacggagtctcttggattcgcca gcctccggggaagggtcttgaatggattggggtgatttggggatcagagactactt actaccagtcatcacttaagtcacgggtcaccatcagcaaagataatagcaagaac

		caagtgtcacttaagctgtcatctgtgaccgccgctgacaccgccgtgtactattg tgccaaacattactattacggagggtcttatgctatggactactggggacagggga ccctggtgactgtctctagccatcaccatcaccaccatcatcac
99790 **CAR6 - Soluble scFv - aa**	920	**MALPVTALLLPLALLLHAARP**eivmtqspatlslspgeratlscrasqdiskylnw yqqkpgqaprlliyhtsrlhsgiparfsgsgsgtdytltisslqpedfavyfcqqg ntlpytfgqgtkleikggggsggggsggggsggggsqvqlqesgpglvkpsetlsl tctvsgvslpdygvswirqppgkglewigviwgsettyyqsslksrvtiskdnskn qvslklssvtaadtavyycakhyyyggsyamdywgqgtlvtvss**hhhhhhhh**
104880 **CAR6 – Full - nt**	921	atggccctccctgtcaccgccctgctgcttccgctggctcttctgctccacgccgc tcggcccgaaattgtgatgacccagtcacccgccactcttagcctttcacccggtg agcgcgcaaccctgtcttgcagagcctcccaagacatctcaaaataccttaattgg tatcaacagaagcccggacaggctcctcgccttctgatctaccacaccagccggct ccattctggaatccctgccaggttcagcggtagcggatctgggaccgactacaccc tcactatcagctcactgcagccagaggacttcgctgtctatttctgtcagcaaggg aacaccctgccctacacctttggacagggcaccaagctcgagattaaaggtggagg tggcagcggaggaggtgggtccggcggtggaggaagcggaggcggagggagccagg tccaactccaagaaagcggaccgggtcttgtgaagccatcagaaactctttcactg acttgtactgtgagcggagtgtctccccgattacggggtgtcttggatcagaca gccaccggggaagggtctggaatggattggagtgatttggggctctgagactactt actaccaatcatccctcaagtcacgcgtcaccatctcaaaggacaactctaagaat caggtgtcactgaaactgtcatctgtgaccgcagccgacaccgccgtgtactattg cgctaagcattactattatggcgggagctacgcaatggattactggggacagggta ctctggtcaccgtgtccagcaccactaccccagcaccgaggccacccaccccggct cctaccatcgcctcccagcctctgtccctgcgtccggaggcatgtagacccgcagc tggtggggccgtgcatacccggggtcttgacttcgcctgcgatatctacatttggg cccctctggctggtacttgcggggtcctgctgctttcactcgtgatcactctttac tgtaagcgcggtcggaagaagctgctgtacatctttaagcaacccttcatgaggcc tgtgcagactactcaagaggaggacggctgttcatgccggttcccagaggaggagg aaggcggctgcgaactgcgcgtgaaattcagccgcagcgcagatgctccagcctac aagcaggggcagaaccagctctacaacgaactcaatcttggtcggagagaggagta cgacgtgctggacaagcggagaggacgggacccagaaatgggcgggaagccgcgca gaaagaatccccaagagggcctgtacaacgagctccaaaaggataagatggcagaa gcctatagcgagattggtatgaaaggggaacgcagaagaggcaaaggccacgacgg actgtaccagggactcagcaccgccaccaaggacacctatgacgctcttcacatgc aggccctgccgcctcgg
104880 **CAR6 – Full – aa**	922	MALPVTALLLPLALLLHAARPeivmtqspatlslspgeratlsc**rasqdiskyln**w yqqkpgqaprlliy**htsrlhs**giparfsgsgsgtdytltisslqpedfavyfc**qqg** **ntlpyt**fgqgtkleikggggsggggsggggsggggsqvqlqesgpglvkpsetlsl tctvsgvslp**dygvs**wirqppgkglewig**viwgsettyyqsslks**rvtiskdnskn qvslklssvtaadtavyycak**hyyyggsyamdy**wgqgtlvtvssttpaprpptpa ptiasqplslrpeacrpaaggavhtrgldfacdiyiwaplagtcgvlllslvitly ckrgrkkllyifkqpfmrpvqttqeedgcscrfpeeeeggcelrvkfsrsadapay kqgqnqlynelnlgrreeydvldkrrgrdpemggkprrknpqeglynelqkdkmae ayseigmkgerrrgkghdglyqglstatkdtydalhmqalppr

CAR 7		
CAR7 scFv domain	923	qvqlqesgpglvkpsetlsltctvsgvslpdygvswirqppgkglewigviwgset tyyssslksrvtiskdnsknqvslklssvtaadtavyycakhyyyggsyamdywgq gtlvtvssggggsggggsggggsggggseivmtqspatlslspgeratlscrasqd iskylnwyqqkpgqaprlliyhtsrlhsgiparfsgsgsgtdytltisslqpedfa vyfcqqgntlpytfgqgtkleik
100796 **CAR7 - Soluble scFv - nt**	924	atggcactgcctgtcactgccctcctgctgcctctggccctccttctgcatgccgc caggccccaagtccagctgcaagagtcaggacccggactggtgaagccgtctgaga ctctctcactgacttgtaccgtcagcggcgtgtccctccccgactacggagtgtca tggatccgccaacctcccgggaaagggcttgaatggattggtgtcatctggggttc tgaaaccacctactactcatcttccctgaagtccagggtgaccatcagcaaggata attccaagaaccaggtcagccttaagctgtcatctgtgaccgctgctgacaccgcc gtgtattactgcgccaagcactactattacggaggaagctacgctatggactattg gggacagggcactctcgtgactgtgagcagcggcggtggagggtctggaggtggag gatccggtggtggtgggtcaggcggaggagggagcgagattgtgatgactcagtca ccagccacccttctctttcacccggcgagagagcaaccctgagctgtagagccag ccaggacatttctaagtacctcaactggtatcagcaaaaaccggggcaggcccctc gcctcctgatctaccatacctcacgccttcactctggtatccccgctcggtttagc ggatcaggatctggtaccgactacactctgaccatttccagcctgcagccagaaga tttcgcagtgtatttctgccagcagggcaataccttccttacaccttcggtcagg gaaccaagctcgaaatcaagcaccatcaccatcatcaccaccat
100796 **CAR7 - Soluble scFv - aa**	925	**MALPVTALLLPLALLLHAARP**qvqlqesgpglvkpsetlsltctvsgvslpdygvs wirqppgkglewigviwgsettyyssslksrvtiskdnsknqvslklssvtaadta vyycakhyyyggsyamdywgqgtlvtvssggggsggggsggggsggggseivmtqs patlslspgeratlscrasqdiskylnwyqqkpgqaprlliyhtsrlhsgiparfs gsgsgtdytltisslqpedfavyfcqqgntlpytfgqgtkleik**hhhhhhhh**
104881 **CAR 7** **Full - nt**	926	atggctctgcccgtgaccgcactcctcctgccactggctctgctgcttcacgccgc tcgcccacaagtccagcttcaagaatcagggcctggtctggtgaagccatctgaga ctctgtccctcacttgcaccgtgagcggagtgtccctcccagactacggagtgagc tggattagacagcctcccggaaagggactggagtggatcggagtgatttggggtag cgaaaccacttactattcatcttccctgaagtcacgggtcaccatttcaaaggata actcaaagaatcaagtgagcctcaagctctcatcagtcaccgccgctgacaccgcc gtgtattactgtgccaagcattactactatggagggtcctacgccatggactactg gggccagggaactctggtcactgtgtcatctggtggaggaggtagcggaggaggcg ggagcggtggaggtggctccggaggtggcggaagcgaaatcgtgatgacccagagc cctgcaaccctgtccctttctcccggggaacgggctacccttttcttgtcgggcatc acaagatatctcaaaatacctcaattggtatcaacagaagccgggacaggccccta ggcttcttatctaccacacctctcgcctgcatagcgggattcccgcacgctttagc gggtctggaagcgggaccgactacactctgaccatctcatctctccagcccgagga cttcgccgtctacttctgccagcagggtaacaccctgccgtacaccttcggccagg gcaccaagcttgagatcaaaaccactactcccgctccaaggccacccacccctgcc ccgaccatcgcctctcagccgctttccctgcgtccggaggcatgtagacccgcagc tggtggggccgtgcatacccggggtcttgacttcgcctgcgatatctacatttggg cccctctggctggtacttgcggggtcctgctgctttcactcgtgatcactctttac tgtaagcgcggtcggaagaagctgctgtacatctttaagcaacccttcatgaggcc

		tgtgcagactactcaagaggaggacggctgttcatgccggttcccagaggaggagg aaggcggctgcgaactgcgcgtgaaattcagccgcagcgcagatgctccagcctac aagcaggggcagaaccagctctacaacgaactcaatcttggtcggagagaggagta cgacgtgctggacaagcggagaggacgggacccagaaatgggcgggaagccgcgca gaaagaatccccaagagggcctgtacaacgagctccaaaaggataagatggcagaa gcctatagcgagattggtatgaaaggggaacgcagaagaggcaaaggccacgacgg actgtaccagggactcagcaccgccaccaaggacacctatgacgctcttcacatgc aggccctgccgcctcgg
104881 **CAR 7** **Full - aa**	927	MALPVTALLLPLALLLHAARPqvqlqesgpglvkpsetlsltctvsgvslp**dygvs** wirqppgkglewig**viwgsettyyssslks**rvtiskdnsknqvslklssvtaadta vyycak**hyyyggsyamdy**wgqgtlvtvssggggsggggsggggsggggseivmtqs patlslspgeratlsc**rasqdiskyln**wyqqkpgqaprlliy**htsrlhs**giparfs gsgsgtdytltisslqpedfavyfc**qqgntlpyt**fgqgtkleikttpaprpptpa ptiasqplslrpeacrpaaggavhtrgldfacdiyiwaplagtcgvlllslvitly ckrgrkkllyifkqpfmrpvqttqeedgcscrfpeeeeggcelrvkfsrsadapay kqgqnqlynelnlgrreeydvldkrrgrdpemggkprrknpqeglynelqkdkmae ayseigmkgerrrgkghdglyqglstatkdtydalhmqalppr
CAR 8		
CAR8 scFv domain	928	qvqlqesgpglvkpsetlsltctvsgvslpdygvswirqppgkglewigviwgset tyyqsslksrvtiskdnsknqvslklssvtaadtavyycakhyyyggsyamdywgq gtlvtvssggggsggggsggggsggggseivmtqspatlslspgeratlscrasqd iskylnwyqqkpgqaprlliyhtsrlhsgiparfsgsgsgtdytltisslqpedfa vyfcqqgntlpytfgqgtkleik
100798 **CAR8 - Soluble scFv - nt**	929	atggcactgcctgtcactgccctcctgctgcctctggccctccttctgcatgccgc caggccccaagtccagctgcaagagtcaggacccggactggtgaagccgtctgaga ctctctcactgacttgtaccgtcagcggcgtgtccctccccgactacggagtgtca tggatccgccaacctcccgggaaagggcttgaatggattggtgtcatctggggttc tgaaaccacctactaccagtcttccctgaagtccagggtgaccatcagcaaggata attccaagaaccaggtcagccttaagctgtcatctgtgaccgctgctgacaccgcc gtgtattactgcgccaagcactactattacggaggaagctacgctatggactattg gggacagggcactctcgtgactgtgagcagcggcggtggagggtctggaggtggag gatccggtggtggtgggtcaggcggaggagggagcgagattgtgatgactcagtca ccagccacccttctctttcacccggcgagagagcaaccctgagctgtagagccag ccaggacatttctaagtacctcaactggtatcagcaaaaaccggggcaggccctc gcctcctgatctaccatacctcacgccttcactctggtatccccgctcggtttagc ggatcaggatctggtaccgactacactctgaccatttccagcctgcagccagaaga tttcgcagtgtatttctgccagcagggcaataccttccttacaccttcggtcagg gaaccaagctcgaaatcaagcaccatcaccatcatcatcaccac
100798 **CAR8 - Soluble scFv - aa**	930	**MALPVTALLLPLALLLHAARP**qvqlqesgpglvkpsetlsltctvsgvslpdygvs wirqppgkglewigviwgsettyyqsslksrvtiskdnsknqvslklssvtaadta vyycakhyyyggsyamdywgqgtlvtvssggggsggggsggggsggggseivmtqs patlslspgeratlscrasqdiskylnwyqqkpgqaprlliyhtsrlhsgiparfs gsgsgtdytltisslqpedfavyfcqqgntlpytfgqgtkleik**hhhhhhhh**

104882 CAR 8 – Full - nt	931	atggctctgcccgtgaccgcactcctcctgccactggctctgctgcttcacgccgc tcgcccacaagtccagcttcaagaatcagggcctggtctggtgaagccatctgaga ctctgtccctcacttgcaccgtgagcggagtgtccctcccagactacggagtgagc tggattagacagcctcccggaaagggactggagtggatcggagtgatttggggtag cgaaaccacttactatcaatcttccctgaagtcacgggtcaccatttcaaaggata actcaaagaatcaagtgagcctcaagctctcatcagtcaccgccgctgacaccgcc gtgtattactgtgccaagcattactactatggagggtcctacgccatggactactg gggccagggaactctggtcactgtgtcatctggtggaggaggtagcggaggaggcg ggagcggtggaggtggctccggaggcggtgggtcagaaatcgtgatgacccagagc cctgcaaccctgtccctttctcccggggaacgggctacccttcttgtcgggcatc acaagatatctcaaaatacctcaattggtatcaacagaagccgggacaggccccta ggcttcttatctaccacacctctcgcctgcatagcgggattcccgcacgctttagc gggtctggaagcgggaccgactacactctgaccatctcatctctccagcccgagga cttcgccgtctacttctgccagcagggtaacaccctgccgtacaccttcggccagg gcaccaagcttgagatcaaaaccactactcccgctccaaggccacccacccctgcc ccgaccatcgcctctcagccgctttccctgcgtccggaggcatgtagacccgcagc tggtggggccgtgcatacccggggtcttgacttcgcctgcgatatctacatttggg cccctctggctggtacttgcggggtcctgctgctttcactcgtgatcactctttac tgtaagcgcggtcggaagaagctgctgtacatctttaagcaacccttcatgaggcc tgtgcagactactcaagaggaggacggctgttcatgccggttcccagaggaggagg aaggcggctgcgaactgcgcgtgaaattcagccgcagcgcagatgctccagcctac aagcaggggcagaaccagctctacaacgaactcaatcttggtcggagagaggagta cgacgtgctggacaagcggagaggacgggacccagaaatgggcgggaagccgcgca gaaagaatccccaagagggcctgtacaacgagctccaaaaggataagatggcagaa gcctatagcgagattggtatgaaaggggaacgcagaagaggcaaaggccacgacgg actgtaccagggactcagcaccgccaccaaggacacctatgacgctcttcacatgc aggccctgccgcctcgg
104882 CAR 8 – Full - aa	932	MALPVTALLLPLALLLHAARPqvqlqesgpglvkpsetlsltctvsgvslp**dygvs** wirqppgkglewig**viwgsettyyqsslks**rvtiskdnsknqvslklssvtaadta vyycak**hyyyggsyamdy**wgqgtlvtvssggggsggggsggggsggggseivmtqs patlslspgeratlsc**rasqdiskyln**wyqqkpgqaprlliy**htsrlhs**giparfs gsgsgtdytltisslqpedfavyfc**qqgntlpyt**fgqgtkleikttpaprpptpa ptiasqplslrpeacrpaaggavhtrgldfacdiyiwaplagtcgvlllslvitly ckrgrkkllyifkqpfmrpvqttqeedgcscrfpeeeeggcelrvkfsrsadapay kqgqnqlynelnlgrreeydvldkrrgrdpemggkprrknpqeglynelqkdkmae ayseigmkgerrrgkghdglyqglstatkdtydalhmqalppr
CAR 9		
CAR9 scFv domain	933	eivmtqspatlslspgeratlscrasqdiskylnwyqqkpgqaprlliyhtsrlhs giparfsgsgsgtdytltisslqpedfavyfcqqgntlpytfgqgtkleikggggs ggggsggggsggggsqvqlqesgpglvkpsetlsltctvsgvslpdygvswirqpp gkglewigviwgsettyynsslksrvtiskdnsknqvslklssvtaadtavyycak hyyyggsyamdywgqgtlvtvss
99789 CAR9 -	934	atggccctcccagtgaccgctctgctgctgcctctcgcacttcttctccatgccgc tcggcctgagatcgtcatgacccaaagccccgctaccctgtccctgtcacccggcg agagggcaacccctttcatgcagggccagccaggacatttctaagtacctcaactgg tatcagcagaagccagggcaggctcctcgcctgctgatctaccacaccagccgcct

Soluble scFv - nt		ccacagcggtatccccgccagattttccgggagcgggtctggaaccgactacaccc tcaccatctcttctctgcagcccgaggatttcgccgtctatttctgccagcagggg aatactctgccgtacaccttcggtcaaggtaccaagctggaaatcaagggaggcgg aggatcaggcggtggcggaagcggaggaggtggctccggaggaggaggttcccaag tgcagcttcaagaatcaggacccggacttgtgaagccatcagaaaccctctccctg acttgtaccgtgtccggtgtgagcctccccgactacggagtctcttggattcgcca gcctccggggaagggtcttgaatggattggggtgatttggggatcagagactactt actacaattcatcacttaagtcacgggtcaccatcagcaaagataatagcaagaac caagtgtcacttaagctgtcatctgtgaccgccgctgacaccgccgtgtactattg tgccaaacattactattacggagggtcttatgctatggactactggggacagggga ccctggtgactgtctctagccatcaccatcaccaccatcatcac
99789 **CAR9 - Soluble scFv - aa**	935	**MALPVTALLLPLALLLHAARP**eivmtqspatlslspgeratlscrasqdiskylnw yqqkpgqaprlliyhtsrlhsgiparfsgsgsgtdytltisslqpedfavyfcqqg ntlpytfgqgtkleikggggsggggsggggsggggsqvqlqesgpglvkpsetlsl tctvsgvslpdygvswirqppgkglewigviwgsettyynsslksrvtiskdnskn qvslklssvtaadtavyycakhyyyggsyamdywgqgtlvtvss**hhhhhhhh**
105974 **CAR 9 – Full - nt**	936	atggccctccctgtcaccgccctgctgcttccgctggctcttctgctccacgccgc tcggcccgaaattgtgatgacccagtcacccgccactcttagcctttcacccggtg agcgcgcaaccctgtcttgcagagcctcccaagacatctcaaaataccttaattgg tatcaacagaagcccggacaggctcctcgccttctgatctaccacaccagccggct ccattctggaatccctgccaggttcagcggtagcggatctgggaccgactacaccc tcactatcagctcactgcagccagaggacttcgctgtctatttctgtcagcaaggg aacaccctgccctacacctttggacagggcaccaagctcgagattaaaggtggagg tggcagcggaggaggtgggtccggcggtggaggaagcggaggcggtgggagccagg tccaactccaagaaagcggaccgggtcttgtgaagccatcagaaactctttcactg acttgtactgtgagcggagtgtctctccccgattacggggtgtcttggatcagaca gccaccggggaagggtctggaatggattggagtgatttggggctctgagactactt actacaactcatccctcaagtcacgcgtcaccatctcaaaggacaactctaagaat caggtgtcactgaaactgtcatctgtgaccgcagccgacaccgccgtgtactattg cgctaagcattactattatggcgggagctacgcaatggattactggggacagggta ctctggtcaccgtgtccagcaccactaccccagcaccgaggccacccaccccggct cctaccatcgcctcccagcctctgtccctgcgtccggaggcatgtagacccgcagc tggtggggccgtgcatacccggggtcttgacttcgcctgcgatatctacatttggg cccctctggctggtacttgcggggtcctgctgctttcactcgtgatcactctttac tgtaagcgcggtcggaagaagctgctgtacatctttaagcaacccttcatgaggcc tgtgcagactactcaagaggaggacggctgttcatgccggttcccagaggaggagg aaggcggctgcgaactgcgcgtgaaattcagccgcagcgcagatgctccagcctac aagcaggggcagaaccagctctacaacgaactcaatcttggtcggagagaggagta cgacgtgctggacaagcggagaggacgggacccagaaatgggcgggaagccgcgca gaaagaatccccaagagggcctgtacaacgagctccaaaaggataagatggcagaa gcctatagcgagattggtatgaaaggggaacgcagaagaggcaaaggccacgacgg actgtaccagggactcagcaccgccaccaaggacacctatgacgctcttcacatgc aggccctgccgcctcgg
105974 **CAR 9 – Full**	937	MALPVTALLLPLALLLHAARPeivmtqspatlslspgeratlsc**rasqdiskyln**w yqqkpgqaprlliy**htsrlhs**giparfsgsgsgtdytltisslqpedfavyfc**qqg** **ntlpyt**fgqgtkleikggggsggggsggggsggggsqvqlqesgpglvkpsetlsl

- aa		tctvsgvslp**dygvs**wirqppgkglewig**viwgsettyynsslks**rvtiskdnskn qvslklssvtaadtavyycak**hyyyggsyamdy**wgqgtlvtvsstttpaprpptpa ptiasqplslrpeacrpaaggavhtrgldfacdiyiwaplagtcgvlllslvitly ckrgrkkllyifkqpfmrpvqttqeedgcscrfpeeeeggcelrvkfsrsadapay kqgqnqlynelnlgrreeydvldkrrgrdpemggkprrknpqeglynelqkdkmae ayseigmkgerrrgkghdglyqglstatkdtydalhmqalppr
CAR10		
CAR10 scFv domain	938	qvqlqesgpglvkpsetlsltctvsgvslpdygvswirqppgkglewigviwgset tyynsslksrvtiskdnsknqvslklssvtaadtavyycakhyyyggsyamdywgq gtlvtvssggggsggggsggggsggggseivmtqspatlslspgeratlscrasqd iskylnwyqqkpgqaprlliyhtsrlhsgiparfsgsgsgtdytltisslqpedfa vyfcqqgntlpytfgqgtkleik
100796 CAR10 - Soluble scFv - nt	939	atggcactgcctgtcactgccctcctgctgcctctggccctccttctgcatgccgc caggccccaagtccagctgcaagagtcaggacccggactggtgaagccgtctgaga ctctctcactgacttgtaccgtcagcggcgtgtccctccccgactacggagtgtca tggatccgccaacctcccgggaaagggcttgaatggattggtgtcatctggggttc tgaaaccacctactacaactcttccctgaagtccagggtgaccatcagcaaggata attccaagaaccaggtcagccttaagctgtcatctgtgaccgctgctgacaccgcc gtgtattactgcgccaagcactactattacggaggaagctacgctatggactattg gggacagggcactctcgtgactgtgagcagcggcggtggagggtctggaggtggag gatccggtggtggtgggtcaggcggaggagggagcgagattgtgatgactcagtca ccagccacccttctctttcacccggcgagagagcaaccctgagctgtagagccag ccaggacatttctaagtacctcaactggtatcagcaaaaaccggggcaggcccctc gcctcctgatctaccatacctcacgccttcactctggtatccccgctcggtttagc ggatcaggatctggtaccgactacactctgaccatttccagcctgcagccagaaga tttcgcagtgtatttctgccagcagggcaataccttccttacaccttcggtcagg gaaccaagctcgaaatcaagcaccatcaccatcatcaccaccat
100796 CAR10 - Soluble scFv - aa	940	**MALPVTALLLPLALLLHAARP**qvqlqesgpglvkpsetlsltctvsgvslpdygvs wirqppgkglewigviwgsettyynsslksrvtiskdnsknqvslklssvtaadta vyycakhyyyggsyamdywgqgtlvtvssggggsggggsggggsggggseivmtqs patlslspgeratlscrasqdiskylnwyqqkpgqaprlliyhtsrlhsgiparfs gsgsgtdytltisslqpedfavyfcqqgntlpytfgqgtkleik**hhhhhhhh**
105975 CAR 10 Full - nt	941	atggccctccctgtcaccgccctgctgcttccgctggctcttctgctccacgccgc tcggcccgaaattgtgatgacccagtcacccgccactcttagcctttcacccggtg agcgcgcaaccctgtcttgcagagcctcccaagacatctcaaaataccttaattgg tatcaacagaagcccggacaggctcctcgccttctgatctaccacaccagccggct ccattctggaatccctgccaggttcagcggtagcggatctgggaccgactacaccc tcactatcagctcactgcagccagaggacttcgctgtctatttctgtcagcaaggg aacaccctgccctacacctttggacagggcaccaagctcgagattaaaggtggagg tggcagcggaggaggtgggtccggcggtggaggaagcggaggcggtgggagccagg tccaactccaagaaagcggaccgggtcttgtgaagccatcagaaactctttcactg acttgtactgtgagcggagtgtctctccccgattacggggtgtcttggatcagaca gccaccggggaagggtctggaatggattggagtgatttggggctctgagactactt actacaactcatccctcaagtcacgcgtcaccatctcaaaggacaactctaagaat

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105975 **CAR 10 Full - aa**	942	MALPVTALLLPLALLLHAARPEIVMTQSPATLSLSPGERATLSC**RASQDISKYLN**W YQQKPGQAPRLLIY**HTSRLHS**GIPARFSGSGSGTDYTLTISSLQPEDFAVYFC**QQG** **NTLPYT**FGQGTKLEIKGGGGSGGGGSGGGGSGGGGSQVQLQESGPGLVKPSETLSL TCTVSGVSLP**DYGVS**WIRQPPGKGLEWIG**VIWGSETTYYNSSLKS**RVTISKDNSKN QVSLKLSSVTAADTAVYYCAK**HYYYGGSYAMDY**WGQGTLVTVSSTTTPAPRPPTPA PTIASQPLSLRPEACRPAAGGAVHTRGLDFACDIYIWAPLAGTCGVLLLSLVITLY CKRGRKKLLYIFKQPFMRPVQTTQEEDGCSCRFPEEEEGGCELRVKFSRSADAPAY KQGQNQLYNELNLGRREEYDVLDKRRGRDPEMGGKPRRKNPQEGLYNELQKDKMAE AYSEIGMKGERRRGKGHDGLYQGLSTATKDTYDALHMQALPPR
CAR11		
CAR11 scFv domain	943	eivmtqspatlslspgeratlscrasqdiskylnwyqqkpgqaprlliyhtsrlhs giparfsgsgsgtdytltisslqpedfavyfcqqgntlpytfgqgtkleikggggs ggggsggggsqvqlqesgpglvkpsetlsltctvsgvslpdygvswirqppgkgle wigviwgsettyynsslksrvtiskdnsknqvslklssvtaadtavyycakhyyyg gsyamdywgqgtlvtvss
103101 **CAR11 - Soluble scFv - nt**	944	Atggccctccctgtcaccgccctgctgcttccgctggctcttctgctccacgccgc tcggcccgaaattgtgatgacccagtcacccgccactcttagcctttcacccggtg agcgcgcaaccctgtcttgcagagcctcccaagacatctcaaaataccttaattgg tatcaacagaagcccggacaggctcctcgccttctgatctaccacaccagccggct ccattctggaatccctgccaggttcagcggtagcggatctgggaccgactacaccc tcactatcagctcactgcagccagaggacttcgctgtctatttctgtcagcaaggg aacaccctgccctacacctttggacagggcaccaagctcgagattaaaggtggagg tggcagcggaggaggtgggtccggcggtggaggaagccaggtccaactccaagaaa gcggaccgggtcttgtgaagccatcagaaactctttcactgacttgtactgtgagc ggagtgtctctccccgattacggggtgtcttggatcagacagccaccggggaaggg tctggaatggattggagtgatttggggctctgagactacttactacaattcatccc tcaagtcacgcgtcaccatctcaaaggacaactctaagaatcaggtgtcactgaaa

		ctgtcatctgtgaccgcagccgacaccgccgtgtactattgcgctaagcattacta ttatggcgggagctacgcaatggattactggggacagggtactctggtcaccgtgt ccagccaccaccatcatcaccatcaccat
103101 CAR11 - Soluble scFv - aa	945	**MALPVTALLLPLALLLHAARP**eivmtqspatlslspgeratlscrasqdiskylnw yqqkpgqaprlliyhtsrlhsgiparfsgsgsgtdytltisslqpedfavyfcqqg ntlpytfgqgtkleikggggsggggsggggsqvqlqesgpglvkpsetlsltctvs gvslpdygvswirqppgkglewigviwgsettyynsslksrvtiskdnsknqvslk lssvtaadtavyycakhyyyggsyamdywgqgtlvtvss**hhhhhhhh**
105976 CAR 11 Full - nt	946	atggctctgcccgtgaccgcactcctcctgccactggctctgctgcttcacgccgc tcgcccacaagtccagcttcaagaatcagggcctggtctggtgaagccatctgaga ctctgtccctcacttgcaccgtgagcggagtgtccctcccagactacggagtgagc tggattagacagcctcccggaaagggactggagtggatcggagtgatttggggtag cgaaaccacttactataactcttccctgaagtcacgggtcaccatttcaaaggata actcaaagaatcaagtgagcctcaagctctcatcagtcaccgccgctgacaccgcc gtgtattactgtgccaagcattactactatggagggtcctacgccatggactactg gggccagggaactctggtcactgtgtcatctggtggaggaggtagcggaggaggcg ggagcggtggaggtggctccggaggtggcggaagcgaaatcgtgatgacccagagc cctgcaaccctgtccctttctcccggggaacgggctacccttcttgtcgggcatc acaagatatctcaaaatacctcaattggtatcaacagaagccgggacaggccccta ggcttcttatctaccacacctctcgcctgcatagcgggattcccgcacgctttagc gggtctggaagcgggaccgactacactctgaccatctcatctctccagcccgagga cttcgccgtctacttctgccagcagggtaacaccctgccgtacaccttcggccagg gcaccaagcttgagatcaaaaccactactcccgctccaaggccacccaccctgcc ccgaccatcgcctctcagccgctttccctgcgtccggaggcatgtagacccgcagc tggtggggccgtgcatacccggggtcttgacttcgcctgcgatatctacatttggg cccctctggctggtacttgcggggtcctgctgctttcactcgtgatcactctttac tgtaagcgcggtcggaagaagctgctgtacatctttaagcaacccttcatgaggcc tgtgcagactactcaagaggaggacggctgttcatgccggttcccagaggaggagg aaggcggctgcgaactgcgcgtgaaattcagccgcagcgcagatgctccagcctac aagcaggggcagaaccagctctacaacgaactcaatcttggtcggagagaggagta cgacgtgctggacaagcggagaggacgggacccagaaatgggcgggaagccgcgca gaaagaatccccaagagggcctgtacaacgagctccaaaaggataagatggcagaa gcctatagcgagattggtatgaaaggggaacgcagaagaggcaaaggccacgacgg actgtaccagggactcagcaccgccaccaaggacacctatgacgctcttcacatgc aggccctgccgcctcgg
105976 CAR 11 Full - aa	947	MALPVTALLLPLALLLHAARPQVQLQESGPGLVKPSETLSLTCTVSGVSLP**DYGVS** WIRQPPGKGLEWIG**VIWGSETTYYNSSLKS**RVTISKDNSKNQVSLKLSSVTAADTA VYYCAK**HYYYGGSYAMDY**WGQGTLVTVSSGGGGSGGGGSGGGGSGGGGSEIVMTQS PATLSLSPGERATLSC**RASQDISKYLN**WYQQKPGQAPRLLIY**HTSRLHS**GIPARFS GSGSGTDYTLTISSLQPEDFAVYFC**QQGNTLPYT**FGQGTKLEIKTTTPAPRPPTPA PTIASQPLSLRPEACRPAAGGAVHTRGLDFACDIYIWAPLAGTCGVLLLSLVITLY CKRGRKKLLYIFKQPFMRPVQTTQEEDGCSCRFPEEEEGGCELRVKFSRSADAPAY KQGQNQLYNELNLGRREEYDVLDKRRGRDPEMGGKPRRKNPQEGLYNELQKDKMAE AYSEIGMKGERRRGKGHDGLYQGLSTATKDTYDALHMQALPPR

CAR12		
CAR12 **scFv domain**	948	qvqlqesgpglvkpsetlsltctvsgvslpdygvswirqppgkglewigviwgset tyynsslksrvtiskdnsknqvslklssvtaadtavyycakhyyyggsyamdywgq gtlvtvssggggsggggsggggseivmtqspatlslspgeratlscrasqdiskyl nwyqqkpgqaprlliyhtsrlhsgiparfsgsgsgtdytltisslqpedfavyfcq qgntlpytfgqgtkleik
103104 **CAR12 - Soluble scFv - nt**	949	atggctctgcccgtgaccgcactcctcctgccactggctctgctgcttcacgccgc tcgcccacaagtccagcttcaagaatcagggcctggtctggtgaagccatctgaga ctctgtccctcacttgcaccgtgagcggagtgtccctcccagactacggagtgagc tggattagacagcctcccggaaagggactggagtggatcggagtgatttggggtag cgaaaccacttactataactcttccctgaagtcacgggtcaccatttcaaaggata actcaaagaatcaagtgagcctcaagctctcatcagtcaccgccgctgacaccgcc gtgtattactgtgccaagcattactactatggagggtcctacgccatggactactg gggccagggaactctggtcactgtgtcatctggtggaggaggtagcggaggaggcg ggagcggtggaggtggctccgaaatcgtgatgacccagagccctgcaaccctgtcc ctttctcccggggaacgggctacccttcttgtcgggcatcacaagatatctcaaa atacctcaattggtatcaacagaagccgggacaggcccctaggcttcttatctacc acacctctcgcctgcatagcgggattcccgcacgctttagcgggtctggaagcggg accgactacactctgaccatctcatctctccagcccgaggacttcgccgtctactt ctgccagcagggtaacaccctgccgtacaccttcggccagggcaccaagcttgaga tcaaacatcaccaccatcatcaccatcac
103104 **CAR12 - Soluble scFv -aa**	950	**MALPVTALLLPLALLLHAARP**qvqlqesgpglvkpsetlsltctvsgvslpdygvs wirqppgkglewigviwgsettyynsslksrvtiskdnsknqvslklssvtaadta vyycakhyyyggsyamdywgqgtlvtvssggggsggggsggggseivmtqspatls lspgeratlscrasqdiskylnwyqqkpgqaprlliyhtsrlhsgiparfsgsgsg tdytltisslqpedfavyfcqqgntlpytfgqgtkleik**hhhhhhhh**
105977 **CAR 12 – Full - nt**	951	atggccctccctgtcaccgccctgctgcttccgctggctcttctgctccacgccgc tcggcccgaaattgtgatgacccagtcacccgccactcttagcctttcacccggtg agcgcgcaaccctgtcttgcagagcctcccaagacatctcaaaataccttaattgg tatcaacagaagcccggacaggctcctcgccttctgatctaccacaccagccggct ccattctggaatccctgccaggttcagcggtagcggatctgggaccgactacaccc tcactatcagctcactgcagccagaggacttcgctgtctatttctgtcagcaaggg aacaccctgccctacacctttggacagggcaccaagctcgagattaaaggtggagg tggcagcggaggaggtgggtccggcggtggaggaagccaggtccaactccaagaaa gcggaccgggtcttgtgaagccatcagaaactctttcactgacttgtactgtgagc ggagtgtctctccccgattacggggtgtcttggatcagacagccaccggggaaggg tctggaatggattggagtgatttggggctctgagactacttactacaactcatccc tcaagtcacgcgtcaccatctcaaaggacaactctaagaatcaggtgtcactgaaa ctgtcatctgtgaccgcagccgacaccgccgtgtactattgcgctaagcattacta ttatggcgggagctacgcaatggattactggggacagggtactctggtcaccgtgt ccagcaccactaccccagcaccgaggccacccaccccggctcctaccatcgcctcc cagcctctgtccctgcgtccggaggcatgtagacccgcagctggtggggccgtgca tacccggggtcttgacttcgcctgcgatatctacatttgggcccctctggctggta

		cttgcggggtcctgctgctttcactcgtgatcactctttactgtaagcgcggtcgg aagaagctgctgtacatctttaagcaacccttcatgaggcctgtgcagactactca agaggaggacggctgttcatgccggttcccagaggaggaggaaggcggctgcgaac tgcgcgtgaaattcagccgcagcgcagatgctccagcctacaagcaggggcagaac cagctctacaacgaactcaatcttggtcggagagaggagtacgacgtgctggacaa gcggagaggacgggacccagaaatgggcgggaagccgcgcagaaagaatccccaag agggcctgtacaacgagctccaaaaggataagatggcagaagcctatagcgagatt ggtatgaaaggggaacgcagaagaggcaaaggccacgacggactgtaccagggact cagcaccgccaccaaggacacctatgacgctcttcacatgcaggccctgccgcctc gg
105977 **CAR 12 – Full - aa**	952	MALPVTALLLPLALLLHAARPEIVMTQSPATLSLSPGERATLSC**RASQDISKYLN**W YQQKPGQAPRLLIY**HTSRLHS**GIPARFSGSGSGTDYTLTISSLQPEDFAVYFC**QQG** **NTLPYT**FGQGTKLEIKGGGGSGGGGSGGGGSQVQLQESGPGLVKPSETLSLTCTVS GVSLP**DYGVS**WIRQPPGKGLEWIG**VIWGSETTYYNSSLKS**RVTISKDNSKNQVSLK LSSVTAADTAVYYCAK**HYYYGGSYAMDY**WGQGTLVTVSSTTTPAPRPPTPAPTIAS QPLSLRPEACRPAAGGAVHTRGLDFACDIYIWAPLAGTCGVLLLSLVITLYCKRGR KKLLYIFKQPFMRPVQTTQEEDGCSCRFPEEEEGGCELRVKFSRSADAPAYKQGQN QLYNELNLGRREEYDVLDKRRGRDPEMGGKPRRKNPQEGLYNELQKDKMAEAYSEI GMKGERRRGKGHDGLYQGLSTATKDTYDALHMQALPPR
CTL019		
CTL019 – Soluble scFv-Histag - nt	953	atggccctgcccgtcaccgctctgctgctgcccttgctctgcttcttcatgcagc aaggccggacatccagatgacccaaaccacctcatccctctctgcctctcttggag acagggtgaccatttcttgtcgcgccagccaggacatcagcaagtatctgaactgg tatcagcagaagccggacggaaccgtgaagctcctgatctaccatacctctcgcct gcatagcggcgtgccctcacgcttctctggaagcggatcaggaaccgattattctc tcactatttcaaatcttgagcaggaagatattgccacctatttctgccagcagggt aataccctgccctacaccttcggaggagggaccaagctcgaaatcaccggtggagg aggcagcggcggtggagggtctggtggaggtggttctgaggtgaagctgcaagaat caggccctggacttgtggcccttcacagtccctgagcgtgacttgcaccgtgtcc ggagtctccctgcccgactacggagtgtcatggatcagacaacctccacggaaagg actggaatggctcggtgtcatctggggtagcgaaactacttactacaattcagccc tcaaaagcaggctgactattatcaaggacaacagcaagtcccaagtctttcttaag atgaactcactccagactgacgacaccgcaatctactattgtgctaagcactacta ctacggaggatcctacgctatggattactggggacaaggtacttccgtcactgtct cttcacaccatcatcaccatcaccatcac
CTL019 – Soluble scFv-Histag - aa	954	**MALPVTALLLPLALLLHAARP**diqmtqttsslsaslgdrvtiscrasqdiskylnw yqqkpdgtvklliyhtsrlhsgvpsrfsgsgsgtdysltisnleqediatyfcqqg ntlpytfgggtkleitggggsggggsggggsevklqesgpglvapsqslsvtctvs gvslpdygvswirqpprkglewlgviwgsettyynsalksrltiikdnsksqvflk mnslqtddtaiyycakhyyyggsyamdywgqgtsvtvss**hhhhhhhh**

CTL019 Full - nt	955	atggccttaccagtgaccgccttgctcctgccgctggccttgctgctccacgccgc caggccggacatccagatgacacagactacatcctccctgtctgcctctctgggag acagagtcaccatcagttgcagggcaagtcaggacattagtaaatatttaaattgg tatcagcagaaaccagatggaactgttaaactcctgatctaccatacatcaagatt acactcaggagtcccatcaaggttcagtggcagtgggtctggaacagattattctc tcaccattagcaacctggagcaagaagatattgccacttacttttgccaacagggt aatacgcttccgtacacgttcggaggggggaccaagctggagatcacaggtggcgg tggctcgggcggtggtgggtcgggtggcggcggatctgaggtgaaactgcaggagt caggacctggcctggtggcgccctcacagagcctgtccgtcacatgcactgtctca ggggtctcattacccgactatggtgtaagctggattcgccagcctccacgaaaggg tctggagtggctgggagtaatatggggtagtgaaaccacatactataattcagctc tcaaatccagactgaccatcatcaaggacaactccaagagccaagttttcttaaaa atgaacagtctgcaaactgatgacacagccatttactactgtgccaaacattatta ctacggtggtagctatgctatggactactggggccaaggaacctcagtcaccgtct cctcaaccacgacgccagcgccgcgaccaccaacaccggcgcccaccatcgcgtcg cagcccctgtccctgcgcccagaggcgtgccggccagcggcgggggcgcagtgca cacgaggggctggacttcgcctgtgatatctacatctgggcgcccttggccggga cttgtggggtccttctcctgtcactggttatcacccttactgcaaacggggcaga aagaaactcctgtatatattcaaacaaccatttatgagaccagtacaaactactca agaggaagatggctgtagctgccgatttccagaagaagaagaaggaggatgtgaac tgagagtgaagttcagcaggagcgcagacgcccccgcgtacaagcagggccagaac cagctctataacgagctcaatctaggacgaagagaggagtacgatgttttggacaa gagacgtggccgggaccctgagatgggggaaagccgagaaggaagaaccctcagg aaggcctgtacaatgaactgcagaaagataagatggcggaggcctacagtgagatt gggatgaaaggcgagcgccggaggggcaaggggcacgatggcctttaccagggtct cagtacagccaccaaggacacctacgacgcccttcacatgcaggccctgcccccctc gc
CTL019 Full - aa	956	MALPVTALLLPLALLLHAARPdiqmtqttsslsaslgdrvtiscrasqdiskylnw yqqkpdgtvklliyhtsrlhsgvpsrfsgsgsgtdysltisnleqediatyfcqqg ntlpytfgggtkleitggggsggggsggggsevklqesgpglvapsqslsvtctvs gvslpdygvswirqpprkglewlgviwgsettyynsalksrltiikdnsksqvflk mnslqtddtaiyycakhyyyggsyamdywgqgtsvtvsstttpaprpptpaptias qplslrpeacrpaaggavhtrgldfacdiyiwaplagtcgvlllslvitlyckrgr kkllyifkqpfmrpvqttqeedgcscrfpeeeeggcelrvkfsrsadapaykqgqn qlynelnlgrreeydvldkrrgrdpemggkprrknpqeglynelqkdkmaeaysei gmkgerrrgkghdglyqglstatkdtydalhmqalppr
CTL019 scFv domain	957	diqmtqttsslsaslgdrvtiscrasqdiskylnwyqqkpdgtvklliyhtsrlhs gvpsrfsgsgsgtdysltisnleqediatyfcqqgntlpytfgggtkleitgggs ggggsggggsevklqesgpglvapsqslsvtctvsgvslpdygvswirqpprkgle wlgviwgsettyynsalksrltiikdnsksqvflkmnslqtddtaiyycakhyyyg gsyamdywgqgtsvtvss

In some embodiments, the CD19 CAR or binding domain includes the amino acid sequence of CTL019, or is encoded by the nucleotide sequence of CTL019 according to Table 9 with or without the

leader sequence or the his tag, or a sequence substantially identical thereto (e.g., at least 85%, 90%, 95% or higher identity).

In some embodiments, the CDRs are defined according to the Kabat numbering scheme, the Chothia numbering scheme, or a combination thereof.

The sequences of humanized CDR sequences of the scFv domains are shown in Table 10A for the heavy chain variable domains and in Table 10B for the light chain variable domains. "ID" stands for the respective SEQ ID NO for each CDR.

Table 10A. Heavy Chain Variable Domain CDRs (according to Kabat)

Candidate	FW	HCDR1	SEQ ID	HCDR2	SEQ ID	HCDR3	SEQ ID
murine_CART19		DYGVS	958	VIWGSETTYYNSALKS	959	HYYYGGSYAMDY	960
humanized_CART19 a	VH4	DYGVS	958	VIWGSETTYY***S***S***S***LKS	961	HYYYGGSYAMDY	960
humanized_CART19 b	VH4	DYGVS	958	VIWGSETTYY***Q***S***S***LKS	962	HYYYGGSYAMDY	960
humanized_CART19 c	VH4	DYGVS	958	VIWGSETTYYNS***S***LKS	963	HYYYGGSYAMDY	960

Table 10B Light Chain Variable Domain CDRs (according to Kabat)

Candidate	FW	LCDR1	SEQ ID	LCDR2	SEQ ID	LCDR3	SEQ ID
murine_CART19		RASQDISKYLN	964	HTSRLHS	965	QQGNTLPYT	966
humanized_CART19 a	VK3	RASQDISKYLN	964	HTSRLHS	965	QQGNTLPYT	966
humanized_CART19 b	VK3	RASQDISKYLN	964	HTSRLHS	965	QQGNTLPYT	966
humanized_CART19 c	VK3	RASQDISKYLN	964	HTSRLHS	965	QQGNTLPYT	966

In one embodiment, the CAR molecule comprises a BCMA CAR molecule described herein, e.g., a BCMA CAR described in US-2016-0046724-A1 or WO2016/014565. In embodiments, the BCMA CAR comprises an amino acid, or has a nucleotide sequence of a CAR molecule, or an antigen binding domain

according to US-2016-0046724-A1, or Table 1 or 16, SEQ ID NO: 271 or SEQ ID NO: 273 of WO2016/014565, incorporated herein by reference, or a sequence substantially identical to any of the aforesaid sequences (e.g., at least 85%, 90%, 95% or more identical to any of the aforesaid BCMA CAR sequences). The amino acid and nucleotide sequences encoding the BCMA CAR molecules and antigen binding domains (*e.g.*, including one, two, three VH CDRs; and one, two, three VL CDRs according to Kabat or Chothia), are specified in WO2016/014565.

In embodiments, the BCMA CAR comprises an anti-BCMA binding domain (e.g., human or humanized anti-BCMA binding domain), a transmembrane domain, and an intracellular signaling domain, and wherein said anti-BCMA binding domain comprises a heavy chain complementary determining region 1 (HC CDR1), a heavy chain complementary determining region 2 (HC CDR2), and a heavy chain complementary determining region 3 (HC CDR3) of any anti-BMCA heavy chain binding domain amino acid sequences listed in Table 11A or 11B, or a sequence at least 85%, 90%, 95% or more identical thereto (e.g., having less than 5, 4, 3, 2 or 1 amino acid substitutions, e.g., conservative substitutions).

In one embodiment, the anti- BCMA binding domain comprises a light chain variable region described herein (e.g., in Table 11A or 11B) and/or a heavy chain variable region described herein (e.g., in Table 11A or 11B), or a sequence at least 85%, 90%, 95% or more identical thereto.

In one embodiment, the encoded anti- BCMA binding domain is a scFv comprising a light chain and a heavy chain of an amino acid sequence of Table 11A or 11B.

In an embodiment, the human or humanized anti-BCMA binding domain (e.g., an scFv) comprises: a light chain variable region comprising an amino acid sequence having at least one, two or three modifications (e.g., substitutions, e.g., conservative substitutions) but not more than 30, 20 or 10 modifications (e.g., substitutions, e.g., conservative substitutions) of an amino acid sequence of a light chain variable region provided in Table 11A or 11B, or a sequence at least 85%, 90%, 95% or more identical thereto; and/or a heavy chain variable region comprising an amino acid sequence having at least one, two or three modifications (e.g., substitutions, e.g., conservative substitutions) but not more than 30, 20 or 10 modifications (e.g., substitutions, e.g., conservative substitutions) of an amino acid sequence of a heavy chain variable region provided in Table 11A or 11B, or a sequence at least 85%, 90%, 95% or more identical thereto.

Table 11A. Amino Acid and Nucleic Acid Sequences of exemplary anti-BCMA scFv domains and BCMA CAR molecules

Name/ Description	SEQ ID NO:	Sequence
139109		
139109- aa **ScFv domain**	967	EVQLVESGGGLVQPGGSLRLSCAVSGFALSNHGMSWVRRAPGKGLEWV SGIVYSGSTYYAASVKGRFTISRDNSRNTLYLQMNSLRPEDTAIYYCS AHGGESDVWGQGTTVTVSSASGGGGSGGRASGGGGSDIQLTQSPSSLS ASVGDRVTITCRASQSISSYLNWYQQKPGKAPKLLIYAASSLQSGVPS RFSGSGSGTDFTLTISSLQPEDFATYYCQQSYSTPYTFGQGTKVEIK
139109- nt **ScFv domain**	968	GAAGTGCAATTGGTGGAATCAGGGGGAGGACTTGTGCAGCCTGGAGGA TCGCTGAGACTGTCATGTGCCGTGTCCGGCTTTGCCCTGTCCAACCAC GGGATGTCCTGGGTCCGCCGCGCGCCTGGAAAGGGCCTCGAATGGGTG TCGGGTATTGTGTACAGCGGTAGCACCTACTATGCCGCATCCGTGAAG GGGAGATTCACCATCAGCCGGGACAACTCCAGGAACACTCTGTACCTC CAAATGAATTCGCTGAGGCCAGAGGACACTGCCATCTACTACTGCTCC GCGCATGGCGGAGAGTCCGACGTCTGGGGACAGGGGACCACCGTGACC GTGTCTAGCGCGTCCGGCGGAGGCGGCAGCGGGGGTCGGGCATCAGGG GGCGGCGGATCGGACATCCAGCTCACCCAGTCCCCGAGCTCGCTGTCC GCCTCCGTGGGAGATCGGGTCACCATCACGTGCCGCGCCAGCCAGTCG ATTTCCTCCTACCTGAACTGGTACCAACAGAAGCCCGGAAAAGCCCCG AAGCTTCTCATCTACGCCGCCTCGAGCCTGCAGTCAGGAGTGCCCTCA CGGTTCTCCGGCTCCGGTTCCGGTACTGATTTCACCCTGACCATTTCC TCCCTGCAACCGGAGGACTTCGCTACTTACTACTGCCAGCAGTCGTAC TCCACCCCCTACACTTTCGGACAAGGCACCAAGGTCGAAATCAAG
139109- aa **VH**	969	EVQLVESGGGLVQPGGSLRLSCAVSGFALSNHGMSWVRRAPGKGLEWV SGIVYSGSTYYAASVKGRFTISRDNSRNTLYLQMNSLRPEDTAIYYCS AHGGESDVWGQGTTVTVSS
139109- aa **VL**	970	DIQLTQSPSSLSASVGDRVTITCRASQSISSYLNWYQQKPGKAPKLLI YAASSLQSGVPSRFSGSGSGTDFTLTISSLQPEDFATYYCQQSYSTPY TFGQGTKVEIK
139109- aa **Full CAR**	971	MALPVTALLLPLALLLHAARPEVQLVESGGGLVQPGGSLRLSCAVSGF ALSNHGMSWVRRAPGKGLEWVSGIVYSGSTYYAASVKGRFTISRDNSR NTLYLQMNSLRPEDTAIYYCSAHGGESDVWGQGTTVTVSSASGGGGSG GRASGGGGSDIQLTQSPSSLSASVGDRVTITCRASQSISSYLNWYQQK PGKAPKLLIYAASSLQSGVPSRFSGSGSGTDFTLTISSLQPEDFATYY CQQSYSTPYTFGQGTKVEIKTTTPAPRPPTPAPTIASQPLSLRPEACR PAAGGAVHTRGLDFACDIYIWAPLAGTCGVLLLSLVITLYCKRGRKKL LYIFKQPFMRPVQTTQEEDGCSCRFPEEEEGGCELRVKFSRSADAPAY KQGQNQLYNELNLGRREEYDVLDKRRGRDPEMGGKPRRKNPQEGLYNE LQKDKMAEAYSEIGMKGERRRGKGHDGLYQGLSTATKDTYDALHMQAL PPR

139109- nt **Full CAR**	972	ATGGCCCTCCCTGTCACCGCCCTGCTGCTTCCGCTGGCTCTTCTGCTC CACGCCGCTCGGCCCGAAGTGCAATTGGTGGAATCAGGGGGAGGACTT GTGCAGCCTGGAGGATCGCTGAGACTGTCATGTGCCGTGTCCGGCTTT GCCCTGTCCAACCACGGGATGTCCTGGGTCCGCCGCGCGCCTGGAAAG GGCCTCGAATGGGTGTCGGGTATTGTGTACAGCGGTAGCACCTACTAT GCCGCATCCGTGAAGGGGAGATTCACCATCAGCCGGGACAACTCCAGG AACACTCTGTACCTCCAAATGAATTCGCTGAGGCCAGAGGACACTGCC ATCTACTACTGCTCCGCGCATGGCGGAGAGTCCGACGTCTGGGGACAG GGGACCACCGTGACCGTGTCTAGCGCGTCCGGCGGAGGCGGCAGCGGG GGTCGGGCATCAGGGGGCGGCGGATCGGACATCCAGCTCACCCAGTCC CCGAGCTCGCTGTCCGCCTCCGTGGGAGATCGGGTCACCATCACGTGC CGCGCCAGCCAGTCGATTTCCTCCTACCTGAACTGGTACCAACAGAAG CCCGGAAAAGCCCCGAAGCTTCTCATCTACGCCGCCTCGAGCCTGCAG TCAGGAGTGCCCTCACGGTTCTCCGGCTCCGGTTCCGGTACTGATTTC ACCCTGACCATTTCCTCCCTGCAACCGGAGGACTTCGCTACTTACTAC TGCCAGCAGTCGTACTCCACCCCCTACACTTTCGGACAAGGCACCAAG GTCGAAATCAAGACCACTACCCCAGCACCGAGGCCACCCACCCCGGCT CCTACCATCGCCTCCCAGCCTCTGTCCCTGCGTCCGGAGGCATGTAGA CCCGCAGCTGGTGGGGCCGTGCATACCCGGGGTCTTGACTTCGCCTGC GATATCTACATTTGGGCCCCTCTGGCTGGTACTTGCGGGGTCCTGCTG CTTTCACTCGTGATCACTCTTTACTGTAAGCGCGGTCGGAAGAAGCTG CTGTACATCTTTAAGCAACCCTTCATGAGGCCTGTGCAGACTACTCAA GAGGAGGACGGCTGTTCATGCCGGTTCCCAGAGGAGGAGGAAGGCGGC TGCGAACTGCGCGTGAAATTCAGCCGCAGCGCAGATGCTCCAGCCTAC AAGCAGGGGCAGAACCAGCTCTACAACGAACTCAATCTTGGTCGGAGA GAGGAGTACGACGTGCTGGACAAGCGGAGAGGACGGGACCCAGAAATG GGCGGGAAGCCGCGCAGAAAGAATCCCCAAGAGGGCCTGTACAACGAG CTCCAAAAGGATAAGATGGCAGAAGCCTATAGCGAGATTGGTATGAAA GGGGAACGCAGAAGAGGCAAAGGCCACGACGGACTGTACCAGGGACTC AGCACCGCCACCAAGGACACCTATGACGCTCTTCACATGCAGGCCCTG CCGCCTCGG
139103		
139103- aa **ScFv domain**	973	QVQLVESGGGLVQPGRSLRLSCAASGFTFSNYAMSWVRQAPGKGLGWV SGISRSGENTYYADSVKGRFTISRDNSKNTLYLQMNSLRDEDTAVYYC ARSPAHYYGGMDVWGQGTTVTVSSASGGGGSGGRASGGGGSDIVLTQS PGTLSLSPGERATLSCRASQSISSSFLAWYQQKPGQAPRLLIYGASRR ATGIPDRFSGSGSGTDFTLTISRLEPEDSAVYYCQQYHSSPSWTFGQG TKLEIK
139103- nt **ScFv domain**	974	CAAGTGCAACTCGTGGAATCTGGTGGAGGACTCGTGCAACCCGGAAGA TCGCTTAGACTGTCGTGTGCCGCCAGCGGGTTCACTTTCTCGAACTAC GCGATGTCCTGGGTCCGCCAGGCACCCGGAAAGGGACTCGGTTGGGTG TCCGGCATTTCCCGGTCCGGCGAAAATACCTACTACGCCGACTCCGTG AAGGGCCGCTTCACCATCTCAAGGGACAACAGCAAAAACACCCTGTAC TTGCAAATGAACTCCCTGCGGGATGAAGATACAGCCGTGTACTATTGC GCCCGGTCGCCTGCCCATTACTACGGCGGAATGGACGTCTGGGGACAG

		GGAACCACTGTGACTGTCAGCAGCGCGTCGGGTGGCGGCGGCTCAGGG GGTCGGGCCTCCGGGGGGGGAGGGTCCGACATCGTGCTGACCCAGTCC CCGGGAACCCTGAGCCTGAGCCCGGGAGAGCGCGCGACCCTGTCATGC CGGGCATCCCAGAGCATTAGCTCCTCCTTTCTCGCCTGGTATCAGCAG AAGCCCGGACAGGCCCCGAGGCTGCTGATCTACGGCGCTAGCAGAAGG GCTACCGGAATCCCAGACCGGTTCTCCGGCTCCGGTTCCGGGACCGAT TTCACCCTTACTATCTCGCGCCTGGAACCTGAGGACTCCGCCGTCTAC TACTGCCAGCAGTACCACTCATCCCCGTCGTGGACGTTCGGACAGGGC ACCAAGCTGGAGATTAAG
139103- aa **VH**	975	QVQLVESGGGLVQPGRSLRLSCAASGFTFSNYAMSWVRQAPGKGLGWV SGISRSGENTYYADSVKGRFTISRDNSKNTLYLQMNSLRDEDTAVYYC ARSPAHYYGGMDVWGQGTTVTVSS
139103- aa **VL**	976	DIVLTQSPGTLSLSPGERATLSCRASQSISSSFLAWYQQKPGQAPRLL IYGASRRATGIPDRFSGSGSGTDFTLTISRLEPEDSAVYYCQQYHSSP SWTFGQGTKLEIK
139103- aa **Full CAR**	977	MALPVTALLLPLALLLHAARPQVQLVESGGGLVQPGRSLRLSCAASGF TFSNYAMSWVRQAPGKGLGWVSGISRSGENTYYADSVKGRFTISRDNS KNTLYLQMNSLRDEDTAVYYCARSPAHYYGGMDVWGQGTTVTVSSASG GGGSGGRASGGGGSDIVLTQSPGTLSLSPGERATLSCRASQSISSSFL AWYQQKPGQAPRLLIYGASRRATGIPDRFSGSGSGTDFTLTISRLEPE DSAVYYCQQYHSSPSWTFGQGTKLEIKTTTPAPRPPTPAPTIASQPLS LRPEACRPAAGGAVHTRGLDFACDIYIWAPLAGTCGVLLLSLVITLYC KRGRKKLLYIFKQPFMRPVQTTQEEDGCSCRFPEEEEGGCELRVKFSR SADAPAYKQGQNQLYNELNLGRREEYDVLDKRRGRDPEMGGKPRRKNP QEGLYNELQKDKMAEAYSEIGMKGERRRGKGHDGLYQGLSTATKDTYD ALHMQALPPR
139103- nt **Full CAR**	978	ATGGCCCTCCCTGTCACCGCCCTGCTGCTTCCGCTGGCTCTTCTGCTC CACGCCGCTCGGCCCCAAGTGCAACTCGTGGAATCTGGTGGAGGACTC GTGCAACCCGGAAGATCGCTTAGACTGTCGTGTGCCGCCAGCGGGTTC ACTTTCTCGAACTACGCGATGTCCTGGGTCCGCCAGGCACCCGGAAAG GGACTCGGTTGGGTGTCCGGCATTTCCCGGTCCGGCGAAAATACCTAC TACGCCGACTCCGTGAAGGGCCGCTTCACCATCTCAAGGGACAACAGC AAAAACACCCTGTACTTGCAAATGAACTCCCTGCGGGATGAAGATACA GCCGTGTACTATTGCGCCCGGTCGCCTGCCCATTACTACGGCGGAATG GACGTCTGGGGACAGGGAACCACTGTGACTGTCAGCAGCGCGTCGGGT GGCGGCGGCTCAGGGGGTCGGGCCTCCGGGGGGGGAGGGTCCGACATC GTGCTGACCCAGTCCCCGGGAACCCTGAGCCTGAGCCCGGGAGAGCGC GCGACCCTGTCATGCCGGGCATCCCAGAGCATTAGCTCCTCCTTTCTC GCCTGGTATCAGCAGAAGCCCGGACAGGCCCCGAGGCTGCTGATCTAC GGCGCTAGCAGAAGGGCTACCGGAATCCCAGACCGGTTCTCCGGCTCC GGTTCCGGGACCGATTTCACCCTTACTATCTCGCGCCTGGAACCTGAG GACTCCGCCGTCTACTACTGCCAGCAGTACCACTCATCCCCGTCGTGG ACGTTCGGACAGGGCACCAAGCTGGAGATTAAGACCACTACCCCAGCA CCGAGGCCACCCACCCCGGCTCCTACCATCGCCTCCCAGCCTCTGTCC

		CTGCGTCCGGAGGCATGTAGACCCGCAGCTGGTGGGGCCGTGCATACC CGGGGTCTTGACTTCGCCTGCGATATCTACATTTGGGCCCCTCTGGCT GGTACTTGCGGGGTCCTGCTGCTTTCACTCGTGATCACTCTTTACTGT AAGCGCGGTCGGAAGAAGCTGCTGTACATCTTTAAGCAACCCTTCATG AGGCCTGTGCAGACTACTCAAGAGGAGGACGGCTGTTCATGCCGGTTC CCAGAGGAGGAGGAAGGCGGCTGCGAACTGCGCGTGAAATTCAGCCGC AGCGCAGATGCTCCAGCCTACAAGCAGGGGCAGAACCAGCTCTACAAC GAACTCAATCTTGGTCGGAGAGAGGAGTACGACGTGCTGGACAAGCGG AGAGGACGGGACCCAGAAATGGGCGGGAAGCCGCGCAGAAAGAATCCC CAAGAGGGCCTGTACAACGAGCTCCAAAAGGATAAGATGGCAGAAGCC TATAGCGAGATTGGTATGAAAGGGGAACGCAGAAGAGGCAAAGGCCAC GACGGACTGTACCAGGGACTCAGCACCGCCACCAAGGACACCTATGAC GCTCTTCACATGCAGGCCCTGCCGCCTCGG
139105		
139105- aa **ScFv domain**	979	QVQLVESGGGLVQPGRSLRLSCAASGFTFDDYAMHWVRQAPGKGLEWV SGISWNSGSIGYADSVKGRFTISRDNAKNSLYLQMNSLRAEDTALYYC SVHSFLAYWGQGTLVTVSSASGGGGSGGRASGGGGSDIVMTQTPLSLP VTPGEPASISCRSSQSLLHSNGYNYLDWYLQKPGQSPQLLIYLGSNRA SGVPDRFSGSGSGTDFTLKISRVEAEDVGVYYCMQALQTPYTFGQGTK VEIK
139105- nt **ScFv domain**	980	CAAGTGCAACTCGTCGAATCCGGTGGAGGTCTGGTCCAACCTGGTAGA AGCCTGAGACTGTCGTGTGCGGCCAGCGGATTCACCTTTGATGACTAT GCTATGCACTGGGTGCGGCAGGCCCCAGGAAAGGGCCTGGAATGGGTG TCGGGAATTAGCTGGAACTCCGGGTCCATTGGCTACGCCGACTCCGTG AAGGGCCGCTTCACCATCTCCCGCGACAACGCAAAGAACTCCCTGTAC TTGCAAATGAACTCGCTCAGGGCTGAGGATACCGCGCTGTACTACTGC TCCGTGCATTCCTTCCTGGCCTACTGGGGACAGGGAACTCTGGTCACC GTGTCGAGCGCCTCCGGCGGCGGGGGCTCGGGTGGACGGGCCTCGGGC GGAGGGGGGTCCGACATCGTGATGACCCAGACCCCGCTGAGCTTGCCC GTGACTCCCGGAGAGCCTGCATCCATCTCCTGCCGGTCATCCCAGTCC CTTCTCCACTCCAACGGATACAACTACCTCGACTGGTACCTCCAGAAG CCGGGACAGAGCCCTCAGCTTCTGATCTACCTGGGGTCAAATAGAGCC TCAGGAGTGCCGGATCGGTTCAGCGGATCTGGTTCGGGAACTGATTTC ACTCTGAAGATTTCCCGCGTGGAAGCCGAGGACGTGGGCGTCTACTAC TGTATGCAGGCGCTGCAGACCCCCTATACCTTCGGCCAAGGGACGAAA GTGGAGATCAAG
139105- aa **VH**	981	QVQLVESGGGLVQPGRSLRLSCAASGFTFDDYAMHWVRQAPGKGLEWV SGISWNSGSIGYADSVKGRFTISRDNAKNSLYLQMNSLRAEDTALYYC SVHSFLAYWGQGTLVTVSS
139105- aa **VL**	982	DIVMTQTPLSLPVTPGEPASISCRSSQSLLHSNGYNYLDWYLQKPGQS PQLLIYLGSNRASGVPDRFSGSGSGTDFTLKISRVEAEDVGVYYCMQA LQTPYTFGQGTKVEIK

139105- aa **Full CAR**	983	MALPVTALLLPLALLLHAARPQVQLVESGGGLVQPGRSLRLSCAASGF TFDDYAMHWVRQAPGKGLEWVSGISWNSGSIGYADSVKGRFTISRDNA KNSLYLQMNSLRAEDTALYYCSVHSFLAYWGQGTLVTVSSASGGGGSG GRASGGGGSDIVMTQTPLSLPVTPGEPASISCRSSQSLLHSNGYNYLD WYLQKPGQSPQLLIYLGSNRASGVPDRFSGSGSGTDFTLKISRVEAED VGVYYCMQALQTPYTFGQGTKVEIKTTTPAPRPPTPAPTIASQPLSLR PEACRPAAGGAVHTRGLDFACDIYIWAPLAGTCGVLLLSLVITLYCKR GRKKLLYIFKQPFMRPVQTTQEEDGCSCRFPEEEEGGCELRVKFSRSA DAPAYKQGQNQLYNELNLGRREEYDVLDKRRGRDPEMGGKPRRKNPQE GLYNELQKDKMAEAYSEIGMKGERRRGKGHDGLYQGLSTATKDTYDAL HMQALPPR
139105- nt **Full CAR**	984	ATGGCCCTCCCTGTCACCGCCCTGCTGCTTCCGCTGGCTCTTCTGCTC CACGCCGCTCGGCCCCAAGTGCAACTCGTCGAATCCGGTGGAGGTCTG GTCCAACCTGGTAGAAGCCTGAGACTGTCGTGTGCGGCCAGCGGATTC ACCTTTGATGACTATGCTATGCACTGGGTGCGGCAGGCCCCAGGAAAG GGCCTGGAATGGGTGTCGGGAATTAGCTGGAACTCCGGGTCCATTGGC TACGCCGACTCCGTGAAGGGCCGCTTCACCATCTCCCGCGACAACGCA AAGAACTCCCTGTACTTGCAAATGAACTCGCTCAGGGCTGAGGATACC GCGCTGTACTACTGCTCCGTGCATTCCTTCCTGGCCTACTGGGGACAG GGAACTCTGGTCACCGTGTCGAGCGCCTCCGGCGGCGGGGGCTCGGGT GGACGGGCCTCGGGCGGAGGGGGGTCCGACATCGTGATGACCCAGACC CCGCTGAGCTTGCCCGTGACTCCCGGAGAGCCTGCATCCATCTCCTGC CGGTCATCCCAGTCCCTTCTCCACTCCAACGGATACAACTACCTCGAC TGGTACCTCCAGAAGCCGGGACAGAGCCCTCAGCTTCTGATCTACCTG GGGTCAAATAGAGCCTCAGGAGTGCCGGATCGGTTCAGCGGATCTGGT TCGGGAACTGATTTCACTCTGAAGATTTCCCGCGTGGAAGCCGAGGAC GTGGGCGTCTACTACTGTATGCAGGCGCTGCAGACCCCCTATACCTTC GGCCAAGGGACGAAAGTGGAGATCAAGACCACTACCCCAGCACCGAGG CCACCCACCCCGGCTCCTACCATCGCCTCCCAGCCTCTGTCCCTGCGT CCGGAGGCATGTAGACCCGCAGCTGGTGGGGCCGTGCATACCCGGGGT CTTGACTTCGCCTGCGATATCTACATTTGGGCCCCTCTGGCTGGTACT TGCGGGGTCCTGCTGCTTTCACTCGTGATCACTCTTTACTGTAAGCGC GGTCGGAAGAAGCTGCTGTACATCTTTAAGCAACCCTTCATGAGGCCT GTGCAGACTACTCAAGAGGAGGACGGCTGTTCATGCCGGTTCCCAGAG GAGGAGGAAGGCGGCTGCGAACTGCGCGTGAAATTCAGCCGCAGCGCA GATGCTCCAGCCTACAAGCAGGGGCAGAACCAGCTCTACAACGAACTC AATCTTGGTCGGAGAGAGGAGTACGACGTGCTGGACAAGCGGAGAGGA CGGGACCCAGAAATGGGCGGGAAGCCGCGCAGAAAGAATCCCCAAGAG GGCCTGTACAACGAGCTCCAAAAGGATAAGATGGCAGAAGCCTATAGC GAGATTGGTATGAAAGGGGAACGCAGAAGAGGCAAAGGCCACGACGGA CTGTACCAGGGACTCAGCACCGCCACCAAGGACACCTATGACGCTCTT CACATGCAGGCCCTGCCGCCTCGG
139111		
139111- aa	985	EVQLLESGGGLVQPGGSLRLSCAVSGFALSNHGMSWVRRAPGKGLEWV SGIVYSGSTYYAASVKGRFTISRDNSRNTLYLQMNSLRPEDTAIYYCS

ScFv domain		AHGGESDVWGQGTTVTVSSASGGGGSGGRASGGGGSDIVMTQTPLSLS VTPGQPASISCKSSQSLLRNDGKTPLYWYLQKAGQPPQLLIYEVSNRF SGVPDRFSGSGSGTDFTLKISRVEAEDVGAYYCMQNIQFPSFGGGTKL EIK
139111- nt **ScFv domain**	986	GAAGTGCAATTGTTGGAATCTGGAGGAGGACTTGTGCAGCCTGGAGGA TCACTGAGACTTTCGTGTGCGGTGTCAGGCTTCGCCCTGAGCAACCAC GGCATGAGCTGGGTGCGGAGAGCCCCGGGGAAGGGTCTGGAATGGGTG TCCGGGATCGTCTACTCCGGTTCAACTTACTACGCCGCAAGCGTGAAG GGTCGCTTCACCATTTCCCGCGATAACTCCCGGAACACCCTGTACCTC CAAATGAACTCCCTGCGGCCCGAGGACACCGCCATCTACTACTGTTCC GCGCATGGAGGAGAGTCCGATGTCTGGGGACAGGGCACTACCGTGACC GTGTCGAGCGCCTCGGGGGGAGGAGGCTCCGGCGGTCGCGCCTCCGGG GGGGGTGGCAGCGACATTGTGATGACGCAGACTCCACTCTCGCTGTCC GTGACCCCGGGACAGCCCGCGTCCATCTCGTGCAAGAGCTCCCAGAGC CTGCTGAGGAACGACGGAAAGACTCCTCTGTATTGGTACCTCCAGAAG GCTGGACAGCCCCCGCAACTGCTCATCTACGAAGTGTCAAATCGCTTC TCCGGGGTGCCGGATCGGTTTTCCGGCTCGGGATCGGGCACCGACTTC ACCCTGAAAATCTCCAGGGTCGAGGCCGAGGACGTGGGAGCCTACTAC TGCATGCAAAACATCCAGTTCCCTTCCTTCGGCGGCGGCACAAAGCTG GAGATTAAG
139111- aa **VH**	987	EVQLLESGGGLVQPGGSLRLSCAVSGFALSNHGMSWVRRAPGKGLEWV SGIVYSGSTYYAASVKGRFTISRDNSRNTLYLQMNSLRPEDTAIYYCS AHGGESDVWGQGTTVTVSS
139111- aa **VL**	988	DIVMTQTPLSLSVTPGQPASISCKSSQSLLRNDGKTPLYWYLQKAGQP PQLLIYEVSNRFSGVPDRFSGSGSGTDFTLKISRVEAEDVGAYYCMQN IQFPSFGGGTKLEIK
139111- aa **Full CAR**	989	MALPVTALLLPLALLLHAARPEVQLLESGGGLVQPGGSLRLSCAVSGF ALSNHGMSWVRRAPGKGLEWVSGIVYSGSTYYAASVKGRFTISRDNSR NTLYLQMNSLRPEDTAIYYCSAHGGESDVWGQGTTVTVSSASGGGGSG GRASGGGGSDIVMTQTPLSLSVTPGQPASISCKSSQSLLRNDGKTPLY WYLQKAGQPPQLLIYEVSNRFSGVPDRFSGSGSGTDFTLKISRVEAED VGAYYCMQNIQFPSFGGGTKLEIKTTTPAPRPPTPAPTIASQPLSLRP EACRPAAGGAVHTRGLDFACDIYIWAPLAGTCGVLLLSLVITLYCKRG RKKLLYIFKQPFMRPVQTTQEEDGCSCRFPEEEEGGCELRVKFSRSAD APAYKQGQNQLYNELNLGRREEYDVLDKRRGRDPEMGGKPRRKNPQEG LYNELQKDKMAEAYSEIGMKGERRRGKGHDGLYQGLSTATKDTYDALH MQALPPR
139111- nt **Full CAR**	990	ATGGCCCTCCCTGTCACCGCCCTGCTGCTTCCGCTGGCTCTTCTGCTC CACGCCGCTCGGCCCGAAGTGCAATTGTTGGAATCTGGAGGAGGACTT GTGCAGCCTGGAGGATCACTGAGACTTTCGTGTGCGGTGTCAGGCTTC GCCCTGAGCAACCACGGCATGAGCTGGGTGCGGAGAGCCCCGGGGAAG GGTCTGGAATGGGTGTCCGGGATCGTCTACTCCGGTTCAACTTACTAC GCCGCAAGCGTGAAGGGTCGCTTCACCATTTCCCGCGATAACTCCCGG

		AACACCCTGTACCTCCAAATGAACTCCCTGCGGCCCGAGGACACCGCC ATCTACTACTGTTCCGCGCATGGAGGAGAGTCCGATGTCTGGGGACAG GGCACTACCGTGACCGTGTCGAGCGCCTCGGGGGGGAGGAGGCTCCGGC GGTCGCGCCTCCGGGGGGGGTGGCAGCGACATTGTGATGACGCAGACT CCACTCTCGCTGTCCGTGACCCCGGGACAGCCCGCGTCCATCTCGTGC AAGAGCTCCCAGAGCCTGCTGAGGAACGACGGAAAGACTCCTCTGTAT TGGTACCTCCAGAAGGCTGGACAGCCCCCGCAACTGCTCATCTACGAA GTGTCAAATCGCTTCTCCGGGGTGCCGGATCGGTTTTCCGGCTCGGGA TCGGGCACCGACTTCACCCTGAAAATCTCCAGGGTCGAGGCCGAGGAC GTGGGAGCCTACTACTGCATGCAAAACATCCAGTTCCCTTCCTTCGGC GGCGGCACAAAGCTGGAGATTAAGACCACTACCCCAGCACCGAGGCCA CCCACCCCGGCTCCTACCATCGCCTCCCAGCCTCTGTCCCTGCGTCCG GAGGCATGTAGACCCGCAGCTGGTGGGGCCGTGCATACCCGGGGTCTT GACTTCGCCTGCGATATCTACATTTGGGCCCCTCTGGCTGGTACTTGC GGGGTCCTGCTGCTTTCACTCGTGATCACTCTTTACTGTAAGCGCGGT CGGAAGAAGCTGCTGTACATCTTTAAGCAACCCTTCATGAGGCCTGTG CAGACTACTCAAGAGGAGGACGGCTGTTCATGCCGGTTCCCAGAGGAG GAGGAAGGCGGCTGCGAACTGCGCGTGAAATTCAGCCGCAGCGCAGAT GCTCCAGCCTACAAGCAGGGGCAGAACCAGCTCTACAACGAACTCAAT CTTGGTCGGAGAGAGGAGTACGACGTGCTGGACAAGCGGAGAGGACGG GACCCAGAAATGGGCGGGAAGCCGCGCAGAAAGAATCCCCAAGAGGGC CTGTACAACGAGCTCCAAAAGGATAAGATGGCAGAAGCCTATAGCGAG ATTGGTATGAAAGGGGAACGCAGAAGAGGCAAAGGCCACGACGGACTG TACCAGGGACTCAGCACCGCCACCAAGGACACCTATGACGCTCTTCAC ATGCAGGCCCTGCCGCCTCGG
139100		
139100- aa **ScFv domain**	991	QVQLVQSGAEVRKTGASVKVSCKASGYIFDNFGINWVRQAPGQGLEWM GWINPKNNNTNYAQKFQGRVTITADESTNTAYMEVSSLRSEDTAVYYC ARGPYYYQSYMDVWGQGTMVTVSSASGGGGSGGRASGGGGSDIVMTQT PLSLPVTPGEPASISCRSSQSLLHSNGYNYLNWYLQKPGQSPQLLIYL GSKRASGVPDRFSGSGSGTDFTLHITRVGAEDVGVYYCMQALQTPYTF GQGTKLEIK
139100- nt **ScFv domain**	992	CAAGTCCAACTCGTCCAGTCCGGCGCAGAAGTCAGAAAAACCGGTGCT AGCGTGAAAGTGTCCTGCAAGGCCTCCGGCTACATTTTCGATAACTTC GGAATCAACTGGGTCAGACAGGCCCCGGGCCAGGGGCTGGAATGGATG GGATGGATCAACCCCAAGAACAACAACACCAACTACGCACAGAAGTTC CAGGGCCGCGTGACTATCACCGCCGATGAATCGACCAATACCGCCTAC ATGGAGGTGTCCTCCCTGCGGTCGGAGGACACTGCCGTGTATTACTGC GCGAGGGGCCCATACTACTACCAAAGCTACATGGACGTCTGGGGACAG GGAACCATGGTGACCGTGTCATCCGCCTCCGGTGGTGGAGGCTCCGGG GGGCGGGCTTCAGGAGGCGGAGGAAGCGATATTGTGATGACCCAGACT CCGCTTAGCCTGCCCGTGACTCCTGGAGAACCGGCCTCCATTTCCTGC CGGTCCTCGCAATCACTCCTGCATTCCAACGGTTACAACTACCTGAAT TGGTACCTCCAGAAGCCTGGCCAGTCGCCCCAGTTGCTGATCTATCTG GGCTCGAAGCGCGCCTCCGGGGTGCCTGACCGGTTTAGCGGATCTGGG

		AGCGGCACGGACTTCACTCTCCACATCACCCGCGTGGGAGCGGAGGAC GTGGGAGTGTACTACTGTATGCAGGCGCTGCAGACTCCGTACACATTC GGACAGGGCACCAAGCTGGAGATCAAG
139100- aa **VH**	993	QVQLVQSGAEVRKTGASVKVSCKASGYIFDNFGINWVRQAPGQGLEWM GWINPKNNNTNYAQKFQGRVTITADESTNTAYMEVSSLRSEDTAVYYC ARGPYYYQSYMDVWGQGTMVTVSS
139100- aa **VL**	994	DIVMTQTPLSLPVTPGEPASISCRSSQSLLHSNGYNYLNWYLQKPGQS PQLLIYLGSKRASGVPDRFSGSGSGTDFTLHITRVGAEDVGVYYCMQA LQTPYTFGQGTKLEIK
139100- aa **Full CAR**	995	MALPVTALLLPLALLLHAARPQVQLVQSGAEVRKTGASVKVSCKASGY IFDNFGINWVRQAPGQGLEWMGWINPKNNNTNYAQKFQGRVTITADES TNTAYMEVSSLRSEDTAVYYCARGPYYYQSYMDVWGQGTMVTVSSASG GGGSGGRASGGGGSDIVMTQTPLSLPVTPGEPASISCRSSQSLLHSNG YNYLNWYLQKPGQSPQLLIYLGSKRASGVPDRFSGSGSGTDFTLHITR VGAEDVGVYYCMQALQTPYTFGQGTKLEIKTTTPAPRPPTPAPTIASQ PLSLRPEACRPAAGGAVHTRGLDFACDIYIWAPLAGTCGVLLLSLVIT LYCKRGRKKLLYIFKQPFMRPVQTTQEEDGCSCRFPEEEEGGCELRVK FSRSADAPAYKQGQNQLYNELNLGRREEYDVLDKRRGRDPEMGGKPRR KNPQEGLYNELQKDKMAEAYSEIGMKGERRRGKGHDGLYQGLSTATKD TYDALHMQALPPR
139100- nt **Full CAR**	996	ATGGCCCTCCCTGTCACCGCCCTGCTGCTTCCGCTGGCTCTTCTGCTC CACGCCGCTCGGCCCCAAGTCCAACTCGTCCAGTCCGGCGCAGAAGTC AGAAAAACCGGTGCTAGCGTGAAAGTGTCCTGCAAGGCCTCCGGCTAC ATTTTCGATAACTTCGGAATCAACTGGGTCAGACAGGCCCCGGGCCAG GGGCTGGAATGGATGGGATGGATCAACCCCAAGAACAACAACACCAAC TACGCACAGAAGTTCCAGGGCCGCGTGACTATCACCGCCGATGAATCG ACCAATACCGCCTACATGGAGGTGTCCTCCCTGCGGTCGGAGGACACT GCCGTGTATTACTGCGCGAGGGGCCCATACTACTACCAAAGCTACATG GACGTCTGGGGACAGGGAACCATGGTGACCGTGTCATCCGCCTCCGGT GGTGGAGGCTCCGGGGGGCGGGCTTCAGGAGGCGGAGGAAGCGATATT GTGATGACCCAGACTCCGCTTAGCCTGCCCGTGACTCCTGGAGAACCG GCCTCCATTTCCTGCCGGTCCTCGCAATCACTCCTGCATTCCAACGGT TACAACTACCTGAATTGGTACCTCCAGAAGCCTGGCCAGTCGCCCCAG TTGCTGATCTATCTGGGCTCGAAGCGCGCCTCCGGGGTGCCTGACCGG TTTAGCGGATCTGGGAGCGGCACGGACTTCACTCTCCACATCACCCGC GTGGGAGCGGAGGACGTGGGAGTGTACTACTGTATGCAGGCGCTGCAG ACTCCGTACACATTCGGACAGGGCACCAAGCTGGAGATCAAGACCACT ACCCCAGCACCGAGGCCACCCACCCCGGCTCCTACCATCGCCTCCCAG CCTCTGTCCCTGCGTCCGGAGGCATGTAGACCCGCAGCTGGTGGGGCC GTGCATACCCGGGGTCTTGACTTCGCCTGCGATATCTACATTTGGGCC CCTCTGGCTGGTACTTGCGGGGTCCTGCTGCTTTCACTCGTGATCACT CTTTACTGTAAGCGCGGTCGGAAGAAGCTGCTGTACATCTTTAAGCAA CCCTTCATGAGGCCTGTGCAGACTACTCAAGAGGAGGACGGCTGTTCA TGCCGGTTCCCAGAGGAGGAGGAAGGCGGCTGCGAACTGCGCGTGAAA

		TTCAGCCGCAGCGCAGATGCTCCAGCCTACAAGCAGGGGCAGAACCAG CTCTACAACGAACTCAATCTTGGTCGGAGAGAGGAGTACGACGTGCTG GACAAGCGGAGAGGACGGGACCCAGAAATGGGCGGGAAGCCGCGCAGA AAGAATCCCCAAGAGGGCCTGTACAACGAGCTCCAAAAGGATAAGATG GCAGAAGCCTATAGCGAGATTGGTATGAAAGGGGAACGCAGAAGAGGC AAAGGCCACGACGGACTGTACCAGGGACTCAGCACCGCCACCAAGGAC ACCTATGACGCTCTTCACATGCAGGCCCTGCCGCCTCGG
139101		
139101- aa **ScFv domain**	997	QVQLQESGGGLVQPGGSLRLSCAASGFTFSSDAMTWVRQAPGKGLEWV SVISGSGGTTYYADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYC AKLDSSGYYYARGPRYWGQGTLVTVSSASGGGGSGGRASGGGGSDIQL TQSPSSLSASVGDRVTITCRASQSISSYLNWYQQKPGKAPKLLIYGAS TLASGVPARFSGSGSGTHFTLTINSLQSEDSATYYCQQSYKRASFGQG TKVEIK
139101- nt **ScFv domain**	998	CAAGTGCAACTTCAAGAATCAGGCGGAGGACTCGTGCAGCCCGGAGGA TCATTGCGGCTCTCGTGCGCCGCCTCGGGCTTCACCTTCTCGAGCGAC GCCATGACCTGGGTCCGCCAGGCCCCGGGGAAGGGGCTGGAATGGGTG TCTGTGATTTCCGGCTCCGGGGGAACTACGTACTACGCCGATTCCGTG AAAGGTCGCTTCACTATCTCCCGGGACAACAGCAAGAACACCCTTTAT CTGCAAATGAATTCCCTCCGCGCCGAGGACACCGCCGTGTACTACTGC GCCAAGCTGGACTCCTCGGGCTACTACTATGCCCGGGGTCCGAGATAC TGGGGACAGGGAACCCTCGTGACCGTGTCCTCCGCGTCCGGCGGAGGA GGGTCGGGAGGGCGGGCCTCCGGCGGCGGCGGTTCGGACATCCAGCTG ACCCAGTCCCCATCCTCACTGAGCGCAAGCGTGGGCGACAGAGTCACC ATTACATGCAGGGCGTCCCAGAGCATCAGCTCCTACCTGAACTGGTAC CAACAGAAGCCTGGAAAGGCTCCTAAGCTGTTGATCTACGGGGCTTCG ACCCTGGCATCCGGGGTGCCCGCGAGGTTTAGCGGAAGCGGTAGCGGC ACTCACTTCACTCTGACCATTAACAGCCTCCAGTCCGAGGATTCAGCC ACTTACTACTGTCAGCAGTCCTACAAGCGGGCCAGCTTCGGACAGGGC ACTAAGGTCGAGATCAAG
139101- aa **VH**	999	QVQLQESGGGLVQPGGSLRLSCAASGFTFSSDAMTWVRQAPGKGLEWV SVISGSGGTTYYADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYC AKLDSSGYYYARGPRYWGQGTLVTVSS
139101- aa **VL**	1000	DIQLTQSPSSLSASVGDRVTITCRASQSISSYLNWYQQKPGKAPKLLI YGASTLASGVPARFSGSGSGTHFTLTINSLQSEDSATYYCQQSYKRAS FGQGTKVEIK
139101- aa **Full CAR**	1001	MALPVTALLLPLALLLHAARPQVQLQESGGGLVQPGGSLRLSCAASGF TFSSDAMTWVRQAPGKGLEWVSVISGSGGTTYYADSVKGRFTISRDNS KNTLYLQMNSLRAEDTAVYYCAKLDSSGYYYARGPRYWGQGTLVTVSS ASGGGGSGGRASGGGGSDIQLTQSPSSLSASVGDRVTITCRASQSISS YLNWYQQKPGKAPKLLIYGASTLASGVPARFSGSGSGTHFTLTINSLQ SEDSATYYCQQSYKRASFGQGTKVEIKTTTPAPRPPTPAPTIASQPLS

		LRPEACRPAAGGAVHTRGLDFACDIYIWAPLAGTCGVLLLSLVITLYC KRGRKKLLYIFKQPFMRPVQTTQEEDGCSCRFPEEEEGGCELRVKFSR SADAPAYKQGQNQLYNELNLGRREEYDVLDKRRGRDPEMGGKPRRKNP QEGLYNELQKDKMAEAYSEIGMKGERRRGKGHDGLYQGLSTATKDTYD ALHMQALPPR
139101- nt **Full CAR**	1002	ATGGCCCTCCCTGTCACCGCCCTGCTGCTTCCGCTGGCTCTTCTGCTC CACGCCGCTCGGCCCCAAGTGCAACTTCAAGAATCAGGCGGAGGACTC GTGCAGCCCGGAGGATCATTGCGGCTCTCGTGCGCCGCCTCGGGCTTC ACCTTCTCGAGCGACGCCATGACCTGGGTCCGCCAGGCCCCGGGGAAG GGGCTGGAATGGGTGTCTGTGATTTCCGGCTCCGGGGGAACTACGTAC TACGCCGATTCCGTGAAAGGTCGCTTCACTATCTCCCGGGACAACAGC AAGAACACCCTTTATCTGCAAATGAATTCCCTCCGCGCCGAGGACACC GCCGTGTACTACTGCGCCAAGCTGGACTCCTCGGGCTACTACTATGCC CGGGGTCCGAGATACTGGGGACAGGGAACCCTCGTGACCGTGTCCTCC GCGTCCGGCGGAGGAGGGTCGGGAGGGCGGGCCTCCGGCGGCGGCGGT TCGGACATCCAGCTGACCCAGTCCCCATCCTCACTGAGCGCAAGCGTG GGCGACAGAGTCACCATTACATGCAGGGCGTCCCAGAGCATCAGCTCC TACCTGAACTGGTACCAACAGAAGCCTGGAAAGGCTCCTAAGCTGTTG ATCTACGGGGCTTCGACCCTGGCATCCGGGGTGCCCGCGAGGTTTAGC GGAAGCGGTAGCGGCACTCACTTCACTCTGACCATTAACAGCCTCCAG TCCGAGGATTCAGCCACTTACTACTGTCAGCAGTCCTACAAGCGGGCC AGCTTCGGACAGGGCACTAAGGTCGAGATCAAGACCACTACCCCAGCA CCGAGGCCACCCACCCCGGCTCCTACCATCGCCTCCCAGCCTCTGTCC CTGCGTCCGGAGGCATGTAGACCCGCAGCTGGTGGGGCCGTGCATACC CGGGGTCTTGACTTCGCCTGCGATATCTACATTTGGGCCCCTCTGGCT GGTACTTGCGGGGTCCTGCTGCTTTCACTCGTGATCACTCTTTACTGT AAGCGCGGTCGGAAGAAGCTGCTGTACATCTTTAAGCAACCCTTCATG AGGCCTGTGCAGACTACTCAAGAGGAGGACGGCTGTTCATGCCGGTTC CCAGAGGAGGAGGAAGGCGGCTGCGAACTGCGCGTGAAATTCAGCCGC AGCGCAGATGCTCCAGCCTACAAGCAGGGGCAGAACCAGCTCTACAAC GAACTCAATCTTGGTCGGAGAGAGGAGTACGACGTGCTGGACAAGCGG AGAGGACGGGACCCAGAAATGGGCGGGAAGCCGCGCAGAAAGAATCCC CAAGAGGGCCTGTACAACGAGCTCCAAAAGGATAAGATGGCAGAAGCC TATAGCGAGATTGGTATGAAAGGGGAACGCAGAAGAGGCAAAGGCCAC GACGGACTGTACCAGGGACTCAGCACCGCCACCAAGGACACCTATGAC GCTCTTCACATGCAGGCCCTGCCGCCTCGG
139102		
139102- aa **ScFv domain**	1003	QVQLVQSGAEVKKPGASVKVSCKASGYTFSNYGITWVRQAPGQGLEWM GWISAYNGNTNYAQKFQGRVTMTRNTSISTAYMELSSLRSEDTAVYYC ARGPYYYYMDVWGKGTMVTVSSASGGGGSGGRASGGGGSEIVMTQSPL SLPVTPGEPASISCRSSQSLLYSNGYNYVDWYLQKPGQSPQLLIYLGS NRASGVPDRFSGSGSGTDFKLQISRVEAEDVGIYYCMQGRQFPYSFGQ GTKVEIK

139102- nt **ScFv domain**	1004	CAAGTCCAACTGGTCCAGAGCGGTGCAGAAGTGAAGAAGCCCGGAGCG AGCGTGAAAGTGTCCTGCAAGGCTTCCGGGTACACCTTCTCCAACTAC GGCATCACTTGGGTGCGCCAGGCCCCGGGACAGGGCCTGGAATGGATG GGGTGGATTTCCGCGTACAACGGCAATACGAACTACGCTCAGAAGTTC CAGGGTAGAGTGACCATGACTAGGAACACCTCCATTTCCACCGCCTAC ATGGAACTGTCCTCCCTGCGGAGCGAGGACACCGCCGTGTACTATTGC GCCCGGGGACCATACTACTACTACATGGATGTCTGGGGGAAGGGGACT ATGGTCACCGTGTCATCCGCCTCGGGAGGCGGCGGATCAGGAGGACGC GCCTCTGGTGGTGGAGGATCGGAGATCGTGATGACCCAGAGCCCTCTC TCCTTGCCCGTGACTCCTGGGGAGCCCGCATCCATTTCATGCCGGAGC TCCCAGTCACTTCTCTACTCCAACGGCTATAACTACGTGGATTGGTAC CTCCAAAAGCCGGGCCAGAGCCCGCAGCTGCTGATCTACCTGGGCTCG AACAGGGCCAGCGGAGTGCCTGACCGGTTCTCCGGGTCGGGAAGCGGG ACCGACTTCAAGCTGCAAATCTCGAGAGTGGAGGCCGAGGACGTGGGA ATCTACTACTGTATGCAGGGCCGCCAGTTTCCGTACTCGTTCGGACAG GGCACCAAAGTGGAAATCAAG
139102- aa **VH**	1005	QVQLVQSGAEVKKPGASVKVSCKASGYTFSNYGITWVRQAPGQGLEWM GWISAYNGNTNYAQKFQGRVTMTRNTSISTAYMELSSLRSEDTAVYYC ARGPYYYYMDVWGKGTMVTVSS
139102- aa **VL**	1006	EIVMTQSPLSLPVTPGEPASISCRSSQSLLYSNGYNYVDWYLQKPGQS PQLLIYLGSNRASGVPDRFSGSGSGTDFKLQISRVEAEDVGIYYCMQG RQFPYSFGQGTKVEIK
139102- aa **Full CAR**	1007	MALPVTALLLPLALLLHAARPQVQLVQSGAEVKKPGASVKVSCKASGY TFSNYGITWVRQAPGQGLEWMGWISAYNGNTNYAQKFQGRVTMTRNTS ISTAYMELSSLRSEDTAVYYCARGPYYYYMDVWGKGTMVTVSSASGGG GSGGRASGGGGSEIVMTQSPLSLPVTPGEPASISCRSSQSLLYSNGYN YVDWYLQKPGQSPQLLIYLGSNRASGVPDRFSGSGSGTDFKLQISRVE AEDVGIYYCMQGRQFPYSFGQGTKVEIKTTTPAPRPPTPAPTIASQPL SLRPEACRPAAGGAVHTRGLDFACDIYIWAPLAGTCGVLLLSLVITLY CKRGRKKLLYIFKQPFMRPVQTTQEEDGCSCRFPEEEEGGCELRVKFS RSADAPAYKQGQNQLYNELNLGRREEYDVLDKRRGRDPEMGGKPRRKN PQEGLYNELQKDKMAEAYSEIGMKGERRRGKGHDGLYQGLSTATKDTY DALHMQALPPR
139102- nt **Full CAR**	1008	ATGGCCCTCCCTGTCACCGCCCTGCTGCTTCCGCTGGCTCTTCTGCTC CACGCCGCTCGGCCCCAAGTCCAACTGGTCCAGAGCGGTGCAGAAGTG AAGAAGCCCGGAGCGAGCGTGAAAGTGTCCTGCAAGGCTTCCGGGTAC ACCTTCTCCAACTACGGCATCACTTGGGTGCGCCAGGCCCCGGGACAG GGCCTGGAATGGATGGGGTGGATTTCCGCGTACAACGGCAATACGAAC TACGCTCAGAAGTTCCAGGGTAGAGTGACCATGACTAGGAACACCTCC ATTTCCACCGCCTACATGGAACTGTCCTCCCTGCGGAGCGAGGACACC GCCGTGTACTATTGCGCCCGGGGACCATACTACTACTACATGGATGTC TGGGGGAAGGGGACTATGGTCACCGTGTCATCCGCCTCGGGAGGCGGC GGATCAGGAGGACGCGCCTCTGGTGGTGGAGGATCGGAGATCGTGATG ACCCAGAGCCCTCTCTCCTTGCCCGTGACTCCTGGGGAGCCCGCATCC

		ATTTCATGCCGGAGCTCCCAGTCACTTCTCTACTCCAACGGCTATAAC TACGTGGATTGGTACCTCCAAAAGCCGGGCCAGAGCCCGCAGCTGCTG ATCTACCTGGGCTCGAACAGGGCCAGCGGAGTGCCTGACCGGTTCTCC GGGTCGGGAAGCGGGACCGACTTCAAGCTGCAAATCTCGAGAGTGGAG GCCGAGGACGTGGGAATCTACTACTGTATGCAGGGCCGCCAGTTTCCG TACTCGTTCGGACAGGGCACCAAAGTGGAAATCAAGACCACTACCCCA GCACCGAGGCCACCCACCCCGGCTCCTACCATCGCCTCCCAGCCTCTG TCCCTGCGTCCGGAGGCATGTAGACCCGCAGCTGGTGGGGCCGTGCAT ACCCGGGGTCTTGACTTCGCCTGCGATATCTACATTTGGGCCCCTCTG GCTGGTACTTGCGGGGTCCTGCTGCTTTCACTCGTGATCACTCTTTAC TGTAAGCGCGGTCGGAAGAAGCTGCTGTACATCTTTAAGCAACCCTTC ATGAGGCCTGTGCAGACTACTCAAGAGGAGGACGGCTGTTCATGCCGG TTCCCAGAGGAGGAGGAAGGCGGCTGCGAACTGCGCGTGAAATTCAGC CGCAGCGCAGATGCTCCAGCCTACAAGCAGGGGCAGAACCAGCTCTAC AACGAACTCAATCTTGGTCGGAGAGAGGAGTACGACGTGCTGGACAAG CGGAGAGGACGGGACCCAGAAATGGGCGGGAAGCCGCGCAGAAAGAAT CCCCAAGAGGGCCTGTACAACGAGCTCCAAAAGGATAAGATGGCAGAA GCCTATAGCGAGATTGGTATGAAAGGGGAACGCAGAAGAGGCAAAGGC CACGACGGACTGTACCAGGGACTCAGCACCGCCACCAAGGACACCTAT GACGCTCTTCACATGCAGGCCCTGCCGCCTCGG
139104		
139104- aa **ScFv domain**	1009	EVQLLETGGGLVQPGGSLRLSCAVSGFALSNHGMSWVRRAPGKGLEWV SGIVYSGSTYYAASVKGRFTISRDNSRNTLYLQMNSLRPEDTAIYYCS AHGGESDVWGQGTTVTVSSASGGGGSGGRASGGGGSEIVLTQSPATLS VSPGESATLSCRASQSVSSNLAWYQQKPGQAPRLLIYGASTRASGIPD RFSGSGSGTDFTLTISSLQAEDVAVYYCQQYGSSLTFGGGTKVEIK
139104- nt **ScFv domain**	1010	GAAGTGCAATTGCTCGAAACTGGAGGAGGTCTGGTGCAACCTGGAGGA TCACTTCGCCTGTCCTGCGCCGTGTCGGGCTTTGCCCTGTCCAACCAT GGAATGAGCTGGGTCCGCCGCGCGCCGGGGAAGGGCCTCGAATGGGTG TCCGGCATCGTCTACTCCGGCTCCACCTACTACGCCGCGTCCGTGAAG GGCCGGTTCACGATTTCACGGGACAACTCGCGGAACACCCTGTACCTC CAAATGAATTCCCTTCGGCCGGAGGATACTGCCATCTACTACTGCTCC GCCCACGGTGGCGAATCCGACGTCTGGGGCCAGGGAACCACCGTGACC GTGTCCAGCGCGTCCGGGGGAGGAGGAAGCGGGGGTAGAGCATCGGGT GGAGGCGGATCAGAGATCGTGCTGACCCAGTCCCCCGCCACCTTGAGC GTGTCACCAGGAGAGTCCGCCACCCTGTCATGCCGCGCCAGCCAGTCC GTGTCCTCCAACCTGGCTTGGTACCAGCAGAAGCCGGGGCAGGCCCCT AGACTCCTGATCTATGGGGCGTCGACCCGGGCATCTGGAATTCCCGAT AGGTTCAGCGGATCGGGCTCGGGCACTGACTTCACTCTGACCATCTCC TCGCTGCAAGCCGAGGACGTGGCTGTGTACTACTGTCAGCAGTACGGA AGCTCCCTGACTTTCGGTGGCGGGACCAAAGTCGAGATTAAG
139104- aa	1011	EVQLLETGGGLVQPGGSLRLSCAVSGFALSNHGMSWVRRAPGKGLEWV SGIVYSGSTYYAASVKGRFTISRDNSRNTLYLQMNSLRPEDTAIYYCS

VH		AHGGESDVWGQGTTVTVSS
139104- aa **VL**	1012	EIVLTQSPATLSVSPGESATLSCRASQSVSSNLAWYQQKPGQAPRLLI YGASTRASGIPDRFSGSGSGTDFTLTISSLQAEDVAVYYCQQYGSSLT FGGGTKVEIK
139104- aa **Full CAR**	1013	MALPVTALLLPLALLLHAARPEVQLLETGGGLVQPGGSLRLSCAVSGF ALSNHGMSWVRRAPGKGLEWVSGIVYSGSTYYAASVKGRFTISRDNSR NTLYLQMNSLRPEDTAIYYCSAHGGESDVWGQGTTVTVSSASGGGGSG GRASGGGGSEIVLTQSPATLSVSPGESATLSCRASQSVSSNLAWYQQK PGQAPRLLIYGASTRASGIPDRFSGSGSGTDFTLTISSLQAEDVAVYY CQQYGSSLTFGGGTKVEIKTTTPAPRPPTPAPTIASQPLSLRPEACRP AAGGAVHTRGLDFACDIYIWAPLAGTCGVLLLSLVITLYCKRGRKKLL YIFKQPFMRPVQTTQEEDGCSCRFPEEEEGGCELRVKFSRSADAPAYK QGQNQLYNELNLGRREEYDVLDKRRGRDPEMGGKPRRKNPQEGLYNEL QKDKMAEAYSEIGMKGERRRGKGHDGLYQGLSTATKDTYDALHMQALP PR
139104- nt **Full CAR**	1014	ATGGCCCTCCCTGTCACCGCCCTGCTGCTTCCGCTGGCTCTTCTGCTC CACGCCGCTCGGCCCGAAGTGCAATTGCTCGAAACTGGAGGAGGTCTG GTGCAACCTGGAGGATCACTTCGCCTGTCCTGCGCCGTGTCGGGCTTT GCCCTGTCCAACCATGGAATGAGCTGGGTCCGCCGCGCGCCGGGGAAG GGCCTCGAATGGGTGTCCGGCATCGTCTACTCCGGCTCCACCTACTAC GCCGCGTCCGTGAAGGGCCGGTTCACGATTTCACGGGACAACTCGCGG AACACCCTGTACCTCCAAATGAATTCCCTTCGGCCGGAGGATACTGCC ATCTACTACTGCTCCGCCCACGGTGGCGAATCCGACGTCTGGGGCCAG GGAACCACCGTGACCGTGTCCAGCGCGTCCGGGGGAGGAGGAAGCGGG GGTAGAGCATCGGGTGGAGGCGGATCAGAGATCGTGCTGACCCAGTCC CCCGCCACCTTGAGCGTGTCACCAGGAGAGTCCGCCACCCTGTCATGC CGCGCCAGCCAGTCCGTGTCCTCCAACCTGGCTTGGTACCAGCAGAAG CCGGGGCAGGCCCCTAGACTCCTGATCTATGGGGCGTCGACCCGGGCA TCTGGAATTCCCGATAGGTTCAGCGGATCGGGCTCGGGCACTGACTTC ACTCTGACCATCTCCTCGCTGCAAGCCGAGGACGTGGCTGTGTACTAC TGTCAGCAGTACGGAAGCTCCCTGACTTTCGGTGGCGGGACCAAAGTC GAGATTAAGACCACTACCCCAGCACCGAGGCCACCCACCCCGGCTCCT ACCATCGCCTCCCAGCCTCTGTCCCTGCGTCCGGAGGCATGTAGACCC GCAGCTGGTGGGGCCGTGCATACCCGGGGTCTTGACTTCGCCTGCGAT ATCTACATTTGGGCCCCTCTGGCTGGTACTTGCGGGGTCCTGCTGCTT TCACTCGTGATCACTCTTTACTGTAAGCGCGGTCGGAAGAAGCTGCTG TACATCTTTAAGCAACCCTTCATGAGGCCTGTGCAGACTACTCAAGAG GAGGACGGCTGTTCATGCCGGTTCCCAGAGGAGGAGGAAGGCGGCTGC GAACTGCGCGTGAAATTCAGCCGCAGCGCAGATGCTCCAGCCTACAAG CAGGGGCAGAACCAGCTCTACAACGAACTCAATCTTGGTCGGAGAGAG GAGTACGACGTGCTGGACAAGCGGAGAGGACGGGACCCAGAAATGGGC GGGAAGCCGCGCAGAAAGAATCCCCAAGAGGGCCTGTACAACGAGCTC CAAAAGGATAAGATGGCAGAAGCCTATAGCGAGATTGGTATGAAAGGG GAACGCAGAAGAGGCAAAGGCCACGACGGACTGTACCAGGGACTCAGC ACCGCCACCAAGGACACCTATGACGCTCTTCACATGCAGGCCCTGCCG

		CCTCGG
139106		
139106- aa **ScFv domain**	1015	EVQLVETGGGLVQPGGSLRLSCAVSGFALSNHGMSWVRRAPGKGLEWV SGIVYSGSTYYAASVKGRFTISRDNSRNTLYLQMNSLRPEDTAIYYCS AHGGESDVWGQGTTVTVSSASGGGGSGGRASGGGGSEIVMTQSPATLS VSPGERATLSCRASQSVSSKLAWYQQKPGQAPRLLMYGASIRATGIPD RFSGSGSGTEFTLTISSLEPEDFAVYYCQQYGSSSWTFGQGTKVEIK
139106- nt **ScFv domain**	1016	GAAGTGCAATTGGTGGAAACTGGAGGAGGACTTGTGCAACCTGGAGGA TCATTGAGACTGAGCTGCGCAGTGTCGGGATTCGCCCTGAGCAACCAT GGAATGTCCTGGGTCAGAAGGGCCCCTGGAAAAGGCCTCGAATGGGTG TCAGGGATCGTGTACTCCGGTTCCACTTACTACGCCGCCTCCGTGAAG GGGCGCTTCACTATCTCACGGGATAACTCCCGCAATACCCTGTACCTC CAAATGAACAGCCTGCGGCCGGAGGATACCGCCATCTACTACTGTTCC GCCCACGGTGGAGAGTCTGACGTCTGGGGCCAGGGAACTACCGTGACC GTGTCCTCCGCGTCCGGCGGTGGAGGGAGCGGCGGCCGCGCCAGCGGC GGCGGAGGCTCCGAGATCGTGATGACCCAGAGCCCCGCTACTCTGTCG GTGTCGCCCGGAGAAAGGGCGACCCTGTCCTGCCGGGCGTCGCAGTCC GTGAGCAGCAAGCTGGCTTGGTACCAGCAGAAGCCGGGCCAGGCACCA CGCCTGCTTATGTACGGTGCCTCCATTCGGGCCACCGGAATCCCGGAC CGGTTCTCGGGGTCGGGGTCCGGTACCGAGTTCACACTGACCATTTCC TCGCTCGAGCCCGAGGACTTTGCCGTCTATTACTGCCAGCAGTACGGC TCCTCCTCATGGACGTTCGGCCAGGGGACCAAGGTCGAAATCAAG
139106- aa **VH**	1017	EVQLVETGGGLVQPGGSLRLSCAVSGFALSNHGMSWVRRAPGKGLEWV SGIVYSGSTYYAASVKGRFTISRDNSRNTLYLQMNSLRPEDTAIYYCS AHGGESDVWGQGTTVTVSS
139106- aa **VL**	1018	EIVMTQSPATLSVSPGERATLSCRASQSVSSKLAWYQQKPGQAPRLLM YGASIRATGIPDRFSGSGSGTEFTLTISSLEPEDFAVYYCQQYGSSSW TFGQGTKVEIK
139106- aa **Full CAR**	1019	MALPVTALLLPLALLLHAARPEVQLVETGGGLVQPGGSLRLSCAVSGF ALSNHGMSWVRRAPGKGLEWVSGIVYSGSTYYAASVKGRFTISRDNSR NTLYLQMNSLRPEDTAIYYCSAHGGESDVWGQGTTVTVSSASGGGGSG GRASGGGGSEIVMTQSPATLSVSPGERATLSCRASQSVSSKLAWYQQK PGQAPRLLMYGASIRATGIPDRFSGSGSGTEFTLTISSLEPEDFAVYY CQQYGSSSWTFGQGTKVEIKTTTPAPRPPTPAPTIASQPLSLRPEACR PAAGGAVHTRGLDFACDIYIWAPLAGTCGVLLLSLVITLYCKRGRKKL LYIFKQPFMRPVQTTQEEDGCSCRFPEEEEGGCELRVKFSRSADAPAY KQGQNQLYNELNLGRREEYDVLDKRRGRDPEMGGKPRRKNPQEGLYNE LQKDKMAEAYSEIGMKGERRRGKGHDGLYQGLSTATKDTYDALHMQAL PPR
139106- nt	1020	ATGGCCCTCCCTGTCACCGCCCTGCTGCTTCCGCTGGCTCTTCTGCTC CACGCCGCTCGGCCCGAAGTGCAATTGGTGGAAACTGGAGGAGGACTT GTGCAACCTGGAGGATCATTGAGACTGAGCTGCGCAGTGTCGGGATTC

Full CAR		GCCCTGAGCAACCATGGAATGTCCTGGGTCAGAAGGGCCCCTGGAAAA GGCCTCGAATGGGTGTCAGGGATCGTGTACTCCGGTTCCACTTACTAC GCCGCCTCCGTGAAGGGGCGCTTCACTATCTCACGGGATAACTCCCGC AATACCCTGTACCTCCAAATGAACAGCCTGCGGCCGGAGGATACCGCC ATCTACTACTGTTCCGCCCACGGTGGAGAGTCTGACGTCTGGGGCCAG GGAACTACCGTGACCGTGTCCTCCGCGTCCGGCGGTGGAGGGAGCGGC GGCCGCGCCAGCGGCGGCGGAGGCTCCGAGATCGTGATGACCCAGAGC CCCGCTACTCTGTCGGTGTCGCCCGGAGAAAGGGCGACCCTGTCCTGC CGGGCGTCGCAGTCCGTGAGCAGCAAGCTGGCTTGGTACCAGCAGAAG CCGGGCCAGGCACCACGCCTGCTTATGTACGGTGCCTCCATTCGGGCC ACCGGAATCCCGGACCGGTTCTCGGGGTCGGGGTCCGGTACCGAGTTC ACACTGACCATTTCCTCGCTCGAGCCCGAGGACTTTGCCGTCTATTAC TGCCAGCAGTACGGCTCCTCCTCATGGACGTTCGGCCAGGGGACCAAG GTCGAAATCAAGACCACTACCCCAGCACCGAGGCCACCCACCCCGGCT CCTACCATCGCCTCCCAGCCTCTGTCCCTGCGTCCGGAGGCATGTAGA CCCGCAGCTGGTGGGGCCGTGCATACCCGGGGTCTTGACTTCGCCTGC GATATCTACATTTGGGCCCCTCTGGCTGGTACTTGCGGGGTCCTGCTG CTTTCACTCGTGATCACTCTTTACTGTAAGCGCGGTCGGAAGAAGCTG CTGTACATCTTTAAGCAACCCTTCATGAGGCCTGTGCAGACTACTCAA GAGGAGGACGGCTGTTCATGCCGGTTCCCAGAGGAGGAGGAAGGCGGC TGCGAACTGCGCGTGAAATTCAGCCGCAGCGCAGATGCTCCAGCCTAC AAGCAGGGGCAGAACCAGCTCTACAACGAACTCAATCTTGGTCGGAGA GAGGAGTACGACGTGCTGGACAAGCGGAGAGGACGGGACCCAGAAATG GGCGGGAAGCCGCGCAGAAAGAATCCCCAAGAGGGCCTGTACAACGAG CTCCAAAAGGATAAGATGGCAGAAGCCTATAGCGAGATTGGTATGAAA GGGGAACGCAGAAGAGGCAAAGGCCACGACGGACTGTACCAGGGACTC AGCACCGCCACCAAGGACACCTATGACGCTCTTCACATGCAGGCCCTG CCGCCTCGG
139107		
139107- aa **ScFv domain**	1021	EVQLVETGGGVVQPGGSLRLSCAVSGFALSNHGMSWVRRAPGKGLEWV SGIVYSGSTYYAASVKGRFTISRDNSRNTLYLQMNSLRPEDTAIYYCS AHGGESDVWGQGTTVTVSSASGGGGSGGRASGGGGSEIVLTQSPGTLS LSPGERATLSCRASQSVGSTNLAWYQQKPGQAPRLLIYDASNRATGIP DRFSGGGSGTDFTLTISRLEPEDFAVYYCQQYGSSPPWTFGQGTKVEI K
139107- nt **ScFv domain**	1022	GAAGTGCAATTGGTGGAGACTGGAGGAGGAGTGGTGCAACCTGGAGGA AGCCTGAGACTGTCATGCGCGGTGTCGGGCTTCGCCCTCTCCAACCAC GGAATGTCCTGGGTCCGCCGGGCCCCTGGGAAAGGACTTGAATGGGTG TCCGGCATCGTGTACTCGGGTTCCACCTACTACGCGGCCTCAGTGAAG GGCCGGTTTACTATTAGCCGCGACAACTCCAGAAACACACTGTACCTC CAAATGAACTCGCTGCGGCCGGAAGATACCGCTATCTACTACTGCTCC GCCCATGGGGGAGAGTCGGACGTCTGGGGACAGGGCACCACTGTCACT GTGTCCAGCGCTTCCGGCGGTGGTGGAAGCGGGGGACGGGCCTCAGGA GGCGGTGGCAGCGAGATTGTGCTGACCCAGTCCCCCGGGACCCTGAGC CTGTCCCCGGGAGAAAGGGCCACCCTCTCCTGTCGGGCATCCCAGTCC

		GTGGGGTCTACTAACCTTGCATGGTACCAGCAGAAGCCCGGCCAGGCC CCTCGCCTGCTGATCTACGACGCGTCCAATAGAGCCACCGGCATCCCG GATCGCTTCAGCGGAGGCGGATCGGGCACCGACTTCACCCTCACCATT TCAAGGCTGGAACCGGAGGACTTCGCCGTGTACTACTGCCAGCAGTAT GGTTCGTCCCCACCCTGGACGTTCGGCCAGGGGACTAAGGTCGAGATC AAG
139107- aa **VH**	1023	EVQLVETGGGVVQPGGSLRLSCAVSGFALSNHGMSWVRRAPGKGLEWV SGIVYSGSTYYAASVKGRFTISRDNSRNTLYLQMNSLRPEDTAIYYCS AHGGESDVWGQGTTVTVSS
139107- aa **VL**	1024	EIVLTQSPGTLSLSPGERATLSCRASQSVGSTNLAWYQQKPGQAPRLL IYDASNRATGIPDRFSGGGSGTDFTLTISRLEPEDFAVYYCQQYGSSP PWTFGQGTKVEIK
139107- aa **Full CAR**	1025	MALPVTALLLPLALLLHAARPEVQLVETGGGVVQPGGSLRLSCAVSGF ALSNHGMSWVRRAPGKGLEWVSGIVYSGSTYYAASVKGRFTISRDNSR NTLYLQMNSLRPEDTAIYYCSAHGGESDVWGQGTTVTVSSASGGGGSG GRASGGGGSEIVLTQSPGTLSLSPGERATLSCRASQSVGSTNLAWYQQ KPGQAPRLLIYDASNRATGIPDRFSGGGSGTDFTLTISRLEPEDFAVY YCQQYGSSPPWTFGQGTKVEIKTTTPAPRPPTPAPTIASQPLSLRPEA CRPAAGGAVHTRGLDFACDIYIWAPLAGTCGVLLLSLVITLYCKRGRK KLLYIFKQPFMRPVQTTQEEDGCSCRFPEEEEGGCELRVKFSRSADAP AYKQGQNQLYNELNLGRREEYDVLDKRRGRDPEMGGKPRRKNPQEGLY NELQKDKMAEAYSEIGMKGERRRGKGHDGLYQGLSTATKDTYDALHMQ ALPPR
139107- nt **Full CAR**	1026	ATGGCCCTCCCTGTCACCGCCCTGCTGCTTCCGCTGGCTCTTCTGCTC CACGCCGCTCGGCCCGAAGTGCAATTGGTGGAGACTGGAGGAGGAGTG GTGCAACCTGGAGGAAGCCTGAGACTGTCATGCGCGGTGTCGGGCTTC GCCCTCTCCAACCACGGAATGTCCTGGGTCCGCCGGGCCCCTGGGAAA GGACTTGAATGGGTGTCCGGCATCGTGTACTCGGGTTCCACCTACTAC GCGGCCTCAGTGAAGGGCCGGTTTACTATTAGCCGCGACAACTCCAGA AACACACTGTACCTCCAAATGAACTCGCTGCGGCCGGAAGATACCGCT ATCTACTACTGCTCCGCCCATGGGGGAGAGTCGGACGTCTGGGGACAG GGCACCACTGTCACTGTGTCCAGCGCTTCCGGCGGTGGTGGAAGCGGG GGACGGGCCTCAGGAGGCGGTGGCAGCGAGATTGTGCTGACCCAGTCC CCCGGGACCCTGAGCCTGTCCCCGGGAGAAAGGGCCACCCTCTCCTGT CGGGCATCCCAGTCCGTGGGGTCTACTAACCTTGCATGGTACCAGCAG AAGCCCGGCCAGGCCCCTCGCCTGCTGATCTACGACGCGTCCAATAGA GCCACCGGCATCCCGGATCGCTTCAGCGGAGGCGGATCGGGCACCGAC TTCACCCTCACCATTTCAAGGCTGGAACCGGAGGACTTCGCCGTGTAC TACTGCCAGCAGTATGGTTCGTCCCCACCCTGGACGTTCGGCCAGGGG ACTAAGGTCGAGATCAAGACCACTACCCCAGCACCGAGGCCACCCACC CCGGCTCCTACCATCGCCTCCCAGCCTCTGTCCCTGCGTCCGGAGGCA TGTAGACCCGCAGCTGGTGGGGCCGTGCATACCCGGGGTCTTGACTTC GCCTGCGATATCTACATTTGGGCCCCTCTGGCTGGTACTTGCGGGGTC CTGCTGCTTTCACTCGTGATCACTCTTTACTGTAAGCGCGGTCGGAAG

		AAGCTGCTGTACATCTTTAAGCAACCCTTCATGAGGCCTGTGCAGACT ACTCAAGAGGAGGACGGCTGTTCATGCCGGTTCCCAGAGGAGGAGGAA GGCGGCTGCGAACTGCGCGTGAAATTCAGCCGCAGCGCAGATGCTCCA GCCTACAAGCAGGGGCAGAACCAGCTCTACAACGAACTCAATCTTGGT CGGAGAGAGGAGTACGACGTGCTGGACAAGCGGAGAGGACGGGACCCA GAAATGGGCGGGAAGCCGCGCAGAAAGAATCCCCAAGAGGGCCTGTAC AACGAGCTCCAAAAGGATAAGATGGCAGAAGCCTATAGCGAGATTGGT ATGAAAGGGGAACGCAGAAGAGGCAAAGGCCACGACGGACTGTACCAG GGACTCAGCACCGCCACCAAGGACACCTATGACGCTCTTCACATGCAG GCCCTGCCGCCTCGG
139108		
139108- aa **ScFv domain**	1027	QVQLVESGGGLVKPGGSLRLSCAASGFTFSDYYMSWIRQAPGKGLEWV SYISSSGSTIYYADSVKGRFTISRDNAKNSLYLQMNSLRAEDTAVYYC ARESGDGMDVWGQGTTVTVSSASGGGGSGGRASGGGGSDIQMTQSPSS LSASVGDRVTITCRASQSISSYLNWYQQKPGKAPKLLIYAASSLQSGV PSRFSGSGSGTDFTLTISSLQPEDFATYYCQQSYTLAFGQGTKVDIK
139108- nt **ScFv domain**	1028	CAAGTGCAACTCGTGGAATCTGGTGGAGGACTCGTGAAACCTGGAGGA TCATTGAGACTGTCATGCGCGGCCTCGGGATTCACGTTCTCCGATTAC TACATGAGCTGGATTCGCCAGGCTCCGGGGAAGGGACTGGAATGGGTG TCCTACATTTCCTCATCCGGCTCCACCATCTACTACGCGGACTCCGTG AAGGGGAGATTCACCATTAGCCGCGATAACGCCAAGAACAGCCTGTAC CTTCAGATGAACTCCCTGCGGGCTGAAGATACTGCCGTCTACTACTGC GCAAGGGAGAGCGGAGATGGGATGGACGTCTGGGGACAGGGTACCACT GTGACCGTGTCGTCGGCCTCCGGCGGAGGGGGTTCGGGTGGAAGGGCC AGCGGCGGCGGAGGCAGCGACATCCAGATGACCCAGTCCCCCTCATCG CTGTCCGCCTCCGTGGGCGACCGCGTCACCATCACATGCCGGGCCTCA CAGTCGATCTCCTCCTACCTCAATTGGTATCAGCAGAAGCCCGGAAAG GCCCCTAAGCTTCTGATCTACGCAGCGTCCTCCCTGCAATCCGGGGTC CCATCTCGGTTCTCCGGCTCGGGCAGCGGTACCGACTTCACTCTGACC ATCTCGAGCCTGCAGCCGGAGGACTTCGCCACTTACTACTGTCAGCAA AGCTACACCCTCGCGTTTGGCCAGGGCACCAAAGTGGACATCAAG
139108- aa **VH**	1029	QVQLVESGGGLVKPGGSLRLSCAASGFTFSDYYMSWIRQAPGKGLEWV SYISSSGSTIYYADSVKGRFTISRDNAKNSLYLQMNSLRAEDTAVYYC ARESGDGMDVWGQGTTVTVSS
139108- aa **VL**	1030	DIQMTQSPSSLSASVGDRVTITCRASQSISSYLNWYQQKPGKAPKLLI YAASSLQSGVPSRFSGSGSGTDFTLTISSLQPEDFATYYCQQSYTLAF GQGTKVDIK
139108- aa **Full CAR**	1031	MALPVTALLLPLALLLHAARPQVQLVESGGGLVKPGGSLRLSCAASGF TFSDYYMSWIRQAPGKGLEWVSYISSSGSTIYYADSVKGRFTISRDNA KNSLYLQMNSLRAEDTAVYYCARESGDGMDVWGQGTTVTVSSASGGGG SGGRASGGGGSDIQMTQSPSSLSASVGDRVTITCRASQSISSYLNWYQ QKPGKAPKLLIYAASSLQSGVPSRFSGSGSGTDFTLTISSLQPEDFAT

		YYCQQSYTLAFGQGTKVDIKTTTPAPRPPTPAPTIASQPLSLRPEACR PAAGGAVHTRGLDFACDIYIWAPLAGTCGVLLLSLVITLYCKRGRKKL LYIFKQPFMRPVQTTQEEDGCSCRFPEEEEGGCELRVKFSRSADAPAY KQGQNQLYNELNLGRREEYDVLDKRRGRDPEMGGKPRRKNPQEGLYNE LQKDKMAEAYSEIGMKGERRRGKGHDGLYQGLSTATKDTYDALHMQAL PPR
139108- nt **Full CAR**	1032	ATGGCCCTCCCTGTCACCGCCCTGCTGCTTCCGCTGGCTCTTCTGCTC CACGCCGCTCGGCCCCAAGTGCAACTCGTGGAATCTGGTGGAGGACTC GTGAAACCTGGAGGATCATTGAGACTGTCATGCGCGGCCTCGGGATTC ACGTTCTCCGATTACTACATGAGCTGGATTCGCCAGGCTCCGGGGAAG GGACTGGAATGGGTGTCCTACATTTCCTCATCCGGCTCCACCATCTAC TACGCGGACTCCGTGAAGGGGAGATTCACCATTAGCCGCGATAACGCC AAGAACAGCCTGTACCTTCAGATGAACTCCCTGCGGGCTGAAGATACT GCCGTCTACTACTGCGCAAGGGAGAGCGGAGATGGGATGGACGTCTGG GGACAGGGTACCACTGTGACCGTGTCGTCGGCCTCCGGCGGAGGGGGT TCGGGTGGAAGGGCCAGCGGCGGCGGAGGCAGCGACATCCAGATGACC CAGTCCCCCTCATCGCTGTCCGCCTCCGTGGGCGACCGCGTCACCATC ACATGCCGGGCCTCACAGTCGATCTCCTCCTACCTCAATTGGTATCAG CAGAAGCCCGGAAAGGCCCCTAAGCTTCTGATCTACGCAGCGTCCTCC CTGCAATCCGGGGTCCCATCTCGGTTCTCCGGCTCGGGCAGCGGTACC GACTTCACTCTGACCATCTCGAGCCTGCAGCCGGAGGACTTCGCCACT TACTACTGTCAGCAAAGCTACACCCTCGCGTTTGGCCAGGGCACCAAA GTGGACATCAAGACCACTACCCCAGCACCGAGGCCACCCACCCCGGCT CCTACCATCGCCTCCCAGCCTCTGTCCCTGCGTCCGGAGGCATGTAGA CCCGCAGCTGGTGGGGCCGTGCATACCCGGGGTCTTGACTTCGCCTGC GATATCTACATTTGGGCCCCTCTGGCTGGTACTTGCGGGGTCCTGCTG CTTTCACTCGTGATCACTCTTTACTGTAAGCGCGGTCGGAAGAAGCTG CTGTACATCTTTAAGCAACCCTTCATGAGGCCTGTGCAGACTACTCAA GAGGAGGACGGCTGTTCATGCCGGTTCCCAGAGGAGGAGGAAGGCGGC TGCGAACTGCGCGTGAAATTCAGCCGCAGCGCAGATGCTCCAGCCTAC AAGCAGGGGCAGAACCAGCTCTACAACGAACTCAATCTTGGTCGGAGA GAGGAGTACGACGTGCTGGACAAGCGGAGAGGACGGGACCCAGAAATG GGCGGGAAGCCGCGCAGAAAGAATCCCCAAGAGGGCCTGTACAACGAG CTCCAAAAGGATAAGATGGCAGAAGCCTATAGCGAGATTGGTATGAAA GGGGAACGCAGAAGAGGCAAAGGCCACGACGGACTGTACCAGGGACTC AGCACCGCCACCAAGGACACCTATGACGCTCTTCACATGCAGGCCCTG CCGCCTCGG
139110		
139110- aa **ScFv domain**	1033	QVQLVQSGGGLVKPGGSLRLSCAASGFTFSDYYMSWIRQAPGKGLEWV SYISSSGNTIYYADSVKGRFTISRDNAKNSLYLQMNSLRAEDTAVYYC ARSTMVREDYWGQGTLVTVSSASGGGGSGGRASGGGGSDIVLTQSPLS LPVTLGQPASISCKSSESLVHNSGKTYLNWFHQRPGQSPRRLIYEVSN RDSGVPDRFTGSGSGTDFTLKISRVEAEDVGVYYCMQGTHWPGTFGQG TKLEIK

139110- nt **ScFv domain**	1034	CAAGTGCAACTGGTGCAAAGCGGAGGAGGATTGGTCAAACCCGGAGGA AGCCTGAGACTGTCATGCGCGGCCTCTGGATTCACCTTCTCCGATTAC TACATGTCATGGATCAGACAGGCCCCGGGGAAGGGCCTCGAATGGGTG TCCTACATCTCGTCCTCCGGGAACACCATCTACTACGCCGACAGCGTG AAGGGCCGCTTTACCATTTCCCGCGACAACGCAAAGAACTCGCTGTAC CTTCAGATGAATTCCCTGCGGGCTGAAGATACCGCGGTGTACTATTGC GCCCGGTCCACTATGGTCCGGGAGGACTACTGGGGACAGGGCACACTC GTGACCGTGTCCAGCGCGAGCGGGGGTGGAGGCAGCGGTGGACGCGCC TCCGGCGGCGGCGGTTCAGACATCGTGCTGACTCAGTCGCCCCTGTCG CTGCCGGTCACCCTGGGCCAACCGGCCTCAATTAGCTGCAAGTCCTCG GAGAGCCTGGTGCACAACTCAGGAAAGACTTACCTGAACTGGTTCCAT CAGCGGCCTGGACAGTCCCCACGGAGGCTCATCTATGAAGTGTCCAAC AGGGATTCGGGGGTGCCCGACCGCTTCACTGGCTCCGGGTCCGGCACC GACTTCACCTTGAAAATCTCCAGAGTGGAAGCCGAGGACGTGGGCGTG TACTACTGTATGCAGGGTACCCACTGGCCTGGAACCTTTGGACAAGGA ACTAAGCTCGAGATTAAG
139110- aa **VH**	1035	QVQLVQSGGGLVKPGGSLRLSCAASGFTFSDYYMSWIRQAPGKGLEWV SYISSSGNTIYYADSVKGRFTISRDNAKNSLYLQMNSLRAEDTAVYYC ARSTMVREDYWGQGTLVTVSS
139110- aa **VL**	1036	DIVLTQSPLSLPVTLGQPASISCKSSESLVHNSGKTYLNWFHQRPGQS PRRLIYEVSNRDSGVPDRFTGSGSGTDFTLKISRVEAEDVGVYYCMQG THWPGTFGQGTKLEIK
139110- aa **Full CAR**	1037	MALPVTALLLPLALLLHAARPQVQLVQSGGGLVKPGGSLRLSCAASGF TFSDYYMSWIRQAPGKGLEWVSYISSSGNTIYYADSVKGRFTISRDNA KNSLYLQMNSLRAEDTAVYYCARSTMVREDYWGQGTLVTVSSASGGGG SGGRASGGGGSDIVLTQSPLSLPVTLGQPASISCKSSESLVHNSGKTY LNWFHQRPGQSPRRLIYEVSNRDSGVPDRFTGSGSGTDFTLKISRVEA EDVGVYYCMQGTHWPGTFGQGTKLEIKTTTPAPRPPTPAPTIASQPLS LRPEACRPAAGGAVHTRGLDFACDIYIWAPLAGTCGVLLLSLVITLYC KRGRKKLLYIFKQPFMRPVQTTQEEDGCSCRFPEEEEGGCELRVKFSR SADAPAYKQGQNQLYNELNLGRREEYDVLDKRRGRDPEMGGKPRRKNP QEGLYNELQKDKMAEAYSEIGMKGERRRGKGHDGLYQGLSTATKDTYD ALHMQALPPR
139110- nt **Full CAR**	1038	ATGGCCCTCCCTGTCACCGCCCTGCTGCTTCCGCTGGCTCTTCTGCTC CACGCCGCTCGGCCCCAAGTGCAACTGGTGCAAAGCGGAGGAGGATTG GTCAAACCCGGAGGAAGCCTGAGACTGTCATGCGCGGCCTCTGGATTC ACCTTCTCCGATTACTACATGTCATGGATCAGACAGGCCCCGGGGAAG GGCCTCGAATGGGTGTCCTACATCTCGTCCTCCGGGAACACCATCTAC TACGCCGACAGCGTGAAGGGCCGCTTTACCATTTCCCGCGACAACGCA AAGAACTCGCTGTACCTTCAGATGAATTCCCTGCGGGCTGAAGATACC GCGGTGTACTATTGCGCCCGGTCCACTATGGTCCGGGAGGACTACTGG GGACAGGGCACACTCGTGACCGTGTCCAGCGCGAGCGGGGGTGGAGGC AGCGGTGGACGCGCCTCCGGCGGCGGCGGTTCAGACATCGTGCTGACT CAGTCGCCCCTGTCGCTGCCGGTCACCCTGGGCCAACCGGCCTCAATT

		AGCTGCAAGTCCTCGGAGAGCCTGGTGCACAACTCAGGAAAGACTTAC CTGAACTGGTTCCATCAGCGGCCTGGACAGTCCCCACGGAGGCTCATC TATGAAGTGTCCAACAGGGATTCGGGGGTGCCCGACCGCTTCACTGGC TCCGGGTCCGGCACCGACTTCACCTTGAAAATCTCCAGAGTGGAAGCC GAGGACGTGGGCGTGTACTACTGTATGCAGGGTACCCACTGGCCTGGA ACCTTTGGACAAGGAACTAAGCTCGAGATTAAGACCACTACCCCAGCA CCGAGGCCACCCACCCCGGCTCCTACCATCGCCTCCCAGCCTCTGTCC CTGCGTCCGGAGGCATGTAGACCCGCAGCTGGTGGGGCCGTGCATACC CGGGGTCTTGACTTCGCCTGCGATATCTACATTTGGGCCCCTCTGGCT GGTACTTGCGGGGTCCTGCTGCTTTCACTCGTGATCACTCTTTACTGT AAGCGCGGTCGGAAGAAGCTGCTGTACATCTTTAAGCAACCCTTCATG AGGCCTGTGCAGACTACTCAAGAGGAGGACGGCTGTTCATGCCGGTTC CCAGAGGAGGAGGAAGGCGGCTGCGAACTGCGCGTGAAATTCAGCCGC AGCGCAGATGCTCCAGCCTACAAGCAGGGGCAGAACCAGCTCTACAAC GAACTCAATCTTGGTCGGAGAGAGGAGTACGACGTGCTGGACAAGCGG AGAGGACGGGACCCAGAAATGGGCGGGAAGCCGCGCAGAAAGAATCCC CAAGAGGGCCTGTACAACGAGCTCCAAAAGGATAAGATGGCAGAAGCC TATAGCGAGATTGGTATGAAAGGGGAACGCAGAAGAGGCAAAGGCCAC GACGGACTGTACCAGGGACTCAGCACCGCCACCAAGGACACCTATGAC GCTCTTCACATGCAGGCCCTGCCGCCTCGG
139112		
139112- aa **ScFv domain**	1039	QVQLVESGGGLVQPGGSLRLSCAVSGFALSNHGMSWVRRAPGKGLEWV SGIVYSGSTYYAASVKGRFTISRDNSRNTLYLQMNSLRPEDTAIYYCS AHGGESDVWGQGTTVTVSSASGGGGSGGRASGGGGSDIRLTQSPSPLS ASVGDRVTITCQASEDINKFLNWYHQTPGKAPKLLIYDASTLQTGVPS RFSGSGSGTDFTLTINSLQPEDIGTYYCQQYESLPLTFGGGTKVEIK
139112- nt **ScFv domain**	1040	CAAGTGCAACTCGTGGAATCTGGTGGAGGACTCGTGCAACCCGGTGGA AGCCTTAGGCTGTCGTGCGCCGTCAGCGGGTTTGCTCTGAGCAACCAT GGAATGTCCTGGGTCCGCCGGGCACCGGGAAAAGGGCTGGAATGGGTG TCCGGCATCGTGTACAGCGGGTCAACCTATTACGCCGCGTCCGTGAAG GGCAGATTCACTATCTCAAGAGACAACAGCCGGAACACCCTGTACTTG CAAATGAATTCCCTGCGCCCCGAGGACACCGCCATCTACTACTGCTCC GCCCACGGAGGAGAGTCGGACGTGTGGGGCCAGGGAACGACTGTGACT GTGTCCAGCGCATCAGGAGGGGGTGGTTCGGGCGGCCGGGCCTCGGGG GGAGGAGGTTCCGACATTCGGCTGACCCAGTCCCCGTCCCCACTGTCG GCCTCCGTCGGCGACCGCGTGACCATCACTTGTCAGGCGTCCGAGGAC ATTAACAAGTTCCTGAACTGGTACCACCAGACCCCTGGAAAGGCCCCC AAGCTGCTGATCTACGATGCCTCGACCCTTCAAACTGGAGTGCCTAGC CGGTTCTCCGGGTCCGGCTCCGGCACTGATTTCACTCTGACCATCAAC TCATTGCAGCCGGAAGATATCGGGACCTACTATTGCCAGCAGTACGAA TCCCTCCCGCTCACATTCGGCGGGGGAACCAAGGTCGAGATTAAG
139112- aa	1041	QVQLVESGGGLVQPGGSLRLSCAVSGFALSNHGMSWVRRAPGKGLEWV SGIVYSGSTYYAASVKGRFTISRDNSRNTLYLQMNSLRPEDTAIYYCS

VH		AHGGESDVWGQGTTVTVSS
139112- aa **VL**	1042	DIRLTQSPSPLSASVGDRVTITCQASEDINKFLNWYHQTPGKAPKLLI YDASTLQTGVPSRFSGSGSGTDFTLTINSLQPEDIGTYYCQQYESLPL TFGGGTKVEIK
139112- aa **Full CAR**	1043	MALPVTALLLPLALLLHAARPQVQLVESGGGLVQPGGSLRLSCAVSGF ALSNHGMSWVRRAPGKGLEWVSGIVYSGSTYYAASVKGRFTISRDNSR NTLYLQMNSLRPEDTAIYYCSAHGGESDVWGQGTTVTVSSASGGGGSG GRASGGGGSDIRLTQSPSPLSASVGDRVTITCQASEDINKFLNWYHQT PGKAPKLLIYDASTLQTGVPSRFSGSGSGTDFTLTINSLQPEDIGTYY CQQYESLPLTFGGGTKVEIKTTTPAPRPPTPAPTIASQPLSLRPEACR PAAGGAVHTRGLDFACDIYIWAPLAGTCGVLLLSLVITLYCKRGRKKL LYIFKQPFMRPVQTTQEEDGCSCRFPEEEEGGCELRVKFSRSADAPAY KQGQNQLYNELNLGRREEYDVLDKRRGRDPEMGGKPRRKNPQEGLYNE LQKDKMAEAYSEIGMKGERRRGKGHDGLYQGLSTATKDTYDALHMQAL PPR
139112- nt **Full CAR**	1044	ATGGCCCTCCCTGTCACCGCCCTGCTGCTTCCGCTGGCTCTTCTGCTC CACGCCGCTCGGCCCCAAGTGCAACTCGTGGAATCTGGTGGAGGACTC GTGCAACCCGGTGGAAGCCTTAGGCTGTCGTGCGCCGTCAGCGGGTTT GCTCTGAGCAACCATGGAATGTCCTGGGTCCGCCGGGCACCGGGAAAA GGGCTGGAATGGGTGTCCGGCATCGTGTACAGCGGGTCAACCTATTAC GCCGCGTCCGTGAAGGGCAGATTCACTATCTCAAGAGACAACAGCCGG AACACCCTGTACTTGCAAATGAATTCCCTGCGCCCCGAGGACACCGCC ATCTACTACTGCTCCGCCCACGGAGGAGAGTCGGACGTGTGGGGCCAG GGAACGACTGTGACTGTGTCCAGCGCATCAGGAGGGGGTGGTTCGGGC GGCCGGGCCTCGGGGGGAGGAGGTTCCGACATTCGGCTGACCCAGTCC CCGTCCCCACTGTCGGCCTCCGTCGGCGACCGCGTGACCATCACTTGT CAGGCGTCCGAGGACATTAACAAGTTCCTGAACTGGTACCACCAGACC CCTGGAAAGGCCCCCAAGCTGCTGATCTACGATGCCTCGACCCTTCAA ACTGGAGTGCCTAGCCGGTTCTCCGGGTCCGGCTCCGGCACTGATTTC ACTCTGACCATCAACTCATTGCAGCCGGAAGATATCGGGACCTACTAT TGCCAGCAGTACGAATCCCTCCCGCTCACATTCGGCGGGGGAACCAAG GTCGAGATTAAGACCACTACCCCAGCACCGAGGCCACCCACCCCGGCT CCTACCATCGCCTCCCAGCCTCTGTCCCTGCGTCCGGAGGCATGTAGA CCCGCAGCTGGTGGGGCCGTGCATACCCGGGGTCTTGACTTCGCCTGC GATATCTACATTTGGGCCCCTCTGGCTGGTACTTGCGGGGTCCTGCTG CTTTCACTCGTGATCACTCTTTACTGTAAGCGCGGTCGGAAGAAGCTG CTGTACATCTTTAAGCAACCCTTCATGAGGCCTGTGCAGACTACTCAA GAGGAGGACGGCTGTTCATGCCGGTTCCCAGAGGAGGAGGAAGGCGGC TGCGAACTGCGCGTGAAATTCAGCCGCAGCGCAGATGCTCCAGCCTAC AAGCAGGGGCAGAACCAGCTCTACAACGAACTCAATCTTGGTCGGAGA GAGGAGTACGACGTGCTGGACAAGCGGAGAGGACGGGACCCAGAAATG GGCGGGAAGCCGCGCAGAAAGAATCCCCAAGAGGGCCTGTACAACGAG CTCCAAAAGGATAAGATGGCAGAAGCCTATAGCGAGATTGGTATGAAA GGGGAACGCAGAAGAGGCAAAGGCCACGACGGACTGTACCAGGGACTC AGCACCGCCACCAAGGACACCTATGACGCTCTTCACATGCAGGCCCTG

		CCGCCTCGG
139113		
139113- aa **ScFv domain**	1045	EVQLVETGGGLVQPGGSLRLSCAVSGFALSNHGMSWVRRAPGKGLEWV SGIVYSGSTYYAASVKGRFTISRDNSRNTLYLQMNSLRPEDTAIYYCS AHGGESDVWGQGTTVTVSSASGGGGSGGRASGGGGSETTLTQSPATLS VSPGERATLSCRASQSVGSNLAWYQQKPGQGPRLLIYGASTRATGIPA RFSGSGSGTEFTLTISSLQPEDFAVYYCQQYNDWLPVTFGQGTKVEIK
139113- nt **ScFv domain**	1046	GAAGTGCAATTGGTGGAAACTGGAGGAGGACTTGTGCAACCTGGAGGA TCATTGCGGCTCTCATGCGCTGTCTCCGGCTTCGCCCTGTCAAATCAC GGGATGTCGTGGGTCAGACGGGCCCCGGGAAAGGGTCTGGAATGGGTG TCGGGGATTGTGTACAGCGGCTCCACCTACTACGCCGCTTCGGTCAAG GGCCGCTTCACTATTTCACGGGACAACAGCCGCAACACCCTCTATCTG CAAATGAACTCTCTCCGCCCGGAGGATACCGCCATCTACTACTGCTCC GCACACGGCGGCGAATCCGACGTGTGGGGACAGGGAACCACTGTCACC GTGTCGTCCGCATCCGGTGGCGGAGGATCGGGTGGCCGGGCCTCCGGG GGCGGCGGCAGCGAGACTACCCTGACCCAGTCCCCTGCCACTCTGTCC GTGAGCCCGGGAGAGAGAGCCACCCTTAGCTGCCGGGCCAGCCAGAGC GTGGGCTCCAACCTGGCCTGGTACCAGCAGAAGCCAGGACAGGGTCCC AGGCTGCTGATCTACGGAGCCTCCACTCGCGCGACCGGCATCCCCGCG AGGTTCTCCGGGTCGGGTTCCGGGACCGAGTTCACCCTGACCATCTCC TCCCTCCAACCGGAGGACTTCGCGGTGTACTACTGTCAGCAGTACAAC GATTGGCTGCCCGTGACATTTGGACAGGGGACGAAGGTGGAAATCAAA
139113- aa **VH**	1047	EVQLVETGGGLVQPGGSLRLSCAVSGFALSNHGMSWVRRAPGKGLEWV SGIVYSGSTYYAASVKGRFTISRDNSRNTLYLQMNSLRPEDTAIYYCS AHGGESDVWGQGTTVTVSS
139113- aa **VL**	1048	ETTLTQSPATLSVSPGERATLSCRASQSVGSNLAWYQQKPGQGPRLLI YGASTRATGIPARFSGSGSGTEFTLTISSLQPEDFAVYYCQQYNDWLP VTFGQGTKVEIK
139113- aa **Full CAR**	1049	MALPVTALLLPLALLLHAARPEVQLVETGGGLVQPGGSLRLSCAVSGF ALSNHGMSWVRRAPGKGLEWVSGIVYSGSTYYAASVKGRFTISRDNSR NTLYLQMNSLRPEDTAIYYCSAHGGESDVWGQGTTVTVSSASGGGGSG GRASGGGGSETTLTQSPATLSVSPGERATLSCRASQSVGSNLAWYQQK PGQGPRLLIYGASTRATGIPARFSGSGSGTEFTLTISSLQPEDFAVYY CQQYNDWLPVTFGQGTKVEIKTTTPAPRPPTPAPTIASQPLSLRPEAC RPAAGGAVHTRGLDFACDIYIWAPLAGTCGVLLLSLVITLYCKRGRKK LLYIFKQPFMRPVQTTQEEDGCSCRFPEEEEGGCELRVKFSRSADAPA YKQGQNQLYNELNLGRREEYDVLDKRRGRDPEMGGKPRRKNPQEGLYN ELQKDKMAEAYSEIGMKGERRRGKGHDGLYQGLSTATKDTYDALHMQA LPPR
139113- nt	1050	ATGGCCCTCCCTGTCACCGCCCTGCTGCTTCCGCTGGCTCTTCTGCTC CACGCCGCTCGGCCCGAAGTGCAATTGGTGGAAACTGGAGGAGGACTT GTGCAACCTGGAGGATCATTGCGGCTCTCATGCGCTGTCTCCGGCTTC

Full CAR		GCCCTGTCAAATCACGGGATGTCGTGGGTCAGACGGGCCCCGGGAAAG GGTCTGGAATGGGTGTCGGGGATTGTGTACAGCGGCTCCACCTACTAC GCCGCTTCGGTCAAGGGCCGCTTCACTATTTCACGGGACAACAGCCGC AACACCCTCTATCTGCAAATGAACTCTCTCCGCCCGGAGGATACCGCC ATCTACTACTGCTCCGCACACGGCGGCGAATCCGACGTGTGGGGACAG GGAACCACTGTCACCGTGTCGTCCGCATCCGGTGGCGGAGGATCGGGT GGCCGGGCCTCCGGGGGCGGCGGCAGCGAGACTACCCTGACCCAGTCC CCTGCCACTCTGTCCGTGAGCCCGGGAGAGAGAGCCACCCTTAGCTGC CGGGCCAGCCAGAGCGTGGGCTCCAACCTGGCCTGGTACCAGCAGAAG CCAGGACAGGGTCCCAGGCTGCTGATCTACGGAGCCTCCACTCGCGCG ACCGGCATCCCCGCGAGGTTCTCCGGGTCGGGTTCCGGGACCGAGTTC ACCCTGACCATCTCCTCCCTCCAACCGGAGGACTTCGCGGTGTACTAC TGTCAGCAGTACAACGATTGGCTGCCCGTGACATTTGGACAGGGGACG AAGGTGGAAATCAAAACCACTACCCCAGCACCGAGGCCACCCACCCCG GCTCCTACCATCGCCTCCCAGCCTCTGTCCCTGCGTCCGGAGGCATGT AGACCCGCAGCTGGTGGGGCCGTGCATACCCGGGGTCTTGACTTCGCC TGCGATATCTACATTTGGGCCCCTCTGGCTGGTACTTGCGGGGTCCTG CTGCTTTCACTCGTGATCACTCTTTACTGTAAGCGCGGTCGGAAGAAG CTGCTGTACATCTTTAAGCAACCCTTCATGAGGCCTGTGCAGACTACT CAAGAGGAGGACGGCTGTTCATGCCGGTTCCCAGAGGAGGAGGAAGGC GGCTGCGAACTGCGCGTGAAATTCAGCCGCAGCGCAGATGCTCCAGCC TACAAGCAGGGGCAGAACCAGCTCTACAACGAACTCAATCTTGGTCGG AGAGAGGAGTACGACGTGCTGGACAAGCGGAGAGGACGGGACCCAGAA ATGGGCGGGAAGCCGCGCAGAAAGAATCCCCAAGAGGGCCTGTACAAC GAGCTCCAAAAGGATAAGATGGCAGAAGCCTATAGCGAGATTGGTATG AAAGGGGAACGCAGAAGAGGCAAAGGCCACGACGGACTGTACCAGGGA CTCAGCACCGCCACCAAGGACACCTATGACGCTCTTCACATGCAGGCC CTGCCGCCTCGG
139114		
139114- aa **ScFv domain**	1051	EVQLVESGGGLVQPGGSLRLSCAVSGFALSNHGMSWVRRAPGKGLEWV SGIVYSGSTYYAASVKGRFTISRDNSRNTLYLQMNSLRPEDTAIYYCS AHGGESDVWGQGTTVTVSSASGGGGSGGRASGGGGSEIVLTQSPGTLS LSPGERATLSCRASQSIGSSSLAWYQQKPGQAPRLLMYGASSRASGIP DRFSGSGSGTDFTLTISRLEPEDFAVYYCQQYAGSPPFTFGQGTKVEI K
139114- nt **ScFv domain**	1052	GAAGTGCAATTGGTGGAATCTGGTGGAGGACTTGTGCAACCTGGAGGA TCACTGAGACTGTCATGCGCGGTGTCCGGTTTTGCCCTGAGCAATCAT GGGATGTCGTGGGTCCGGCGCGCCCCCGGAAAGGGTCTGGAATGGGTG TCGGGTATCGTCTACTCCGGGAGCACTTACTACGCCGCGAGCGTGAAG GGCCGCTTCACCATTTCCCGCGATAACTCCCGCAACACCCTGTACTTG CAAATGAACTCGCTCCGGCCTGAGGACACTGCCATCTACTACTGCTCC GCACACGGAGGAGAATCCGACGTGTGGGGCCAGGGAACTACCGTGACC GTCAGCAGCGCCTCCGGCGGCGGGGGCTCAGGCGGACGGGCTAGCGGC GGCGGTGGCTCCGAGATCGTGCTGACCCAGTCGCCTGGCACTCTCTCG CTGAGCCCCGGGGAAAGGGCAACCCTGTCCTGTCGGGCCAGCCAGTCC

		ATTGGATCATCCTCCCTCGCCTGGTATCAGCAGAAACCGGGACAGGCT CCGCGGCTGCTTATGTATGGGGCCAGCTCAAGAGCCTCCGGCATTCCC GACCGGTTCTCCGGGTCCGGTTCCGGCACCGATTTCACCCTGACTATC TCGAGGCTGGAGCCAGAGGACTTCGCCGTGTACTACTGCCAGCAGTAC GCGGGGTCCCCGCCGTTCACGTTCGGACAGGGAACCAAGGTCGAGATC AAG
139114- aa **VH**	1053	EVQLVESGGGLVQPGGSLRLSCAVSGFALSNHGMSWVRRAPGKGLEWV SGIVYSGSTYYAASVKGRFTISRDNSRNTLYLQMNSLRPEDTAIYYCS AHGGESDVWGQGTTVTVSS
139114- aa **VL**	1054	EIVLTQSPGTLSLSPGERATLSCRASQSIGSSSLAWYQQKPGQAPRLL MYGASSRASGIPDRFSGSGSGTDFTLTISRLEPEDFAVYYCQQYAGSP PFTFGQGTKVEIK
139114- aa **Full CAR**	1055	MALPVTALLLPLALLLHAARPEVQLVESGGGLVQPGGSLRLSCAVSGF ALSNHGMSWVRRAPGKGLEWVSGIVYSGSTYYAASVKGRFTISRDNSR NTLYLQMNSLRPEDTAIYYCSAHGGESDVWGQGTTVTVSSASGGGGSG GRASGGGGSEIVLTQSPGTLSLSPGERATLSCRASQSIGSSSLAWYQQ KPGQAPRLLMYGASSRASGIPDRFSGSGSGTDFTLTISRLEPEDFAVY YCQQYAGSPPFTFGQGTKVEIKTTTPAPRPPTPAPTIASQPLSLRPEA CRPAAGGAVHTRGLDFACDIYIWAPLAGTCGVLLLSLVITLYCKRGRK KLLYIFKQPFMRPVQTTQEEDGCSCRFPEEEEGGCELRVKFSRSADAP AYKQGQNQLYNELNLGRREEYDVLDKRRGRDPEMGGKPRRKNPQEGLY NELQKDKMAEAYSEIGMKGERRRGKGHDGLYQGLSTATKDTYDALHMQ ALPPR
139114- nt **Full CAR**	1056	ATGGCCCTCCCTGTCACCGCCCTGCTGCTTCCGCTGGCTCTTCTGCTC CACGCCGCTCGGCCCGAAGTGCAATTGGTGGAATCTGGTGGAGGACTT GTGCAACCTGGAGGATCACTGAGACTGTCATGCGCGGTGTCCGGTTTT GCCCTGAGCAATCATGGGATGTCGTGGGTCCGGCGCGCCCCCGGAAAG GGTCTGGAATGGGTGTCGGGTATCGTCTACTCCGGGAGCACTTACTAC GCCGCGAGCGTGAAGGGCCGCTTCACCATTTCCCGCGATAACTCCCGC AACACCCTGTACTTGCAAATGAACTCGCTCCGGCCTGAGGACACTGCC ATCTACTACTGCTCCGCACACGGAGGAGAATCCGACGTGTGGGGCCAG GGAACTACCGTGACCGTCAGCAGCGCCTCCGGCGGCGGGGGCTCAGGC GGACGGGCTAGCGGCGGCGGTGGCTCCGAGATCGTGCTGACCCAGTCG CCTGGCACTCTCTCGCTGAGCCCCGGGGAAAGGGCAACCCTGTCCTGT CGGGCCAGCCAGTCCATTGGATCATCCTCCCTCGCCTGGTATCAGCAG AAACCGGGACAGGCTCCGCGGCTGCTTATGTATGGGGCCAGCTCAAGA GCCTCCGGCATTCCCGACCGGTTCTCCGGGTCCGGTTCCGGCACCGAT TTCACCCTGACTATCTCGAGGCTGGAGCCAGAGGACTTCGCCGTGTAC TACTGCCAGCAGTACGCGGGGTCCCCGCCGTTCACGTTCGGACAGGGA ACCAAGGTCGAGATCAAGACCACTACCCCAGCACCGAGGCCACCCACC CCGGCTCCTACCATCGCCTCCCAGCCTCTGTCCCTGCGTCCGGAGGCA TGTAGACCCGCAGCTGGTGGGGCCGTGCATACCCGGGGTCTTGACTTC GCCTGCGATATCTACATTTGGGCCCCTCTGGCTGGTACTTGCGGGGTC CTGCTGCTTTCACTCGTGATCACTCTTTACTGTAAGCGCGGTCGGAAG

		AAGCTGCTGTACATCTTTAAGCAACCCTTCATGAGGCCTGTGCAGACT ACTCAAGAGGAGGACGGCTGTTCATGCCGGTTCCCAGAGGAGGAGGAA GGCGGCTGCGAACTGCGCGTGAAATTCAGCCGCAGCGCAGATGCTCCA GCCTACAAGCAGGGGCAGAACCAGCTCTACAACGAACTCAATCTTGGT CGGAGAGAGGAGTACGACGTGCTGGACAAGCGGAGAGGACGGGACCCA GAAATGGGCGGGAAGCCGCGCAGAAAGAATCCCCAAGAGGGCCTGTAC AACGAGCTCCAAAAGGATAAGATGGCAGAAGCCTATAGCGAGATTGGT ATGAAAGGGGAACGCAGAAGAGGCAAAGGCCACGACGGACTGTACCAG GGACTCAGCACCGCCACCAAGGACACCTATGACGCTCTTCACATGCAG GCCCTGCCGCCTCGG
149362		
149362-aa ScFv domain	1057	QVQLQESGPGLVKPSETLSLTCTVSGGSISSSYYYWGWIRQPPGKGLE WIGSIYYSGSAYYNPSLKSRVTISVDTSKNQFSLRLSSVTAADTAVYY CARHWQEWPDAFDIWGQGTMVTVSSGGGGSGGGGSGGGGSETTLTQSP AFMSATPGDKVIISCKASQDIDDAMNWYQQKPGEAPLFIIQSATSPVP GIPPRFSGSGFGTDFSLTINNIESEDAAYYFCLQHDNFPLTFGQGTKL EIK
149362-nt ScFv domain	1058	CAAGTGCAGCTTCAGGAAAGCGGACCGGGCCTGGTCAAGCCATCCGAA ACTCTCTCCCTGACTTGCACTGTGTCTGGCGGTTCCATCTCATCGTCG TACTACTACTGGGGCTGGATTAGGCAGCCGCCCGGAAAGGGACTGGAG TGGATCGGAAGCATCTACTATTCCGGCTCGGCGTACTACAACCCTAGC CTCAAGTCGAGAGTGACCATCTCCGTGGATACCTCCAAGAACCAGTTT TCCCTGCGCCTGAGCTCCGTGACCGCCGCTGACACCGCCGTGTACTAC TGTGCTCGGCATTGGCAGGAATGGCCCGATGCCTTCGACATTTGGGGC CAGGGCACTATGGTCACTGTGTCATCCGGGGGTGGAGGCAGCGGGGGA GGAGGGTCCGGGGGGGGAGGTTCAGAGACAACCTTGACCCAGTCACCC GCATTCATGTCCGCCACTCCGGGAGACAAGGTCATCATCTCGTGCAAA GCGTCCCAGGATATCGACGATGCCATGAATTGGTACCAGCAGAAGCCT GGCGAAGCGCCGCTGTTCATTATCCAATCCGCAACCTCGCCCGTGCCT GGAATCCCACCGCGGTTCAGCGGCAGCGGTTTCGGAACCGACTTTTCC CTGACCATTAACAACATTGAGTCCGAGGACGCCGCCTACTACTTCTGC CTGCAACACGACAACTTCCCTCTCACGTTCGGCCAGGGAACCAAGCTG GAAATCAAG
149362-aa VH	1059	QVQLQESGPGLVKPSETLSLTCTVSGGSISSSYYYWGWIRQPPGKGLE WIGSIYYSGSAYYNPSLKSRVTISVDTSKNQFSLRLSSVTAADTAVYY CARHWQEWPDAFDIWGQGTMVTVSS
149362-aa VL	1060	ETTLTQSPAFMSATPGDKVIISCKASQDIDDAMNWYQQKPGEAPLFII QSATSPVPGIPPRFSGSGFGTDFSLTINNIESEDAAYYFCLQHDNFPL TFGQGTKLEIK
149362-aa Full CAR	1061	MALPVTALLLPLALLLHAARPQVQLQESGPGLVKPSETLSLTCTVSGG SISSSYYYWGWIRQPPGKGLEWIGSIYYSGSAYYNPSLKSRVTISVDT SKNQFSLRLSSVTAADTAVYYCARHWQEWPDAFDIWGQGTMVTVSSGG

		GGSGGGGSGGGGSETTLTQSPAFMSATPGDKVIISCKASQDIDDAMNW YQQKPGEAPLFIIQSATSPVPGIPPRFSGSGFGTDFSLTINNIESEDA AYYFCLQHDNFPLTFGQGTKLEIKTTTPAPRPPTPAPTIASQPLSLRP EACRPAAGGAVHTRGLDFACDIYIWAPLAGTCGVLLLSLVITLYCKRG RKKLLYIFKQPFMRPVQTTQEEDGCSCRFPEEEEGGCELRVKFSRSAD APAYKQGQNQLYNELNLGRREEYDVLDKRRGRDPEMGGKPRRKNPQEG LYNELQKDKMAEAYSEIGMKGERRRGKGHDGLYQGLSTATKDTYDALH MQALPPR
149362-nt **Full CAR**	1062	ATGGCCCTCCCTGTCACCGCCCTGCTGCTTCCGCTGGCTCTTCTGCTC CACGCCGCTCGGCCCCAAGTGCAGCTTCAGGAAAGCGGACCGGGCCTG GTCAAGCCATCCGAAACTCTCTCCCTGACTTGCACTGTGTCTGGCGGT TCCATCTCATCGTCGTACTACTACTGGGGCTGGATTAGGCAGCCGCCC GGAAAGGGACTGGAGTGGATCGGAAGCATCTACTATTCCGGCTCGGCG TACTACAACCCTAGCCTCAAGTCGAGAGTGACCATCTCCGTGGATACC TCCAAGAACCAGTTTTCCCTGCGCCTGAGCTCCGTGACCGCCGCTGAC ACCGCCGTGTACTACTGTGCTCGGCATTGGCAGGAATGGCCCGATGCC TTCGACATTTGGGGCCAGGGCACTATGGTCACTGTGTCATCCGGGGGT GGAGGCAGCGGGGGAGGAGGGTCCGGGGGGGGAGGTTCAGAGACAACC TTGACCCAGTCACCCGCATTCATGTCCGCCACTCCGGGAGACAAGGTC ATCATCTCGTGCAAAGCGTCCCAGGATATCGACGATGCCATGAATTGG TACCAGCAGAAGCCTGGCGAAGCGCCGCTGTTCATTATCCAATCCGCA ACCTCGCCCGTGCCTGGAATCCCACCGCGGTTCAGCGGCAGCGGTTTC GGAACCGACTTTTCCCTGACCATTAACAACATTGAGTCCGAGGACGCC GCCTACTACTTCTGCCTGCAACACGACAACTTCCCTCTCACGTTCGGC CAGGGAACCAAGCTGGAAATCAAGACCACTACCCCAGCACCGAGGCCA CCCACCCCGGCTCCTACCATCGCCTCCCAGCCTCTGTCCCTGCGTCCG GAGGCATGTAGACCCGCAGCTGGTGGGGCCGTGCATACCCGGGGTCTT GACTTCGCCTGCGATATCTACATTTGGGCCCCTCTGGCTGGTACTTGC GGGGTCCTGCTGCTTTCACTCGTGATCACTCTTTACTGTAAGCGCGGT CGGAAGAAGCTGCTGTACATCTTTAAGCAACCCTTCATGAGGCCTGTG CAGACTACTCAAGAGGAGGACGGCTGTTCATGCCGGTTCCCAGAGGAG GAGGAAGGCGGCTGCGAACTGCGCGTGAAATTCAGCCGCAGCGCAGAT GCTCCAGCCTACAAGCAGGGGCAGAACCAGCTCTACAACGAACTCAAT CTTGGTCGGAGAGAGGAGTACGACGTGCTGGACAAGCGGAGAGGACGG GACCCAGAAATGGGCGGGAAGCCGCGCAGAAAGAATCCCCAAGAGGGC CTGTACAACGAGCTCCAAAAGGATAAGATGGCAGAAGCCTATAGCGAG ATTGGTATGAAAGGGGAACGCAGAAGAGGCAAAGGCCACGACGGACTG TACCAGGGACTCAGCACCGCCACCAAGGACACCTATGACGCTCTTCAC ATGCAGGCCCTGCCGCCTCGG
149363		
149363-aa **ScFv domain**	1063	VNLRESGPALVKPTQTLTLTCTFSGFSLRTSGMCVSWIRQPPGKALEW LARIDWDEDKFYSTSLKTRLTISKDTSDNQVVLRMTNMDPADTATYYC ARSGAGGTSATAFDIWGPGTMVTVSSGGGGSGGGGSGGGGSDIQMTQS PSSLSASVGDRVTITCRASQDIYNNLAWFQLKPGSAPRSLMYAANKSQ SGVPSRFSGSASGTDFTLTISSLQPEDFATYYCQHYYRFPYSFGQGTK

		LEIK
149363-nt ScFv domain	1064	CAAGTCAATCTGCGCGAATCCGGCCCCGCCTTGGTCAAGCCTACCCAG ACCCTCACTCTGACCTGTACTTTCTCCGGCTTCTCCCTGCGGACTTCC GGGATGTGCGTGTCCTGGATCAGACAGCCTCCGGGAAAGGCCCTGGAG TGGCTCGCTCGCATTGACTGGGATGAGGACAAGTTCTACTCCACCTCA CTCAAGACCAGGCTGACCATCAGCAAAGATACCTCTGACAACCAAGTG GTGCTCCGCATGACCAACATGGACCCAGCCGACACTGCCACTTACTAC TGCGCGAGGAGCGGAGCGGGCGGAACCTCCGCCACCGCCTTCGATATT TGGGGCCCGGGTACCATGGTCACCGTGTCAAGCGGAGGAGGGGGGTCC GGGGGCGGCGGTTCCGGGGGAGGCGGATCGGACATTCAGATGACTCAG TCACCATCGTCCCTGAGCGCTAGCGTGGGCGACAGAGTGACAATCACT TGCCGGGCATCCCAGGACATCTATAACAACCTTGCGTGGTTCCAGCTG AAGCCTGGTTCCGCACCGCGGTCACTTATGTACGCCGCCAACAAGAGC CAGTCGGGAGTGCCGTCCCGGTTTTCCGGTTCGGCCTCGGGAACTGAC TTCACCCTGACGATCTCCAGCCTGCAACCCGAGGATTTCGCCACCTAC TACTGCCAGCACTACTACCGCTTTCCCTACTCGTTCGGACAGGGAACC AAGCTGGAAATCAAG
149363-aa VH	1065	QVNLRESGPALVKPTQTLTLTCTFSGFSLRTSGMCVSWIRQPPGKALE WLARIDWDEDKFYSTSLKTRLTISKDTSDNQVVLRMTNMDPADTATYY CARSGAGGTSATAFDIWGPGTMVTVSS
149363-aa VL	1066	DIQMTQSPSSLSASVGDRVTITCRASQDIYNNLAWFQLKPGSAPRSLM YAANKSQSGVPSRFSGSASGTDFTLTISSLQPEDFATYYCQHYYRFPY SFGQGTKLEIK
149363-aa Full CAR	1067	MALPVTALLLPLALLLHAARPQVNLRESGPALVKPTQTLTLTCTFSGF SLRTSGMCVSWIRQPPGKALEWLARIDWDEDKFYSTSLKTRLTISKDT SDNQVVLRMTNMDPADTATYYCARSGAGGTSATAFDIWGPGTMVTVSS GGGGSGGGGSGGGGSDIQMTQSPSSLSASVGDRVTITCRASQDIYNNL AWFQLKPGSAPRSLMYAANKSQSGVPSRFSGSASGTDFTLTISSLQPE DFATYYCQHYYRFPYSFGQGTKLEIKTTTPAPRPPTPAPTIASQPLSL RPEACRPAAGGAVHTRGLDFACDIYIWAPLAGTCGVLLLSLVITLYCK RGRKKLLYIFKQPFMRPVQTTQEEDGCSCRFPEEEEGGCELRVKFSRS ADAPAYKQGQNQLYNELNLGRREEYDVLDKRRGRDPEMGGKPRRKNPQ EGLYNELQKDKMAEAYSEIGMKGERRRGKGHDGLYQGLSTATKDTYDA LHMQALPPR
149363-nt Full CAR	1068	ATGGCCCTCCCTGTCACCGCCCTGCTGCTTCCGCTGGCTCTTCTGCTC CACGCCGCTCGGCCCCAAGTCAATCTGCGCGAATCCGGCCCCGCCTTG GTCAAGCCTACCCAGACCCTCACTCTGACCTGTACTTTCTCCGGCTTC TCCCTGCGGACTTCCGGGATGTGCGTGTCCTGGATCAGACAGCCTCCG GGAAAGGCCCTGGAGTGGCTCGCTCGCATTGACTGGGATGAGGACAAG TTCTACTCCACCTCACTCAAGACCAGGCTGACCATCAGCAAAGATACC TCTGACAACCAAGTGGTGCTCCGCATGACCAACATGGACCCAGCCGAC ACTGCCACTTACTACTGCGCGAGGAGCGGAGCGGGCGGAACCTCCGCC ACCGCCTTCGATATTTGGGGCCCGGGTACCATGGTCACCGTGTCAAGC

		GGAGGAGGGGGGTCCGGGGGCGGCGGTTCCGGGGGAGGCGGATCGGAC ATTCAGATGACTCAGTCACCATCGTCCCTGAGCGCTAGCGTGGGCGAC AGAGTGACAATCACTTGCCGGGCATCCCAGGACATCTATAACAACCTT GCGTGGTTCCAGCTGAAGCCTGGTTCCGCACCGCGGTCACTTATGTAC GCCGCCAACAAGAGCCAGTCGGGAGTGCCGTCCCGGTTTTCCGGTTCG GCCTCGGGAACTGACTTCACCCTGACGATCTCCAGCCTGCAACCCGAG GATTTCGCCACCTACTACTGCCAGCACTACTACCGCTTTCCCTACTCG TTCGGACAGGGAACCAAGCTGGAAATCAAGACCACTACCCCAGCACCG AGGCCACCCACCCCGGCTCCTACCATCGCCTCCCAGCCTCTGTCCCTG CGTCCGGAGGCATGTAGACCCGCAGCTGGTGGGGCCGTGCATACCCGG GGTCTTGACTTCGCCTGCGATATCTACATTTGGGCCCCTCTGGCTGGT ACTTGCGGGGTCCTGCTGCTTTCACTCGTGATCACTCTTTACTGTAAG CGCGGTCGGAAGAAGCTGCTGTACATCTTTAAGCAACCCTTCATGAGG CCTGTGCAGACTACTCAAGAGGAGGACGGCTGTTCATGCCGGTTCCCA GAGGAGGAGGAAGGCGGCTGCGAACTGCGCGTGAAATTCAGCCGCAGC GCAGATGCTCCAGCCTACAAGCAGGGGCAGAACCAGCTCTACAACGAA CTCAATCTTGGTCGGAGAGAGGAGTACGACGTGCTGGACAAGCGGAGA GGACGGGACCCAGAAATGGGCGGGAAGCCGCGCAGAAAGAATCCCCAA GAGGGCCTGTACAACGAGCTCCAAAAGGATAAGATGGCAGAAGCCTAT AGCGAGATTGGTATGAAAGGGGAACGCAGAAGAGGCAAAGGCCACGAC GGACTGTACCAGGGACTCAGCACCGCCACCAAGGACACCTATGACGCT CTTCACATGCAGGCCCTGCCGCCTCGG
149364		
149364-aa ScFv domain	1069	EVQLVESGGGLVKPGGSLRLSCAASGFTFSSYSMNWVRQAPGKGLEWV SSISSSSSYIYYADSVKGRFTISRDNAKNSLYLQMNSLRAEDTAVYYC AKTIAAVYAFDIWGQGTTVTVSSGGGGSGGGGSGGGGSEIVLTQSPLS LPVTPEEPASISCRSSQSLLHSNGYNYLDWYLQKPGQSPQLLIYLGSN RASGVPDRFSGSGSGTDFTLKISRVEAEDVGVYYCMQALQTPYTFGQG TKLEIK
149364-nt ScFv domain	1070	GAAGTGCAGCTTGTCGAATCCGGGGGGGGACTGGTCAAGCCGGGCGGA TCACTGAGACTGTCCTGCGCCGCGAGCGGCTTCACGTTCTCCTCCTAC TCCATGAACTGGGTCCGCCAAGCCCCCGGGAAGGGACTGGAATGGGTG TCCTCTATCTCCTCGTCGTCGTCCTACATCTACTACGCCGACTCCGTG AAGGGAAGATTCACCATTTCCCGCGACAACGCAAAGAACTCACTGTAC TTGCAAATGAACTCACTCCGGGCCGAAGATACTGCTGTGTACTATTGC GCCAAGACTATTGCCGCCGTCTACGCTTTCGACATCTGGGGCCAGGGA ACCACCGTGACTGTGTCGTCCGGTGGTGGTGGCTCGGGCGGAGGAGGA AGCGGCGGCGGGGGGTCCGAGATTGTGCTGACCCAGTCGCCACTGAGC CTCCCTGTGACCCCCGAGGAACCCGCCAGCATCAGCTGCCGGTCCAGC CAGTCCCTGCTCCACTCCAACGGATACAATTACCTCGATTGGTACCTT CAGAAGCCTGGACAAAGCCCGCAGCTGCTCATCTACTTGGGATCAAAC CGCGCGTCAGGAGTGCCTGACCGGTTCTCCGGCTCGGGCAGCGGTACC GATTTCACCCTGAAAATCTCCAGGGTGGAGGCAGAGGACGTGGGAGTG TATTACTGTATGCAGGCGCTGCAGACTCCGTACACATTTGGGCAGGGC

		ACCAAGCTGGAGATCAAG
149364-aa VH	1071	EVQLVESGGGLVKPGGSLRLSCAASGFTFSSYSMNWVRQAPGKGLEWV SSISSSSSYIYYADSVKGRFTISRDNAKNSLYLQMNSLRAEDTAVYYC AKTIAAVYAFDIWGQGTTVTVSS
149364-aa VL	1072	EIVLTQSPLSLPVTPEEPASISCRSSQSLLHSNGYNYLDWYLQKPGQS PQLLIYLGSNRASGVPDRFSGSGSGTDFTLKISRVEAEDVGVYYCMQA LQTPYTFGQGTKLEIK
149364-aa Full CAR	1073	MALPVTALLLPLALLLHAARPEVQLVESGGGLVKPGGSLRLSCAASGF TFSSYSMNWVRQAPGKGLEWVSSISSSSSYIYYADSVKGRFTISRDNA KNSLYLQMNSLRAEDTAVYYCAKTIAAVYAFDIWGQGTTVTVSSGGGG SGGGGSGGGGSEIVLTQSPLSLPVTPEEPASISCRSSQSLLHSNGYNY LDWYLQKPGQSPQLLIYLGSNRASGVPDRFSGSGSGTDFTLKISRVEA EDVGVYYCMQALQTPYTFGQGTKLEIKTTTPAPRPPTPAPTIASQPLS LRPEACRPAAGGAVHTRGLDFACDIYIWAPLAGTCGVLLLSLVITLYC KRGRKKLLYIFKQPFMRPVQTTQEEDGCSCRFPEEEEGGCELRVKFSR SADAPAYKQGQNQLYNELNLGRREEYDVLDKRRGRDPEMGGKPRRKNP QEGLYNELQKDKMAEAYSEIGMKGERRRGKGHDGLYQGLSTATKDTYD ALHMQALPPR
149364-nt **Full CAR**	1074	ATGGCCCTCCCTGTCACCGCCCTGCTGCTTCCGCTGGCTCTTCTGCTC CACGCCGCTCGGCCCGAAGTGCAGCTTGTCGAATCCGGGGGGGGACTG GTCAAGCCGGGCGGATCACTGAGACTGTCCTGCGCCGCGAGCGGCTTC ACGTTCTCCTCCTACTCCATGAACTGGGTCCGCCAAGCCCCCGGGAAG GGACTGGAATGGGTGTCCTCTATCTCCTCGTCGTCGTCCTACATCTAC TACGCCGACTCCGTGAAGGGAAGATTCACCATTTCCCGCGACAACGCA AAGAACTCACTGTACTTGCAAATGAACTCACTCCGGGCCGAAGATACT GCTGTGTACTATTGCGCCAAGACTATTGCCGCCGTCTACGCTTTCGAC ATCTGGGGCCAGGGAACCACCGTGACTGTGTCGTCCGGTGGTGGTGGC TCGGGCGGAGGAGGAAGCGGCGGCGGGGGGTCCGAGATTGTGCTGACC CAGTCGCCACTGAGCCTCCCTGTGACCCCCGAGGAACCCGCCAGCATC AGCTGCCGGTCCAGCCAGTCCCTGCTCCACTCCAACGGATACAATTAC CTCGATTGGTACCTTCAGAAGCCTGGACAAAGCCCGCAGCTGCTCATC TACTTGGGATCAAACCGCGCGTCAGGAGTGCCTGACCGGTTCTCCGGC TCGGGCAGCGGTACCGATTTCACCCTGAAAATCTCCAGGGTGGAGGCA GAGGACGTGGGAGTGTATTACTGTATGCAGGCGCTGCAGACTCCGTAC ACATTTGGGCAGGGCACCAAGCTGGAGATCAAGACCACTACCCCAGCA CCGAGGCCACCCACCCCGGCTCCTACCATCGCCTCCCAGCCTCTGTCC CTGCGTCCGGAGGCATGTAGACCCGCAGCTGGTGGGGCCGTGCATACC CGGGGTCTTGACTTCGCCTGCGATATCTACATTTGGGCCCCTCTGGCT GGTACTTGCGGGGTCCTGCTGCTTTCACTCGTGATCACTCTTTACTGT AAGCGCGGTCGGAAGAAGCTGCTGTACATCTTTAAGCAACCCTTCATG AGGCCTGTGCAGACTACTCAAGAGGAGGACGGCTGTTCATGCCGGTTC CCAGAGGAGGAGGAAGGCGGCTGCGAACTGCGCGTGAAATTCAGCCGC AGCGCAGATGCTCCAGCCTACAAGCAGGGGCAGAACCAGCTCTACAAC GAACTCAATCTTGGTCGGAGAGAGGAGTACGACGTGCTGGACAAGCGG

		AGAGGACGGGACCCAGAAATGGGCGGGAAGCCGCGCAGAAAGAATCCC CAAGAGGGCCTGTACAACGAGCTCCAAAAGGATAAGATGGCAGAAGCC TATAGCGAGATTGGTATGAAAGGGGAACGCAGAAGAGGCAAAGGCCAC GACGGACTGTACCAGGGACTCAGCACCGCCACCAAGGACACCTATGAC GCTCTTCACATGCAGGCCCTGCCGCCTCGG
149365		
149365-aa ScFv domain	1075	EVQLVESGGGLVKPGGSLRLSCAASGFTFSDYYMSWIRQAPGKGLEWV SYISSSGSTIYYADSVKGRFTISRDNAKNSLYLQMNSLRAEDTAVYYC ARDLRGAFDIWGQGTMVTVSSGGGGSGGGGSGGGGSSYVLTQSPSVSA APGYTATISCGGNNIGTKSVHWYQQKPGQAPLLVIRDDSVRPSKIPGR FSGSNSGNMATLTISGVQAGDEADFYCQVWDSDSEHVVFGGGTKLTVL
149365-nt ScFv domain	1076	GAAGTCCAGCTCGTGGAGTCCGGCGGAGGCCTTGTGAAGCCTGGAGGT TCGCTGAGACTGTCCTGCGCCGCCTCCGGCTTCACCTTCTCCGACTAC TACATGTCCTGGATCAGACAGGCCCCGGGAAAGGGCCTGGAATGGGTG TCCTACATCTCGTCATCGGGCAGCACTATCTACTACGCGGACTCAGTG AAGGGGCGGTTCACCATTTCCCGGGATAACGCGAAGAACTCGCTGTAT CTGCAAATGAACTCACTGAGGGCCGAGGACACCGCCGTGTACTACTGC GCCCGCGATCTCCGCGGGGCATTTGACATCTGGGGACAGGGAACCATG GTCACAGTGTCCAGCGGAGGGGGAGGATCGGGTGGCGGAGGTTCCGGG GGTGGAGGCTCCTCCTACGTGCTGACTCAGAGCCCAAGCGTCAGCGCT GCGCCCGGTTACACGGCAACCATCTCCTGTGGCGGAAACAACATTGGG ACCAAGTCTGTGCACTGGTATCAGCAGAAGCCGGGCCAAGCTCCCCTG TTGGTGATCCGCGATGACTCCGTGCGGCCTAGCAAAATTCCGGGACGG TTCTCCGGCTCCAACAGCGGCAATATGGCCACTCTCACCATCTCGGGA GTGCAGGCCGGAGATGAAGCCGACTTCTACTGCCAAGTCTGGGACTCA GACTCCGAGCATGTGGTGTTCGGGGGCGGAACCAAGCTGACTGTGCTC
149365-aa VH	1077	EVQLVESGGGLVKPGGSLRLSCAASGFTFSDYYMSWIRQAPGKGLEWV SYISSSGSTIYYADSVKGRFTISRDNAKNSLYLQMNSLRAEDTAVYYC ARDLRGAFDIWGQGTMVTVSS
149365-aa VL	1078	SYVLTQSPSVSAAPGYTATISCGGNNIGTKSVHWYQQKPGQAPLLVIR DDSVRPSKIPGRFSGSNSGNMATLTISGVQAGDEADFYCQVWDSDSEH VVFGGGTKLTVL
149365-aa Full CAR	1079	MALPVTALLLPLALLLHAARPEVQLVESGGGLVKPGGSLRLSCAASGF TFSDYYMSWIRQAPGKGLEWVSYISSSGSTIYYADSVKGRFTISRDNA KNSLYLQMNSLRAEDTAVYYCARDLRGAFDIWGQGTMVTVSSGGGGSG GGGSGGGGSSYVLTQSPSVSAAPGYTATISCGGNNIGTKSVHWYQQKP GQAPLLVIRDDSVRPSKIPGRFSGSNSGNMATLTISGVQAGDEADFYC QVWDSDSEHVVFGGGTKLTVLTTTPAPRPPTPAPTIASQPLSLRPEAC RPAAGGAVHTRGLDFACDIYIWAPLAGTCGVLLLSLVITLYCKRGRKK LLYIFKQPFMRPVQTTQEEDGCSCRFPEEEEGGCELRVKFSRSADAPA YKQGQNQLYNELNLGRREEYDVLDKRRGRDPEMGGKPRRKNPQEGLYN ELQKDKMAEAYSEIGMKGERRRGKGHDGLYQGLSTATKDTYDALHMQA

		LPPR
149365-nt **Full CAR**	1080	ATGGCCCTCCCTGTCACCGCCCTGCTGCTTCCGCTGGCTCTTCTGCTC CACGCCGCTCGGCCCGAAGTCCAGCTCGTGGAGTCCGGCGGAGGCCTT GTGAAGCCTGGAGGTTCGCTGAGACTGTCCTGCGCCGCCTCCGGCTTC ACCTTCTCCGACTACTACATGTCCTGGATCAGACAGGCCCCGGGAAAG GGCCTGGAATGGGTGTCCTACATCTCGTCATCGGGCAGCACTATCTAC TACGCGGACTCAGTGAAGGGGCGGTTCACCATTTCCCGGGATAACGCG AAGAACTCGCTGTATCTGCAAATGAACTCACTGAGGGCCGAGGACACC GCCGTGTACTACTGCGCCCGCGATCTCCGCGGGGCATTTGACATCTGG GGACAGGGAACCATGGTCACAGTGTCCAGCGGAGGGGGAGGATCGGGT GGCGGAGGTTCCGGGGGTGGAGGCTCCTCCTACGTGCTGACTCAGAGC CCAAGCGTCAGCGCTGCGCCCGGTTACACGGCAACCATCTCCTGTGGC GGAAACAACATTGGGACCAAGTCTGTGCACTGGTATCAGCAGAAGCCG GGCCAAGCTCCCCTGTTGGTGATCCGCGATGACTCCGTGCGGCCTAGC AAAATTCCGGGACGGTTCTCCGGCTCCAACAGCGGCAATATGGCCACT CTCACCATCTCGGGAGTGCAGGCCGGAGATGAAGCCGACTTCTACTGC CAAGTCTGGGACTCAGACTCCGAGCATGTGGTGTTCGGGGGCGGAACC AAGCTGACTGTGCTCACCACTACCCCAGCACCGAGGCCACCCACCCCG GCTCCTACCATCGCCTCCCAGCCTCTGTCCCTGCGTCCGGAGGCATGT AGACCCGCAGCTGGTGGGGCCGTGCATACCCGGGGTCTTGACTTCGCC TGCGATATCTACATTTGGGCCCCTCTGGCTGGTACTTGCGGGGTCCTG CTGCTTTCACTCGTGATCACTCTTTACTGTAAGCGCGGTCGGAAGAAG CTGCTGTACATCTTTAAGCAACCCTTCATGAGGCCTGTGCAGACTACT CAAGAGGAGGACGGCTGTTCATGCCGGTTCCCAGAGGAGGAGGAAGGC GGCTGCGAACTGCGCGTGAAATTCAGCCGCAGCGCAGATGCTCCAGCC TACAAGCAGGGGCAGAACCAGCTCTACAACGAACTCAATCTTGGTCGG AGAGAGGAGTACGACGTGCTGGACAAGCGGAGAGGACGGGACCCAGAA ATGGGCGGGAAGCCGCGCAGAAAGAATCCCCAAGAGGGCCTGTACAAC GAGCTCCAAAAGGATAAGATGGCAGAAGCCTATAGCGAGATTGGTATG AAAGGGGAACGCAGAAGAGGCAAAGGCCACGACGGACTGTACCAGGGA CTCAGCACCGCCACCAAGGACACCTATGACGCTCTTCACATGCAGGCC CTGCCGCCTCGG
149366		
149366-aa **ScFv domain**	1081	QVQLVQSGAEVKKPGASVKVSCKPSGYTVTSHYIHWVRRAPGQGLEWM GMINPSGGVTAYSQTLQGRVTMTSDTSSSTVYMELSSLRSEDTAMYYC AREGSGSGWYFDFWGRGTLVTVSSGGGGSGGGGSGGGGSSYVLTQPPS VSVSPGQTASITCSGDGLSKKYVSWYQQKAGQSPVVLISRDKERPSGI PDRFSGSNSADTATLTISGTQAMDEADYYCQAWDDTTVVFGGGTKLTV L
149366-nt **ScFv domain**	1082	CAAGTGCAGCTGGTGCAGAGCGGGGCCGAAGTCAAGAAGCCGGGAGCC TCCGTGAAAGTGTCCTGCAAGCCTTCGGGATACACCGTGACCTCCCAC TACATTCATTGGGTCCGCCGCGCCCCCGGCCAAGGACTCGAGTGGATG GGCATGATCAACCCTAGCGGCGGAGTGACCGCGTACAGCCAGACGCTG CAGGGACGCGTGACTATGACCTCGGATACCTCCTCCTCCACCGTCTAT

		ATGGAACTGTCCAGCCTGCGGTCCGAGGATACCGCCATGTACTACTGC GCCCGGGAAGGATCAGGCTCCGGGTGGTATTTCGACTTCTGGGGAAGA GGCACCCTCGTGACTGTGTCATCTGGGGGAGGGGGTTCCGGTGGTGGC GGATCGGGAGGAGGCGGTTCATCCTACGTGCTGACCCAGCCACCCTCC GTGTCCGTGAGCCCCGGCCAGACTGCATCGATTACATGTAGCGGCGAC GGCCTCTCCAAGAAATACGTGTCGTGGTACCAGCAGAAGGCCGGACAG AGCCCGGTGGTGCTGATCTCAAGAGATAAGGAGCGGCCTAGCGGAATC CCGGACAGGTTCTCGGGTTCCAACTCCGCGGACACTGCTACTCTGACC ATCTCGGGGACCCAGGCTATGGACGAAGCCGATTACTACTGCCAAGCC TGGGACGACACTACTGTCGTGTTTGGAGGGGCACCAAGTTGACCGTC CTT
149366-aa VH	1083	QVQLVQSGAEVKKPGASVKVSCKPSGYTVTSHYIHWVRRAPGQGLEWM GMINPSGGVTAYSQTLQGRVTMTSDTSSSTVYMELSSLRSEDTAMYYC AREGSGSGWYFDFWGRGTLVTVSS
149366-aa VL	1084	SYVLTQPPSVSVSPGQTASITCSGDGLSKKYVSWYQQKAGQSPVVLIS RDKERPSGIPDRFSGSNSADTATLTISGTQAMDEADYYCQAWDDTTVV FGGGTKLTVL
149366-aa Full CAR	1085	MALPVTALLLPLALLLHAARPQVQLVQSGAEVKKPGASVKVSCKPSGY TVTSHYIHWVRRAPGQGLEWMGMINPSGGVTAYSQTLQGRVTMTSDTS SSTVYMELSSLRSEDTAMYYCAREGSGSGWYFDFWGRGTLVTVSSGGG GSGGGGSGGGGSSYVLTQPPSVSVSPGQTASITCSGDGLSKKYVSWYQ QKAGQSPVVLISRDKERPSGIPDRFSGSNSADTATLTISGTQAMDEAD YYCQAWDDTTVVFGGGTKLTVLTTTPAPRPPTPAPTIASQPLSLRPEA CRPAAGGAVHTRGLDFACDIYIWAPLAGTCGVLLLSLVITLYCKRGRK KLLYIFKQPFMRPVQTTQEEDGCSCRFPEEEEGGCELRVKFSRSADAP AYKQGQNQLYNELNLGRREEYDVLDKRRGRDPEMGGKPRRKNPQEGLY NELQKDKMAEAYSEIGMKGERRRGKGHDGLYQGLSTATKDTYDALHMQ ALPPR
149366-nt **Full CAR**	1086	ATGGCCCTCCCTGTCACCGCCCTGCTGCTTCCGCTGGCTCTTCTGCTC CACGCCGCTCGGCCCCAAGTGCAGCTGGTGCAGAGCGGGGCCGAAGTC AAGAAGCCGGGAGCCTCCGTGAAAGTGTCCTGCAAGCCTTCGGGATAC ACCGTGACCTCCCACTACATTCATTGGGTCCGCCGCGCCCCCGGCCAA GGACTCGAGTGGATGGGCATGATCAACCCTAGCGGCGGAGTGACCGCG TACAGCCAGACGCTGCAGGGACGCGTGACTATGACCTCGGATACCTCC TCCTCCACCGTCTATATGGAACTGTCCAGCCTGCGGTCCGAGGATACC GCCATGTACTACTGCGCCCGGGAAGGATCAGGCTCCGGGTGGTATTTC GACTTCTGGGGAAGAGGCACCCTCGTGACTGTGTCATCTGGGGGAGGG GGTTCCGGTGGTGGCGGATCGGGAGGAGGCGGTTCATCCTACGTGCTG ACCCAGCCACCCTCCGTGTCCGTGAGCCCCGGCCAGACTGCATCGATT ACATGTAGCGGCGACGGCCTCTCCAAGAAATACGTGTCGTGGTACCAG CAGAAGGCCGGACAGAGCCCGGTGGTGCTGATCTCAAGAGATAAGGAG CGGCCTAGCGGAATCCCGGACAGGTTCTCGGGTTCCAACTCCGCGGAC ACTGCTACTCTGACCATCTCGGGGACCCAGGCTATGGACGAAGCCGAT TACTACTGCCAAGCCTGGGACGACACTACTGTCGTGTTTGGAGGGGGC

		ACCAAGTTGACCGTCCTTACCACTACCCCAGCACCGAGGCCACCCACC CCGGCTCCTACCATCGCCTCCCAGCCTCTGTCCCTGCGTCCGGAGGCA TGTAGACCCGCAGCTGGTGGGGCCGTGCATACCCGGGGTCTTGACTTC GCCTGCGATATCTACATTTGGGCCCCTCTGGCTGGTACTTGCGGGGTC CTGCTGCTTTCACTCGTGATCACTCTTTACTGTAAGCGCGGTCGGAAG AAGCTGCTGTACATCTTTAAGCAACCCTTCATGAGGCCTGTGCAGACT ACTCAAGAGGAGGACGGCTGTTCATGCCGGTTCCCAGAGGAGGAGGAA GGCGGCTGCGAACTGCGCGTGAAATTCAGCCGCAGCGCAGATGCTCCA GCCTACAAGCAGGGGCAGAACCAGCTCTACAACGAACTCAATCTTGGT CGGAGAGAGGAGTACGACGTGCTGGACAAGCGGAGAGGACGGGACCCA GAAATGGGCGGGAAGCCGCGCAGAAAGAATCCCCAAGAGGGCCTGTAC AACGAGCTCCAAAAGGATAAGATGGCAGAAGCCTATAGCGAGATTGGT ATGAAAGGGGAACGCAGAAGAGGCAAAGGCCACGACGGACTGTACCAG GGACTCAGCACCGCCACCAAGGACACCTATGACGCTCTTCACATGCAG GCCCTGCCGCCTCGG
149367		
149367-aa ScFv domain	1087	QVQLQESGPGLVKPSQTLSLTCTVSGGSISSGGYYWSWIRQHPGKGLE WIGYIYYSGSTYYNPSLKSRVTISVDTSKNQFSLKLSSVTAADTAVYY CARAGIAARLRGAFDIWGQGTMVTVSSGGGGSGGGGSGGGGSDIVMTQ SPSSVSASVGDRVIITCRASQGIRNWLAWYQQKPGKAPNLLIYAASNL QSGVPSRFSGSGSGADFTLTISSLQPEDVATYYCQKYNSAPFTFGPGT KVDIK
149367-nt ScFv domain	1088	CAAGTGCAGCTTCAGGAGAGCGGCCCGGGACTCGTGAAGCCGTCCCAG ACCCTGTCCCTGACTTGCACCGTGTCGGGAGGAAGCATCTCGAGCGGA GGCTACTATTGGTCGTGGATTCGGCAGCACCCTGGAAAGGGCCTGGAA TGGATCGGCTACATCTACTACTCCGGCTCGACCTACTACAACCCATCG CTGAAGTCCAGAGTGACAATCTCAGTGGACACGTCCAAGAATCAGTTC AGCCTGAAGCTCTCTTCCGTGACTGCGGCCGACACCGCCGTGTACTAC TGCGCACGCGCTGGAATTGCCGCCCGGCTGAGGGGTGCCTTCGACATT TGGGGACAGGGCACCATGGTCACCGTGTCCTCCGGCGGCGGAGGTTCC GGGGGTGGAGGCTCAGGAGGAGGGGGGTCCGACATCGTCATGACTCAG TCGCCCTCAAGCGTCAGCGCGTCCGTCGGGGACAGAGTGATCATCACC TGTCGGGCGTCCCAGGGAATTCGCAACTGGCTGGCCTGGTATCAGCAG AAGCCCGGAAAGGCCCCCAACCTGTTGATCTACGCCGCCTCAAACCTC CAATCCGGGGTGCCGAGCCGCTTCAGCGGCTCCGGTTCGGGTGCCGAT TTCACTCTGACCATCTCCTCCCTGCAACCTGAAGATGTGGCTACCTAC TACTGCCAAAAGTACAACTCCGCACCTTTTACTTTCGGACCGGGGACC AAAGTGGACATTAAG
149367-aa VH	1089	QVQLQESGPGLVKPSQTLSLTCTVSGGSISSGGYYWSWIRQHPGKGLE WIGYIYYSGSTYYNPSLKSRVTISVDTSKNQFSLKLSSVTAADTAVYY CARAGIAARLRGAFDIWGQGTMVTVSS
149367-aa	1090	DIVMTQSPSSVSASVGDRVIITCRASQGIRNWLAWYQQKPGKAPNLLI YAASNLQSGVPSRFSGSGSGADFTLTISSLQPEDVATYYCQKYNSAPF

VL		TFGPGTKVDIK
149367-aa Full CAR	1091	MALPVTALLLPLALLLHAARPQVQLQESGPGLVKPSQTLSLTCTVSGG SISSGGYYWSWIRQHPGKGLEWIGYIYYSGSTYYNPSLKSRVTISVDT SKNQFSLKLSSVTAADTAVYYCARAGIAARLRGAFDIWGQGTMVTVSS GGGGSGGGGSGGGGSDIVMTQSPSSVSASVGDRVIITCRASQGIRNWL AWYQQKPGKAPNLLIYAASNLQSGVPSRFSGSGSGADFTLTISSLQPE DVATYYCQKYNSAPFTFGPGTKVDIKTTTPAPRPPTPAPTIASQPLSL RPEACRPAAGGAVHTRGLDFACDIYIWAPLAGTCGVLLLSLVITLYCK RGRKKLLYIFKQPFMRPVQTTQEEDGCSCRFPEEEEGGCELRVKFSRS ADAPAYKQGQNQLYNELNLGRREEYDVLDKRRGRDPEMGGKPRRKNPQ EGLYNELQKDKMAEAYSEIGMKGERRRGKGHDGLYQGLSTATKDTYDA LHMQALPPR
149367-nt **Full CAR**	1092	ATGGCCCTCCCTGTCACCGCCCTGCTGCTTCCGCTGGCTCTTCTGCTC CACGCCGCTCGGCCCCAAGTGCAGCTTCAGGAGAGCGGCCCGGGACTC GTGAAGCCGTCCCAGACCCTGTCCCTGACTTGCACCGTGTCGGGAGGA AGCATCTCGAGCGGAGGCTACTATTGGTCGTGGATTCGGCAGCACCCT GGAAAGGGCCTGGAATGGATCGGCTACATCTACTACTCCGGCTCGACC TACTACAACCCATCGCTGAAGTCCAGAGTGACAATCTCAGTGGACACG TCCAAGAATCAGTTCAGCCTGAAGCTCTCTTCCGTGACTGCGGCCGAC ACCGCCGTGTACTACTGCGCACGCGCTGGAATTGCCGCCCGGCTGAGG GGTGCCTTCGACATTTGGGGACAGGGCACCATGGTCACCGTGTCCTCC GGCGGCGGAGGTTCCGGGGGTGGAGGCTCAGGAGGAGGGGGGTCCGAC ATCGTCATGACTCAGTCGCCCTCAAGCGTCAGCGCGTCCGTCGGGGAC AGAGTGATCATCACCTGTCGGGCGTCCCAGGGAATTCGCAACTGGCTG GCCTGGTATCAGCAGAAGCCCGGAAAGGCCCCCAACCTGTTGATCTAC GCCGCCTCAAACCTCCAATCCGGGGTGCCGAGCCGCTTCAGCGGCTCC GGTTCGGGTGCCGATTTCACTCTGACCATCTCCTCCCTGCAACCTGAA GATGTGGCTACCTACTACTGCCAAAAGTACAACTCCGCACCTTTTACT TTCGGACCGGGGACCAAAGTGGACATTAAGACCACTACCCCAGCACCG AGGCCACCCACCCCGGCTCCTACCATCGCCTCCCAGCCTCTGTCCCTG CGTCCGGAGGCATGTAGACCCGCAGCTGGTGGGGCCGTGCATACCCGG GGTCTTGACTTCGCCTGCGATATCTACATTTGGGCCCCTCTGGCTGGT ACTTGCGGGGTCCTGCTGCTTTCACTCGTGATCACTCTTTACTGTAAG CGCGGTCGGAAGAAGCTGCTGTACATCTTTAAGCAACCCTTCATGAGG CCTGTGCAGACTACTCAAGAGGAGGACGGCTGTTCATGCCGGTTCCCA GAGGAGGAGGAAGGCGGCTGCGAACTGCGCGTGAAATTCAGCCGCAGC GCAGATGCTCCAGCCTACAAGCAGGGGCAGAACCAGCTCTACAACGAA CTCAATCTTGGTCGGAGAGAGGAGTACGACGTGCTGGACAAGCGGAGA GGACGGGACCCAGAAATGGGCGGGAAGCCGCGCAGAAAGAATCCCCAA GAGGGCCTGTACAACGAGCTCCAAAAGGATAAGATGGCAGAAGCCTAT AGCGAGATTGGTATGAAAGGGGAACGCAGAAGAGGCAAAGGCCACGAC GGACTGTACCAGGGACTCAGCACCGCCACCAAGGACACCTATGACGCT CTTCACATGCAGGCCCTGCCGCCTCGG
149368		

149368-aa ScFv domain	1093	QVQLVQSGAEVKKPGSSVKVSCKASGGTFSSYAISWVRQAPGQGLEWM GGIIPIFGTANYAQKFQGRVTITADESTSTAYMELSSLRSEDTAVYYC ARRGGYQLLRWDVGLLRSAFDIWGQGTMVTVSSGGGGSGGGGSGGGGS SYVLTQPPSVSVAPGQTARITCGGNNIGSKSVHWYQQKPGQAPVLVLY GKNNRPSGVPDRFSGSRSGTTASLTITGAQAEDEADYYCSSRDSSGDH LRVFGTGTKVTVL
149368-nt ScFv domain	1094	CAAGTGCAGCTGGTCCAGTCGGGCGCCGAGGTCAAGAAGCCCGGGAGC TCTGTGAAAGTGTCCTGCAAGGCCTCCGGGGGCACCTTTAGCTCCTAC GCCATCTCCTGGGTCCGCCAAGCACCGGGTCAAGGCCTGGAGTGGATG GGGGGAATTATCCCTATCTTCGGCACTGCCAACTACGCCCAGAAGTTC CAGGGACGCGTGACCATTACCGCGGACGAATCCACCTCCACCGCTTAT ATGGAGCTGTCCAGCTTGCGCTCGGAAGATACCGCCGTGTACTACTGC GCCCGGAGGGGTGGATACCAGCTGCTGAGATGGGACGTGGGCCTCCTG CGGTCGGCGTTCGACATCTGGGGCCAGGGCACTATGGTCACTGTGTCC AGCGGAGGAGGCGGATCGGGAGGCGGCGGATCAGGGGGAGGCGGTTCC AGCTACGTGCTTACTCAACCCCCTTCGGTGTCCGTGGCCCCGGGACAG ACCGCCAGAATCACTTGCGGAGGAAACAACATTGGGTCCAAGAGCGTG CATTGGTACCAGCAGAAGCCAGGACAGGCCCCTGTGCTGGTGCTCTAC GGGAAGAACAATCGGCCCAGCGGAGTGCCGGACAGGTTCTCGGGTTCA CGCTCCGGTACAACCGCTTCACTGACTATCACCGGGGCCCAGGCAGAG GATGAAGCGGACTACTACTGTTCCTCCCGGGATTCATCCGGCGACCAC CTCCGGGTGTTCGGAACCGGAACGAAGGTCACCGTGCTG
149368-aa VH	1095	QVQLVQSGAEVKKPGSSVKVSCKASGGTFSSYAISWVRQAPGQGLEWM GGIIPIFGTANYAQKFQGRVTITADESTSTAYMELSSLRSEDTAVYYC ARRGGYQLLRWDVGLLRSAFDIWGQGTMVTVSS
149368-aa VL	1096	SYVLTQPPSVSVAPGQTARITCGGNNIGSKSVHWYQQKPGQAPVLVLY GKNNRPSGVPDRFSGSRSGTTASLTITGAQAEDEADYYCSSRDSSGDH LRVFGTGTKVTVL
149368-aa Full CAR	1097	MALPVTALLLPLALLLHAARPQVQLVQSGAEVKKPGSSVKVSCKASGG TFSSYAISWVRQAPGQGLEWMGGIIPIFGTANYAQKFQGRVTITADES TSTAYMELSSLRSEDTAVYYCARRGGYQLLRWDVGLLRSAFDIWGQGT MVTVSSGGGGSGGGGSGGGGSSYVLTQPPSVSVAPGQTARITCGGNNI GSKSVHWYQQKPGQAPVLVLYGKNNRPSGVPDRFSGSRSGTTASLTIT GAQAEDEADYYCSSRDSSGDHLRVFGTGTKVTVLTTTPAPRPPTPAPT IASQPLSLRPEACRPAAGGAVHTRGLDFACDIYIWAPLAGTCGVLLLS LVITLYCKRGRKKLLYIFKQPFMRPVQTTQEEDGCSCRFPEEEEGGCE LRVKFSRSADAPAYKQGQNQLYNELNLGRREEYDVLDKRRGRDPEMGG KPRRKNPQEGLYNELQKDKMAEAYSEIGMKGERRRGKGHDGLYQGLST ATKDTYDALHMQALPPR
149368-nt Full CAR	1098	ATGGCCCTCCCTGTCACCGCCCTGCTGCTTCCGCTGGCTCTTCTGCTC CACGCCGCTCGGCCCCAAGTGCAGCTGGTCCAGTCGGGCGCCGAGGTC AAGAAGCCCGGGAGCTCTGTGAAAGTGTCCTGCAAGGCCTCCGGGGGC ACCTTTAGCTCCTACGCCATCTCCTGGGTCCGCCAAGCACCGGGTCAA

		GGCCTGGAGTGGATGGGGGGAATTATCCCTATCTTCGGCACTGCCAAC TACGCCCAGAAGTTCCAGGGACGCGTGACCATTACCGCGGACGAATCC ACCTCCACCGCTTATATGGAGCTGTCCAGCTTGCGCTCGGAAGATACC GCCGTGTACTACTGCGCCCGGAGGGGTGGATACCAGCTGCTGAGATGG GACGTGGGCCTCCTGCGGTCGGCGTTCGACATCTGGGGCCAGGGCACT ATGGTCACTGTGTCCAGCGGAGGAGGCGGATCGGGAGGCGGCGGATCA GGGGGAGGCGGTTCCAGCTACGTGCTTACTCAACCCCCTTCGGTGTCC GTGGCCCCGGGACAGACCGCCAGAATCACTTGCGGAGGAAACAACATT GGGTCCAAGAGCGTGCATTGGTACCAGCAGAAGCCAGGACAGGCCCCT GTGCTGGTGCTCTACGGGAAGAACAATCGGCCCAGCGGAGTGCCGGAC AGGTTCTCGGGTTCACGCTCCGGTACAACCGCTTCACTGACTATCACC GGGGCCCAGGCAGAGGATGAAGCGGACTACTACTGTTCCTCCCGGGAT TCATCCGGCGACCACCTCCGGGTGTTCGGAACCGGAACGAAGGTCACC GTGCTGACCACTACCCCAGCACCGAGGCCACCCACCCCGGCTCCTACC ATCGCCTCCCAGCCTCTGTCCCTGCGTCCGGAGGCATGTAGACCCGCA GCTGGTGGGGCCGTGCATACCCGGGGTCTTGACTTCGCCTGCGATATC TACATTTGGGCCCCTCTGGCTGGTACTTGCGGGGTCCTGCTGCTTTCA CTCGTGATCACTCTTTACTGTAAGCGCGGTCGGAAGAAGCTGCTGTAC ATCTTTAAGCAACCCTTCATGAGGCCTGTGCAGACTACTCAAGAGGAG GACGGCTGTTCATGCCGGTTCCCAGAGGAGGAGGAAGGCGGCTGCGAA CTGCGCGTGAAATTCAGCCGCAGCGCAGATGCTCCAGCCTACAAGCAG GGGCAGAACCAGCTCTACAACGAACTCAATCTTGGTCGGAGAGAGGAG TACGACGTGCTGGACAAGCGGAGAGGACGGGACCCAGAAATGGGCGGG AAGCCGCGCAGAAAGAATCCCCAAGAGGGCCTGTACAACGAGCTCCAA AAGGATAAGATGGCAGAAGCCTATAGCGAGATTGGTATGAAAGGGGAA CGCAGAAGAGGCAAAGGCCACGACGGACTGTACCAGGGACTCAGCACC GCCACCAAGGACACCTATGACGCTCTTCACATGCAGGCCCTGCCGCCT CGG
149369		
149369-aa ScFv domain	1099	EVQLQQSGPGLVKPSQTLSLTCAISGDSVSSNSAAWNWIRQSPSRGLE WLGRTYYRSKWYSFYAISLKSRIIINPDTSKNQFSLQLKSVTPEDTAV YYCARSSPEGLFLYWFDPWGQGTLVTVSSGGDGSGGGGSGGGGSSSEL TQDPAVSVALGQTIRITCQGDSLGNYYATWYQQKPGQAPVLVIYGTNN RPSGIPDRFSASSSGNTASLTITGAQAEDEADYYCNSRDSSGHHLLFG TGTKVTVL
149369-nt ScFv domain	1100	GAAGTGCAGCTCCAACAGTCAGGACCGGGGCTCGTGAAGCCATCCCAG ACCCTGTCCCTGACTTGTGCCATCTCGGGAGATAGCGTGTCATCGAAC TCCGCCGCCTGGAACTGGATTCGGCAGAGCCCGTCCCGCGGACTGGAG TGGCTTGGAAGGACCTACTACCGGTCCAAGTGGTACTCTTTCTACGCG ATCTCGCTGAAGTCCCGCATTATCATTAACCCTGATACCTCCAAGAAT CAGTTCTCCCTCCAACTGAAATCCGTCACCCCCGAGGACACAGCAGTG TATTACTGCGCACGGAGCAGCCCCGAAGGACTGTTCCTGTATTGGTTT GACCCCTGGGGCCAGGGGACTCTTGTGACCGTGTCGAGCGGCGGAGAT GGGTCCGGTGGCGGTGGTTCGGGGGGCGGCGGATCATCATCCGAACTG ACCCAGGACCCGGCTGTGTCCGTGGCGCTGGGACAAACCATCCGCATT

		ACGTGCCAGGGAGACTCCCTGGGCAACTACTACGCCACTTGGTACCAG CAGAAGCCGGGCCAAGCCCCTGTGTTGGTCATCTACGGGACCAACAAC AGACCTTCCGGCATCCCCGACCGGTTCAGCGCTTCGTCCTCCGGCAAC ACTGCCAGCCTGACCATCACTGGAGCGCAGGCCGAAGATGAGGCCGAC TACTACTGCAACAGCAGAGACTCCTCGGGTCATCACCTCTTGTTCGGA ACTGGAACCAAGGTCACCGTGCTG
149369-aa VH	1101	EVQLQQSGPGLVKPSQTLSLTCAISGDSVSSNSAAWNWIRQSPSRGLE WLGRTYYRSKWYSFYAISLKSRIIINPDTSKNQFSLQLKSVTPEDTAV YYCARSSPEGLFLYWFDPWGQGTLVTVSS
149369-aa VL	1102	SSELTQDPAVSVALGQTIRITCQGDSLGNYYATWYQQKPGQAPVLVIY GTNNRPSGIPDRFSASSSGNTASLTITGAQAEDEADYYCNSRDSSGHH LLFGTGTKVTVL
149369-aa Full CAR	1103	MALPVTALLLPLALLLHAARPEVQLQQSGPGLVKPSQTLSLTCAISGD SVSSNSAAWNWIRQSPSRGLEWLGRTYYRSKWYSFYAISLKSRIIINP DTSKNQFSLQLKSVTPEDTAVYYCARSSPEGLFLYWFDPWGQGTLVTV SSGGDGSGGGGSGGGGSSSELTQDPAVSVALGQTIRITCQGDSLGNYY ATWYQQKPGQAPVLVIYGTNNRPSGIPDRFSASSSGNTASLTITGAQA EDEADYYCNSRDSSGHHLLFGTGTKVTVLTTTPAPRPPTPAPTIASQP LSLRPEACRPAAGGAVHTRGLDFACDIYIWAPLAGTCGVLLLSLVITL YCKRGRKKLLYIFKQPFMRPVQTTQEEDGCSCRFPEEEEGGCELRVKF SRSADAPAYKQGQNQLYNELNLGRREEYDVLDKRRGRDPEMGGKPRRK NPQEGLYNELQKDKMAEAYSEIGMKGERRRGKGHDGLYQGLSTATKDT YDALHMQALPPR
149369-nt **Full CAR**	1104	ATGGCCCTCCCTGTCACCGCCCTGCTGCTTCCGCTGGCTCTTCTGCTC CACGCCGCTCGGCCCGAAGTGCAGCTCCAACAGTCAGGACCGGGGCTC GTGAAGCCATCCCAGACCCTGTCCCTGACTTGTGCCATCTCGGGAGAT AGCGTGTCATCGAACTCCGCCGCCTGGAACTGGATTCGGCAGAGCCCG TCCCGCGGACTGGAGTGGCTTGGAAGGACCTACTACCGGTCCAAGTGG TACTCTTTCTACGCGATCTCGCTGAAGTCCCGCATTATCATTAACCCT GATACCTCCAAGAATCAGTTCTCCCTCCAACTGAAATCCGTCACCCCC GAGGACACAGCAGTGTATTACTGCGCACGGAGCAGCCCCGAAGGACTG TTCCTGTATTGGTTTGACCCCTGGGGCCAGGGGACTCTTGTGACCGTG TCGAGCGGCGGAGATGGGTCCGGTGGCGGTGGTTCGGGGGGCGGCGGA TCATCATCCGAACTGACCCAGGACCCGGCTGTGTCCGTGGCGCTGGGA CAAACCATCCGCATTACGTGCCAGGGAGACTCCCTGGGCAACTACTAC GCCACTTGGTACCAGCAGAAGCCGGGCCAAGCCCCTGTGTTGGTCATC TACGGGACCAACAACAGACCTTCCGGCATCCCCGACCGGTTCAGCGCT TCGTCCTCCGGCAACACTGCCAGCCTGACCATCACTGGAGCGCAGGCC GAAGATGAGGCCGACTACTACTGCAACAGCAGAGACTCCTCGGGTCAT CACCTCTTGTTCGGAACTGGAACCAAGGTCACCGTGCTGACCACTACC CCAGCACCGAGGCCACCCACCCCCGGCTCCTACCATCGCCTCCCAGCCT CTGTCCCTGCGTCCGGAGGCATGTAGACCCGCAGCTGGTGGGGCCGTG CATACCCGGGGTCTTGACTTCGCCTGCGATATCTACATTTGGGCCCCT CTGGCTGGTACTTGCGGGGTCCTGCTGCTTTCACTCGTGATCACTCTT

		TACTGTAAGCGCGGTCGGAAGAAGCTGCTGTACATCTTTAAGCAACCC TTCATGAGGCCTGTGCAGACTACTCAAGAGGAGGACGGCTGTTCATGC CGGTTCCCAGAGGAGGAGGAAGGCGGCTGCGAACTGCGCGTGAAATTC AGCCGCAGCGCAGATGCTCCAGCCTACAAGCAGGGGCAGAACCAGCTC TACAACGAACTCAATCTTGGTCGGAGAGAGGAGTACGACGTGCTGGAC AAGCGGAGAGGACGGGACCCAGAAATGGGCGGGAAGCCGCGCAGAAAG AATCCCCAAGAGGGCCTGTACAACGAGCTCCAAAAGGATAAGATGGCA GAAGCCTATAGCGAGATTGGTATGAAAGGGGAACGCAGAAGAGGCAAA GGCCACGACGGACTGTACCAGGGACTCAGCACCGCCACCAAGGACACC TATGACGCTCTTCACATGCAGGCCCTGCCGCCTCGG
BCMA_EBB-C1978-A4		
BCMA_EBB-C1978-A4 - aa **ScFv domain**	1105	EVQLVESGGGLVQPGGSLRLSCAASGFTFSSYAMSWVRQAPGKGLEWV SAISGSGGSTYYADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYC AKVEGSGSLDYWGQGTLVTVSSGGGGSGGGGSGGGGSEIVMTQSPGTL SLSPGERATLSCRASQSVSSAYLAWYQQKPGQPPRLLISGASTRATGI PDRFGGSGSGTDFTLTISRLEPEDFAVYYCQHYGSSFNGSSLFTFGQG TRLEIK
BCMA_EBB-C1978-A4 - nt **ScFv domain**	1106	GAAGTGCAGCTCGTGGAGTCAGGAGGCGGCCTGGTCCAGCCGGGAGGG TCCCTTAGACTGTCATGCGCCGCAAGCGGATTCACTTTCTCCTCCTAT GCCATGAGCTGGGTCCGCCAAGCCCCCGGAAAGGGACTGGAATGGGTG TCCGCCATCTCGGGGTCTGGAGGCTCAACTTACTACGCTGACTCCGTG AAGGGACGGTTCACCATTAGCCGCGACAACTCCAAGAACACCCTCTAC CTCCAAATGAACTCCCTGCGGGCCGAGGATACCGCCGTCTACTACTGC GCCAAAGTGGAAGGTTCAGGATCGCTGGACTACTGGGGACAGGGTACT CTCGTGACCGTGTCATCGGGCGGAGGAGGTTCCGGCGGTGGCGGCTCC GGCGGCGGAGGGTCGGAGATCGTGATGACCCAGAGCCCTGGTACTCTG AGCCTTTCGCCGGGAGAAAGGGCCACCCTGTCCTGCCGCGCTTCCCAA TCCGTGTCCTCCGCGTACTTGGCGTGGTACCAGCAGAAGCCGGGACAG CCCCCTCGGCTGCTGATCAGCGGGGCCAGCACCCGGGCAACCGGAATC CCAGACAGATTCGGGGGTTCCGGCAGCGGCACAGATTTCACCCTGACT ATTTCGAGGTTGGAGCCCGAGGACTTTGCGGTGTATTACTGTCAGCAC TACGGGTCGTCCTTTAATGGCTCCAGCCTGTTCACGTTCGGACAGGGG ACCCGCCTGGAAATCAAG
BCMA_EBB-C1978-A4 - aa **VH**	1107	EVQLVESGGGLVQPGGSLRLSCAASGFTFSSYAMSWVRQAPGKGLEWV SAISGSGGSTYYADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYC AKVEGSGSLDYWGQGTLVTVSS
BCMA_EBB-C1978-A4 - aa	1108	EIVMTQSPGTLSLSPGERATLSCRASQSVSSAYLAWYQQKPGQPPRLL ISGASTRATGIPDRFGGSGSGTDFTLTISRLEPEDFAVYYCQHYGSSF NGSSLFTFGQGTRLEIK

VL		
BCMA_EBB-C1978-A4 - aa **Full CART**	1109	MALPVTALLLPLALLLHAARPEVQLVESGGGLVQPGGSLRLSCAASGF TFSSYAMSWVRQAPGKGLEWVSAISGSGGSTYYADSVKGRFTISRDNS KNTLYLQMNSLRAEDTAVYYCAKVEGSGSLDYWGQGTLVTVSSGGGGS GGGGSGGGGSEIVMTQSPGTLSLSPGERATLSCRASQSVSSAYLAWYQ QKPGQPPRLLISGASTRATGIPDRFGGSGSGTDFTLTISRLEPEDFAV YYCQHYGSSFNGSSLFTFGQGTRLEIKTTTPAPRPPTPAPTIASQPLS LRPEACRPAAGGAVHTRGLDFACDIYIWAPLAGTCGVLLLSLVITLYC KRGRKKLLYIFKQPFMRPVQTTQEEDGCSCRFPEEEEGGCELRVKFSR SADAPAYKQGQNQLYNELNLGRREEYDVLDKRRGRDPEMGGKPRRKNP QEGLYNELQKDKMAEAYSEIGMKGERRRGKGHDGLYQGLSTATKDTYD ALHMQALPPR
BCMA_EBB-C1978-A4 - nt **Full CART**	1110	ATGGCCCTCCCTGTCACCGCCCTGCTGCTTCCGCTGGCTCTTCTGCTC CACGCCGCTCGGCCCGAAGTGCAGCTCGTGGAGTCAGGAGGCGGCCTG GTCCAGCCGGGAGGGTCCCTTAGACTGTCATGCGCCGCAAGCGGATTC ACTTTCTCCTCCTATGCCATGAGCTGGGTCCGCCAAGCCCCCGGAAAG GGACTGGAATGGGTGTCCGCCATCTCGGGGTCTGGAGGCTCAACTTAC TACGCTGACTCCGTGAAGGGACGGTTCACCATTAGCCGCGACAACTCC AAGAACACCCTCTACCTCCAAATGAACTCCCTGCGGGCCGAGGATACC GCCGTCTACTACTGCGCCAAAGTGGAAGGTTCAGGATCGCTGGACTAC TGGGGACAGGGTACTCTCGTGACCGTGTCATCGGGCGGAGGAGGTTCC GGCGGTGGCGGCTCCGGCGGCGGAGGGTCGGAGATCGTGATGACCCAG AGCCCTGGTACTCTGAGCCTTTCGCCGGGAGAAAGGGCCACCCTGTCC TGCCGCGCTTCCCAATCCGTGTCCTCCGCGTACTTGGCGTGGTACCAG CAGAAGCCGGGACAGCCCCCTCGGCTGCTGATCAGCGGGGCCAGCACC CGGGCAACCGGAATCCCAGACAGATTCGGGGGTTCCGGCAGCGGCACA GATTTCACCCTGACTATTTCGAGGTTGGAGCCCGAGGACTTTGCGGTG TATTACTGTCAGCACTACGGGTCGTCCTTTAATGGCTCCAGCCTGTTC ACGTTCGGACAGGGGACCCGCCTGGAAATCAAGACCACTACCCCAGCA CCGAGGCCACCCACCCCGGCTCCTACCATCGCCTCCCAGCCTCTGTCC CTGCGTCCGGAGGCATGTAGACCCGCAGCTGGTGGGGCCGTGCATACC CGGGGTCTTGACTTCGCCTGCGATATCTACATTTGGGCCCCTCTGGCT GGTACTTGCGGGGTCCTGCTGCTTTCACTCGTGATCACTCTTTACTGT AAGCGCGGTCGGAAGAAGCTGCTGTACATCTTTAAGCAACCCTTCATG AGGCCTGTGCAGACTACTCAAGAGGAGGACGGCTGTTCATGCCGGTTC CCAGAGGAGGAGGAAGGCGGCTGCGAACTGCGCGTGAAATTCAGCCGC AGCGCAGATGCTCCAGCCTACAAGCAGGGGCAGAACCAGCTCTACAAC GAACTCAATCTTGGTCGGAGAGAGGAGTACGACGTGCTGGACAAGCGG AGAGGACGGGACCCAGAAATGGGCGGGAAGCCGCGCAGAAAGAATCCC CAAGAGGGCCTGTACAACGAGCTCCAAAAGGATAAGATGGCAGAAGCC TATAGCGAGATTGGTATGAAAGGGGAACGCAGAAGAGGCAAAGGCCAC GACGGACTGTACCAGGGACTCAGCACCGCCACCAAGGACACCTATGAC GCTCTTCACATGCAGGCCCTGCCGCCTCGG
BCMA_EBB-C1978-G1		

BCMA_EBB-C1978-G1 - aa **ScFv domain**	1111	EVQLVETGGGLVQPGGSLRLSCAASGITFSRYPMSWVRQAPGKGLEWVSGISD SGVSTYYADSAKGRFTISRDNSKNTLFLQMSSLRDEDTAVYYCVTRAGSEASDI WGQGTMVTVSSGGGGSGGGGSGGGGSEIVLTQSPATLSLSPGERATLSCRAS QSVSNSLAWYQQKPGQAPRLLIYDASSRATGIPDRFSGSGSGTDFTLTISRLEPE DFAIYYCQQFGTSSGLTFGGGTKLEIK
BCMA_EBB-C1978-G1 - nt **ScFv domain**	1112	GAAGTGCAACTGGTGGAAACCGGTGGCGGCCTGGTGCAGCCTGGAGGA TCATTGAGGCTGTCATGCGCGGCCAGCGGTATTACCTTCTCCCGGTAC CCCATGTCCTGGGTCAGACAGGCCCCGGGGAAAGGGCTTGAATGGGTG TCCGGGATCTCGGACTCCGGTGTCAGCACTTACTACGCCGACTCCGCC AAGGGACGCTTCACCATTTCCCGGGACAACTCGAAGAACACCCTGTTC CTCCAAATGAGCTCCCTCCGGGACGAGGATACTGCAGTGTACTACTGC GTGACCCGCGCCGGGTCCGAGGCGTCTGACATTTGGGGACAGGGCACT ATGGTCACCGTGTCGTCCGGCGGAGGGGCTCGGGAGGCGGTGGCAGC GGAGGAGGAGGGTCCGAGATCGTGCTGACCCAATCCCCGGCCACCCTC TCGCTGAGCCCTGGAGAAAGGGCAACCTTGTCCTGTCGCGCGAGCCAG TCCGTGAGCAACTCCCTGGCCTGGTACCAGCAGAAGCCCGGACAGGCT CCGAGACTTCTGATCTACGACGCTTCGAGCCGGGCCACTGGAATCCCC GACCGCTTTTCGGGGTCCGGCTCAGGAACCGATTTCACCCTGACAATC TCACGGCTGGAGCCAGAGGATTTCGCCATCTATTACTGCCAGCAGTTC GGTACTTCCTCCGGCCTGACTTTCGGAGGCGGCACGAAGCTCGAAATC AAG
BCMA_EBB-C1978-G1 - aa **VH**	1113	EVQLVETGGGLVQPGGSLRLSCAASGITFSRYPMSWVRQAPGKGLEWV SGISDSGVSTYYADSAKGRFTISRDNSKNTLFLQMSSLRDEDTAVYYC VTRAGSEASDIWGQGTMVTVSS
BCMA_EBB-C1978-G1 - aa **VL**	1114	EIVLTQSPATLSLSPGERATLSCRASQSVSNSLAWYQQKPGQAPRLLI YDASSRATGIPDRFSGSGSGTDFTLTISRLEPEDFAIYYCQQFGTSSG LTFGGGTKLEIK
BCMA_EBB-C1978-G1 - aa **Full CART**	1115	MALPVTALLLPLALLLHAARPEVQLVETGGGLVQPGGSLRLSCAASGITFSRYPM SWVRQAPGKGLEWVSGISDSGVSTYYADSAKGRFTISRDNSKNTLFLQMSSLR DEDTAVYYCVTRAGSEASDIWGQGTMVTVSSGGGGSGGGGSGGGGSEIVLTQ SPATLSLSPGERATLSCRASQSVSNSLAWYQQKPGQAPRLLIYDASSRATGIPDR FSGSGSGTDFTLTISRLEPEDFAIYYCQQFGTSSGLTFGGGTKLEIKTTTPAPRPPT PAPTIASQPLSLRPEACRPAAGGAVHTRGLDFACDIYIWAPLAGTCGVLLLSLVIT LYCKRGRKKLLYIFKQPFMRPVQTTQEEDGCSCRFPEEEEGGCELRVKFSRSADA PAYKQGQNQLYNELNLGRREEYDVLDKRRGRDPEMGGKPRRKNPQEGLYNEL QKDKMAEAYSEIGMKGERRRGKGHDGLYQGLSTATKDTYDALHMQALPPR
BCMA_EBB-C1978-G1 -	1116	ATGGCCCTCCCTGTCACCGCCCTGCTGCTTCCGCTGGCTCTTCTGCTC CACGCCGCTCGGCCCGAAGTGCAACTGGTGGAAACCGGTGGCGGCCTG

nt **Full CART**		GTGCAGCCTGGAGGATCATTGAGGCTGTCATGCGCGGCCAGCGGTATT ACCTTCTCCCGGTACCCCATGTCCTGGGTCAGACAGGCCCCGGGGAAA GGGCTTGAATGGGTGTCCGGGATCTCGGACTCCGGTGTCAGCACTTAC TACGCCGACTCCGCCAAGGGACGCTTCACCATTTCCCGGGACAACTCG AAGAACACCCTGTTCCTCCAAATGAGCTCCCTCCGGGACGAGGATACT GCAGTGTACTACTGCGTGACCCGCGCCGGGTCCGAGGCGTCTGACATT TGGGGACAGGGCACTATGGTCACCGTGTCGTCCGGCGGAGGGGCTCG GGAGGCGGTGGCAGCGGAGGAGGAGGGTCCGAGATCGTGCTGACCCAA TCCCCGGCCACCCTCTCGCTGAGCCCTGGAGAAAGGGCAACCTTGTCC TGTCGCGCGAGCCAGTCCGTGAGCAACTCCCTGGCCTGGTACCAGCAG AAGCCCGGACAGGCTCCGAGACTTCTGATCTACGACGCTTCGAGCCGG GCCACTGGAATCCCCGACCGCTTTTCGGGGTCCGGCTCAGGAACCGAT TTCACCCTGACAATCTCACGGCTGGAGCCAGAGGATTTCGCCATCTAT TACTGCCAGCAGTTCGGTACTTCCTCCGGCCTGACTTTCGGAGGCGGC ACGAAGCTCGAAATCAAGACCACTACCCCAGCACCGAGGCCACCCACC CCGGCTCCTACCATCGCCTCCCAGCCTCTGTCCCTGCGTCCGGAGGCA TGTAGACCCGCAGCTGGTGGGGCCGTGCATACCCGGGGTCTTGACTTC GCCTGCGATATCTACATTTGGGCCCCTCTGGCTGGTACTTGCGGGGTC CTGCTGCTTTCACTCGTGATCACTCTTTACTGTAAGCGCGGTCGGAAG AAGCTGCTGTACATCTTTAAGCAACCCTTCATGAGGCCTGTGCAGACT ACTCAAGAGGAGGACGGCTGTTCATGCCGGTTCCCAGAGGAGGAGGAA GGCGGCTGCGAACTGCGCGTGAAATTCAGCCGCAGCGCAGATGCTCCA GCCTACAAGCAGGGGCAGAACCAGCTCTACAACGAACTCAATCTTGGT CGGAGAGAGGAGTACGACGTGCTGGACAAGCGGAGAGGACGGGACCCA GAAATGGGCGGGAAGCCGCGCAGAAAGAATCCCCAAGAGGGCCTGTAC AACGAGCTCCAAAAGGATAAGATGGCAGAAGCCTATAGCGAGATTGGT ATGAAAGGGGAACGCAGAAGAGGCAAAGGCCACGACGGACTGTACCAG GGACTCAGCACCGCCACCAAGGACACCTATGACGCTCTTCACATGCAG GCCCTGCCGCCTCGG
BCMA_EBB-C1979-C1		
BCMA_EBB-C1979-C1 - aa **ScFv domain**	1117	QVQLVESGGGLVQPGGSLRLSCAASGFTFSSYAMSWVRQAPGKGLEWV SAISGSGGSTYYADSVKGRFTISRDNAKNSLYLQMNSLRAEDTAIYYC ARATYKRELRYYYGMDVWGQGTMVTVSSGGGGSGGGGSGGGGSEIVMT QSPGTVSLSPGERATLSCRASQSVSSSFLAWYQQKPGQAPRLLIYGAS SRATGIPDRFSGSGSGTDFTLTISRLEPEDSAVYYCQQYHSSPSWTFG QGTRLEIK
BCMA_EBB-C1979-C1 - nt **ScFv domain**	1118	CAAGTGCAGCTCGTGGAATCGGGTGGCGGACTGGTGCAGCCGGGGGGC TCACTTAGACTGTCCTGCGCGGCCAGCGGATTCACTTTCTCCTCCTAC GCCATGTCCTGGGTCAGACAGGCCCCTGGAAAGGGCCTGGAATGGGTG TCCGCAATCAGCGGCAGCGGCGGCTCGACCTATTACGCGGATTCAGTG AAGGGCAGATTCACCATTTCCCGGGACAACGCCAAGAACTCCTTGTAC CTTCAAATGAACTCCCTCCGCGCGGAAGATACCGCAATCTACTACTGC GCTCGGGCCACTTACAAGAGGGAACTGCGCTACTACTACGGGATGGAC GTCTGGGGCCAGGGAACCATGGTCACCGTGTCCAGCGGAGGAGGAGGA TCGGGAGGAGGCGGTAGCGGGGGTGGAGGGTCGGAGATCGTGATGACC

		CAGTCCCCCGGCACTGTGTCGCTGTCCCCCGGCGAACGGGCCACCCTG TCATGTCGGGCCAGCCAGTCAGTGTCGTCAAGCTTCCTCGCCTGGTAC CAGCAGAAACCGGGACAAGCTCCCCGCCTGCTGATCTACGGAGCCAGC AGCCGGGCCACCGGTATTCCTGACCGGTTCTCCGGTTCGGGGTCCGGG ACCGACTTTACTCTGACTATCTCTCGCCTCGAGCCAGAGGACTCCGCC GTGTATTACTGCCAGCAGTACCACTCCTCCCCGTCCTGGACGTTCGGA CAGGGCACAAGGCTGGAGATTAAG
BCMA_EBB-C1979-C1 - aa **VH**	1119	QVQLVESGGGLVQPGGSLRLSCAASGFTFSSYAMSWVRQAPGKGLEWV SAISGSGGSTYYADSVKGRFTISRDNAKNSLYLQMNSLRAEDTAIYYC ARATYKRELRYYYGMDVWGQGTMVTVSS
BCMA_EBB-C1979-C1 - aa **VL**	1120	EIVMTQSPGTVSLSPGERATLSCRASQSVSSSFLAWYQQKPGQAPRLL IYGASSRATGIPDRFSGSGSGTDFTLTISRLEPEDSAVYYCQQYHSSP SWTFGQGTRLEIK
BCMA_EBB-C1979-C1 - aa **Full CART**	1121	MALPVTALLLPLALLLHAARPQVQLVESGGGLVQPGGSLRLSCAASGF TFSSYAMSWVRQAPGKGLEWVSAISGSGGSTYYADSVKGRFTISRDNA KNSLYLQMNSLRAEDTAIYYCARATYKRELRYYYGMDVWGQGTMVTVS SGGGGSGGGGSGGGGSEIVMTQSPGTVSLSPGERATLSCRASQSVSSS FLAWYQQKPGQAPRLLIYGASSRATGIPDRFSGSGSGTDFTLTISRLE PEDSAVYYCQQYHSSPSWTFGQGTRLEIKTTTPAPRPPTPAPTIASQP LSLRPEACRPAAGGAVHTRGLDFACDIYIWAPLAGTCGVLLLSLVITL YCKRGRKKLLYIFKQPFMRPVQTTQEEDGCSCRFPEEEEGGCELRVKF SRSADAPAYKQGQNQLYNELNLGRREEYDVLDKRRGRDPEMGGKPRRK NPQEGLYNELQKDKMAEAYSEIGMKGERRRGKGHDGLYQGLSTATKDT YDALHMQALPPR
BCMA_EBB-C1979-C1 - nt **Full CART**	1122	ATGGCCCTCCCTGTCACCGCCCTGCTGCTTCCGCTGGCTCTTCTGCTC CACGCCGCTCGGCCCCAAGTGCAGCTCGTGGAATCGGGTGGCGGACTG GTGCAGCCGGGGGGCTCACTTAGACTGTCCTGCGCGGCCAGCGGATTC ACTTTCTCCTCCTACGCCATGTCCTGGGTCAGACAGGCCCCTGGAAAG GGCCTGGAATGGGTGTCCGCAATCAGCGGCAGCGGCGGCTCGACCTAT TACGCGGATTCAGTGAAGGGCAGATTCACCATTTCCCGGGACAACGCC AAGAACTCCTTGTACCTTCAAATGAACTCCCTCCGCGCGGAAGATACC GCAATCTACTACTGCGCTCGGGCCACTTACAAGAGGGAACTGCGCTAC TACTACGGGATGGACGTCTGGGGCCAGGGAACCATGGTCACCGTGTCC AGCGGAGGAGGAGGATCGGGAGGAGGCGGTAGCGGGGGTGGAGGGTCG GAGATCGTGATGACCCAGTCCCCCGGCACTGTGTCGCTGTCCCCCGGC GAACGGGCCACCCTGTCATGTCGGGCCAGCCAGTCAGTGTCGTCAAGC TTCCTCGCCTGGTACCAGCAGAAACCGGGACAAGCTCCCCGCCTGCTG ATCTACGGAGCCAGCAGCCGGGCCACCGGTATTCCTGACCGGTTCTCC GGTTCGGGGTCCGGGACCGACTTTACTCTGACTATCTCTCGCCTCGAG CCAGAGGACTCCGCCGTGTATTACTGCCAGCAGTACCACTCCTCCCCG

		TCCTGGACGTTCGGACAGGGCACAAGGCTGGAGATTAAGACCACTACC CCAGCACCGAGGCCACCCACCCCGGCTCCTACCATCGCCTCCCAGCCT CTGTCCCTGCGTCCGGAGGCATGTAGACCCGCAGCTGGTGGGGCCGTG CATACCCGGGGTCTTGACTTCGCCTGCGATATCTACATTTGGGCCCCT CTGGCTGGTACTTGCGGGGTCCTGCTGCTTTCACTCGTGATCACTCTT TACTGTAAGCGCGGTCGGAAGAAGCTGCTGTACATCTTTAAGCAACCC TTCATGAGGCCTGTGCAGACTACTCAAGAGGAGGACGGCTGTTCATGC CGGTTCCCAGAGGAGGAGGAAGGCGGCTGCGAACTGCGCGTGAAATTC AGCCGCAGCGCAGATGCTCCAGCCTACAAGCAGGGGCAGAACCAGCTC TACAACGAACTCAATCTTGGTCGGAGAGAGGAGTACGACGTGCTGGAC AAGCGGAGAGGACGGGACCCAGAAATGGGCGGGAAGCCGCGCAGAAAG AATCCCCAAGAGGGCCTGTACAACGAGCTCCAAAAGGATAAGATGGCA GAAGCCTATAGCGAGATTGGTATGAAAGGGGAACGCAGAAGAGGCAAA GGCCACGACGGACTGTACCAGGGACTCAGCACCGCCACCAAGGACACC TATGACGCTCTTCACATGCAGGCCCTGCCGCCTCGG
BCMA_EBB-C1978-C7		
BCMA_EBB-C1978-C7 - aa **ScFv domain**	1123	EVQLVETGGGLVQPGGSLRLSCAASGFTFSSYAMSWVRQAPGKGLEWV SAISGSGGSTYYADSVKGRFTISRDNSKNTLYLQMNTLKAEDTAVYYC ARATYKRELRYYYGMDVWGQGTTVTVSSGGGGSGGGGSGGGGSEIVLT QSPSTLSLSPGESATLSCRASQSVSTTFLAWYQQKPGQAPRLLIYGSS NRATGIPDRFSGSGSGTDFTLTIRRLEPEDFAVYYCQQYHSSPSWTFG QGTKVEIK
BCMA_EBB-C1978-C7 - nt **ScFv domain**	1124	GAGGTGCAGCTTGTGGAAACCGGTGGCGGACTGGTGCAGCCCGGAGGA AGCCTCAGGCTGTCCTGCGCCGCGTCCGGCTTCACCTTCTCCTCGTAC GCCATGTCCTGGGTCCGCCAGGCCCCCGGAAAGGGCCTGGAATGGGTG TCCGCCATCTCTGGAAGCGGAGGTTCCACGTACTACGCGGACAGCGTC AAGGGAAGGTTCACAATCTCCCGCGATAATTCGAAGAACACTCTGTAC CTTCAAATGAACACCCTGAAGGCCGAGGACACTGCTGTGTACTACTGC GCACGGGCCACCTACAAGAGAGAGCTCCGGTACTACTACGGAATGGAC GTCTGGGGCCAGGGAACTACTGTGACCGTGTCCTCGGGAGGGGGTGGC TCCGGGGGGGGCGGCTCCGGCGGAGGCGGTTCCGAGATTGTGCTGACC CAGTCACCTTCAACTCTGTCGCTGTCCCCGGGAGAGAGCGCTACTCTG AGCTGCCGGGCCAGCCAGTCCGTGTCCACCACCTTCCTCGCCTGGTAT CAGCAGAAGCCGGGGCAGGCACCACGGCTCTTGATCTACGGGTCAAGC AACAGAGCGACCGGAATTCCTGACCGCTTCTCGGGGAGCGGTTCAGGC ACCGACTTCACCCTGACTATCCGGCGCCTGGAACCCGAAGATTTCGCC GTGTATTACTGTCAACAGTACCACTCCTCGCCGTCCTGGACCTTTGGC CAAGGAACCAAAGTGGAAATCAAG
BCMA_EBB-C1978-C7 - aa **VH**	1125	EVQLVETGGGLVQPGGSLRLSCAASGFTFSSYAMSWVRQAPGKGLEWV SAISGSGGSTYYADSVKGRFTISRDNSKNTLYLQMNTLKAEDTAVYYC ARATYKRELRYYYGMDVWGQGTTVTVSS

BCMA_EBB-C1978-C7 - aa **VL**	1126	EIVLTQSPSTLSLSPGESATLSCRASQSVSTTFLAWYQQKPGQAPRLL IYGSSNRATGIPDRFSGSGSGTDFTLTIRRLEPEDFAVYYCQQYHSSP SWTFGQGTKVEIK
BCMA_EBB-C1978-C7 - aa **Full CART**	1127	MALPVTALLLPLALLLHAARPEVQLVETGGGLVQPGGSLRLSCAASGF TFSSYAMSWVRQAPGKGLEWVSAISGSGGSTYYADSVKGRFTISRDNS KNTLYLQMNTLKAEDTAVYYCARATYKRELRYYYGMDVWGQGTTVTVS SGGGGSGGGGSGGGGSEIVLTQSPSTLSLSPGESATLSCRASQSVSTT FLAWYQQKPGQAPRLLIYGSSNRATGIPDRFSGSGSGTDFTLTIRRLE PEDFAVYYCQQYHSSPSWTFGQGTKVEIKTTTPAPRPPTPAPTIASQP LSLRPEACRPAAGGAVHTRGLDFACDIYIWAPLAGTCGVLLLSLVITL YCKRGRKKLLYIFKQPFMRPVQTTQEEDGCSCRFPEEEEGGCELRVKF SRSADAPAYKQGQNQLYNELNLGRREEYDVLDKRRGRDPEMGGKPRRK NPQEGLYNELQKDKMAEAYSEIGMKGERRRGKGHDGLYQGLSTATKDT YDALHMQALPPR
BCMA_EBB-C1978-C7 - nt **Full CART**	1128	ATGGCCCTCCCTGTCACCGCCCTGCTGCTTCCGCTGGCTCTTCTGCTC CACGCCGCTCGGCCCGAGGTGCAGCTTGTGGAAACCGGTGGCGGACTG GTGCAGCCCGGAGGAAGCCTCAGGCTGTCCTGCGCCGCGTCCGGCTTC ACCTTCTCCTCGTACGCCATGTCCTGGGTCCGCCAGGCCCCCGGAAAG GGCCTGGAATGGGTGTCCGCCATCTCTGGAAGCGGAGGTTCCACGTAC TACGCGGACAGCGTCAAGGGAAGGTTCACAATCTCCCGCGATAATTCG AAGAACACTCTGTACCTTCAAATGAACACCCTGAAGGCCGAGGACACT GCTGTGTACTACTGCGCACGGGCCACCTACAAGAGAGAGCTCCGGTAC TACTACGGAATGGACGTCTGGGGCCAGGGAACTACTGTGACCGTGTCC TCGGGAGGGGGTGGCTCCGGGGGGGGCGGCTCCGGCGGAGGCGGTTCC GAGATTGTGCTGACCCAGTCACCTTCAACTCTGTCGCTGTCCCCGGGA GAGAGCGCTACTCTGAGCTGCCGGGCCAGCCAGTCCGTGTCCACCACC TTCCTCGCCTGGTATCAGCAGAAGCCGGGGCAGGCACCACGGCTCTTG ATCTACGGGTCAAGCAACAGAGCGACCGGAATTCCTGACCGCTTCTCG GGGAGCGGTTCAGGCACCGACTTCACCCTGACTATCCGGCGCCTGGAA CCCGAAGATTTCGCCGTGTATTACTGTCAACAGTACCACTCCTCGCCG TCCTGGACCTTTGGCCAAGGAACCAAAGTGGAAATCAAGACCACTACC CCAGCACCGAGGCCACCCACCCCGGCTCCTACCATCGCCTCCCAGCCT CTGTCCCTGCGTCCGGAGGCATGTAGACCCGCAGCTGGTGGGGCCGTG CATACCCGGGGTCTTGACTTCGCCTGCGATATCTACATTTGGGCCCCT CTGGCTGGTACTTGCGGGGTCCTGCTGCTTTCACTCGTGATCACTCTT TACTGTAAGCGCGGTCGGAAGAAGCTGCTGTACATCTTTAAGCAACCC TTCATGAGGCCTGTGCAGACTACTCAAGAGGAGGACGGCTGTTCATGC CGGTTCCCAGAGGAGGAGGAAGGCGGCTGCGAACTGCGCGTGAAATTC AGCCGCAGCGCAGATGCTCCAGCCTACAAGCAGGGGCAGAACCAGCTC TACAACGAACTCAATCTTGGTCGGAGAGAGGAGTACGACGTGCTGGAC AAGCGGAGAGGACGGGACCCAGAAATGGGCGGGAAGCCGCGCAGAAAG AATCCCCAAGAGGGCCTGTACAACGAGCTCCAAAAGGATAAGATGGCA GAAGCCTATAGCGAGATTGGTATGAAAGGGGAACGCAGAAGAGGCAAA GGCCACGACGGACTGTACCAGGGACTCAGCACCGCCACCAAGGACACC

		TATGACGCTCTTCACATGCAGGCCCTGCCGCCTCGG
BCMA_EBB-C1978-D10		
BCMA_EBB-C1978-D10 - aa **ScFv domain**	1129	EVQLVETGGGLVQPGRSLRLSCAASGFTFDDYAMHWVRQAPGKGLEWV SGISWNSGSIGYADSVKGRFTISRDNAKNSLYLQMNSLRDEDTAVYYC ARVGKAVPDVWGQGTTVTVSSGGGGSGGGGSGGGGSDIVMTQTPSSLS ASVGDRVTITCRASQSISSYLNWYQQKPGKAPKLLIYAASSLQSGVPS RFSGSGSGTDFTLTISSLQPEDFATYYCQQSYSTPYSFGQGTRLEIK
BCMA_EBB-C1978-D10-nt **ScFv domain**	1130	GAAGTGCAGCTCGTGGAAACTGGAGGTGGACTCGTGCAGCCTGGACGG TCGCTGCGGCTGAGCTGCGCTGCATCCGGCTTCACCTTCGACGATTAT GCCATGCACTGGGTCAGACAGGCGCCAGGGAAGGGACTTGAGTGGGTG TCCGGTATCAGCTGGAATAGCGGCTCAATCGGATACGCGGACTCCGTG AAGGGAAGGTTCACCATTTCCCGCGACAACGCCAAGAACTCCCTGTAC TTGCAAATGAACAGCCTCCGGGATGAGGACACTGCCGTGTACTACTGC GCCCGCGTCGGAAAAGCTGTGCCCGACGTCTGGGGCCAGGGAACCACT GTGACCGTGTCCAGCGGCGGGGGTGGATCGGGCGGTGGAGGGTCCGGT GGAGGGGGCTCAGATATTGTGATGACCCAGACCCCCTCGTCCCTGTCC GCCTCGGTCGGCGACCGCGTGACTATCACATGTAGAGCCTCGCAGAGC ATCTCCAGCTACCTGAACTGGTATCAGCAGAAGCCGGGGAAGGCCCCG AAGCTCCTGATCTACGCGGCATCATCACTGCAATCGGGAGTGCCGAGC CGGTTTTCCGGGTCCGGCTCCGGCACCGACTTCACGCTGACCATTTCT TCCCTGCAACCCGAGGACTTCGCCACTTACTACTGCCAGCAGTCCTAC TCCACCCCTTACTCCTTCGGCCAAGGAACCAGGCTGGAAATCAAG
BCMA_EBB-C1978-D10 - aa **VH**	1131	EVQLVETGGGLVQPGRSLRLSCAASGFTFDDYAMHWVRQAPGKGLEWV SGISWNSGSIGYADSVKGRFTISRDNAKNSLYLQMNSLRDEDTAVYYC ARVGKAVPDVWGQGTTVTVSS
BCMA_EBB-C1978-D10-aa **VL**	1132	DIVMTQTPSSLSASVGDRVTITCRASQSISSYLNWYQQKPGKAPKLLI YAASSLQSGVPSRFSGSGSGTDFTLTISSLQPEDFATYYCQQSYSTPY SFGQGTRLEIK
BCMA_EBB-C1978-D10 - aa **Full CART**	1133	MALPVTALLLPLALLLHAARPEVQLVETGGGLVQPGRSLRLSCAASGF TFDDYAMHWVRQAPGKGLEWVSGISWNSGSIGYADSVKGRFTISRDNA KNSLYLQMNSLRDEDTAVYYCARVGKAVPDVWGQGTTVTVSSGGGGSG GGGSGGGGSDIVMTQTPSSLSASVGDRVTITCRASQSISSYLNWYQQK PGKAPKLLIYAASSLQSGVPSRFSGSGSGTDFTLTISSLQPEDFATYY CQQSYSTPYSFGQGTRLEIKTTTPAPRPPTPAPTIASQPLSLRPEACR PAAGGAVHTRGLDFACDIYIWAPLAGTCGVLLLSLVITLYCKRGRKKL LYIFKQPFMRPVQTTQEEDGCSCRFPEEEEGGCELRVKFSRSADAPAY KQGQNQLYNELNLGRREEYDVLDKRRGRDPEMGGKPRRKNPQEGLYNE LQKDKMAEAYSEIGMKGERRRGKGHDGLYQGLSTATKDTYDALHMQAL

		PPR
BCMA_EBB-C1978-D10 - nt **Full CART**	1134	ATGGCCCTCCCTGTCACCGCCCTGCTGCTTCCGCTGGCTCTTCTGCTC CACGCCGCTCGGCCCGAAGTGCAGCTCGTGGAAACTGGAGGTGGACTC GTGCAGCCTGGACGGTCGCTGCGGCTGAGCTGCGCTGCATCCGGCTTC ACCTTCGACGATTATGCCATGCACTGGGTCAGACAGGCGCCAGGGAAG GGACTTGAGTGGGTGTCCGGTATCAGCTGGAATAGCGGCTCAATCGGA TACGCGGACTCCGTGAAGGGAAGGTTCACCATTTCCCGCGACAACGCC AAGAACTCCCTGTACTTGCAAATGAACAGCCTCCGGGATGAGGACACT GCCGTGTACTACTGCGCCCGCGTCGGAAAAGCTGTGCCCGACGTCTGG GGCCAGGGAACCACTGTGACCGTGTCCAGCGGCGGGGGTGGATCGGGC GGTGGAGGGTCCGGTGGAGGGGGCTCAGATATTGTGATGACCCAGACC CCCTCGTCCCTGTCCGCCTCGGTCGGCGACCGCGTGACTATCACATGT AGAGCCTCGCAGAGCATCTCCAGCTACCTGAACTGGTATCAGCAGAAG CCGGGGAAGGCCCCGAAGCTCCTGATCTACGCGGCATCATCACTGCAA TCGGGAGTGCCGAGCCGGTTTTCCGGGTCCGGCTCCGGCACCGACTTC ACGCTGACCATTTCTTCCCTGCAACCCGAGGACTTCGCCACTTACTAC TGCCAGCAGTCCTACTCCACCCCTTACTCCTTCGGCCAAGGAACCAGG CTGGAAATCAAGACCACTACCCCAGCACCGAGGCCACCCACCCCGGCT CCTACCATCGCCTCCCAGCCTCTGTCCCTGCGTCCGGAGGCATGTAGA CCCGCAGCTGGTGGGGCCGTGCATACCCGGGGTCTTGACTTCGCCTGC GATATCTACATTTGGGCCCCTCTGGCTGGTACTTGCGGGGTCCTGCTG CTTTCACTCGTGATCACTCTTTACTGTAAGCGCGGTCGGAAGAAGCTG CTGTACATCTTTAAGCAACCCTTCATGAGGCCTGTGCAGACTACTCAA GAGGAGGACGGCTGTTCATGCCGGTTCCCAGAGGAGGAGGAAGGCGGC TGCGAACTGCGCGTGAAATTCAGCCGCAGCGCAGATGCTCCAGCCTAC AAGCAGGGGCAGAACCAGCTCTACAACGAACTCAATCTTGGTCGGAGA GAGGAGTACGACGTGCTGGACAAGCGGAGAGGACGGGACCCAGAAATG GGCGGGAAGCCGCGCAGAAAGAATCCCCAAGAGGGCCTGTACAACGAG CTCCAAAAGGATAAGATGGCAGAAGCCTATAGCGAGATTGGTATGAAA GGGGAACGCAGAAGAGGCAAAGGCCACGACGGACTGTACCAGGGACTC AGCACCGCCACCAAGGACACCTATGACGCTCTTCACATGCAGGCCCTG CCGCCTCGG
BCMA_EBB-C1979-C12		
BCMA_EBB-C1979-C12-aa **ScFv domain**	1135	EVQLVESGGGLVQPGRSLRLSCTASGFTFDDYAMHWVRQRPGKGLEWV ASINWKGNSLAYGDSVKGRFAISRDNAKNTVFLQMNSLRTEDTAVYYC ASHQGVAYYNYAMDVWGRGTLVTVSSGGGGSGGGGSGGGGSEIVLTQS PGTLSLSPGERATLSCRATQSIGSSFLAWYQQRPGQAPRLLIYGASQR ATGIPDRFSGRGSGTDFTLTISRVEPEDSAVYYCQHYESSPSWTFGQG TKVEIK
BCMA_EBB-C1979-C12 - nt	1136	GAAGTGCAGCTCGTGGAGAGCGGGGGAGGATTGGTGCAGCCCGGAAGG TCCCTGCGGCTCTCCTGCACTGCGTCTGGCTTCACCTTCGACGACTAC GCGATGCACTGGGTCAGACAGCGCCCGGGAAAGGGCCTGGAATGGGTC GCCTCAATCAACTGGAAGGGAAACTCCCTGGCCTATGGCGACAGCGTG AAGGGCCGCTTCGCCATTTCGCGCGACAACGCCAAGAACACCGTGTTT

ScFv domain		CTGCAAATGAATTCCCTGCGGACCGAGGATACCGCTGTGTACTACTGC GCCAGCCACCAGGGCGTGGCATACTATAACTACGCCATGGACGTGTGG GGAAGAGGGACGCTCGTCACCGTGTCCTCCGGGGGCGGTGGATCGGGT GGAGGAGGAAGCGGTGGCGGGGGCAGCGAAATCGTGCTGACTCAGAGC CCGGGAACTCTTTCACTGTCCCCGGGAGAACGGGCCACTCTCTCGTGC CGGGCCACCCAGTCCATCGGCTCCTCCTTCCTTGCCTGGTACCAGCAG AGGCCAGGACAGGCGCCCCGCCTGCTGATCTACGGTGCTTCCCAACGC GCCACTGGCATTCCTGACCGGTTCAGCGGCAGAGGGTCGGGAACCGAT TTCACACTGACCATTTCCCGGGTGGAGCCCGAAGATTCGGCAGTCTAC TACTGTCAGCATTACGAGTCCTCCCCTTCATGGACCTTCGGTCAAGGG ACCAAAGTGGAGATCAAG
BCMA_EBB-C1979-C12 - aa **VH**	1137	EVQLVESGGGLVQPGRSLRLSCTASGFTFDDYAMHWVRQRPGKGLEWV ASINWKGNSLAYGDSVKGRFAISRDNAKNTVFLQMNSLRTEDTAVYYC ASHQGVAYYNYAMDVWGRGTLVTVSS
BCMA_EBB-C1979-C12 - aa **VL**	1138	EIVLTQSPGTLSLSPGERATLSCRATQSIGSSFLAWYQQRPGQAPRLL IYGASQRATGIPDRFSGRGSGTDFTLTISRVEPEDSAVYYCQHYESSP SWTFGQGTKVEIK
BCMA_EBB-C1979-C12 - aa **Full CART**	1139	MALPVTALLLPLALLLHAARPEVQLVESGGGLVQPGRSLRLSCTASGF TFDDYAMHWVRQRPGKGLEWVASINWKGNSLAYGDSVKGRFAISRDNA KNTVFLQMNSLRTEDTAVYYCASHQGVAYYNYAMDVWGRGTLVTVSSG GGGSGGGGSGGGGSEIVLTQSPGTLSLSPGERATLSCRATQSIGSSFL AWYQQRPGQAPRLLIYGASQRATGIPDRFSGRGSGTDFTLTISRVEPE DSAVYYCQHYESSPSWTFGQGTKVEIKTTTPAPRPPTPAPTIASQPLS LRPEACRPAAGGAVHTRGLDFACDIYIWAPLAGTCGVLLLSLVITLYC KRGRKKLLYIFKQPFMRPVQTTQEEDGCSCRFPEEEEGGCELRVKFSR SADAPAYKQGQNQLYNELNLGRREEYDVLDKRRGRDPEMGGKPRRKNP QEGLYNELQKDKMAEAYSEIGMKGERRRGKGHDGLYQGLSTATKDTYD ALHMQALPPR
BCMA_EBB-C1979-C12 - nt **Full CART**	1140	ATGGCCCTCCCTGTCACCGCCCTGCTGCTTCCGCTGGCTCTTCTGCTC CACGCCGCTCGGCCCGAAGTGCAGCTCGTGGAGAGCGGGGGAGGATTG GTGCAGCCCGGAAGGTCCCTGCGGCTCTCCTGCACTGCGTCTGGCTTC ACCTTCGACGACTACGCGATGCACTGGGTCAGACAGCGCCCGGGAAAG GGCCTGGAATGGGTCGCCTCAATCAACTGGAAGGGAAACTCCCTGGCC TATGGCGACAGCGTGAAGGGCCGCTTCGCCATTTCGCGCGACAACGCC AAGAACACCGTGTTTCTGCAAATGAATTCCCTGCGGACCGAGGATACC GCTGTGTACTACTGCGCCAGCCACCAGGGCGTGGCATACTATAACTAC GCCATGGACGTGTGGGGAAGAGGGACGCTCGTCACCGTGTCCTCCGGG GGCGGTGGATCGGGTGGAGGAGGAAGCGGTGGCGGGGGCAGCGAAATC GTGCTGACTCAGAGCCCGGGAACTCTTTCACTGTCCCCGGGAGAACGG GCCACTCTCTCGTGCCGGGCCACCCAGTCCATCGGCTCCTCCTTCCTT

		GCCTGGTACCAGCAGAGGCCAGGACAGGCGCCCCGCCTGCTGATCTAC GGTGCTTCCCAACGCGCCACTGGCATTCCTGACCGGTTCAGCGGCAGA GGGTCGGGAACCGATTTCACACTGACCATTTCCCGGGTGGAGCCCGAA GATTCGGCAGTCTACTACTGTCAGCATTACGAGTCCTCCCCTTCATGG ACCTTCGGTCAAGGGACCAAAGTGGAGATCAAGACCACTACCCCAGCA CCGAGGCCACCCACCCCGGCTCCTACCATCGCCTCCCAGCCTCTGTCC CTGCGTCCGGAGGCATGTAGACCCGCAGCTGGTGGGGCCGTGCATACC CGGGGTCTTGACTTCGCCTGCGATATCTACATTTGGGCCCCTCTGGCT GGTACTTGCGGGGTCCTGCTGCTTTCACTCGTGATCACTCTTTACTGT AAGCGCGGTCGGAAGAAGCTGCTGTACATCTTTAAGCAACCCTTCATG AGGCCTGTGCAGACTACTCAAGAGGAGGACGGCTGTTCATGCCGGTTC CCAGAGGAGGAGGAAGGCGGCTGCGAACTGCGCGTGAAATTCAGCCGC AGCGCAGATGCTCCAGCCTACAAGCAGGGGCAGAACCAGCTCTACAAC GAACTCAATCTTGGTCGGAGAGAGGAGTACGACGTGCTGGACAAGCGG AGAGGACGGGACCCAGAAATGGGCGGGAAGCCGCGCAGAAAGAATCCC CAAGAGGGCCTGTACAACGAGCTCCAAAAGGATAAGATGGCAGAAGCC TATAGCGAGATTGGTATGAAAGGGGAACGCAGAAGAGGCAAAGGCCAC GACGGACTGTACCAGGGACTCAGCACCGCCACCAAGGACACCTATGAC GCTCTTCACATGCAGGCCCTGCCGCCTCGG
BCMA_EBB-C1980-G4		
BCMA_EBB-C1980-G4- aa **ScFv domain**	1141	EVQLVESGGGLVQPGGSLRLSCAASGFTFSSYAMSWVRQAPGKGLEWV SAISGSGGSTYYADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYC AKVVRDGMDVWGQGTTVTVSSGGGGSGGGGSGGGGSEIVLTQSPATLS LSPGERATLSCRASQSVSSSYLAWYQQKPGQAPRLLIYGASSRATGIP DRFSGNGSGTDFTLTISRLEPEDFAVYYCQQYGSPPRFTFGPGTKVDI K
BCMA_EBB-C1980-G4- nt **ScFv domain**	1142	GAGGTGCAGTTGGTCGAAAGCGGGGGCGGGCTTGTGCAGCCTGGCGGA TCACTGCGGCTGTCCTGCGCGGCATCAGGCTTCACGTTTTCTTCCTAC GCCATGTCCTGGGTGCGCCAGGCCCCTGGAAAGGGACTGGAATGGGTG TCCGCGATTTCGGGGTCCGGCGGGAGCACCTACTACGCCGATTCCGTG AAGGGCCGCTTCACTATCTCGCGGGACAACTCCAAGAACACCCTCTAC CTCCAAATGAATAGCCTGCGGGCCGAGGATACCGCCGTCTACTATTGC GCTAAGGTCGTGCGCGACGGAATGGACGTGTGGGGACAGGGTACCACC GTGACAGTGTCCTCGGGGGGAGGCGGTAGCGGCGGAGGAGGAAGCGGT GGTGGAGGTTCCGAGATTGTGCTGACTCAATCACCCGCGACCCTGAGC CTGTCCCCCGGCGAAAGGGCCACTCTGTCCTGTCGGGCCAGCCAATCA GTCTCCTCCTCGTACCTGGCCTGGTACCAGCAGAAGCCAGGACAGGCT CCGAGACTCCTTATCTATGGCGCATCCTCCCGCGCCACCGGAATCCCG GATAGGTTCTCGGGAAACGGATCGGGGACCGACTTCACTCTCACCATC TCCCGGCTGGAACCGGAGGACTTCGCCGTGTACTACTGCCAGCAGTAC GGCAGCCCGCCTAGATTCACTTTCGGCCCCGGCACCAAAGTGGACATC AAG
BCMA_EBB-	1143	EVQLVESGGGLVQPGGSLRLSCAASGFTFSSYAMSWVRQAPGKGLEWV SAISGSGGSTYYADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYC

C1980-G4- aa **VH**		AKVVRDGMDVWGQGTTVTVSS
BCMA_EBB-C1980-G4- aa **VL**	1144	EIVLTQSPATLSLSPGERATLSCRASQSVSSSYLAWYQQKPGQAPRLLIYGASSRATGIPDRFSGNGSGTDFTLTISRLEPEDFAVYYCQQYGSPPRFTFGPGTKVDIK
BCMA_EBB-C1980-G4- aa **Full CART**	1145	MALPVTALLLPLALLLHAARPEVQLVESGGGLVQPGGSLRLSCAASGFTFSSYAMSWVRQAPGKGLEWVSAISGSGGSTYYADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCAKVVRDGMDVWGQGTTVTVSSGGGGSGGGGSGGGGSEIVLTQSPATLSLSPGERATLSCRASQSVSSSYLAWYQQKPGQAPRLLIYGASSRATGIPDRFSGNGSGTDFTLTISRLEPEDFAVYYCQQYGSPPRFTFGPGTKVDIKTTTPAPRPPTPAPTIASQPLSLRPEACRPAAGGAVHTRGLDFACDIYIWAPLAGTCGVLLLSLVITLYCKRGRKKLLYIFKQPFMRPVQTTQEEDGCSCRFPEEEEGGCELRVKFSRSADAPAYKQGQNQLYNELNLGRREEYDVLDKRRGRDPEMGGKPRRKNPQEGLYNELQKDKMAEAYSEIGMKGERRRGKGHDGLYQGLSTATKDTYDALHMQALPPR
BCMA_EBB-C1980-G4- nt **Full CART**	1146	ATGGCCCTCCCTGTCACCGCCCTGCTGCTTCCGCTGGCTCTTCTGCTCCACGCCGCTCGGCCCGAGGTGCAGTTGGTCGAAAGCGGGGGCGGGCTTGTGCAGCCTGGCGGATCACTGCGGCTGTCCTGCGCGGCATCAGGCTTCACGTTTTCTTCCTACGCCATGTCCTGGGTGCGCCAGGCCCCTGGAAAGGGACTGGAATGGGTGTCCGCGATTTCGGGGTCCGGCGGGAGCACCTACTACGCCGATTCCGTGAAGGGCCGCTTCACTATCTCGCGGGACAACTCCAAGAACACCCTCTACCTCCAAATGAATAGCCTGCGGGCCGAGGATACCGCCGTCTACTATTGCGCTAAGGTCGTGCGCGACGGAATGGACGTGTGGGGACAGGGTACCACCGTGACAGTGTCCTCGGGGGGGAGGCGGTAGCGGCGGAGGAGGAAGCGGTGGTGGAGGTTCCGAGATTGTGCTGACTCAATCACCCGCGACCCTGAGCCTGTCCCCCGGCGAAAGGGCCACTCTGTCCTGTCGGGCCAGCCAATCAGTCTCCTCCTCGTACCTGGCCTGGTACCAGCAGAAGCCAGGACAGGCTCCGAGACTCCTTATCTATGGCGCATCCTCCCGCGCCACCGGAATCCCGGATAGGTTCTCGGGAAACGGATCGGGGACCGACTTCACTCTCACCATCTCCCGGCTGGAACCGGAGGACTTCGCCGTGTACTACTGCCAGCAGTACGGCAGCCCGCCTAGATTCACTTTCGGCCCCGGCACCAAAGTGGACATCAAGACCACTACCCCAGCACCGAGGCCACCCACCCCGGCTCCTACCATCGCCTCCCAGCCTCTGTCCCTGCGTCCGGAGGCATGTAGACCCGCAGCTGGTGGGGCCGTGCATACCCGGGGTCTTGACTTCGCCTGCGATATCTACATTTGGGCCCCTCTGGCTGGTACTTGCGGGGTCCTGCTGCTTTCACTCGTGATCACTCTTTACTGTAAGCGCGGTCGGAAGAAGCTGCTGTACATCTTTAAGCAACCCTTCATGAGGCCTGTGCAGACTACTCAAGAGGAGGACGGCTGTTCATGCCGGTTCCCAGAGGAGGAGGAAGGCGGCTGCGAACTGCGCGTGAAATTCAGCCGCAGCGCAGATGCTCCAGCCTACAAGCAGGGGCAGAACCAGCTCTACAACGAACTCAATCTTGGTCGGAGAGAGGAGTACGACGTGCTGGACAAGCGGAGAGGACGGGACCCAGAAATGGGCGGGAAGCCGCGCAGAAAGAATCCCCAAGAGGGCCTGTAC

		AACGAGCTCCAAAAGGATAAGATGGCAGAAGCCTATAGCGAGATTGGT ATGAAAGGGGAACGCAGAAGAGGCAAAGGCCACGACGGACTGTACCAG GGACTCAGCACCGCCACCAAGGACACCTATGACGCTCTTCACATGCAG GCCCTGCCGCCTCGG
BCMA_EBB-C1980-D2		
BCMA_EBB-C1980-D2- aa **ScFv domain**	1147	EVQLLESGGGLVQPGGSLRLSCAASGFTFSSYAMSWVRQAPGKGLEWV SAISGSGGSTYYADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYC AKIPQTGTFDYWGQGTLVTVSSGGGGSGGGGSGGGGSEIVLTQSPGTL SLSPGERATLSCRASQSVSSSYLAWYQQRPGQAPRLLIYGASSRATGI PDRFSGSGSGTDFTLTISRLEPEDFAVYYCQHYGSSPSWTFGQGTRLE IK
BCMA_EBB-C1980-D2- nt **ScFv domain**	1148	GAAGTGCAGCTGCTGGAGTCCGGCGGTGGATTGGTGCAACCGGGGGGA TCGCTCAGACTGTCCTGTGCGGCGTCAGGCTTCACCTTCTCGAGCTAC GCCATGTCATGGGTCAGACAGGCCCCTGGAAAGGGTCTGGAATGGGTG TCCGCCATTTCCGGGAGCGGGGGATCTACATACTACGCCGATAGCGTG AAGGGCCGCTTCACCATTTCCCGGGACAACTCCAAGAACACTCTCTAT CTGCAAATGAACTCCCTCCGCGCTGAGGACACTGCCGTGTACTACTGC GCCAAAATCCCTCAGACCGGCACCTTCGACTACTGGGGACAGGGGACT CTGGTCACCGTCAGCAGCGGTGGCGGAGGTTCGGGGGGAGGAGGAAGC GGCGGCGGAGGGTCCGAGATTGTGCTGACCCAGTCACCCGGCACTTTG TCCCTGTCGCCTGGAGAAAGGGCCACCCTTTCCTGCCGGGCATCCCAA TCCGTGTCCTCCTCGTACCTGGCCTGGTACCAGCAGAGGCCCGGACAG GCCCCACGGCTTCTGATCTACGGAGCAAGCAGCCGCGCGACCGGTATC CCGGACCGGTTTTCGGGCTCGGGCTCAGGAACTGACTTCACCCTCACC ATCTCCCGCCTGGAACCCGAAGATTTCGCTGTGTATTACTGCCAGCAC TACGGCAGCTCCCCGTCCTGGACGTTCGGCCAGGGAACTCGGCTGGAG ATCAAG
BCMA_EBB-C1980-D2- aa **VH**	1149	EVQLLESGGGLVQPGGSLRLSCAASGFTFSSYAMSWVRQAPGKGLEWV SAISGSGGSTYYADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYC AKIPQTGTFDYWGQGTLVTVSS
BCMA_EBB-C1980-D2- aa **VL**	1150	EIVLTQSPGTLSLSPGERATLSCRASQSVSSSYLAWYQQRPGQAPRLL IYGASSRATGIPDRFSGSGSGTDFTLTISRLEPEDFAVYYCQHYGSSP SWTFGQGTRLEIK
BCMA_EBB-C1980-D2- aa **Full CART**	1151	MALPVTALLLPLALLLHAARPEVQLLESGGGLVQPGGSLRLSCAASGF TFSSYAMSWVRQAPGKGLEWVSAISGSGGSTYYADSVKGRFTISRDNS KNTLYLQMNSLRAEDTAVYYCAKIPQTGTFDYWGQGTLVTVSSGGGGS GGGGSGGGGSEIVLTQSPGTLSLSPGERATLSCRASQSVSSSYLAWYQ QRPGQAPRLLIYGASSRATGIPDRFSGSGSGTDFTLTISRLEPEDFAV YYCQHYGSSPSWTFGQGTRLEIKTTTPAPRPPTPAPTIASQPLSLRPE ACRPAAGGAVHTRGLDFACDIYIWAPLAGTCGVLLLSLVITLYCKRGR

		KKLLYIFKQPFMRPVQTTQEEDGCSCRFPEEEEGGCELRVKFSRSADA PAYKQGQNQLYNELNLGRREEYDVLDKRRGRDPEMGGKPRRKNPQEGL YNELQKDKMAEAYSEIGMKGERRRGKGHDGLYQGLSTATKDTYDALHM QALPPR
BCMA_EBB-C1980-D2- nt **Full CART**	1152	ATGGCCCTCCCTGTCACCGCCCTGCTGCTTCCGCTGGCTCTTCTGCTC CACGCCGCTCGGCCCGAAGTGCAGCTGCTGGAGTCCGGCGGTGGATTG GTGCAACCGGGGGGATCGCTCAGACTGTCCTGTGCGGCGTCAGGCTTC ACCTTCTCGAGCTACGCCATGTCATGGGTCAGACAGGCCCCTGGAAAG GGTCTGGAATGGGTGTCCGCCATTTCCGGGAGCGGGGGATCTACATAC TACGCCGATAGCGTGAAGGGCCGCTTCACCATTTCCCGGGACAACTCC AAGAACACTCTCTATCTGCAAATGAACTCCCTCCGCGCTGAGGACACT GCCGTGTACTACTGCGCCAAAATCCCTCAGACCGGCACCTTCGACTAC TGGGGACAGGGGACTCTGGTCACCGTCAGCAGCGGTGGCGGAGGTTCG GGGGGAGGAGGAAGCGGCGGCGGAGGGTCCGAGATTGTGCTGACCCAG TCACCCGGCACTTTGTCCCTGTCGCCTGGAGAAAGGGCCACCCTTTCC TGCCGGGCATCCCAATCCGTGTCCTCCTCGTACCTGGCCTGGTACCAG CAGAGGCCCGGACAGGCCCCACGGCTTCTGATCTACGGAGCAAGCAGC CGCGCGACCGGTATCCCGGACCGGTTTTCGGGCTCGGGCTCAGGAACT GACTTCACCCTCACCATCTCCCGCCTGGAACCCGAAGATTTCGCTGTG TATTACTGCCAGCACTACGGCAGCTCCCCGTCCTGGACGTTCGGCCAG GGAACTCGGCTGGAGATCAAGACCACTACCCCAGCACCGAGGCCACCC ACCCCGGCTCCTACCATCGCCTCCCAGCCTCTGTCCCTGCGTCCGGAG GCATGTAGACCCGCAGCTGGTGGGGCCGTGCATACCCGGGGTCTTGAC TTCGCCTGCGATATCTACATTTGGGCCCCTCTGGCTGGTACTTGCGGG GTCCTGCTGCTTTCACTCGTGATCACTCTTTACTGTAAGCGCGGTCGG AAGAAGCTGCTGTACATCTTTAAGCAACCCTTCATGAGGCCTGTGCAG ACTACTCAAGAGGAGGACGGCTGTTCATGCCGGTTCCCAGAGGAGGAG GAAGGCGGCTGCGAACTGCGCGTGAAATTCAGCCGCAGCGCAGATGCT CCAGCCTACAAGCAGGGGCAGAACCAGCTCTACAACGAACTCAATCTT GGTCGGAGAGAGGAGTACGACGTGCTGGACAAGCGGAGAGGACGGGAC CCAGAAATGGGCGGGAAGCCGCGCAGAAAGAATCCCCAAGAGGGCCTG TACAACGAGCTCCAAAAGGATAAGATGGCAGAAGCCTATAGCGAGATT GGTATGAAAGGGGAACGCAGAAGAGGCAAAGGCCACGACGGACTGTAC CAGGGACTCAGCACCGCCACCAAGGACACCTATGACGCTCTTCACATG CAGGCCCTGCCGCCTCGG
BCMA_EBB-C1978-A10		
BCMA_EBB-C1978-A10-aa **ScFv domain**	1153	EVQLVETGGGLVQPGGSLRLSCAASGFTFSSYAMSWVRQAPGKGLEWV SAISGSGGSTYYADSVKGRFTMSRENDKNSVFLQMNSLRVEDTGVYYC ARANYKRELRYYYGMDVWGQGTMVTVSSGGGGSGGGGSGGGGSEIVMT QSPGTLSLSPGESATLSCRASQRVASNYLAWYQHKPGQAPSLLISGAS SRATGVPDRFSGSGSGTDFTLAISRLEPEDSAVYYCQHYDSSPSWTFG QGTKVEIK
BCMA_EBB-C1978-A10-	1154	GAAGTGCAACTGGTGGAAACCGGTGGAGGACTCGTGCAGCCTGGCGGC AGCCTCCGGCTGAGCTGCGCCGCTTCGGGATTCACCTTTTCCTCCTAC

nt **ScFv domain**		GCGATGTCTTGGGTCAGACAGGCCCCCGGAAAGGGGCTGGAATGGGTG TCAGCCATCTCCGGCTCCGGCGGATCAACGTACTACGCCGACTCCGTG AAAGGCCGGTTCACCATGTCGCGCGAGAATGACAAGAACTCCGTGTTC CTGCAAATGAACTCCCTGAGGGTGGAGGACACCGGAGTGTACTATTGT GCGCGCGCCAACTACAAGAGAGAGCTGCGGTACTACTACGGAATGGAC GTCTGGGGACAGGGAACTATGGTGACCGTGTCATCCGGTGGAGGGGGA AGCGGCGGTGGAGGCAGCGGGGGCGGGGGTTCAGAAATTGTCATGACC CAGTCCCCGGGAACTCTTTCCCTCTCCCCCGGGGAATCCGCGACTTTG TCCTGCCGGGCCAGCCAGCGCGTGGCCTCGAACTACCTCGCATGGTAC CAGCATAAGCCAGGCCAAGCCCCTTCCCTGCTGATTTCCGGGGCTAGC AGCCGCGCCACTGGCGTGCCGGATAGGTTCTCGGGAAGCGGCTCGGGT ACCGATTTCACCCTGGCAATCTCGCGGCTGGAACCGGAGGATTCGGCC GTGTACTACTGCCAGCACTATGACTCATCCCCCTCCTGGACATTCGGA CAGGGCACCAAGGTCGAGATCAAG
BCMA_EBB-C1978-A10-aa **VH**	1155	EVQLVETGGGLVQPGGSLRLSCAASGFTFSSYAMSWVRQAPGKGLEWV SAISGSGGSTYYADSVKGRFTMSRENDKNSVFLQMNSLRVEDTGVYYC ARANYKRELRYYYGMDVWGQGTMVTVSS
BCMA_EBB-C1978-A10-aa **VL**	1156	EIVMTQSPGTLSLSPGESATLSCRASQRVASNYLAWYQHKPGQAPSLL ISGASSRATGVPDRFSGSGSGTDFTLAISRLEPEDSAVYYCQHYDSSP SWTFGQGTKVEIK
BCMA_EBB-C1978-A10-aa **Full CART**	1157	MALPVTALLLPLALLLHAARPEVQLVETGGGLVQPGGSLRLSCAASGF TFSSYAMSWVRQAPGKGLEWVSAISGSGGSTYYADSVKGRFTMSREND KNSVFLQMNSLRVEDTGVYYCARANYKRELRYYYGMDVWGQGTMVTVS SGGGGSGGGGSGGGGSEIVMTQSPGTLSLSPGESATLSCRASQRVASN YLAWYQHKPGQAPSLLISGASSRATGVPDRFSGSGSGTDFTLAISRLE PEDSAVYYCQHYDSSPSWTFGQGTKVEIKTTTPAPRPPTPAPTIASQP LSLRPEACRPAAGGAVHTRGLDFACDIYIWAPLAGTCGVLLLSLVITL YCKRGRKKLLYIFKQPFMRPVQTTQEEDGCSCRFPEEEEGGCELRVKF SRSADAPAYKQGQNQLYNELNLGRREEYDVLDKRRGRDPEMGGKPRRK NPQEGLYNELQKDKMAEAYSEIGMKGERRRGKGHDGLYQGLSTATKDT YDALHMQALPPR
BCMA_EBB-C1978-A10-nt **Full CART**	1158	ATGGCCCTCCCTGTCACCGCCCTGCTGCTTCCGCTGGCTCTTCTGCTC CACGCCGCTCGGCCCGAAGTGCAACTGGTGGAAACCGGTGGAGGACTC GTGCAGCCTGGCGGCAGCCTCCGGCTGAGCTGCGCCGCTTCGGGATTC ACCTTTTCCTCCTACGCGATGTCTTGGGTCAGACAGGCCCCCGGAAAG GGGCTGGAATGGGTGTCAGCCATCTCCGGCTCCGGCGGATCAACGTAC TACGCCGACTCCGTGAAAGGCCGGTTCACCATGTCGCGCGAGAATGAC AAGAACTCCGTGTTCCTGCAAATGAACTCCCTGAGGGTGGAGGACACC GGAGTGTACTATTGTGCGCGCGCCAACTACAAGAGAGAGCTGCGGTAC TACTACGGAATGGACGTCTGGGGACAGGGAACTATGGTGACCGTGTCA

		TCCGGTGGAGGGGGAAGCGGCGGTGGAGGCAGCGGGGGCGGGGGTTCA GAAATTGTCATGACCCAGTCCCCGGGAACTCTTTCCCTCTCCCCCGGG GAATCCGCGACTTTGTCCTGCCGGGCCAGCCAGCGCGTGGCCTCGAAC TACCTCGCATGGTACCAGCATAAGCCAGGCCAAGCCCCTTCCCTGCTG ATTTCCGGGGCTAGCAGCCGCGCCACTGGCGTGCCGGATAGGTTCTCG GGAAGCGGCTCGGGTACCGATTTCACCCTGGCAATCTCGCGGCTGGAA CCGGAGGATTCGGCCGTGTACTACTGCCAGCACTATGACTCATCCCCC TCCTGGACATTCGGACAGGGCACCAAGGTCGAGATCAAGACCACTACC CCAGCACCGAGGCCACCCACCCCGGCTCCTACCATCGCCTCCCAGCCT CTGTCCCTGCGTCCGGAGGCATGTAGACCCGCAGCTGGTGGGGCCGTG CATACCCGGGGTCTTGACTTCGCCTGCGATATCTACATTTGGGCCCCT CTGGCTGGTACTTGCGGGGTCCTGCTGCTTTCACTCGTGATCACTCTT TACTGTAAGCGCGGTCGGAAGAAGCTGCTGTACATCTTTAAGCAACCC TTCATGAGGCCTGTGCAGACTACTCAAGAGGAGGACGGCTGTTCATGC CGGTTCCCAGAGGAGGAGGAAGGCGGCTGCGAACTGCGCGTGAAATTC AGCCGCAGCGCAGATGCTCCAGCCTACAAGCAGGGGCAGAACCAGCTC TACAACGAACTCAATCTTGGTCGGAGAGAGGAGTACGACGTGCTGGAC AAGCGGAGAGGACGGGACCCAGAAATGGGCGGGAAGCCGCGCAGAAAG AATCCCCAAGAGGGCCTGTACAACGAGCTCCAAAAGGATAAGATGGCA GAAGCCTATAGCGAGATTGGTATGAAAGGGGAACGCAGAAGAGGCAAA GGCCACGACGGACTGTACCAGGGACTCAGCACCGCCACCAAGGACACC TATGACGCTCTTCACATGCAGGCCCTGCCGCCTCGG
BCMA_EBB-C1978-D4		
BCMA_EBB-C1978-D4- aa **ScFv domain**	1159	EVQLLETGGGLVQPGGSLRLSCAASGFSFSSYAMSWVRQAPGKGLEWV SAISGSGGSTYYADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYC AKALVGATGAFDIWGQGTLVTVSSGGGGSGGGGSGGGGSEIVLTQSPG TLSLSPGERATLSCRASQSLSSNFLAWYQQKPGQAPGLLIYGASNWAT GTPDRFSGSGSGTDFTLTITRLEPEDFAVYYCQYYGTSPMYTFGQGTK VEIK
BCMA_EBB-C1978-D4- nt **ScFv domain**	1160	GAAGTGCAGCTGCTCGAAACCGGTGGAGGGCTGGTGCAGCCAGGGGGC TCCCTGAGGCTTTCATGCGCCGCTAGCGGATTCTCCTTCTCCTCTTAC GCCATGTCGTGGGTCCGCCAAGCCCCTGGAAAAGGCCTGGAATGGGTG TCCGCGATTTCCGGGAGCGGAGGTTCGACCTATTACGCCGACTCCGTG AAGGGCCGCTTTACCATCTCCCGGGATAACTCCAAGAACACTCTGTAC CTCCAAATGAACTCGCTGAGAGCCGAGGACACCGCCGTGTATTACTGC GCGAAGGCGCTGGTCGGCGCGACTGGGGCATTCGACATCTGGGGACAG GGAACTCTTGTGACCGTGTCGAGCGGAGGCGGCGGCTCCGGCGGAGGA GGGAGCGGGGGCGGTGGTTCCGAAATCGTGTTGACTCAGTCCCCGGGA ACCCTGAGCTTGTCACCCGGGGAGCGGGCCACTCTCTCCTGTCGCGCC TCCCAATCGCTCTCATCCAATTTCCTGGCCTGGTACCAGCAGAAGCCC GGACAGGCCCCGGGCCTGCTCATCTACGGCGCTTCAAACTGGGCAACG GGAACCCCTGATCGGTTCAGCGGAAGCGGATCGGGTACTGACTTTACC CTGACCATCACCAGACTGGAACCGGAGGACTTCGCCGTGTACTACTGC CAGTACTACGGCACCTCCCCCATGTACACATTCGGACAGGGTACCAAG

		GTCGAGATTAAG
BCMA_EBB-C1978-D4- aa **VH**	1161	EVQLLETGGGLVQPGGSLRLSCAASGFSFSSYAMSWVRQAPGKGLEWV SAISGSGGSTYYADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYC AKALVGATGAFDIWGQGTLVTVSS
BCMA_EBB-C1978-D4- aa **VL**	1162	EIVLTQSPGTLSLSPGERATLSCRASQSLSSNFLAWYQQKPGQAPGLL IYGASNWATGTPDRFSGSGSGTDFTLTITRLEPEDFAVYYCQYYGTSP MYTFGQGTKVEIK
BCMA_EBB-C1978-D4- aa **Full CART**	1163	MALPVTALLLPLALLLHAARPEVQLLETGGGLVQPGGSLRLSCAASGF SFSSYAMSWVRQAPGKGLEWVSAISGSGGSTYYADSVKGRFTISRDNS KNTLYLQMNSLRAEDTAVYYCAKALVGATGAFDIWGQGTLVTVSSGGG GSGGGGSGGGGSEIVLTQSPGTLSLSPGERATLSCRASQSLSSNFLAW YQQKPGQAPGLLIYGASNWATGTPDRFSGSGSGTDFTLTITRLEPEDF AVYYCQYYGTSPMYTFGQGTKVEIKTTTPAPRPPTPAPTIASQPLSLR PEACRPAAGGAVHTRGLDFACDIYIWAPLAGTCGVLLLSLVITLYCKR GRKKLLYIFKQPFMRPVQTTQEEDGCSCRFPEEEEGGCELRVKFSRSA DAPAYKQGQNQLYNELNLGRREEYDVLDKRRGRDPEMGGKPRRKNPQE GLYNELQKDKMAEAYSEIGMKGERRRGKGHDGLYQGLSTATKDTYDAL HMQALPPR
BCMA_EBB-C1978-D4- nt **Full CART**	1164	ATGGCCCTCCCTGTCACCGCCCTGCTGCTTCCGCTGGCTCTTCTGCTC CACGCCGCTCGGCCCGAAGTGCAGCTGCTCGAAACCGGTGGAGGGCTG GTGCAGCCAGGGGGCTCCCTGAGGCTTTCATGCGCCGCTAGCGGATTC TCCTTCTCCTCTTACGCCATGTCGTGGGTCCGCCAAGCCCCTGGAAAA GGCCTGGAATGGGTGTCCGCGATTTCCGGGAGCGGAGGTTCGACCTAT TACGCCGACTCCGTGAAGGGCCGCTTTACCATCTCCCGGGATAACTCC AAGAACACTCTGTACCTCCAAATGAACTCGCTGAGAGCCGAGGACACC GCCGTGTATTACTGCGCGAAGGCGCTGGTCGGCGCGACTGGGGCATTC GACATCTGGGGACAGGGAACTCTTGTGACCGTGTCGAGCGGAGGCGGC GGCTCCGGCGGAGGAGGGAGCGGGGGCGGTGGTTCCGAAATCGTGTTG ACTCAGTCCCCGGGAACCCTGAGCTTGTCACCCGGGGAGCGGGCCACT CTCTCCTGTCGCGCCTCCCAATCGCTCTCATCCAATTTCCTGGCCTGG TACCAGCAGAAGCCCGGACAGGCCCCGGGCCTGCTCATCTACGGCGCT TCAAACTGGGCAACGGGAACCCCTGATCGGTTCAGCGGAAGCGGATCG GGTACTGACTTTACCCTGACCATCACCAGACTGGAACCGGAGGACTTC GCCGTGTACTACTGCCAGTACTACGGCACCTCCCCCATGTACACATTC GGACAGGGTACCAAGGTCGAGATTAAGACCACTACCCCAGCACCGAGG CCACCCACCCCCGGCTCCTACCATCGCCTCCCAGCCTCTGTCCCTGCGT CCGGAGGCATGTAGACCCGCAGCTGGTGGGGCCGTGCATACCCGGGGT CTTGACTTCGCCTGCGATATCTACATTTGGGCCCCTCTGGCTGGTACT TGCGGGGTCCTGCTGCTTTCACTCGTGATCACTCTTTACTGTAAGCGC GGTCGGAAGAAGCTGCTGTACATCTTTAAGCAACCCTTCATGAGGCCT GTGCAGACTACTCAAGAGGAGGACGGCTGTTCATGCCGGTTCCCAGAG GAGGAGGAAGGCGGCTGCGAACTGCGCGTGAAATTCAGCCGCAGCGCA

		GATGCTCCAGCCTACAAGCAGGGGCAGAACCAGCTCTACAACGAACTC AATCTTGGTCGGAGAGAGGAGTACGACGTGCTGGACAAGCGGAGAGGA CGGGACCCAGAAATGGGCGGGAAGCCGCGCAGAAAGAATCCCCAAGAG GGCCTGTACAACGAGCTCCAAAAGGATAAGATGGCAGAAGCCTATAGC GAGATTGGTATGAAAGGGGAACGCAGAAGAGGCAAAGGCCACGACGGA CTGTACCAGGGACTCAGCACCGCCACCAAGGACACCTATGACGCTCTT CACATGCAGGCCCTGCCGCCTCGG
BCMA_EBB-C1980-A2		
BCMA_EBB-C1980-A2- aa **ScFv domain**	1165	EVQLLESGGGLVQPGGSLRLSCAASGFTFSSYAMSWVRQAPGKGLEWV SAISGSGGSTYYADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYC VLWFGEGFDPWGQGTLVTVSSGGGGSGGGGSGGGGSDIVLTQSPLSLP VTPGEPASISCRSSQSLLHSNGYNYLDWYLQKPGQSPQLLIYLGSNRA SGVPDRFSGSGSGTDFTLKISRVEAEDVGVYYCMQALQTPLTFGGGTK VDIK
BCMA_EBB-C1980-A2- nt **ScFv domain**	1166	GAAGTGCAGCTGCTTGAGAGCGGTGGAGGTCTGGTGCAGCCCGGGGGA TCACTGCGCCTGTCCTGTGCCGCGTCCGGTTTCACTTTCTCCTCGTAC GCCATGTCGTGGGTCAGACAGGCACCGGGAAAGGGACTGGAATGGGTG TCAGCCATTTCGGGTTCGGGGGGCAGCACCTACTACGCTGACTCCGTG AAGGGCCGGTTCACCATTTCCCGCGACAACTCCAAGAACACCTTGTAC CTCCAAATGAACTCCCTGCGGGCCGAAGATACCGCCGTGTATTACTGC GTGCTGTGGTTCGGAGAGGGATTCGACCCGTGGGGACAAGGAACACTC GTGACTGTGTCATCCGGCGGAGGCGGCAGCGGTGGCGGCGGTTCCGGC GGCGGCGGATCTGACATCGTGTTGACCCAGTCCCCTCTGAGCCTGCCG GTCACTCCTGGCGAACCAGCCAGCATCTCCTGCCGGTCGAGCCAGTCC CTCCTGCACTCCAATGGGTACAACTACCTCGATTGGTATCTGCAAAAG CCGGGCCAGAGCCCCCAGCTGCTGATCTACCTTGGGTCAAACCGCGCT TCCGGGGTGCCTGATAGATTCTCCGGGTCCGGGAGCGGAACCGACTTT ACCCTGAAAATCTCGAGGGTGGAGGCCGAGGACGTCGGAGTGTACTAC TGCATGCAGGCGCTCCAGACTCCCCTGACCTTCGGAGGAGGAACGAAG GTCGACATCAAGA
BCMA_EBB-C1980-A2- aa **VH**	1167	EVQLLESGGGLVQPGGSLRLSCAASGFTFSSYAMSWVRQAPGKGLEWV SAISGSGGSTYYADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYC VLWFGEGFDPWGQGTLVTVSS
BCMA_EBB-C1980-A2- aa **VL**	1168	DIVLTQSPLSLPVTPGEPASISCRSSQSLLHSNGYNYLDWYLQKPGQS PQLLIYLGSNRASGVPDRFSGSGSGTDFTLKISRVEAEDVGVYYCMQA LQTPLTFGGGTKVDIK
BCMA_EBB-C1980-A2- aa **Full CART**	1169	MALPVTALLLPLALLLHAARPEVQLLESGGGLVQPGGSLRLSCAASGF TFSSYAMSWVRQAPGKGLEWVSAISGSGGSTYYADSVKGRFTISRDNS KNTLYLQMNSLRAEDTAVYYCVLWFGEGFDPWGQGTLVTVSSGGGGSG GGGSGGGGSDIVLTQSPLSLPVTPGEPASISCRSSQSLLHSNGYNYLD

		WYLQKPGQSPQLLIYLGSNRASGVPDRFSGSGSGTDFTLKISRVEAED VGVYYCMQALQTPLTFGGGTKVDIKTTTPAPRPPTPAPTIASQPLSLR PEACRPAAGGAVHTRGLDFACDIYIWAPLAGTCGVLLLSLVITLYCKR GRKKLLYIFKQPFMRPVQTTQEEDGCSCRFPEEEEGGCELRVKFSRSA DAPAYKQGQNQLYNELNLGRREEYDVLDKRRGRDPEMGGKPRRKNPQE GLYNELQKDKMAEAYSEIGMKGERRRGKGHDGLYQGLSTATKDTYDAL HMQALPPR
BCMA_EBB-C1980-A2- nt **Full CART**	1170	ATGGCCCTCCCTGTCACCGCCCTGCTGCTTCCGCTGGCTCTTCTGCTC CACGCCGCTCGGCCCGAAGTGCAGCTGCTTGAGAGCGGTGGAGGTCTG GTGCAGCCCGGGGGATCACTGCGCCTGTCCTGTGCCGCGTCCGGTTTC ACTTTCTCCTCGTACGCCATGTCGTGGGTCAGACAGGCACCGGGAAAG GGACTGGAATGGGTGTCAGCCATTTCGGGTTCGGGGGGCAGCACCTAC TACGCTGACTCCGTGAAGGGCCGGTTCACCATTTCCCGCGACAACTCC AAGAACACCTTGTACCTCCAAATGAACTCCCTGCGGGCCGAAGATACC GCCGTGTATTACTGCGTGCTGTGGTTCGGAGAGGGATTCGACCCGTGG GGACAAGGAACACTCGTGACTGTGTCATCCGGCGGAGGCGGCAGCGGT GGCGGCGGTTCCGGCGGCGGCGGATCTGACATCGTGTTGACCCAGTCC CCTCTGAGCCTGCCGGTCACTCCTGGCGAACCAGCCAGCATCTCCTGC CGGTCGAGCCAGTCCCTCCTGCACTCCAATGGGTACAACTACCTCGAT TGGTATCTGCAAAAGCCGGGCCAGAGCCCCCAGCTGCTGATCTACCTT GGGTCAAACCGCGCTTCCGGGGTGCCTGATAGATTCTCCGGGTCCGGG AGCGGAACCGACTTTACCCTGAAAATCTCGAGGGTGGAGGCCGAGGAC GTCGGAGTGTACTACTGCATGCAGGCGCTCCAGACTCCCCTGACCTTC GGAGGAGGAACGAAGGTCGACATCAAGACCACTACCCCAGCACCGAGG CCACCCACCCCGGCTCCTACCATCGCCTCCCAGCCTCTGTCCCTGCGT CCGGAGGCATGTAGACCCGCAGCTGGTGGGGCCGTGCATACCCGGGGT CTTGACTTCGCCTGCGATATCTACATTTGGGCCCCTCTGGCTGGTACT TGCGGGGTCCTGCTGCTTTCACTCGTGATCACTCTTTACTGTAAGCGC GGTCGGAAGAAGCTGCTGTACATCTTTAAGCAACCCTTCATGAGGCCT GTGCAGACTACTCAAGAGGAGGACGGCTGTTCATGCCGGTTCCCAGAG GAGGAGGAAGGCGGCTGCGAACTGCGCGTGAAATTCAGCCGCAGCGCA GATGCTCCAGCCTACAAGCAGGGGCAGAACCAGCTCTACAACGAACTC AATCTTGGTCGGAGAGAGGAGTACGACGTGCTGGACAAGCGGAGAGGA CGGGACCCAGAAATGGGCGGGAAGCCGCGCAGAAAGAATCCCCAAGAG GGCCTGTACAACGAGCTCCAAAAGGATAAGATGGCAGAAGCCTATAGC GAGATTGGTATGAAAGGGGAACGCAGAAGAGGCAAAGGCCACGACGGA CTGTACCAGGGACTCAGCACCGCCACCAAGGACACCTATGACGCTCTT CACATGCAGGCCCTGCCGCCTCGG
BCMA_EBB-C1981-C3		
BCMA_EBB-C1981-C3- aa **ScFv domain**	1171	QVQLVESGGGLVQPGGSLRLSCAASGFTFSSYAMSWVRQAPGKGLEWV SAISGSGGSTYYADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYC AKVGYDSSGYYRDYYGMDVWGQGTTVTVSSGGGGSGGGGSGGGGSEIV LTQSPGTLSLSPGERATLSCRASQSVSSSYLAWYQQKPGQAPRLLIYG TSSRATGISDRFSGSGSGTDFTLTISRLEPEDFAVYYCQHYGNSPPKF

		TFGPGTKLEIK
BCMA_EBB-C1981-C3- nt **ScFv domain**	1172	CAAGTGCAGCTCGTGGAGTCAGGCGGAGGACTGGTGCAGCCCGGGGGC TCCCTGAGACTTTCCTGCGCGGCATCGGGTTTTACCTTCTCCTCCTAT GCTATGTCCTGGGTGCGCCAGGCCCCGGGAAAGGGACTGGAATGGGTG TCCGCAATCAGCGGTAGCGGGGGCTCAACATACTACGCCGACTCCGTC AAGGGTCGCTTCACTATTTCCCGGGACAACTCCAAGAATACCCTGTAC CTCCAAATGAACAGCCTCAGGGCCGAGGATACTGCCGTGTACTACTGC GCCAAAGTCGGATACGATAGCTCCGGTTACTACCGGGACTACTACGGA ATGGACGTGTGGGGACAGGGCACCACCGTGACCGTGTCAAGCGGCGGA GGCGGTTCAGGAGGGGGAGGCTCCGGCGGTGGAGGGTCCGAAATCGTC CTGACTCAGTCGCCTGGCACTCTGTCGTTGTCCCCGGGGGAGCGCGCT ACCCTGTCGTGTCGGGCGTCGCAGTCCGTGTCGAGCTCCTACCTCGCG TGGTACCAGCAGAAGCCCGGACAGGCCCCTAGACTTCTGATCTACGGC ACTTCTTCACGCGCCACCGGGATCAGCGACAGGTTCAGCGGCTCCGGC TCCGGGACCGACTTCACCCTGACCATTAGCCGGCTGGAGCCTGAAGAT TTCGCCGTGTATTACTGCCAACACTACGGAAACTCGCCGCCAAAGTTC ACGTTCGGACCCGGAACCAAGCTGGAAATCAAG
BCMA_EBB-C1981-C3- aa **VH**	1173	QVQLVESGGGLVQPGGSLRLSCAASGFTFSSYAMSWVRQAPGKGLEWV SAISGSGGSTYYADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYC AKVGYDSSGYYRDYYGMDVWGQGTTVTVSS
BCMA_EBB-C1981-C3- aa **VL**	1174	EIVLTQSPGTLSLSPGERATLSCRASQSVSSSYLAWYQQKPGQAPRLL IYGTSSRATGISDRFSGSGSGTDFTLTISRLEPEDFAVYYCQHYGNSP PKFTFGPGTKLEIK
BCMA_EBB-C1981-C3- aa **Full CART**	1175	MALPVTALLLPLALLLHAARPQVQLVESGGGLVQPGGSLRLSCAASGF TFSSYAMSWVRQAPGKGLEWVSAISGSGGSTYYADSVKGRFTISRDNS KNTLYLQMNSLRAEDTAVYYCAKVGYDSSGYYRDYYGMDVWGQGTTVT VSSGGGGSGGGGSGGGGSEIVLTQSPGTLSLSPGERATLSCRASQSVS SSYLAWYQQKPGQAPRLLIYGTSSRATGISDRFSGSGSGTDFTLTISR LEPEDFAVYYCQHYGNSPPKFTFGPGTKLEIKTTTPAPRPPTPAPTIA SQPLSLRPEACRPAAGGAVHTRGLDFACDIYIWAPLAGTCGVLLLSLV ITLYCKRGRKKLLYIFKQPFMRPVQTTQEEDGCSCRFPEEEEGGCELR VKFSRSADAPAYKQGQNQLYNELNLGRREEYDVLDKRRGRDPEMGGKP RRKNPQEGLYNELQKDKMAEAYSEIGMKGERRRGKGHDGLYQGLSTAT KDTYDALHMQALPPR
BCMA_EBB-C1981-C3- nt **Full CART**	1176	ATGGCCCTCCCTGTCACCGCCCTGCTGCTTCCGCTGGCTCTTCTGCTC CACGCCGCTCGGCCCCAAGTGCAGCTCGTGGAGTCAGGCGGAGGACTG GTGCAGCCCGGGGGCTCCCTGAGACTTTCCTGCGCGGCATCGGGTTTT ACCTTCTCCTCCTATGCTATGTCCTGGGTGCGCCAGGCCCCGGGAAAG GGACTGGAATGGGTGTCCGCAATCAGCGGTAGCGGGGGCTCAACATAC

		TACGCCGACTCCGTCAAGGGTCGCTTCACTATTTCCCGGGACAACTCC AAGAATACCCTGTACCTCCAAATGAACAGCCTCAGGGCCGAGGATACT GCCGTGTACTACTGCGCCAAAGTCGGATACGATAGCTCCGGTTACTAC CGGGACTACTACGGAATGGACGTGTGGGGACAGGGCACCACCGTGACC GTGTCAAGCGGCGGAGGCGGTTCAGGAGGGGGAGGCTCCGGCGGTGGA GGGTCCGAAATCGTCCTGACTCAGTCGCCTGGCACTCTGTCGTTGTCC CCGGGGGAGCGCGCTACCCTGTCGTGTCGGGCGTCGCAGTCCGTGTCG AGCTCCTACCTCGCGTGGTACCAGCAGAAGCCCGGACAGGCCCCTAGA CTTCTGATCTACGGCACTTCTTCACGCGCCACCGGGATCAGCGACAGG TTCAGCGGCTCCGGCTCCGGGACCGACTTCACCCTGACCATTAGCCGG CTGGAGCCTGAAGATTTCGCCGTGTATTACTGCCAACACTACGGAAAC TCGCCGCCAAAGTTCACGTTCGGACCCGGAACCAAGCTGGAAATCAAG ACCACTACCCCAGCACCGAGGCCACCCACCCCGGCTCCTACCATCGCC TCCCAGCCTCTGTCCCTGCGTCCGGAGGCATGTAGACCCGCAGCTGGT GGGGCCGTGCATACCCGGGGTCTTGACTTCGCCTGCGATATCTACATT TGGGCCCCTCTGGCTGGTACTTGCGGGGTCCTGCTGCTTTCACTCGTG ATCACTCTTTACTGTAAGCGCGGTCGGAAGAAGCTGCTGTACATCTTT AAGCAACCCTTCATGAGGCCTGTGCAGACTACTCAAGAGGAGGACGGC TGTTCATGCCGGTTCCCAGAGGAGGAGGAAGGCGGCTGCGAACTGCGC GTGAAATTCAGCCGCAGCGCAGATGCTCCAGCCTACAAGCAGGGGCAG AACCAGCTCTACAACGAACTCAATCTTGGTCGGAGAGAGGAGTACGAC GTGCTGGACAAGCGGAGAGGACGGGACCCAGAAATGGGCGGGAAGCCG CGCAGAAAGAATCCCCAAGAGGGCCTGTACAACGAGCTCCAAAAGGAT AAGATGGCAGAAGCCTATAGCGAGATTGGTATGAAAGGGGAACGCAGA AGAGGCAAAGGCCACGACGGACTGTACCAGGGACTCAGCACCGCCACC AAGGACACCTATGACGCTCTTCACATGCAGGCCCTGCCGCCTCGG
BCMA_EBB-C1978-G4		
BCMA_EBB-C1978-G4- aa **ScFv domain**	1177	EVQLVESGGGLVQPGGSLRLSCAASGFTFSSYAMSWVRQAPGKGLEWV SAISGSGGSTYYADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYC AKMGWSSGYLGAFDIWGQGTTVTVSSGGGGSGGGGSGGGGSEIVLTQS PGTLSLSPGERATLSCRASQSVASSFLAWYQQKPGQAPRLLIYGASGR ATGIPDRFSGSGSGTDFTLTISRLEPEDFAVYYCQHYGGSPRLTFGGG TKVDIK
BCMA_EBB-C1978-G4- nt **ScFv domain**	1178	GAAGTCCAACTGGTGGAGTCCGGGGGAGGGCTCGTGCAGCCCGGAGGC AGCCTTCGGCTGTCGTGCGCCGCCTCCGGGTTCACGTTCTCATCCTAC GCGATGTCGTGGGTCAGACAGGCACCAGGAAAGGGACTGGAATGGGTG TCCGCCATTAGCGGCTCCGGCGGTAGCACCTACTATGCCGACTCAGTG AAGGGAAGGTTCACTATCTCCCGCGACAACAGCAAGAACACCCTGTAC CTCCAAATGAACTCTCTGCGGGCCGAGGATACCGCGGTGTACTATTGC GCCAAGATGGGTTGGTCCAGCGGATACTTGGGAGCCTTCGACATTTGG GGACAGGGCACTACTGTGACCGTGTCCTCCGGGGGTGGCGGATCGGGA GGCGGCGGCTCGGGTGGAGGGGGTTCCGAAATCGTGTTGACCCAGTCA CCGGGAACCCTCTCGCTGTCCCCGGGAGAACGGGCTACACTGTCATGT AGAGCGTCCCAGTCCGTGGCTTCCTCGTTCCTGGCCTGGTACCAGCAG AAGCCGGGACAGGCACCCCGCCTGCTCATCTACGGAGCCAGCGGCCGG

		GCGACCGGCATCCCTGACCGCTTCTCCGGTTCCGGCTCGGGCACCGAC TTTACTCTGACCATTAGCAGGCTTGAGCCCGAGGATTTTGCCGTGTAC TACTGCCAACACTACGGGGGGAGCCCTCGCCTGACCTTCGGAGGCGGA ACTAAGGTCGATATCAAAA
BCMA_EBB-C1978-G4- aa **VH**	1179	EVQLVESGGGLVQPGGSLRLSCAASGFTFSSYAMSWVRQAPGKGLEWV SAISGSGGSTYYADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYC AKMGWSSGYLGAFDIWGQGTTVTVSS
BCMA_EBB-C1978-G4- aa **VL**	1180	EIVLTQSPGTLSLSPGERATLSCRASQSVASSFLAWYQQKPGQAPRLL IYGASGRATGIPDRFSGSGSGTDFTLTISRLEPEDFAVYYCQHYGGSP RLTFGGGTKVDIK
BCMA_EBB-C1978-G4- aa **Full CART**	1181	MALPVTALLLPLALLLHAARPEVQLVESGGGLVQPGGSLRLSCAASGF TFSSYAMSWVRQAPGKGLEWVSAISGSGGSTYYADSVKGRFTISRDNS KNTLYLQMNSLRAEDTAVYYCAKMGWSSGYLGAFDIWGQGTTVTVSSG GGGSGGGGSGGGGSEIVLTQSPGTLSLSPGERATLSCRASQSVASSFL AWYQQKPGQAPRLLIYGASGRATGIPDRFSGSGSGTDFTLTISRLEPE DFAVYYCQHYGGSPRLTFGGGTKVDIKTTTPAPRPPTPAPTIASQPLS LRPEACRPAAGGAVHTRGLDFACDIYIWAPLAGTCGVLLLSLVITLYC KRGRKKLLYIFKQPFMRPVQTTQEEDGCSCRFPEEEEGGCELRVKFSR SADAPAYKQGQNQLYNELNLGRREEYDVLDKRRGRDPEMGGKPRRKNP QEGLYNELQKDKMAEAYSEIGMKGERRRGKGHDGLYQGLSTATKDTYD ALHMQALPPR
BCMA_EBB-C1978-G4- nt **Full CART**	1182	ATGGCCCTCCCTGTCACCGCCCTGCTGCTTCCGCTGGCTCTTCTGCTC CACGCCGCTCGGCCCGAAGTCCAACTGGTGGAGTCCGGGGGAGGCTC GTGCAGCCCGGAGGCAGCCTTCGGCTGTCGTGCGCCGCCTCCGGGTTC ACGTTCTCATCCTACGCGATGTCGTGGGTCAGACAGGCACCAGGAAAG GGACTGGAATGGGTGTCCGCCATTAGCGGCTCCGGCGGTAGCACCTAC TATGCCGACTCAGTGAAGGGAAGGTTCACTATCTCCCGCGACAACAGC AAGAACACCCTGTACCTCCAAATGAACTCTCTGCGGGCCGAGGATACC GCGGTGTACTATTGCGCCAAGATGGGTTGGTCCAGCGGATACTTGGGA GCCTTCGACATTTGGGGACAGGGCACTACTGTGACCGTGTCCTCCGGG GGTGGCGGATCGGGAGGCGGCGGCTCGGGTGGAGGGGGTTCCGAAATC GTGTTGACCCAGTCACCGGGAACCCTCTCGCTGTCCCCGGGAGAACGG GCTACACTGTCATGTAGAGCGTCCCAGTCCGTGGCTTCCTCGTTCCTG GCCTGGTACCAGCAGAAGCCGGGACAGGCACCCCGCCTGCTCATCTAC GGAGCCAGCGGCCGGGCGACCGGCATCCCTGACCGCTTCTCCGGTTCC GGCTCGGGCACCGACTTTACTCTGACCATTAGCAGGCTTGAGCCCGAG GATTTTGCCGTGTACTACTGCCAACACTACGGGGGGAGCCCTCGCCTG ACCTTCGGAGGCGGAACTAAGGTCGATATCAAAACCACTACCCCAGCA CCGAGGCCACCCACCCCGGCTCCTACCATCGCCTCCCAGCCTCTGTCC CTGCGTCCGGAGGCATGTAGACCCGCAGCTGGTGGGGCCGTGCATACC CGGGGTCTTGACTTCGCCTGCGATATCTACATTTGGGCCCCTCTGGCT GGTACTTGCGGGGTCCTGCTGCTTTCACTCGTGATCACTCTTTACTGT

		AAGCGCGGTCGGAAGAAGCTGCTGTACATCTTTAAGCAACCCTTCATG AGGCCTGTGCAGACTACTCAAGAGGAGGACGGCTGTTCATGCCGGTTC CCAGAGGAGGAGGAAGGCGGCTGCGAACTGCGCGTGAAATTCAGCCGC AGCGCAGATGCTCCAGCCTACAAGCAGGGGCAGAACCAGCTCTACAAC GAACTCAATCTTGGTCGGAGAGAGGAGTACGACGTGCTGGACAAGCGG AGAGGACGGGACCCAGAAATGGGCGGGAAGCCGCGCAGAAAGAATCCC CAAGAGGGCCTGTACAACGAGCTCCAAAAGGATAAGATGGCAGAAGCC TATAGCGAGATTGGTATGAAAGGGGAACGCAGAAGAGGCAAAGGCCAC GACGGACTGTACCAGGGACTCAGCACCGCCACCAAGGACACCTATGAC GCTCTTCACATGCAGGCCCTGCCGCCTCGG

Table 11B. Additional exemplary BCMA CAR sequences

Name	Sequence	SEQ ID NO:
A7D12.2 VH	QIQLVQSGPDLKKPGETVKLSCKASGYTFTNFGMNWVKQAPGKGFKWMAWINTYTGESYFA DDFKGRFAFSVETSATTAYLQINNLKTEDTATYFCARGEIYYGYDGGFAYWGQGTLVTVSA	1183
A7D12.2 VL	DVVMTQSHRFMSTSVGDRVSITCRASQDVNTAVSWYQQKPGQSPKLLIFSASYRYTGVPDR FTGSGSGADFTLTISSVQAEDLAVYYCQQHYSTPWTFGGGTKLDIK	1184
A7D12.2 scFv domain	QIQLVQSGPDLKKPGETVKLSCKASGYTFTNFGMNWVKQAPGKGFKWMAWINTYTGESYFA DDFKGRFAFSVETSATTAYLQINNLKTEDTATYFCARGEIYYGYDGGFAYWGQGTLVTVSA GGGGSGGGGSGGGGSDVVMTQSHRFMSTSVGDRVSITCRASQDVNTAVSWYQQKPGQSPKL LIFSASYRYTGVPDRFTGSGSGADFTLTISSVQAEDLAVYYCQQHYSTPWTFGGGTKLDIK	1185
A7D12.2 Full CART	QIQLVQSGPDLKKPGETVKLSCKASGYTFTNFGMNWVKQAPGKGFKWMAWINTYTGESYFA DDFKGRFAFSVETSATTAYLQINNLKTEDTATYFCARGEIYYGYDGGFAYWGQGTLVTVSA GGGGSGGGGSGGGGSDVVMTQSHRFMSTSVGDRVSITCRASQDVNTAVSWYQQKPGQSPKL LIFSASYRYTGVPDRFTGSGSGADFTLTISSVQAEDLAVYYCQQHYSTPWTFGGGTKLDIK TTTPAPRPPTPAPTIASQPLSLRPEACRPAAGGAVHTRGLDFACDIYIWAPLAGTCGVLLL SLVITLYCKRGRKKLLYIFKQPFMRPVQTTQEEDGCSCRFPEEEEGGCELRVKFSRSADAP AYKQGQNQLYNELNLGRREEYDVLDKRRGRDPEMGGKPRRKNPQEGLYNELQKDKMAEAYS EIGMKGERRRGKGHDGLYQGLSTATKDTYDALHMQALPPR	1186
C11D5.3 VH	QIQLVQSGPELKKPGETVKISCKASGYTFTDYSINWVKRAPGKGLKWMGWINTETREPAYA YDFRGRFAFSLETSASTAYLQINNLKYEDTATYFCALDYSYAMDYWGQGTSVTVSS	1187
C11D5.3 VL	DIVLTQSPASLAMSLGKRATISCRASESVSVIGAHLIHWYQQKPGQPPKLLIYLASNLETG VPARFSGSGSGTDFTLTIDPVEEDDVAIYSCLQSRIFPRTFGGGTKLEIK	1188
C11D5.3 scFv domain	QIQLVQSGPELKKPGETVKISCKASGYTFTDYSINWVKRAPGKGLKWMGWINTETREPAYA YDFRGRFAFSLETSASTAYLQINNLKYEDTATYFCALDYSYAMDYWGQGTSVTVSSGGGGS GGGGSGGGGSQIQLVQSGPELKKPGETVKISCKASGYTFTDYSINWVKRAPGKGLKWMGWI NTETREPAYAYDFRGRFAFSLETSASTAYLQINNLKYEDTATYFCALDYSYAMDYWGQGTS VTVSS	1189
C11D5.3 Full CART	QIQLVQSGPELKKPGETVKISCKASGYTFTDYSINWVKRAPGKGLKWMGWINTETREPAYA YDFRGRFAFSLETSASTAYLQINNLKYEDTATYFCALDYSYAMDYWGQGTSVTVSSGGGGS GGGGSGGGGSQIQLVQSGPELKKPGETVKISCKASGYTFTDYSINWVKRAPGKGLKWMGWI NTETREPAYAYDFRGRFAFSLETSASTAYLQINNLKYEDTATYFCALDYSYAMDYWGQGTS VTVSSTTTPAPRPPTPAPTIASQPLSLRPEACRPAAGGAVHTRGLDFACDIYIWAPLAGTC	1190

	GVLLLSLVITLYCKRGRKKLLYIFKQPFMRPVQTTQEEDGCSCRFPEEEEGGCELRVKFSR SADAPAYKQGQNQLYNELNLGRREEYDVLDKRRGRDPEMGGKPRRKNPQEGLYNELQKDKM AEAYSEIGMKGERRRGKGHDGLYQGLSTATKDTYDALHMQALPPR	
C12A3.2 VH	QIQLVQSGPELKKPGETVKISCKASGYTFRHYSMNWVKQAPGKGLKWMGRINTESGVPIYA DDFKGRFAFSVETSASTAYLVINNLKDEDTASYFCSNDYLYSLDFWGQGTALTVSS	1191
C12A3.2 VL	DIVLTQSPPSLAMSLGKRATISCRASESVTILGSHLIYWYQQKPGQPPTLLIQLASNVQTG VPARFSGSGSRTDFTLTIDPVEEDDVAVYYCLQSRTIPRTFGGGTKLEIK	1192
C12A3.2 scFv domain	QIQLVQSGPELKKPGETVKISCKASGYTFRHYSMNWVKQAPGKGLKWMGRINTESGVPIYA DDFKGRFAFSVETSASTAYLVINNLKDEDTASYFCSNDYLYSLDFWGQGTALTVSSGGGGS GGGGSGGGGSDIVLTQSPPSLAMSLGKRATISCRASESVTILGSHLIYWYQQKPGQPPTLL IQLASNVQTGVPARFSGSGSRTDFTLTIDPVEEDDVAVYYCLQSRTIPRTFGGGTKLEIK	1193
C12A3.2 Full CART	QIQLVQSGPELKKPGETVKISCKASGYTFRHYSMNWVKQAPGKGLKWMGRINTESGVPIYA DDFKGRFAFSVETSASTAYLVINNLKDEDTASYFCSNDYLYSLDFWGQGTALTVSSGGGGS GGGGSGGGGSDIVLTQSPPSLAMSLGKRATISCRASESVTILGSHLIYWYQQKPGQPPTLL IQLASNVQTGVPARFSGSGSRTDFTLTIDPVEEDDVAVYYCLQSRTIPRTFGGGTKLEIKT TTPAPRPPTPAPTIASQPLSLRPEACRPAAGGAVHTRGLDFACDIYIWAPLAGTCGVLLLS LVITLYCKRGRKKLLYIFKQPFMRPVQTTQEEDGCSCRFPEEEEGGCELRVKFSRSADAPA YKQGQNQLYNELNLGRREEYDVLDKRRGRDPEMGGKPRRKNPQEGLYNELQKDKMAEAYSE IGMKGERRRGKGHDGLYQGLSTATKDTYDALHMQALPPR	1194
C13F12.1 VH	QIQLVQSGPELKKPGETVKISCKASGYTFTHYSMNWVKQAPGKGLKWMGRINTETGEPLYA DDFKGRFAFSLETSASTAYLVINNLKNEDTATFFCSNDYLYSCDYWGQGTTLTVSS	1195
C13F12.1 VL	DIVLTQSPPSLAMSLGKRATISCRASESVTILGSHLIYWYQQKPGQPPTLLIQLASNVQTG VPARFSGSGSRTDFTLTIDPVEEDDVAVYYCLQSRTIPRTFGGGTKLEIK	1196
C13F12.1 scFv domain	QIQLVQSGPELKKPGETVKISCKASGYTFTHYSMNWVKQAPGKGLKWMGRINTETGEPLYA DDFKGRFAFSLETSASTAYLVINNLKNEDTATFFCSNDYLYSCDYWGQGTTLTVSSGGGGS GGGGSGGGGSDIVLTQSPPSLAMSLGKRATISCRASESVTILGSHLIYWYQQKPGQPPTLL IQLASNVQTGVPARFSGSGSRTDFTLTIDPVEEDDVAVYYCLQSRTIPRTFGGGTKLEIK	1197
C13F12.1 Full CART	QIQLVQSGPELKKPGETVKISCKASGYTFTHYSMNWVKQAPGKGLKWMGRINTETGEPLYA DDFKGRFAFSLETSASTAYLVINNLKNEDTATFFCSNDYLYSCDYWGQGTTLTVSSGGGGS GGGGSGGGGSDIVLTQSPPSLAMSLGKRATISCRASESVTILGSHLIYWYQQKPGQPPTLL IQLASNVQTGVPARFSGSGSRTDFTLTIDPVEEDDVAVYYCLQSRTIPRTFGGGTKLEIKT TTPAPRPPTPAPTIASQPLSLRPEACRPAAGGAVHTRGLDFACDIYIWAPLAGTCGVLLLS LVITLYCKRGRKKLLYIFKQPFMRPVQTTQEEDGCSCRFPEEEEGGCELRVKFSRSADAPA YKQGQNQLYNELNLGRREEYDVLDKRRGRDPEMGGKPRRKNPQEGLYNELQKDKMAEAYSE IGMKGERRRGKGHDGLYQGLSTATKDTYDALHMQALPPR	1198

Exemplary BCMA CAR constructs disclose herein comprise an scFv (e.g., a scFv as disclosed in Tables 11A or 11B, optionally preceded with an optional leader sequence (e.g., SEQ ID NO: 2 and SEQ ID NO: 3 or 1938 for exemplary leader amino acid and nucleotide sequences, respectively). The sequences of the scFv fragments (e.g., an ScFv from any of SEQ ID NOs: 967-1182, e.g., SEQ ID NOs: 967, 973, 979, 985, 991, 997, 1003, 1009, 1015, 1021, 1027, 1033, 1039, 1045, 1051, 1057, 1063, 1069, 1075, 1081, 1087, 1093, 1099, 1105, 1111, 1117, 1123, 1129, 1135, 1141, 1147, 1153, 1159, 1165, 1171, 1177, not including the optional leader sequence) are provided herein in Tables 11A or 11B. The BCMA CAR construct can further include an optional hinge domain, e.g., a CD8 hinge domain (e.g., including the

amino acid sequence of SEQ ID NO: 4 or encoded by a nucleic acid sequence of SEQ ID NO: 5); a transmembrane domain, e.g., a CD8 transmembrane domain (e.g., including the amino acid sequence of SEQ ID NO: 12 or encoded by the nucleotide sequence of SEQ ID NO: 13 or 1939); an intracellular domain, e.g., a 4-1BB intracellular domain (e.g., including the amino acid sequence of SEQ ID NO: 14 or encoded by the nucleotide sequence of SEQ ID NO: 15 or 1940; and a functional signaling domain, e.g., a CD3 zeta domain (e.g., including amino acid sequence of SEQ ID NO: 18 or 20, or encoded by the nucleotide sequence of SEQ ID NO: 19, 1941, or 21). In certain embodiments, the domains are contiguous with and in the same reading frame to form a single fusion protein. In other embodiments, the domain are in separate polypeptides, e.g., as in an RCAR molecule as described herein.

In certain embodiments, the full length BCMA CAR molecule includes the amino acid sequence of, or is encoded by the nucleotide sequence of, BCMA-1, BCMA-2, BCMA-3, BCMA-4, BCMA-5, BCMA-6, BCMA-7, BCMA-8, BCMA-9, BCMA-10, BCMA-11, BCMA-12, BCMA-13, BCMA-14, BCMA-15, 149362, 149363, 149364, 149365, 149366, 149367, 149368, 149369, BCMA_EBB-C1978-A4, BCMA_EBB-C1978-G1, BCMA_EBB-C1979-C1, BCMA_EBB-C1978-C7, BCMA_EBB-C1978-D10, BCMA_EBB-C1979-C12, BCMA_EBB-C1980-G4, BCMA_EBB-C1980-D2, BCMA_EBB-C1978-A10, BCMA_EBB-C1978-D4, BCMA_EBB-C1980-A2, BCMA_EBB-C1981-C3, BCMA_EBB-C1978-G4, A7D12.2, C11D5.3, C12A3.2, or C13F12.1 provided in Table 11A or 11B, or a sequence substantially (e.g., 85%, 95-99% or higher) identical thereto.

In certain embodiments, the BCMA CAR molecule, or the anti-BCMA antigen binding domain, includes the scFv amino acid sequence of BCMA-1, BCMA-2, BCMA-3, BCMA-4, BCMA-5, BCMA-6, BCMA-7, BCMA-8, BCMA-9, BCMA-10, BCMA-11, BCMA-12, BCMA-13, BCMA-14, BCMA-15, 149362, 149363, 149364, 149365, 149366, 149367, 149368, 149369, BCMA_EBB-C1978-A4, BCMA_EBB-C1978-G1, BCMA_EBB-C1979-C1, BCMA_EBB-C1978-C7, BCMA_EBB-C1978-D10, BCMA_EBB-C1979-C12, BCMA_EBB-C1980-G4, BCMA_EBB-C1980-D2, BCMA_EBB-C1978-A10, BCMA_EBB-C1978-D4, BCMA_EBB-C1980-A2, BCMA_EBB-C1981-C3, BCMA_EBB-C1978-G4, A7D12.2, C11D5.3, C12A3.2, or C13F12.1 provided in Table 11A or 11B (with or without the leader sequence), or a sequence substantially identical (e.g., 85%, 95-99% or higher identical, or up to 20, 15, 10, 8, 6, 5, 4, 3, 2, or 1 amino acid changes, e.g., substitutions (e.g., conservative substitutions)) to any of the aforesaid sequences.

In certain embodiments, the BCMA CAR molecule, or the anti-BCMA antigen binding domain, includes the heavy chain variable region and/or the light chain variable region of BCMA-1, BCMA-2,

BCMA-3, BCMA-4, BCMA-5, BCMA-6, BCMA-7, BCMA-8, BCMA-9, BCMA-10, BCMA-11, BCMA-12, BCMA-13, BCMA-14, BCMA-15, 149362, 149363, 149364, 149365, 149366, 149367, 149368, 149369, BCMA_EBB-C1978-A4, BCMA_EBB-C1978-G1, BCMA_EBB-C1979-C1, BCMA_EBB-C1978-C7, BCMA_EBB-C1978-D10, BCMA_EBB-C1979-C12, BCMA_EBB-C1980-G4, BCMA_EBB-C1980-D2, BCMA_EBB-C1978-A10, BCMA_EBB-C1978-D4, BCMA_EBB-C1980-A2, BCMA_EBB-C1981-C3, BCMA_EBB-C1978-G4, A7D12.2, C11D5.3, C12A3.2, or C13F12.1 provided in Table 11A or 11B, or a sequence substantially identical (e.g., 85%, 95-99% or higher identical, or up to 20, 15, 10, 8, 6, 5, 4, 3, 2, or 1 amino acid changes, e.g., substitutions (e.g., conservative substitutions)) to any of the aforesaid sequences.

In certain embodiments, the BCMA CAR molecule, or the anti-BCMA antigen binding domain, includes one, two or three CDRs from the heavy chain variable region (e.g., HCDR1, HCDR2 and/or HCDR3), provided in Table 12; and/or one, two or three CDRs from the light chain variable region (e.g., LCDR1, LCDR2 and/or LCDR3) of BCMA-1, BCMA-2, BCMA-3, BCMA-4, BCMA-5, BCMA-6, BCMA-7, BCMA-8, BCMA-9, BCMA-10, BCMA-11, BCMA-12, BCMA-13, BCMA-14, BCMA-15, 149362, 149363, 149364, 149365, 149366, 149367, 149368, 149369, BCMA_EBB-C1978-A4, BCMA_EBB-C1978-G1, BCMA_EBB-C1979-C1, BCMA_EBB-C1978-C7, BCMA_EBB-C1978-D10, BCMA_EBB-C1979-C12, BCMA_EBB-C1980-G4, BCMA_EBB-C1980-D2, BCMA_EBB-C1978-A10, BCMA_EBB-C1978-D4, BCMA_EBB-C1980-A2, BCMA_EBB-C1981-C3, BCMA_EBB-C1978-G4, A7D12.2, C11D5.3, C12A3.2, or C13F12.1, provided in Table 13; or a sequence substantially identical (e.g., 85%, 95-99% or higher identical, or up to 20, 15, 10, 8, 6, 5, 4, 3, 2, or 1 amino acid changes, e.g., substitutions (e.g., conservative substitutions)) to any of the aforesaid sequences.

In certain embodiments, the BCMA CAR molecule, or the anti-BCMA antigen binding domain, includes one, two or three CDRs from the heavy chain variable region (e.g., HCDR1, HCDR2 and/or HCDR3), provided in Table 14; and/or one, two or three CDRs from the light chain variable region (e.g., LCDR1, LCDR2 and/or LCDR3) of BCMA-1, BCMA-2, BCMA-3, BCMA-4, BCMA-5, BCMA-6, BCMA-7, BCMA-8, BCMA-9, BCMA-10, BCMA-11, BCMA-12, BCMA-13, BCMA-14, BCMA-15, 149362, 149363, 149364, 149365, 149366, 149367, 149368, 149369, BCMA_EBB-C1978-A4, BCMA_EBB-C1978-G1, BCMA_EBB-C1979-C1, BCMA_EBB-C1978-C7, BCMA_EBB-C1978-D10, BCMA_EBB-C1979-C12, BCMA_EBB-C1980-G4, BCMA_EBB-C1980-D2, BCMA_EBB-C1978-A10, BCMA_EBB-C1978-D4, BCMA_EBB-C1980-A2, BCMA_EBB-C1981-C3, BCMA_EBB-C1978-G4, A7D12.2, C11D5.3, C12A3.2, or C13F12.1, provided in Table 15; or a sequence substantially identical (e.g., 85%, 95-99% or higher

identical, or up to 20, 15, 10, 8, 6, 5, 4, 3, 2, or 1 amino acid changes, e.g., substitutions (e.g., conservative substitutions)) to any of the aforesaid sequences.

In certain embodiments, the BCMA CAR molecule, or the anti-BCMA antigen binding domain, includes one, two or three CDRs from the heavy chain variable region (e.g., HCDR1, HCDR2 and/or HCDR3), provided in Table 16; and/or one, two or three CDRs from the light chain variable region (e.g., LCDR1, LCDR2 and/or LCDR3) of BCMA-1, BCMA-2, BCMA-3, BCMA-4, BCMA-5, BCMA-6, BCMA-7, BCMA-8, BCMA-9, BCMA-10, BCMA-11, BCMA-12, BCMA-13, BCMA-14, BCMA-15, 149362, 149363, 149364, 149365, 149366, 149367, 149368, 149369, BCMA_EBB-C1978-A4, BCMA_EBB-C1978-G1, BCMA_EBB-C1979-C1, BCMA_EBB-C1978-C7, BCMA_EBB-C1978-D10, BCMA_EBB-C1979-C12, BCMA_EBB-C1980-G4, BCMA_EBB-C1980-D2, BCMA_EBB-C1978-A10, BCMA_EBB-C1978-D4, BCMA_EBB-C1980-A2, BCMA_EBB-C1981-C3, BCMA_EBB-C1978-G4, A7D12.2, C11D5.3, C12A3.2, or C13F12.1, provided in Table 17; or a sequence substantially identical (e.g., 85%, 95-99% or higher identical, or up to 20, 15, 10, 8, 6, 5, 4, 3, 2, or 1 amino acid changes, e.g., substitutions (e.g., conservative substitutions)) to any of the aforesaid sequences.

The sequences of human CDR sequences of the scFv domains are shown in Tables 12, 14, and 16 for the heavy chain variable domains and in Tables 13, 15, and 17 for the light chain variable domains.

"ID" stands for the respective SEQ ID NO for each CDR.

Table 12: Heavy Chain Variable Domain CDRs according to the Kabat numbering scheme (Kabat et al. (1991), "Sequences of Proteins of Immunological Interest," 5th Ed. Public Health Service, National Institutes of Health, Bethesda, MD)

Candidate	HCDR1	ID	HCDR2	ID	HCDR3	ID
139109	NHGMS	1199	GIVYSGSTYYAASVKG	1239	HGGESDV	1279
139103	NYAMS	1200	GISRSGENTYYADSVKG	1240	SPAHYYGGMDV	1280
139105	DYAMH	1201	GISWNSGSIGYADSVKG	1241	HSFLAY	1281
139111	NHGMS	1202	GIVYSGSTYYAASVKG	1242	HGGESDV	1282
139100	NFGIN	1203	WINPKNNNTNYAQKFQG	1243	GPYYYQSYMDV	1283
139101	SDAMT	1204	VISGSGGTTYYADSVKG	1244	LDSSGYYYARGPRY	1284
139102	NYGIT	1205	WISAYNGNTNYAQKFQG	1245	GPYYYYMDV	1285

139104	NHGMS	1206	GIVYSGSTYYAASVKG	1246	HGGESDV	1286
139106	NHGMS	1207	GIVYSGSTYYAASVKG	1247	HGGESDV	1287
139107	NHGMS	1208	GIVYSGSTYYAASVKG	1248	HGGESDV	1288
139108	DYYMS	1209	YISSSGSTIYYADSVKG	1249	ESGDGMDV	1289
139110	DYYMS	1210	YISSSGNTIYYADSVKG	1250	STMVREDY	1290
139112	NHGMS	1211	GIVYSGSTYYAASVKG	1251	HGGESDV	1291
139113	NHGMS	1212	GIVYSGSTYYAASVKG	1252	HGGESDV	1292
139114	NHGMS	1213	GIVYSGSTYYAASVKG	1253	HGGESDV	1293
149362	SSYYYWG	1214	SIYYSGSAYYNPSLKS	1254	HWQEWPDAFDI	1294
149363	TSGMCVS	1215	RIDWDEDKFYSTSLKT	1255	SGAGGTSATAFDI	1295
149364	SYSMN	1216	SISSSSSYIYYADSVKG	1256	TIAAVYAFDI	1296
149365	DYYMS	1217	YISSSGSTIYYADSVKG	1257	DLRGAFDI	1297
149366	SHYIH	1218	MINPSGGVTAYSQTLQG	1258	EGSGSGWYFDF	1298
149367	SGGYYWS	1219	YIYYSGSTYYNPSLKS	1259	AGIAARLRGAFDI	1299
149368	SYAIS	1220	GIIPIFGTANYAQKFQG	1260	RGGYQLLRWDVGLLR SAFDI	1300
149369	SNSAAWN	1221	RTYYRSKWYSFYAISLKS	1261	SSPEGLFLYWFDP	1301
BCMA_EBB-C1978-A4	SYAMS	1222	AISGSGGSTYYADSVKG	1262	VEGSGSLDY	1302
BCMA_EBB-C1978-G1	RYPMS	1223	GISDSGVSTYYADSAKG	1263	RAGSEASDI	1303
BCMA_EBB-C1979-C1	SYAMS	1224	AISGSGGSTYYADSVKG	1264	ATYKRELRYYYGMDV	1304
BCMA_EBB-C1978-C7	SYAMS	1225	AISGSGGSTYYADSVKG	1265	ATYKRELRYYYGMDV	1305
BCMA_EBB-C1978-D10	DYAMH	1226	GISWNSGSIGYADSVKG	1266	VGKAVPDV	1306
BCMA_EBB-C1979-C12	DYAMH	1227	SINWKGNSLAYGDSVKG	1267	HQGVAYYNYAMDV	1307
BCMA_EBB-C1980-G4	SYAMS	1228	AISGSGGSTYYADSVKG	1268	VVRDGMDV	1308
BCMA_EBB-	SYAMS	1229	AISGSGGSTYYADSVKG	1269	IPQTGTFDY	1309

C1980-D2						
BCMA_EBB-C1978-A10	SYAMS	1230	AISGSGGSTYYADSVKG	1270	ANYKRELRYYYGMDV	1310
BCMA_EBB-C1978-D4	SYAMS	1231	AISGSGGSTYYADSVKG	1271	ALVGATGAFDI	1311
BCMA_EBB-C1980-A2	SYAMS	1232	AISGSGGSTYYADSVKG	1272	WFGEGFDP	1312
BCMA_EBB-C1981-C3	SYAMS	1233	AISGSGGSTYYADSVKG	1273	VGYDSSGYYRDYYGMDV	1313
BCMA_EBB-C1978-G4	SYAMS	1234	AISGSGGSTYYADSVKG	1274	MGWSSGYLGAFDI	1314
A7D12.2	NFGMN	1235	WINTYTGESYFADDFKG	1275	GEIYYGYDGGFAY	1315
C11D5.3	DYSIN	1236	WINTETREPAYAYDFRG	1276	DYSYAMDY	1316
C12A3.2	HYSMN	1237	RINTESGVPIYADDFKG	1277	DYLYSLDF	1317
C13F12.1	HYSMN	1238	RINTETGEPLYADDFKG	1278	DYLYSCDY	1318

Table 13: Light Chain Variable Domain CDRs according to the Kabat numbering scheme (Kabat et al. (1991), “Sequences of Proteins of Immunological Interest,” 5th Ed. Public Health Service, National Institutes of Health, Bethesda, MD)

Candidate	**LCDR1**	**ID**	**LCDR2**	**ID**	**LCDR3**	**ID**
139109	RASQSISSYLN	1319	AASSLQS	1359	QQSYSTPYT	1399
139103	RASQSISSSFLA	1320	GASRRAT	1360	QQYHSSPSWT	1400
139105	RSSQSLLHSNGYNYLD	1321	LGSNRAS	1361	MQALQTPYT	1401
139111	KSSQSLLRNDGKTPLY	1322	EVSNRFS	1362	MQNIQFPS	1402
139100	RSSQSLLHSNGYNYLN	1323	LGSKRAS	1363	MQALQTPYT	1403
139101	RASQSISSYLN	1324	GASTLAS	1364	QQSYKRAS	1404
139102	RSSQSLLYSNGYNYVD	1325	LGSNRAS	1365	MQGRQFPYS	1405
139104	RASQSVSSNLA	1326	GASTRAS	1366	QQYGSSLT	1406
139106	RASQSVSSKLA	1327	GASIRAT	1367	QQYGSSSWT	1407
139107	RASQSVGSTNLA	1328	DASNRAT	1368	QQYGSSPPWT	1408
139108	RASQSISSYLN	1329	AASSLQS	1369	QQSYTLA	1409

139110	KSSESLVHNSGKTYLN	1330	EVSNRDS	1370	MQGTHWPGT	1410
139112	QASEDINKFLN	1331	DASTLQT	1371	QQYESLPLT	1411
139113	RASQSVGSNLA	1332	GASTRAT	1372	QQYNDWLPVT	1412
139114	RASQSIGSSSLA	1333	GASSRAS	1373	QQYAGSPPFT	1413
149362	KASQDIDDAMN	1334	SATSPVP	1374	LQHDNFPLT	1414
149363	RASQDIYNNLA	1335	AANKSQS	1375	QHYYRFPYS	1415
149364	RSSQSLLHSNGYNYLD	1336	LGSNRAS	1376	MQALQTPYT	1416
149365	GGNNIGTKSVH	1337	DDSVRPS	1377	QVWDSDSEHVV	1417
149366	SGDGLSKKYVS	1338	RDKERPS	1378	QAWDDTTVV	1418
149367	RASQGIRNWLA	1339	AASNLQS	1379	QKYNSAPFT	1419
149368	GGNNIGSKSVH	1340	GKNNRPS	1380	SSRDSSGDHLRV	1420
149369	QGDSLGNYYAT	1341	GTNNRPS	1381	NSRDSSGHHLL	1421
BCMA_EBB-C1978-A4	RASQSVSSAYLA	1342	GASTRAT	1382	QHYGSSFNGSSLFT	1422
BCMA_EBB-C1978-G1	RASQSVSNSLA	1343	DASSRAT	1383	QQFGTSSGLT	1423
BCMA_EBB-C1979-C1	RASQSVSSSFLA	1344	GASSRAT	1384	QQYHSSPSWT	1424
BCMA_EBB-C1978-C7	RASQSVSTTFLA	1345	GSSNRAT	1385	QQYHSSPSWT	1425
BCMA_EBB-C1978-D10	RASQSISSYLN	1346	AASSLQS	1386	QQSYSTPYS	1426
BCMA_EBB-C1979-C12	RATQSIGSSFLA	1347	GASQRAT	1387	QHYESSPSWT	1427
BCMA_EBB-C1980-G4	RASQSVSSSYLA	1348	GASSRAT	1388	QQYGSPPRFT	1428
BCMA_EBB-C1980-D2	RASQSVSSSYLA	1349	GASSRAT	1389	QHYGSSPSWT	1429
BCMA_EBB-C1978-A10	RASQRVASNYLA	1350	GASSRAT	1390	QHYDSSPSWT	1430
BCMA_EBB-C1978-D4	RASQSLSSNFLA	1351	GASNWAT	1391	QYYGTSPMYT	1431
BCMA_EBB-	RSSQSLLHSNGYNYLD	1352	LGSNRAS	1392	MQALQTPLT	1432

C1980-A2						
BCMA_EBB-C1981-C3	RASQSVSSSYLA	1353	GTSSRAT	1393	QHYGNSPPKFT	1433
BCMA_EBB-C1978-G4	RASQSVASSFLA	1354	GASGRAT	1394	QHYGGSPRLT	1434
A7D12.2	RASQDVNTAVS	1355	SASYRYT	1395	QQHYSTPWT	1435
C11D5.3	RASESVSVIGAHLIH	1356	LASNLET	1396	LQSRIFPRT	1436
C12A3.2	RASESVTILGSHLIY	1357	LASNVQT	1397	LQSRTIPRT	1437
C13F12.1	RASESVTILGSHLIY	1358	LASNVQT	1398	LQSRTIPRT	1438

Table 14: Heavy Chain Variable Domain CDRs according to the Chothia numbering scheme (Al-Lazikani et al., (1997) JMB 273,927-948)

Candidate	HCDR1	ID	HCDR2	ID	HCDR3	ID
139109	GFALSNH	1439	VYSGS	1479	HGGESDV	1519
139103	GFTFSNY	1440	SRSGEN	1480	SPAHYYGGMDV	1520
139105	GFTFDDY	1441	SWNSGS	1481	HSFLAY	1521
139111	GFALSNH	1442	VYSGS	1482	HGGESDV	1522
139100	GYIFDNF	1443	NPKNNN	1483	GPYYYQSYMDV	1523
139101	GFTFSSD	1444	SGSGGT	1484	LDSSGYYYARGPRY	1524
139102	GYTFSNY	1445	SAYNGN	1485	GPYYYYMDV	1525
139104	GFALSNH	1446	VYSGS	1486	HGGESDV	1526
139106	GFALSNH	1447	VYSGS	1487	HGGESDV	1527
139107	GFALSNH	1448	VYSGS	1488	HGGESDV	1528
139108	GFTFSDY	1449	SSSGST	1489	ESGDGMDV	1529
139110	GFTFSDY	1450	SSSGNT	1490	STMVREDY	1530
139112	GFALSNH	1451	VYSGS	1491	HGGESDV	1531
139113	GFALSNH	1452	VYSGS	1492	HGGESDV	1532
139114	GFALSNH	1453	VYSGS	1493	HGGESDV	1533
149362	GGSISSSYY	1454	YYSGS	1494	HWQEWPDAFDI	1534

149363	GFSLRTSGM	1455	DWDED	1495	SGAGGTSATAFDI	1535
149364	GFTFSSY	1456	SSSSSY	1496	TIAAVYAFDI	1536
149365	GFTFSDY	1457	SSSGST	1497	DLRGAFDI	1537
149366	GYTVTSH	1458	NPSGGV	1498	EGSGSGWYFDF	1538
149367	GGSISSGGY	1459	YYSGS	1499	AGIAARLRGAFDI	1539
149368	GGTFSSY	1460	IPIFGT	1500	RGGYQLLRWDVGLLRSAFDI	1540
149369	GDSVSSNSA	1461	YYRSKWY	1501	SSPEGLFLYWFDP	1541
BCMA_EBB-C1978-A4	GFTFSSY	1462	SGSGGS	1502	VEGSGSLDY	1542
BCMA_EBB-C1978-G1	GITFSRY	1463	SDSGVS	1503	RAGSEASDI	1543
BCMA_EBB-C1979-C1	GFTFSSY	1464	SGSGGS	1504	ATYKRELRYYYGMDV	1544
BCMA_EBB-C1978-C7	GFTFSSY	1465	SGSGGS	1505	ATYKRELRYYYGMDV	1545
BCMA_EBB-C1978-D10	GFTFDDY	1466	SWNSGS	1506	VGKAVPDV	1546
BCMA_EBB-C1979-C12	GFTFDDY	1467	NWKGNS	1507	HQGVAYYNYAMDV	1547
BCMA_EBB-C1980-G4	GFTFSSY	1468	SGSGGS	1508	VVRDGMDV	1548
BCMA_EBB-C1980-D2	GFTFSSY	1469	SGSGGS	1509	IPQTGTFDY	1549
BCMA_EBB-C1978-A10	GFTFSSY	1470	SGSGGS	1510	ANYKRELRYYYGMDV	1550
BCMA_EBB-C1978-D4	GFSFSSY	1471	SGSGGS	1511	ALVGATGAFDI	1551
BCMA_EBB-C1980-A2	GFTFSSY	1472	SGSGGS	1512	WFGEGFDP	1552
BCMA_EBB-C1981-C3	GFTFSSY	1473	SGSGGS	1513	VGYDSSGYYRDYYGMDV	1553
BCMA_EBB-C1978-G4	GFTFSSY	1474	SGSGGS	1514	MGWSSGYLGAFDI	1554

A7D12.2	GYTFTNF	1475	NTYTGE	1515	GEIYYGYDGGFAY	1555
C11D5.3	GYTFTDY	1476	NTETRE	1516	DYSYAMDY	1556
C12A3.2	GYTFRHY	1477	NTESGV	1517	DYLYSLDF	1557
C13F12.1	GYTFTHY	1478	NTETGE	1518	DYLYSCDY	1558

Table 15: Light Chain Variable Domain CDRs according to the Chothia numbering scheme (Al-Lazikani et al., (1997) JMB 273,927-948)

Candidate	LCDR1	ID	LCDR2	ID	LCDR3	ID
139109	SQSISSY	1559	AAS	1599	SYSTPY	1639
139103	SQSISSSF	1560	GAS	1600	YHSSPSW	1640
139105	SQSLLHSNGYNY	1561	LGS	1601	ALQTPY	1641
139111	SQSLLRNDGKTP	1562	EVS	1602	NIQFP	1642
139100	SQSLLHSNGYNY	1563	LGS	1603	ALQTPY	1643
139101	SQSISSY	1564	GAS	1604	SYKRA	1644
139102	SQSLLYSNGYNY	1565	LGS	1605	GRQFPY	1645
139104	SQSVSSN	1566	GAS	1606	YGSSL	1646
139106	SQSVSSK	1567	GAS	1607	YGSSSW	1647
139107	SQSVGSTN	1568	DAS	1608	YGSSPPW	1648
139108	SQSISSY	1569	AAS	1609	SYTL	1649
139110	SESLVHNSGKTY	1570	EVS	1610	GTHWPG	1650
139112	SEDINKF	1571	DAS	1611	YESLPL	1651
139113	SQSVGSN	1572	GAS	1612	YNDWLPV	1652
139114	SQSIGSSS	1573	GAS	1613	YAGSPPF	1653
149362	SQDIDDA	1574	SAT	1614	HDNFPL	1654
149363	SQDIYNN	1575	AAN	1615	YYRFPY	1655
149364	SQSLLHSNGYNY	1576	LGS	1616	ALQTPY	1656
149365	NNIGTKS	1577	DDS	1617	WDSDSEHV	1657
149366	DGLSKKY	1578	RDK	1618	WDDTTV	1658

149367	SQGIRNW	1579	AAS	1619	YNSAPF	1659
149368	NNIGSKS	1580	GKN	1620	RDSSGDHLR	1660
149369	DSLGNYY	1581	GTN	1621	RDSSGHHL	1661
BCMA_EBB-C1978-A4	SQSVSSAY	1582	GAS	1622	YGSSFNGSSLF	1662
BCMA_EBB-C1978-G1	SQSVSNS	1583	DAS	1623	FGTSSGL	1663
BCMA_EBB-C1979-C1	SQSVSSSF	1584	GAS	1624	YHSSPSW	1664
BCMA_EBB-C1978-C7	SQSVSTTF	1585	GSS	1625	YHSSPSW	1665
BCMA_EBB-C1978-D10	SQSISSY	1586	AAS	1626	SYSTPY	1666
BCMA_EBB-C1979-C12	TQSIGSSF	1587	GAS	1627	YESSPSW	1667
BCMA_EBB-C1980-G4	SQSVSSSY	1588	GAS	1628	YGSPPRF	1668
BCMA_EBB-C1980-D2	SQSVSSSY	1589	GAS	1629	YGSSPSW	1669
BCMA_EBB-C1978-A10	SQRVASNY	1590	GAS	1630	YDSSPSW	1670
BCMA_EBB-C1978-D4	SQSLSSNF	1591	GAS	1631	YGTSPMY	1671
BCMA_EBB-C1980-A2	SQSLLHSNGYNY	1592	LGS	1632	ALQTPL	1672
BCMA_EBB-C1981-C3	SQSVSSSY	1593	GTS	1633	YGNSPPKF	1673
BCMA_EBB-C1978-G4	SQSVASSF	1594	GAS	1634	YGGSPRL	1674
A7D12.2	SQDVNTA	1595	SAS	1635	HYSTPW	1675
C11D5.3	SESVSVIGAHL	1596	LAS	1636	SRIFPR	1676
C12A3.2	SESVTILGSHL	1597	LAS	1637	SRTIPR	1677
C13F12.1	SESVTILGSHL	1598	LAS	1638	SRTIPR	1678

Table 16. Heavy Chain Variable Domain CDRs according to a combination of the Kabat numbering scheme (Kabat et al. (1991), "Sequences of Proteins of Immunological Interest," 5th Ed. Public Health Service, National Institutes of Health, Bethesda, MD) and the Chothia numbering scheme (Al-Lazikani et al., (1997) JMB 273,927-948).

Candidate	HCDR1	ID	HCDR2	ID	HCDR3	ID
139109	GFALSNHGMS	1679	GIVYSGSTYYAASVKG	1719	HGGESDV	1759
139103	GFTFSNYAMS	1680	GISRSGENTYYADSVKG	1720	SPAHYYGGMDV	1760
139105	GFTFDDYAMH	1681	GISWNSGSIGYADSVKG	1721	HSFLAY	1761
139111	GFALSNHGMS	1682	GIVYSGSTYYAASVKG	1722	HGGESDV	1762
139100	GYIFDNFGIN	1683	WINPKNNNTNYAQKFQG	1723	GPYYYQSYMDV	1763
139101	GFTFSSDAMT	1684	VISGSGGTTYYADSVKG	1724	LDSSGYYYARGPRY	1764
139102	GYTFSNYGIT	1685	WISAYNGNTNYAQKFQG	1725	GPYYYYMDV	1765
139104	GFALSNHGMS	1686	GIVYSGSTYYAASVKG	1726	HGGESDV	1766
139106	GFALSNHGMS	1687	GIVYSGSTYYAASVKG	1727	HGGESDV	1767
139107	GFALSNHGMS	1688	GIVYSGSTYYAASVKG	1728	HGGESDV	1768
139108	GFTFSDYYMS	1689	YISSSGSTIYYADSVKG	1729	ESGDGMDV	1769
139110	GFTFSDYYMS	1690	YISSSGNTIYYADSVKG	1730	STMVREDY	1770
139112	GFALSNHGMS	1691	GIVYSGSTYYAASVKG	1731	HGGESDV	1771
139113	GFALSNHGMS	1692	GIVYSGSTYYAASVKG	1732	HGGESDV	1772
139114	GFALSNHGMS	1693	GIVYSGSTYYAASVKG	1733	HGGESDV	1773
149362	GGSISSSYYYWG	1694	SIYYSGSAYYNPSLKS	1734	HWQEWPDAFDI	1774
149363	GFSLRTSGMCVS	1695	RIDWDEDKFYSTSLKT	1735	SGAGGTSATAFDI	1775
149364	GFTFSSYSMN	1696	SISSSSSYIYYADSVKG	1736	TIAAVYAFDI	1776
149365	GFTFSDYYMS	1697	YISSSGSTIYYADSVKG	1737	DLRGAFDI	1777

149366	GYTVTSHYIH	1698	MINPSGGVTAYSQTLQG	1738	EGSGSGWYFDF	1778
149367	GGSISSGGYYWS	1699	YIYYSGSTYYNPSLKS	1739	AGIAARLRGAFDI	1779
149368	GGTFSSYAIS	1700	GIIPIFGTANYAQKFQG	1740	RGGYQLLRWDVGLLRSAFDI	1780
149369	GDSVSSNSAAWN	1701	RTYYRSKWYSFYAISLKS	1741	SSPEGLFLYWFDP	1781
BCMA_EBB-C1978-A4	GFTFSSYAMS	1702	AISGSGGSTYYADSVKG	1742	VEGSGSLDY	1782
BCMA_EBB-C1978-G1	GITFSRYPMS	1703	GISDSGVSTYYADSAKG	1743	RAGSEASDI	1783
BCMA_EBB-C1979-C1	GFTFSSYAMS	1704	AISGSGGSTYYADSVKG	1744	ATYKRELRYYYGMDV	1784
BCMA_EBB-C1978-C7	GFTFSSYAMS	1705	AISGSGGSTYYADSVKG	1745	ATYKRELRYYYGMDV	1785
BCMA_EBB-C1978-D10	GFTFDDYAMH	1706	GISWNSGSIGYADSVKG	1746	VGKAVPDV	1786
BCMA_EBB-C1979-C12	GFTFDDYAMH	1707	SINWKGNSLAYGDSVKG	1747	HQGVAYYNYAMDV	1787
BCMA_EBB-C1980-G4	GFTFSSYAMS	1708	AISGSGGSTYYADSVKG	1748	VVRDGMDV	1788
BCMA_EBB-C1980-D2	GFTFSSYAMS	1709	AISGSGGSTYYADSVKG	1749	IPQTGTFDY	1789
BCMA_EBB-C1978-A10	GFTFSSYAMS	1710	AISGSGGSTYYADSVKG	1750	ANYKRELRYYYGMDV	1790
BCMA_EBB-C1978-D4	GFSFSSYAMS	1711	AISGSGGSTYYADSVKG	1751	ALVGATGAFDI	1791
BCMA_EBB-C1980-A2	GFTFSSYAMS	1712	AISGSGGSTYYADSVKG	1752	WFGEGFDP	1792
BCMA_EBB-C1981-C3	GFTFSSYAMS	1713	AISGSGGSTYYADSVKG	1753	VGYDSSGYYRDYYGMDV	1793
BCMA_EBB-C1978-G4	GFTFSSYAMS	1714	AISGSGGSTYYADSVKG	1754	MGWSSGYLGAFDI	1794
A7D12.2	GYTFTNFGMN	1715	WINTYTGESYFADDFKG	1755	GEIYYGYDGGFAY	1795
C11D5.3	GYTFTDYSIN	1716	WINTETREPAYAYDF	1756	DYSYAMDY	1796

			RG			
C12A3.2	GYTFRHYSMN	1717	RINTESGVPIYADDFKG	1757	DYLYSLDF	1797
C13F12.1	GYTFTHYSMN	1718	RINTETGEPLYADDFKG	1758	DYLYSCDY	1798

Table 17. Light Chain Variable Domain CDRs according to a combination of the Kabat numbering scheme (Kabat et al. (1991), "Sequences of Proteins of Immunological Interest," 5th Ed. Public Health Service, National Institutes of Health, Bethesda, MD) and the Chothia numbering scheme (Al-Lazikani et al., (1997) JMB 273,927-948).

Candidate	LCDR1	ID	LCDR2	ID	LCDR3	ID
139109	RASQSISSYLN	1799	AASSLQS	1839	QQSYSTPYT	1879
139103	RASQSISSSFLA	1800	GASRRAT	1840	QQYHSSPSWT	1880
139105	RSSQSLLHSNGYNYLD	1801	LGSNRAS	1841	MQALQTPYT	1881
139111	KSSQSLLRNDGKTPLY	1802	EVSNRFS	1842	MQNIQFPS	1882
139100	RSSQSLLHSNGYNYLN	1803	LGSKRAS	1843	MQALQTPYT	1883
139101	RASQSISSYLN	1804	GASTLAS	1844	QQSYKRAS	1884
139102	RSSQSLLYSNGYNYVD	1805	LGSNRAS	1845	MQGRQFPYS	1885
139104	RASQSVSSNLA	1806	GASTRAS	1846	QQYGSSLT	1886
139106	RASQSVSSKLA	1807	GASIRAT	1847	QQYGSSSWT	1887
139107	RASQSVGSTNLA	1808	DASNRAT	1848	QQYGSSPPWT	1888
139108	RASQSISSYLN	1809	AASSLQS	1849	QQSYTLA	1889
139110	KSSESLVHNSGKTYLN	1810	EVSNRDS	1850	MQGTHWPGT	1890
139112	QASEDINKFLN	1811	DASTLQT	1851	QQYESLPLT	1891
139113	RASQSVGSNLA	1812	GASTRAT	1852	QQYNDWLPVT	1892
139114	RASQSIGSSSLA	1813	GASSRAS	1853	QQYAGSPPFT	1893
149362	KASQDIDDAMN	1814	SATSPVP	1854	LQHDNFPLT	1894
149363	RASQDIYNNLA	1815	AANKSQS	1855	QHYYRFPYS	1895
149364	RSSQSLLHSNGYNYLD	1816	LGSNRAS	1856	MQALQTPYT	1896
149365	GGNNIGTKSVH	1817	DDSVRPS	1857	QVWDSDSEHVV	1897

149366	SGDGLSKKYVS	1818	RDKERPS	1858	QAWDDTTVV	1898
149367	RASQGIRNWLA	1819	AASNLQS	1859	QKYNSAPFT	1899
149368	GGNNIGSKSVH	1820	GKNNRPS	1860	SSRDSSGDHLRV	1900
149369	QGDSLGNYYAT	1821	GTNNRPS	1861	NSRDSSGHHLL	1901
BCMA_EBB-C1978-A4	RASQSVSSAYLA	1822	GASTRAT	1862	QHYGSSFNGSSLFT	1902
BCMA_EBB-C1978-G1	RASQSVSNSLA	1823	DASSRAT	1863	QQFGTSSGLT	1903
BCMA_EBB-C1979-C1	RASQSVSSSFLA	1824	GASSRAT	1864	QQYHSSPSWT	1904
BCMA_EBB-C1978-C7	RASQSVSTTFLA	1825	GSSNRAT	1865	QQYHSSPSWT	1905
BCMA_EBB-C1978-D10	RASQSISSYLN	1826	AASSLQS	1866	QQSYSTPYS	1906
BCMA_EBB-C1979-C12	RATQSIGSSFLA	1827	GASQRAT	1867	QHYESSPSWT	1907
BCMA_EBB-C1980-G4	RASQSVSSSYLA	1828	GASSRAT	1868	QQYGSPPRFT	1908
BCMA_EBB-C1980-D2	RASQSVSSSYLA	1829	GASSRAT	1869	QHYGSSPSWT	1909
BCMA_EBB-C1978-A10	RASQRVASNYLA	1830	GASSRAT	1870	QHYDSSPSWT	1910
BCMA_EBB-C1978-D4	RASQSLSSNFLA	1831	GASNWAT	1871	QYYGTSPMYT	1911
BCMA_EBB-C1980-A2	RSSQSLLHSNGYNYLD	1832	LGSNRAS	1872	MQALQTPLT	1912
BCMA_EBB-C1981-C3	RASQSVSSSYLA	1833	GTSSRAT	1873	QHYGNSPPKFT	1913
BCMA_EBB-C1978-G4	RASQSVASSFLA	1834	GASGRAT	1874	QHYGGSPRLT	1914
A7D12.2	RASQDVNTAVS	1835	SASYRYT	1875	QQHYSTPWT	1915
C11D5.3	RASESVSVIGAHLIH	1836	LASNLET	1876	LQSRIFPRT	1916
C12A3.2	RASESVTILGSHLIY	1837	LASNVQT	1877	LQSRTIPRT	1917
C13F12.1	RASESVTILGSHLIY	1838	LASNVQT	1878	LQSRTIPRT	1918

In certain embodiments, the CAR molecule described herein (*e.g.*, the CAR nucleic acid or the CAR polypeptide) or a BCMA binding domain includes:

(1) one, two, or three light chain (LC) CDRs chosen from one of the following:

(i) a LC CDR1 of SEQ ID NO: 1320, LC CDR2 of SEQ ID NO: 1360 and LC CDR3 of SEQ ID NO: 1400 of BCMA-4 CAR (139103);

(ii) a LC CDR1 of SEQ ID NO: 1319, LC CDR2 of SEQ ID NO: 1359 and LC CDR3 of SEQ ID NO: 1399 of BCMA-10 CAR (139109);

(iii) a LC CDR1 of SEQ ID NO: 1331, LC CDR2 of SEQ ID NO: 1371and LC CDR3 of SEQ ID NO: 1411of BCMA-13 CAR (139112); or

(iv) a LC CDR1 of SEQ ID NO: 1333, LC CDR2 of SEQ ID NO: 1373 and LC CDR3 of SEQ ID NO: 1413 of BCMA-15 CAR (139114), and/or

(2) one, two, or three heavy chain (HC) CDRs from one of the following:

(i) a HC CDR1 of SEQ ID NO: 1200, HC CDR2 of SEQ ID NO: 1240 and HC CDR3 of SEQ ID NO: 1280 of BCMA-4 CAR (139103);

(ii) a HC CDR1 of SEQ ID NO: 1199, HC CDR2 of SEQ ID NO: 1239 and HC CDR3 of SEQ ID NO: 1279 of BCMA-10 CAR (139109);

(iii) a HC CDR1 of SEQ ID NO: 1121, HC CDR2 of SEQ ID NO: 1251 and HC CDR3 of SEQ ID NO: 1291 of BCMA-13 CAR (139112); or

(iv) a HC CDR1 of SEQ ID NO: 1213, HC CDR2 of SEQ ID NO: 1253 and HC CDR3 of SEQ ID NO: 1293 of BCMA-15 (139114).

In certain embodiments, the CAR molecule described herein (*e.g.*, the CAR nucleic acid or the CAR polypeptide) includes:

(1) one, two, or three light chain (LC) CDRs chosen from one of the following:

(i) a LC CDR1 of SEQ ID NO: 1560, LC CDR2 of SEQ ID NO: 1600 and LC CDR3 of SEQ ID NO: 1640 of BCMA-4 CAR (139103);

(ii) a LC CDR1 of SEQ ID NO: 1559, LC CDR2 of SEQ ID NO: 1599 and LC CDR3 of SEQ ID NO: 1639 of BCMA-10 CAR (139109);

(iii) a LC CDR1 of SEQ ID NO: 1571, LC CDR2 of SEQ ID NO: 1611 and LC CDR3 of SEQ ID NO: 1651 of BCMA-13 CAR (139112); or

(iv) a LC CDR1 of SEQ ID NO: 1573, LC CDR2 of SEQ ID NO: 1613 and LC CDR3 of SEQ ID NO: 1653 of BCMA-15 CAR (139114); and/or

(2) one, two, or three heavy chain (HC) CDRs chosen from one of the following:

(i) a HC CDR1 of SEQ ID NO: 1440, HC CDR2 of SEQ ID NO: 1480 and HC CDR3 of SEQ ID NO: 1520 of BCMA-4 CAR (139103);

(ii) a HC CDR1 of SEQ ID NO: 1439, HC CDR2 of SEQ ID NO: 1479 and HC CDR3 of SEQ ID NO: 1519 of BCMA-10 CAR (139109);

(iii) a HC CDR1 of SEQ ID NO: 1451, HC CDR2 of SEQ ID NO: 1491 and HC CDR3 of SEQ ID NO: 1531 of BCMA-13 CAR (139112); or

(iv) a HC CDR1 of SEQ ID NO: 1453, HC CDR2 of SEQ ID NO: 1493 and HC CDR3 of SEQ ID NO: 1533 of BCMA-15 CAR (139114).

In certain embodiments, the CAR molecule described herein (*e.g.*, the CAR nucleic acid or the CAR polypeptide) includes:

(1) one, two, or three light chain (LC) CDRs chosen from one of the following:

(i) a LC CDR1 of SEQ ID NO: 1800 LC CDR2 of SEQ ID NO: 1840 and LC CDR3 of SEQ ID NO: 1880 of BCMA-4 CAR (139103);

(ii) a LC CDR1 of SEQ ID NO: 1799, LC CDR2 of SEQ ID NO: 1839 and LC CDR3 of SEQ ID NO: 1879 of BCMA-10 CAR (139109);

(iii) a LC CDR1 of SEQ ID NO: 1811, LC CDR2 of SEQ ID NO: 1851 and LC CDR3 of SEQ ID NO: 1891 of BCMA-13 CAR (139112); or

(iv) a LC CDR1 of SEQ ID NO: 1813, LC CDR2 of SEQ ID NO: 1853 and LC CDR3 of SEQ ID NO: 1893 of BCMA-15 CAR (139114); and/or

(2) one, two, or three heavy chain (HC) CDRs chosen from one of the following:

(i) a HC CDR1 of SEQ ID NO: 1680, HC CDR2 of SEQ ID NO: 1720 and HC CDR3 of SEQ ID NO: 1760 of BCMA-4 CAR (139103);

(ii) a HC CDR1 of SEQ ID NO: 1679, HC CDR2 of SEQ ID NO: 1719 and HC CDR3 of SEQ ID NO: 1759 of BCMA-10 CAR (139109);

(iii) a HC CDR1 of SEQ ID NO: 1691, HC CDR2 of SEQ ID NO: 1731 and HC CDR3 of SEQ ID NO: 1771 of BCMA-13 CAR (139112);

(iv) a HC CDR1 of SEQ ID NO: 1693, HC CDR2 of SEQ ID NO: 1733 and HC CDR3 of SEQ ID NO: 1773 of BCMA-15 CAR (139114).

Exemplary components of the CAR molecules:

Leader (amino acid sequence) (SEQ ID NO: 1919)

MALPVTALLLPLALLLHAARP

leader (nucleic acid sequence) (SEQ ID NO: 1920)

ATGGCCCTGCCTGTGACAGCCCTGCTGCTGCCTCTGGCTCTGCTGCTGCATGCCGCTAGACCC

CD8 hinge (amino acid sequence) (SEQ ID NO: 1921)

TTTPAPRPPTPAPTIASQPLSLRPEACRPAAGGAVHTRGLDFACD

CD8 hinge (nucleic acid sequence) (SEQ ID NO: 1922)

ACCACGACGCCAGCGCCGCGACCACCAACACCGGCGCCCACCATCGCGTCGCAGCCCCTGTCCCTGCGCCCAGAG
GCGTGCCGGCCAGCGGCGGGGGGCGCAGTGCACACGAGGGGGCTGGACTTCGCCTGTGAT

CD8 transmembrane (amino acid sequence) (SEQ ID NO: 1923)

IYIWAPLAGTCGVLLLSLVITLYC

CD8 transmembrane (nucleic acid sequence) (SEQ ID NO: 1924)

ATCTACATCTGGGCGCCCTTGGCCGGGACTTGTGGGGTCCTTCTCCTGTCACTGGTTATCACCCTTTACTGC

4-1BB Intracellular domain (amino acid sequence) (SEQ ID NO: 1925)

KRGRKKLLYIFKQPFMRPVQTTQEEDGCSCRFPEEEEGGCEL

4-1BB Intracellular domain (nucleic acid sequence) (SEQ ID NO: 1926)

AAACGGGGCAGAAAGAAACTCCTGTATATATTCAAACAACCATTTATGAGACCAGTACAAACTACTCAAGAGGAA
GATGGCTGTAGCTGCCGATTTCCAGAAGAAGAAGAAGGAGGATGTGAACTG

CD28 Intracellular domain (amino acid sequence) (SEQ ID NO: 1927)

RSKRSRLLHSDYMNMTPRRPGPTRKHYQPYAPPRDFAAYRS (SEQ ID NO: 1927)

CD28 Intracellular domain (nucleotide sequence) (SEQ ID NO: 1928)

AGGAGTAAGAGGAGCAGGCTCCTGCACAGTGACTACATGAACATGACTCCCCGCCGCCCCGGGCCCACCCGC AAGCATTACCAGCCCTATGCCCCACCACGCGACTTCGCAGCCTATCGCTCC (SEQ ID NO: 1928)

ICOS Intracellular domain (amino acid sequence) (SEQ ID NO: 1929)

T K K K Y S S S V H D P N G E Y M F M R A V N T A K K S R L T D V T L (SEQ ID NO: 1929)

ICOS Intracellular domain (nucleotide sequence) (SEQ ID NO: 1930)

ACAAAAAAGAAGTATTCATCCAGTGTGCACGACCCTAACGGTGAATACATGTTCATGAGAGCAGTGAACACA GCCAAAAAATCCAGACTCACAGATGTGACCCTA (SEQ ID NO: 1930)

CD3 zeta domain (amino acid sequence) (SEQ ID NO: 1931)

RVKFSRSADAPAYKQGQNQLYNELNLGRREEYDVLDKRRGRDPEMGGKPRRKNPQEGLYNELQKDKMAEAYSEIGM KGERRRGKGHDGLYQGLSTATKDTYDALHMQALPPR

CD3 zeta (nucleic acid sequence) (SEQ ID NO: 1932)

AGAGTGAAGTTCAGCAGGAGCGCAGACGCCCCCGCGTACAAGCAGGGCCAGAACCAGCTCTATAACGAGCTCAA TCTAGGACGAAGAGAGGAGTACGATGTTTTGGACAAGAGACGTGGCCGGGACCCTGAGATGGGGGGAAAGCCG AGAAGGAAGAACCCTCAGGAAGGCCTGTACAATGAACTGCAGAAAGATAAGATGGCGGAGGCCTACAGTGAGAT TGGGATGAAAGGCGAGCGCCGGAGGGGCAAGGGGCACGATGGCCTTTACCAGGGTCTCAGTACAGCCACCAAG GACACCTACGACGCCCTTCACATGCAGGCCCTGCCCCCTCGC

CD3 zeta domain (amino acid sequence; NCBI Reference NM_000734.3) (SEQ ID NO: 1933)

RVKFSRSADAPAYQQGQNQLYNELNLGRREEYDVLDKRRGRDPEMGGKPRRKNPQEGLYNELQKDKMAEAYSEIGM KGERRRGKGHDGLYQGLSTATKDTYDALHMQALPPR

CD3 zeta (nucleic acid sequence; NCBI Reference Sequence NM_000734.3); (SEQ ID NO: 1934)

AGAGTGAAGTTCAGCAGGAGCGCAGACGCCCCCGCGTACCAGCAGGGCCAG

AACCAGCTCTATAACGAGCTCAATCTAGGACGAAGAGAGGAGTACGATGTTT

TGGACAAGAGACGTGGCCGGGACCCTGAGATGGGGGGAAAGCCGAGAAGGA

AGAACCCTCAGGAAGGCCTGTACAATGAACTGCAGAAAGATAAGATGGCGG

AGGCCTACAGTGAGATTGGGATGAAAGGCGAGCGCCGGAGGGGCAAGGGGC

ACGATGGCCTTTACCAGGGTCTCAGTACAGCCACCAAGGACACCTACGACGC

CCTTCACATGCAGGCCCTGCCCCCTCGC

IgG4 Hinge (amino acid sequence) (SEQ ID NO: 1935)

ESKYGPPCPPCPAPEFLGGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSQEDPEVQFNWYVDGVEVHNAKTKPREEQF
NSTYRVVSVLTVLHQDWLNGKEYKCKVSNKGLPSSIEKTISKAKGQPREPQVYTLPPSQEEMTKNQVSLTCLVKGFYPSD
IAVEWESNGQPENNYKTTPPVLDSDGSFFLYSRLTVDKSRWQEGNVFSCSVMHEALHNHYTQKSLSLSLGKM

IgG4 Hinge (nucleotide sequence) (SEQ ID NO: 1936)

GAGAGCAAGTACGGCCCTCCCTGCCCCCCTTGCCCTGCCCCCGAGTTCCTGGGCGGACCCAGCGTGTTCCTGTTCC
CCCCCAAGCCCAAGGACACCCTGATGATCAGCCGGACCCCCGAGGTGACCTGTGTGGTGGTGGACGTGTCCCAGG
AGGACCCCGAGGTCCAGTTCAACTGGTACGTGGACGGCGTGGAGGTGCACAACGCCAAGACCAAGCCCCGGGAG
GAGCAGTTCAATAGCACCTACCGGGTGGTGTCCGTGCTGACCGTGCTGCACCAGGACTGGCTGAACGGCAAGGA
ATACAAGTGTAAGGTGTCCAACAAGGGCCTGCCCAGCAGCATCGAGAAAACCATCAGCAAGGCCAAGGGCCAGC
CTCGGGAGCCCCAGGTGTACACCCTGCCCCCTAGCCAAGAGGAGATGACCAAGAACCAGGTGTCCCTGACCTGCC
TGGTGAAGGGCTTCTACCCCAGCGACATCGCCGTGGAGTGGGAGAGCAACGGCCAGCCCGAGAACAACTACAAG
ACCACCCCCCCTGTGCTGGACAGCGACGGCAGCTTCTTCCTGTACAGCCGGCTGACCGTGGACAAGAGCCGGTGG
CAGGAGGGCAACGTCTTTAGCTGCTCCGTGATGCACGAGGCCCTGCACAACCACTACACCCAGAAGAGCCTGAGC
CTGTCCCTGGGCAAGATG

In an embodiment, the CAR molecule comprises a mesothelin CAR described herein, e.g., a mesothelin CAR described in WO 2015/090230, incorporated herein by reference. In embodiments, the mesothelin CAR comprises an amino acid, or has a nucleotide sequence shown in WO 2015/090230 incorporated herein by reference, or a sequence substantially identical to any of the aforesaid sequences (e.g., at least 85%, 90%, 95% or more identical to any of the aforesaid mesothelin CAR sequences). In one embodiment, the CAR molecule comprises a mesothelin CAR, or an antigen binding domain according to Tables 2-3 of WO 2015/090230, incorporated herein by reference, or a sequence substantially identical thereto (e.g., at least 85%, 90%, 95% or more identical thereto). The amino acid and nucleotide sequences encoding the mesothelin CAR molecules and antigen binding domains (e.g., including one, two, three VH CDRs; and one, two, three VL CDRs according to Kabat or Chothia), are specified in WO 2015/090230.

In an embodiment, the CAR molecule comprises a CLL1 CAR described herein, e.g., a CLL1 CAR described in US2016/0051651A1, incorporated herein by reference. In embodiments, the CLL1 CAR

comprises an amino acid, or has a nucleotide sequence shown in US2016/0051651A1, incorporated herein by reference, or a sequence substantially identical to any of the aforesaid sequences (e.g., at least 85%, 90%, 95% or more identical to any of the aforesaid CLL1 CAR sequences).

In other embodiments, the CLL1 CAR includes a CAR molecule, or an antigen binding domain according to Table 2 of WO2016/014535, incorporated herein by reference, or a sequence substantially identical to any of the aforesaid sequences (e.g., at least 85%, 90%, 95% or more identical to any of the aforesaid CLL1 CAR sequences). The amino acid and nucleotide sequences encoding the CLL-1 CAR molecules and antigen binding domains (e.g., including one, two, three VH CDRs; and one, two, three VL CDRs according to Kabat or Chothia), are specified in WO2016/014535.

In an embodiment, the CAR molecule comprises a CD33 CAR described herein, e.g., a CD33 CAR described in US2016/0096892A1, incorporated herein by reference. In embodiments, the CD33 CAR comprises an amino acid, or has a nucleotide sequence shown in US2016/0096892A1, incorporated herein by reference, or a sequence substantially identical to any of the aforesaid sequences (e.g., at least 85%, 90%, 95% or more identical to any of the aforesaid CD33 CAR sequences). In other embodiments, the CD33 CAR CAR or antigen binding domain thereof can include a CAR molecule (e.g., any of CAR33-1 to CAR-33-9), or an antigen binding domain according to Table 2 or 9 of WO2016/014576, incorporated herein by reference, or a sequence substantially identical to any of the aforesaid sequences (e.g., at least 85%, 90%, 95% or more identical to any of the aforesaid CD33 CAR sequences). The amino acid and nucleotide sequences encoding the CD33 CAR molecules and antigen binding domains (e.g., including one, two, three VH CDRs; and one, two, three VL CDRs according to Kabat or Chothia), are specified in WO2016/014576.

In embodiments, the CAR molecule comprises a CD123 CAR described herein, e.g., a CD123 CAR described in US2014/0322212A1 or US2016/0068601A1, both incorporated herein by reference. In embodiments, the CD123 CAR comprises an amino acid, or has a nucleotide sequence shown in US2014/0322212A1 or US2016/0068601A1, both incorporated herein by reference, or a sequence substantially identical to any of the aforesaid sequences (e.g., at least 85%, 90%, 95% or more identical to any of the aforesaid CD123 CAR sequences). In one embodiment, the CAR molecule comprises a CD123 CAR (*e.g.,* any of the CAR1-CAR8), or an antigen binding domain according to Tables 1-2 of WO 2014/130635, incorporated herein by reference, or a sequence substantially identical thereto (e.g., at least 85%, 90%, 95% or more identical to any of the aforesaid CD123 CAR sequences). The amino acid and nucleotide sequences encoding the CD123 CAR molecules and antigen binding domains (e.g.,

including one, two, three VH CDRs; and one, two, three VL CDRs according to Kabat or Chothia), are specified in WO 2014/130635.

In other embodiments, the CAR molecule comprises a CD123 CAR comprises a CAR molecule (e.g., any of the CAR123-1 to CAR123-4 and hzCAR123-1 to hzCAR123-32), or an antigen binding domain according to Tables 2, 6, and 9 of WO2016/028896, incorporated herein by reference, or a sequence substantially identical thereto (e.g., at least 85%, 90%, 95% or more identical to any of the aforesaid CD123 CAR sequences). The amino acid and nucleotide sequences encoding the CD123 CAR molecules and antigen binding domains (e.g., including one, two, three VH CDRs; and one, two, three VL CDRs according to Kabat or Chothia), are specified in WO2016/028896.

In an embodiment, the CAR molecule comprises an EGFRvIII CAR molecule described herein, e.g., an EGFRvIII CAR described US2014/0322275A1, incorporated herein by reference. In embodiments, the EGFRvIII CAR comprises an amino acid, or has a nucleotide sequence shown in US2014/0322275A1, incorporated herein by reference, or a sequence substantially identical to any of the aforesaid sequences (e.g., at least 85%, 90%, 95% or more identical to any of the aforesaid EGFRvIII CAR sequences). In one embodiment, the CAR molecule comprises an EGFRvIII CAR, or an antigen binding domain according to Table 2 or SEQ ID NO:11 of WO 2014/130657, incorporated herein by reference, or a sequence substantially identical thereto (e.g., at least 85%, 90%, 95% or more identical thereto). The amino acid and nucleotide sequences encoding the EGFRvIII CAR molecules and antigen binding domains (*e.g.*, including one, two, three VH CDRs; and one, two, three VL CDRs according to Kabat or Chothia), are specified in WO 2014/130657.

In other embodiments, the CAR molecule comprises an a GFR ALPHA-4 CAR, e.g., can include a CAR molecule, or an antigen binding domain according to Table 2 of WO2016/025880, incorporated herein by reference, or a sequence substantially identical to any of the aforesaid sequences (e.g., at least 85%, 90%, 95% or more identical to any of the aforesaid GFR ALPHA-4 sequences). The amino acid and nucleotide sequences encoding the GFR ALPHA-4 CAR molecules and antigen binding domains (e.g., including one, two, three VH CDRs; and one, two, three VL CDRs according to Kabat or Chothia), are specified in WO2016/025880.

Inhibitory domains

In an embodiment, the vector comprises a nucleic acid sequence that encodes a CAR, e.g., a CAR described herein, and a nucleic acid sequence that encodes an inhibitory molecule comprising: an inhKIR cytoplasmic domain; a transmembrane domain, e.g., a KIR transmembrane domain; and an inhibitor

cytoplasmic domain, e.g., an ITIM domain, e.g., an inhKIR ITIM domain. In an embodiment the inhibitory molecule is a naturally occurring inhKIR, or a sequence sharing at least 50, 60, 70, 80, 85, 90, 95, or 99% homology with, or that differs by no more than 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 15, or 20 residues from, a naturally occurring inhKIR.

In an embodiment, the nucleic acid sequence that encodes an inhibitory molecule comprises: a SLAM family cytoplasmic domain; a transmembrane domain, e.g., a SLAM family transmembrane domain; and an inhibitor cytoplasmic domain, e.g., a SLAM family domain, e.g., an SLAM family ITIM domain. In an embodiment the inhibitory molecule is a naturally occurring SLAM family member, or a sequence sharing at least 50, 60, 70, 80, 85, 90, 95, or 99% homology with, or that differs by no more than 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 15, or 20 residues from, a naturally occurring SLAM family member.

In one embodiment, the vector is an *in vitro* transcribed vector, e.g., a vector that transcribes RNA of a nucleic acid molecule described herein. In one embodiment, the nucleic acid sequence in the vector further comprises a poly(A) tail, e.g., a poly A tail. In one embodiment, the nucleic acid sequence in the vector further comprises a 3'UTR, e.g., a 3' UTR described herein, e.g., comprising at least one repeat of a 3'UTR derived from human beta-globulin. In one embodiment, the nucleic acid sequence in the vector further comprises promoter, e.g., a T2A promoter.

Promoters

In one embodiment, the vector further comprises a promoter. In some embodiments, the promoter is chosen from an EF-1 promoter, a CMV IE gene promoter, an EF-1α promoter, an ubiquitin C promoter, or a phosphoglycerate kinase (PGK) promoter. In one embodiment, the promoter is an EF-1 promoter. In one embodiment, the EF-1 promoter comprises a sequence of SEQ ID NO: 1.

Host cells

As noted above, in some aspects the invention pertains to a cell, e.g., an immune effector cell, (e.g., a population of cells, e.g., a population of immune effector cells) comprising a nucleic acid molecule, a chimeric polypeptide molecule, or a vector as described herein.

In certain aspects of the present disclosure, immune effector cells, e.g., T cells, can be obtained from a unit of blood collected from a subject using any number of techniques known to the skilled artisan, such as Ficoll™ separation. In one preferred aspect, cells from the circulating blood of an individual are obtained by apheresis. The apheresis product typically contains lymphocytes, including T cells, monocytes, granulocytes, B cells, other nucleated white blood cells, red blood cells, and platelets. In one aspect, the cells collected by apheresis may be washed to remove the plasma fraction and,

optionally, to place the cells in an appropriate buffer or media for subsequent processing steps. In one embodiment, the cells are washed with phosphate buffered saline (PBS). In an alternative embodiment, the wash solution lacks calcium and may lack magnesium or may lack many if not all divalent cations.

Initial activation steps in the absence of calcium can lead to magnified activation. As those of ordinary skill in the art would readily appreciate a washing step may be accomplished by methods known to those in the art, such as by using a semi-automated "flow-through" centrifuge (for example, the Cobe 2991 cell processor, the Baxter CytoMate, or the Haemonetics Cell Saver 5) according to the manufacturer's instructions. After washing, the cells may be resuspended in a variety of biocompatible buffers, such as, for example, Ca-free, Mg-free PBS, PlasmaLyte A, or other saline solution with or without buffer. Alternatively, the undesirable components of the apheresis sample may be removed and the cells directly resuspended in culture media.

It is recognized that the methods of the application can utilize culture media conditions comprising 5% or less, for example 2%, human AB serum, and employ known culture media conditions and compositions, for example those described in Smith *et al*., "Ex vivo expansion of human T cells for adoptive immunotherapy using the novel Xeno-free CTS Immune Cell Serum Replacement" *Clinical & Translational Immunology* (2015) 4, e31; doi:10.1038/cti.2014.31. The culture media may additionally include one or more, e.g., one, LSD1 inhibitor(s) as described herein.

In one aspect, T cells are isolated from peripheral blood lymphocytes by lysing the red blood cells and depleting the monocytes, for example, by centrifugation through a PERCOLL^{TM} gradient or by counterflow centrifugal elutriation. Once isolated, the cells may be contacted, e.g., ex vivo, with an LSD1 inhibitor as described herein.

The methods described herein can include, e.g., selection of a specific subpopulation of immune effector cells, e.g., T cells, that are a T regulatory cell-depleted population, CD25+ depleted cells, using, e.g., a negative selection technique, e.g., described herein. Preferably, the population of T regulatory depleted cells contains less than 30%, 25%, 20%, 15%, 10%, 5%, 4%, 3%, 2%, 1% of CD25+ cells. Either before or after selection, the cells may be contacted, e.g., ex vivo, with an LSD1 inhibitor as described herein.

In one embodiment, T regulatory cells, e.g., CD25+ T cells, are removed from the population using an anti-CD25 antibody, or fragment thereof, or a CD25-binding ligand, IL-2. In one embodiment, the anti-CD25 antibody, or fragment thereof, or CD25-binding ligand is conjugated to a substrate, e.g., a

bead, or is otherwise coated on a substrate, e.g., a bead. In one embodiment, the anti-CD25 antibody, or fragment thereof, is conjugated to a substrate as described herein.

In one embodiment, the T regulatory cells, e.g., CD25+ T cells, are removed from the population using CD25 depletion reagent from Miltenyi™. In one embodiment, the ratio of cells to CD25 depletion reagent is 1e7 cells to 20 uL, or 1e7 cells to15 uL, or 1e7 cells to 10 uL, or 1e7 cells to 5 uL, or 1e7 cells to 2.5 uL, or 1e7 cells to 1.25 uL. In one embodiment, e.g., for T regulatory cells, e.g., CD25+ depletion, greater than 500 million cells/ml is used. In a further aspect, a concentration of cells of 600, 700, 800, or 900 million cells/ml is used.

In one embodiment, the population of immune effector cells to be depleted includes about 6 x 10^9 CD25+ T cells. In other aspects, the population of immune effector cells to be depleted include about 1 x 10^9 to 1x 10^{10} CD25+ T cell, and any integer value in between. In one embodiment, the resulting population T regulatory depleted cells has 2 x 10^9 T regulatory cells, e.g., CD25+ cells, or less (e.g., 1 x 10^9, 5 x 10^8, 1 x 10^8, 5 x 10^7, 1 x 10^7, or less CD25+ cells).

In one embodiment, the T regulatory cells, e.g., CD25+ cells, are removed from the population using the CliniMAC system with a depletion tubing set, such as, e.g., tubing 162-01. In one embodiment, the CliniMAC system is run on a depletion setting such as, e.g., DEPLETION2.1.

Without wishing to be bound by a particular theory, decreasing the level of negative regulators of immune cells (e.g., decreasing the number of unwanted immune cells, e.g., T_{REG} cells), in a subject prior to apheresis or during manufacturing of a CAR-expressing cell product can reduce the risk of subject relapse. For example, methods of depleting T_{REG} cells are known in the art. Methods of decreasing T_{REG} cells include, but are not limited to, cyclophosphamide, anti-GITR antibody (an anti-GITR antibody described herein), CD25-depletion, and combinations thereof. These methods may be combined with the methods of manufacture that include contacting the population of immune effector cells with an LSD1 inhibitor as described herein.

In some embodiments, the manufacturing methods comprise reducing the number of (e.g., depleting) T_{REG} cells prior to manufacturing of the CAR-expressing cell. For example, manufacturing methods comprise contacting the sample, e.g., the apheresis sample, with an anti-GITR antibody and/or an anti-CD25 antibody (or fragment thereof, or a CD25-binding ligand), e.g., to deplete T_{REG} cells prior to manufacturing of the CAR-expressing cell (e.g., T cell, NK cell) product. These methods may be combined with the methods of manufacture that include contacting the population of immune effector cells with an LSD1 inhibitor as described herein.

In an embodiment, a subject is pre-treated with one or more therapies that reduce T_{REG} cells prior to collection of cells for CAR-expressing cell product manufacturing, thereby reducing the risk of subject relapse to CAR-expressing cell treatment. In an embodiment, methods of decreasing T_{REG} cells include, but are not limited to, administration to the subject of one or more of cyclophosphamide, anti-GITR antibody, CD25-depletion, or a combination thereof. Administration of one or more of cyclophosphamide, anti-GITR antibody, CD25-depletion, or a combination thereof, can occur before, during or after an infusion of the CAR-expressing cell product. These methods may be combined with the methods of manufacture that include contacting the population of immune effector cells with an LSD1 inhibitor as described herein.

In an embodiment, a subject is pre-treated with cyclophosphamide prior to collection of cells for CAR-expressing cell product manufacturing, thereby reducing the risk of subject relapse to CAR-expressing cell treatment. In an embodiment, a subject is pre-treated with an anti-GITR antibody prior to collection of cells for CAR-expressing cell product manufacturing, thereby reducing the risk of subject relapse to CAR-expressing cell treatment. These methods may be combined with the methods of manufacture that include contacting the population of immune effector cells with an LSD1 inhibitor as described herein.

In one embodiment, the population of cells to be removed are neither the regulatory T cells or tumor cells, but cells that otherwise negatively affect the expansion and/or function of CART cells, e.g. cells expressing CD14, CD11b, CD33, CD15, or other markers expressed by potentially immune suppressive cells. In one embodiment, such cells are envisioned to be removed concurrently with regulatory T cells and/or tumor cells, or following said depletion, or in another order.

The methods described herein can include more than one selection step, e.g., more than one depletion step. Enrichment of a T cell population by negative selection can be accomplished, e.g., with a combination of antibodies directed to surface markers unique to the negatively selected cells. One method is cell sorting and/or selection via negative magnetic immunoadherence or flow cytometry that uses a cocktail of monoclonal antibodies directed to cell surface markers present on the cells negatively selected. For example, to enrich for CD4+ cells by negative selection, a monoclonal antibody cocktail can include antibodies to CD14, CD20, CD11b, CD16, HLA-DR, and CD8. These steps may be combined with the methods of manufacture that include contacting the population of immune effector cells with an LSD1 inhibitor as described herein.

The methods described herein can further include removing cells from the population which express a tumor antigen, e.g., a tumor antigen that does not comprise CD25, e.g., CD19, CD30, CD38, CD123, CD20, CD14 or CD11b, to thereby provide a population of T regulatory depleted, e.g., CD25+ depleted, and tumor antigen depleted cells that are suitable for expression of a CAR, e.g., a CAR described herein. In one embodiment, tumor antigen expressing cells are removed simultaneously with the T regulatory, e.g., CD25+ cells. For example, an anti-CD25 antibody, or fragment thereof, and an anti-tumor antigen antibody, or fragment thereof, can be attached to the same substrate, e.g., bead, which can be used to remove the cells or an anti-CD25 antibody, or fragment thereof, or the anti-tumor antigen antibody, or fragment thereof, can be attached to separate beads, a mixture of which can be used to remove the cells. In other embodiments, the removal of T regulatory cells, e.g., CD25+ cells, and the removal of the tumor antigen expressing cells is sequential, and can occur, e.g., in either order. These steps may be combined with the methods of manufacture that include contacting the population of immune effector cells with an LSD1 inhibitor as described herein.

Also provided are methods that include removing cells from the population which express a check point inhibitor, e.g., a check point inhibitor described herein, e.g., one or more of PD1+ cells, LAG3+ cells, and TIM3+ cells, to thereby provide a population of T regulatory depleted, e.g., CD25+ depleted cells, and check point inhibitor depleted cells, e.g., PD1+, LAG3+ and/or TIM3+ depleted cells. Exemplary check point inhibitors include B7-H1, B7-1, CD160, P1H, 2B4, PD1, TIM3, CEACAM (e.g., CEACAM-1, CEACAM-3 and/or CEACAM-5), LAG3, TIGIT, CTLA-4, BTLA and LAIR1. In one embodiment, check point inhibitor expressing cells are removed simultaneously with the T regulatory, e.g., CD25+ cells. For example, an anti-CD25 antibody, or fragment thereof, and an anti-check point inhibitor antibody, or fragment thereof, can be attached to the same bead which can be used to remove the cells, or an anti-CD25 antibody, or fragment thereof, and the anti-check point inhibitor antibody, or fragment there, can be attached to separate beads, a mixture of which can be used to remove the cells. In other embodiments, the removal of T regulatory cells, e.g., CD25+ cells, and the removal of the check point inhibitor expressing cells is sequential, and can occur, e.g., in either order. These steps may be combined with the methods of manufacture that include contacting the population of immune effector cells with an LSD1 inhibitor as described herein.

Methods described herein can include a positive selection step. For example, T cells can isolated by incubation with anti-CD3/anti-CD28 (e.g., 3x28)-conjugated beads, such as DYNABEADS® M-450 CD3/CD28 T, for a time period sufficient for positive selection of the desired T cells. In one

embodiment, the time period is about 30 minutes. In a further embodiment, the time period ranges from 30 minutes to 36 hours or longer and all integer values there between. In a further embodiment, the time period is at least 1, 2, 3, 4, 5, or 6 hours. In yet another embodiment, the time period is 10 to 24 hours, e.g., 24 hours. Longer incubation times may be used to isolate T cells in any situation where there are few T cells as compared to other cell types, such in isolating tumor infiltrating lymphocytes (TIL) from tumor tissue or from immunocompromised individuals. Further, use of longer incubation times can increase the efficiency of capture of CD8+ T cells. Thus, by simply shortening or lengthening the time T cells are allowed to bind to the CD3/CD28 beads and/or by increasing or decreasing the ratio of beads to T cells (as described further herein), subpopulations of T cells can be preferentially selected for or against at culture initiation or at other time points during the process. Additionally, by increasing or decreasing the ratio of anti-CD3 and/or anti-CD28 antibodies on the beads or other surface, subpopulations of T cells can be preferentially selected for or against at culture initiation or at other desired time points. These steps may be combined with the methods of manufacture that include contacting the population of immune effector cells with an LSD1 inhibitor as described herein.

In one embodiment, a T cell population can be selected that expresses one or more of IFN-$^{\gamma}$, TNFα, IL-17A, IL-2, IL-3, IL-4, GM-CSF, IL-10, IL-13, granzyme B, and perforin, or other appropriate molecules, e.g., other cytokines. Methods for screening for cell expression can be determined, e.g., by the methods described in PCT Publication No.: WO 2013/126712.

For isolation of a desired population of cells by positive or negative selection, the concentration of cells and surface (e.g., particles such as beads) can be varied. In certain aspects, it may be desirable to significantly decrease the volume in which beads and cells are mixed together (e.g., increase the concentration of cells), to ensure maximum contact of cells and beads. For example, in one aspect, a concentration of 10 billion cells/ml, 9 billion/ml, 8 billion/ml, 7 billion/ml, 6 billion/ml, or 5 billion/ml is used. In one aspect, a concentration of 1 billion cells/ml is used. In yet one aspect, a concentration of cells from 75, 80, 85, 90, 95, or 100 million cells/ml is used. In further aspects, concentrations of 125 or 150 million cells/ml can be used.

Using high concentrations can result in increased cell yield, cell activation, and cell expansion. Further, use of high cell concentrations allows more efficient capture of cells that may weakly express target antigens of interest, such as CD28-negative T cells, or from samples where there are many tumor cells present (e.g., leukemic blood, tumor tissue, etc.). Such populations of cells may have therapeutic

value and would be desirable to obtain. For example, using high concentration of cells allows more efficient selection of CD8+ T cells that normally have weaker CD28 expression.

In a related aspect, it may be desirable to use lower concentrations of cells. By significantly diluting the mixture of T cells and surface (e.g., particles such as beads), interactions between the particles and cells are minimized. This selects for cells that express high amounts of desired antigens to be bound to the particles. For example, CD4+ T cells express higher levels of CD28 and are more efficiently captured than CD8+ T cells in dilute concentrations. In one aspect, the concentration of cells used is 5 x 10^6/ml. In other aspects, the concentration used can be from about 1 x 10^5/ml to 1 x 10^6/ml, and any integer value in between.

In other aspects, the cells may be incubated on a rotator for varying lengths of time at varying speeds at either 2-10°C or at room temperature.

T cells for stimulation can also be frozen after a washing step. Wishing not to be bound by theory, the freeze and subsequent thaw step provide a more uniform product by removing granulocytes and to some extent monocytes in the cell population. After the washing step that removes plasma and platelets, the cells may be suspended in a freezing solution. While many freezing solutions and parameters are known in the art and will be useful in this context, one method involves using PBS containing 20% DMSO and 8% human serum albumin, or culture media containing 10% Dextran 40 and 5% Dextrose, 20% Human Serum Albumin and 7.5% DMSO, or 31.25% Plasmalyte-A, 31.25% Dextrose 5%, 0.45% NaCl, 10% Dextran 40 and 5% Dextrose, 20% Human Serum Albumin, and 7.5% DMSO or other suitable cell freezing media containing for example, Hespan and PlasmaLyte A, the cells then are frozen to -80°C at a rate of 1° per minute and stored in the vapor phase of a liquid nitrogen storage tank. Other methods of controlled freezing may be used as well as uncontrolled freezing immediately at -20° C or in liquid nitrogen.

In certain aspects, cryopreserved cells are thawed and washed as described herein and allowed to rest for one hour at room temperature prior to activation using the methods of the present invention.

Also contemplated in the context of the invention is the collection of blood samples or apheresis product from a subject at a time period prior to when the expanded cells as described herein might be needed. As such, the source of the cells to be expanded can be collected at any time point necessary, and desired cells, such as T cells, can be isolated and frozen for later use in immune effector cell therapy for any number of diseases or conditions that would benefit from immune effector cell therapy, such as those described herein. In one aspect a blood sample or an apheresis is taken from a generally healthy

subject. In certain aspects, a blood sample or an apheresis is taken from a generally healthy subject who is at risk of developing a disease, but who has not yet developed a disease, and the cells of interest are isolated and frozen for later use. In certain aspects, the T cells may be expanded, frozen, and used at a later time. In certain aspects, samples are collected from a patient shortly after diagnosis of a particular disease as described herein but prior to any treatments. In a further aspect, the cells are isolated from a blood sample or an apheresis from a subject prior to any number of relevant treatment modalities, including but not limited to treatment with agents such as natalizumab, efalizumab, antiviral agents, chemotherapy, radiation, immunosuppressive agents, such as cyclosporin, azathioprine, methotrexate, mycophenolate, and FK506, antibodies, or other immunoablative agents such as CAMPATH, anti-CD3 antibodies, cytoxan, fludarabine, cyclosporin, FK506, rapamycin, mycophenolic acid, steroids, FR901228, and irradiation.

In a further aspect of the present invention, T cells are obtained from a patient directly following treatment that leaves the subject with functional T cells. In this regard, it has been observed that following certain cancer treatments, in particular treatments with drugs that damage the immune system, shortly after treatment during the period when patients would normally be recovering from the treatment, the quality of T cells obtained may be optimal or improved for their ability to expand ex vivo. Likewise, following ex vivo manipulation using the methods described herein, these cells may be in a preferred state for enhanced engraftment and in vivo expansion. Thus, it is contemplated within the context of the present invention to collect blood cells, including T cells, dendritic cells, or other cells of the hematopoietic lineage, during this recovery phase. Further, in certain aspects, mobilization (for example, mobilization with GM-CSF) and conditioning regimens can be used to create a condition in a subject wherein repopulation, recirculation, regeneration, and/or expansion of particular cell types is favored, especially during a defined window of time following therapy. Illustrative cell types include T cells, B cells, dendritic cells, and other cells of the immune system.

In one embodiment, the immune effector cells expressing a CAR molecule, e.g., a CAR molecule described herein, are obtained from a subject that has received an LSD1 inhibitor. In an embodiment, the population of immune effector cells, e.g., T cells, to be engineered to express a CAR, are harvested after a sufficient time, or after sufficient dosing of the LSD1 inhibitor, such that the level of PD1 negative immune effector cells, e.g., T cells, or the ratio of PD1 negative immune effector cells, e.g., T cells/ PD1 positive immune effector cells, e.g., T cells, in the subject or harvested from the subject has been, at least transiently, increased.

In other embodiments, population of immune effector cells, e.g., T cells, which have, or will be engineered to express a CAR, can be treated ex vivo by contact with an amount of an LSD1 inhibitor that increases the number of PD1 negative immune effector cells, e.g., T cells or increases the ratio of PD1 negative immune effector cells, e.g., T cells/ PD1 positive immune effector cells, e.g., T cells.

In one embodiment, a T cell population is diaglycerol kinase (DGK)-deficient. DGK-deficient cells include cells that do not express DGK RNA or protein, or have reduced or inhibited DGK activity. DGK-deficient cells can be generated by genetic approaches, e.g., administering RNA-interfering agents, e.g., siRNA, shRNA, miRNA, to reduce or prevent DGK expression. Alternatively, DGK-deficient cells can be generated by treatment with DGK inhibitors described herein.

In one embodiment, a T cell population is Ikaros-deficient. Ikaros-deficient cells include cells that do not express Ikaros RNA or protein, or have reduced or inhibited Ikaros activity, Ikaros-deficient cells can be generated by genetic approaches, e.g., administering RNA-interfering agents, e.g., siRNA, shRNA, miRNA, to reduce or prevent Ikaros expression. Alternatively, Ikaros-deficient cells can be generated by treatment with Ikaros inhibitors, e.g., lenalidomide.

In embodiments, a T cell population is DGK-deficient and Ikaros-deficient, e.g., does not express DGK and Ikaros, or has reduced or inhibited DGK and Ikaros activity. Such DGK and Ikaros-deficient cells can be generated by any of the methods described herein.

In an embodiment, the NK cells are obtained from the subject. In another embodiment, the NK cells are an NK cell line, e.g., NK-92 cell line (Conkwest).

Additional Expressed Agents

In another embodiment, a CAR-expressing immune effector cell described herein can further express another agent, e.g., an agent which enhances the activity of a CAR-expressing cell. For example, in one embodiment, the agent can be an agent which inhibits an inhibitory molecule. Examples of inhibitory molecules include PD-1, PD-L1, CTLA-4, TIM-3, CEACAM (e.g., CEACAM-1, CEACAM-3 and/or CEACAM-5), LAG-3, VISTA, BTLA, TIGIT, LAIR1, CD160, 2B4 and TGF beta, e.g., as described herein. In one embodiment, the agent that inhibits an inhibitory molecule comprises a first polypeptide, e.g., an inhibitory molecule, associated with a second polypeptide that provides a positive signal to the cell, e.g., an intracellular signaling domain described herein. In one embodiment, the agent comprises a first polypeptide, e.g., of an inhibitory molecule such as PD-1, PD-L1, CTLA-4, TIM-3, CEACAM (e.g., CEACAM-1, CEACAM-3 and/or CEACAM-5), LAG-3, VISTA, BTLA, TIGIT, LAIR1, CD160, 2B4 or TGF beta, or a fragment of any of these, and a second polypeptide which is an intracellular signaling domain described

herein (e.g., comprising a costimulatory domain (e.g., 41BB, CD27 or CD28, e.g., as described herein) and/or a primary signaling domain (e.g., a CD3 zeta signaling domain described herein). In one embodiment, the agent comprises a first polypeptide of PD-1 or a fragment thereof, and a second polypeptide of an intracellular signaling domain described herein (e.g., a CD28, CD27, OX40 or 4-IBB signaling domain described herein and/or a CD3 zeta signaling domain described herein).

In one embodiment, the CAR-expressing immune effector cell described herein can further comprise a second CAR, e.g., a second CAR that includes a different antigen binding domain, e.g., to the same target (e.g., a target described above) or a different target. In one embodiment, the second CAR includes an antigen binding domain to a target expressed on the same cancer cell type as the target of the first CAR. In one embodiment, the CAR-expressing immune effector cell comprises a first CAR that targets a first antigen and includes an intracellular signaling domain having a costimulatory signaling domain but not a primary signaling domain, and a second CAR that targets a second, different, antigen and includes an intracellular signaling domain having a primary signaling domain but not a costimulatory signaling domain.

While not wishing to be bound by theory, placement of a costimulatory signaling domain, e.g., 4-1BB, CD28, CD27 or OX-40, onto the first CAR, and the primary signaling domain, e.g., CD3 zeta, on the second CAR can limit the CAR activity to cells where both targets are expressed. In one embodiment, the CAR expressing immune effector cell comprises a first CAR that includes an antigen binding domain that targets, e.g., a target described above, a transmembrane domain and a costimulatory domain and a second CAR that targets an antigen other than antigen targeted by the first CAR (e.g., an antigen expressed on the same cancer cell type as the first target) and includes an antigen binding domain, a transmembrane domain and a primary signaling domain. In another embodiment, the CAR expressing immune effector cell comprises a first CAR that includes an antigen binding domain that targets, e.g., a target described above, a transmembrane domain and a primary signaling domain and a second CAR that targets an antigen other than antigen targeted by the first CAR (e.g., an antigen expressed on the same cancer cell type as the first target) and includes an antigen binding domain to the antigen, a transmembrane domain and a costimulatory signaling domain.

In one embodiment, the CAR-expressing immune effector cell comprises a CAR described herein, e.g., a CAR to a target described above, and an inhibitory CAR. In one embodiment, the inhibitory CAR comprises an antigen binding domain that binds an antigen found on normal cells but not cancer cells, e.g., normal cells that also express the target. In one embodiment, the inhibitory CAR

comprises the antigen binding domain, a transmembrane domain and an intracellular domain of an inhibitory molecule. For example, the intracellular domain of the inhibitory CAR can be an intracellular domain of PD1, PD-L1, CTLA-4, TIM-3, CEACAM (e.g., CEACAM-1, CEACAM-3 and/or CEACAM-5), LAG-3, VISTA, BTLA, TIGIT, LAIR1, CD160, 2B4 or TGF beta.

In one embodiment, an immune effector cell (e.g., T cell, NK cell) comprises a first CAR comprising an antigen binding domain that binds to a tumor antigen as described herein, and a second CAR comprising a PD1 extracellular domain or a fragment thereof.

In one embodiment, the cell further comprises an inhibitory molecule as described above.

In one embodiment, the second CAR in the cell is an inhibitory CAR, wherein the inhibitory CAR comprises an antigen binding domain, a transmembrane domain, and an intracellular domain of an inhibitory molecule. The inhibitory molecule can be chosen from one or more of: PD1, PD-L1, CTLA-4, TIM-3, LAG-3, VISTA, BTLA, TIGIT, LAIR1, CD160, 2B4, TGF beta, CEACAM-1, CEACAM-3, and CEACAM-5. In one embodiment, the second CAR molecule comprises the extracellular domain of PD1 or a fragment thereof.

In embodiments, the second CAR molecule in the cell further comprises an intracellular signaling domain comprising a primary signaling domain and/or an intracellular signaling domain.

In other embodiments, the intracellular signaling domain in the cell comprises a primary signaling domain comprising the functional domain of CD3 zeta and a costimulatory signaling domain comprising the functional domain of 4-1BB.

In one embodiment, the second CAR molecule in the cell comprises the amino acid sequence of SEQ ID NO: 26.

In certain embodiments, the antigen binding domain of the first CAR molecule comprises a scFv and the antigen binding domain of the second CAR molecule does not comprise a scFv. For example, the antigen binding domain of the first CAR molecule comprises a scFv and the antigen binding domain of the second CAR molecule comprises a camelid VHH domain.

Split CAR

In some embodiments, the CAR-expressing cell uses a split CAR. The split CAR approach is described in more detail in publications WO2014/055442 and WO2014/055657. Briefly, a split CAR

system comprises a cell expressing a first CAR having a first antigen binding domain and a costimulatory domain (e.g., 41BB), and the cell also expresses a second CAR having a second antigen binding domain and an intracellular signaling domain (e.g., CD3 zeta). When the cell encounters the first antigen, the costimulatory domain is activated, and the cell proliferates. When the cell encounters the second antigen, the intracellular signaling domain is activated and cell-killing activity begins. Thus, the CAR-expressing cell is only fully activated in the presence of both antigens.

Multiple CAR expression

In one aspect, the CAR-expressing cell described herein can further comprise a second CAR, e.g., a second CAR that includes a different antigen binding domain, e.g., to the same target or a different target (e.g., a target other than a cancer associated antigen described herein or a different cancer associated antigen described herein). In one embodiment, the second CAR includes an antigen binding domain to a target expressed the same cancer cell type as the cancer associated antigen. In one embodiment, the CAR-expressing cell comprises a first CAR that targets a first antigen and includes an intracellular signaling domain having a costimulatory signaling domain but not a primary signaling domain, and a second CAR that targets a second, different, antigen and includes an intracellular signaling domain having a primary signaling domain but not a costimulatory signaling domain. While not wishing to be bound by theory, placement of a costimulatory signaling domain, e.g., 4-1BB, CD28, CD27 or OX-40, onto the first CAR, and the primary signaling domain, e.g.,CD3 zeta, on the second CAR can limit the CAR activity to cells where both targets are expressed. In one embodiment, the CAR expressing cell comprises a first cancer associated antigen CAR that includes an antigen binding domain that binds a target antigen described herein, a transmembrane domain and a costimulatory domain and a second CAR that targets a different target antigen (e.g., an antigen expressed on that same cancer cell type as the first target antigen) and includes an antigen binding domain, a transmembrane domain and a primary signaling domain. In another embodiment, the CAR expressing cell comprises a first CAR that includes an antigen binding domain that binds a target antigen described herein, a transmembrane domain and a primary signaling domain and a second CAR that targets an antigen other than the first target antigen (e.g., an antigen expressed on the same cancer cell type as the first target antigen) and includes an antigen binding domain to the antigen, a transmembrane domain and a costimulatory signaling domain.

In some embodiments, the claimed invention comprises a first and second CAR, wherein the antigen binding domain of one of said first CAR said second CAR does not comprise a variable light

domain and a variable heavy domain. In some embodiments, the antigen binding domain of one of said first CAR said second CAR is an scFv, and the other is not an scFv. In some embodiments, the antigen binding domain of one of said first CAR said second CAR comprises a single VH domain, e.g., a camelid, shark, or lamprey single VH domain, or a single VH domain derived from a human or mouse sequence. In some embodiments, the antigen binding domain of one of said first CAR said second CAR comprises a nanobody. In some embodiments, the antigen binding domain of one of said first CAR said second CAR comprises a camelid VHH domain.

Allogeneic cells

In embodiments described herein, the immune effector cell can be an allogeneic immune effector cell, e.g., T cell or NK cell. For example, the cell can be an allogeneic T cell, e.g., an allogeneic T cell lacking expression of a functional T cell receptor (TCR) and/or human leukocyte antigen (HLA), e.g., HLA class I and/or HLA class II.

A T cell lacking a functional TCR can be, e.g., engineered such that it does not express any functional TCR on its surface, engineered such that it does not express one or more subunits that comprise a functional TCR or engineered such that it produces very little functional TCR on its surface. Alternatively, the T cell can express a substantially impaired TCR, e.g., by expression of mutated or truncated forms of one or more of the subunits of the TCR. The term “substantially impaired TCR” means that this TCR will not elicit an adverse immune reaction in a host.

A T cell described herein can be, e.g., engineered such that it does not express a functional HLA on its surface. For example, a T cell described herein, can be engineered such that cell surface expression HLA, e.g., HLA class 1 and/or HLA class II, is downregulated.

In some embodiments, the T cell can lack a functional TCR and a functional HLA, e.g., HLA class I and/or HLA class II.

Modified T cells that lack expression of a functional TCR and/or HLA can be obtained by any suitable means, including a knock out or knock down of one or more subunit of TCR or HLA. For example, the T cell can include a knock down of TCR and/or HLA using siRNA, shRNA, clustered regularly interspaced short palindromic repeats (CRISPR) transcription-activator like effector nuclease (TALEN), or zinc finger endonuclease (ZFN).

In some embodiments, the allogeneic cell can be a cell which does not express or expresses at low levels an inhibitory molecule, e.g. by any method described herein. For example, the cell can be a

cell that does not express or expresses at low levels an inhibitory molecule, e.g., that can decrease the ability of a CAR-expressing cell to mount an immune effector response. Examples of inhibitory molecules include PD1, PD-L1, CTLA4, TIM3, CEACAM (e.g., CEACAM-1, CEACAM-3 and/or CEACAM-5), LAG3, VISTA, BTLA, TIGIT, LAIR1, CD160, 2B4 and TGF beta. Inhibition of an inhibitory molecule, e.g., by inhibition at the DNA, RNA or protein level, can optimize a CAR-expressing cell performance. In embodiments, an inhibitory nucleic acid, e.g., an inhibitory nucleic acid, e.g., a dsRNA, e.g., an siRNA or shRNA, a clustered regularly interspaced short palindromic repeats (CRISPR), a transcription-activator like effector nuclease (TALEN), or a zinc finger endonuclease (ZFN), e.g., as described herein, can be used.

siRNA and shRNA to inhibit TCR or HLA

In some embodiments, TCR expression and/or HLA expression can be inhibited using siRNA or shRNA that targets a nucleic acid encoding a TCR and/or HLA in a T cell.

CRISPR Systems to inhibit TCR or HLA

"CRISPR system to TCR and/or HLA" or "CRISPR to inhibit TCR and/or HLA" as used herein refers to a CRISPR system (e.g., a CRISPR/Cas9 system) comprising one or more guide RNA molecules comprising a targeting domain complementary to target sequence within a gene for a component of the TCR and/or HLA or beta-2 microglobulin (B2M), and a RNA-guided endonuclease (e.g., a Cas, e.g., a Cas9).

Artificial CRISPR/Cas systems can be generated which inhibit TCR and/or HLA, using technology known in the art, e.g., that are described in U.S. Publication No. 20140068797, and Cong (2013) Science 339: 819-823. Other artificial CRISPR/Cas systems that are known in the art may also be generated which inhibit TCR and/or HLA, e.g., that described in Tsai (2014) Nature Biotechnol., 32:6 569-576, U.S. Patent No.: 8,871,445; 8,865,406; 8,795,965; 8,771,945; and 8,697,359.

TALEN to inhibit TCR and/or HLA

"TALEN genome editing system to HLA and/or TCR" or "TALEN to inhibit HLA and/or TCR" refers to a transcription activator-like effector nuclease, an artificial nuclease which can be used to edit the HLA and/or TCR gene.

TALENs are produced artificially by fusing a TAL effector DNA binding domain to a DNA cleavage domain. Transcription activator-like effects (TALEs) can be engineered to bind any desired DNA sequence, including a portion of the HLA or TCR gene. By combining an engineered TALE with a DNA cleavage domain, a restriction enzyme can be produced which is specific to any desired DNA sequence,

including a HLA or TCR sequence. These can then be introduced into a cell, wherein they can be used for genome editing. Boch (2011) *Nature Biotech*. 29: 135-6; and Boch et al. (2009) *Science* 326: 1509-12; Moscou et al. (2009) *Science* 326: 3501.

TALENs specific to sequences in HLA or TCR can be constructed using any method known in the art, including various schemes using modular components. Zhang et al. (2011) *Nature Biotech.* 29: 149-53; Geibler et al. (2011) *PLoS ONE* 6: e19509.

Zinc finger nuclease to inhibit HLA and/or TCR

"ZFN genome editing system to HLA and/or TCR" or "ZFN to inhibit HLA and/or TCR" refer to a zinc finger nuclease, an artificial nuclease which can be used to edit the HLA and/or TCR gene.

ZFNs specific to sequences in HLA AND/OR TCR can be constructed using any method known in the art. See, e.g., Provasi (2011) Nature Med. 18: 807-815; Torikai (2013) Blood 122: 1341-1349; Cathomen et al. (2008) *Mol. Ther.* 16: 1200-7; Guo et al. (2010) *J. Mol. Biol.* 400: 96; U.S. Patent Publication 2011/0158957; and U.S. Patent Publication 2012/0060230.

Telomerase expression

While not wishing to be bound by any particular theory, in some embodiments, a therapeutic T cell has short term persistence in a patient, due to shortened telomeres in the T cell; accordingly, transfection with a telomerase gene can lengthen the telomeres of the T cell and improve persistence of the T cell in the patient. See Carl June, "Adoptive T cell therapy for cancer in the clinic", Journal of Clinical Investigation, 117:1466-1476 (2007). Thus, in an embodiment, an immune effector cell, e.g., a T cell, ectopically expresses a telomerase subunit, e.g., the catalytic subunit of telomerase, e.g., TERT, e.g., hTERT. In some aspects, this disclosure provides a method of producing a CAR-expressing cell, comprising contacting a cell with a nucleic acid encoding a telomerase subunit, e.g., the catalytic subunit of telomerase, e.g., TERT, e.g., hTERT. The cell may be contacted with the nucleic acid before, simultaneous with, or after being contacted with a construct encoding a CAR.

Expansion and Activation

Immune effector cells such as T cells may be activated and expanded generally using methods as described, for example, in U.S. Patents 6,352,694; 6,534,055; 6,905,680; 6,692,964; 5,858,358; 6,887,466; 6,905,681; 7,144,575; 7,067,318; 7,172,869; 7,232,566; 7,175,843; 5,883,223; 6,905,874; 6,797,514; 6,867,041; and U.S. Patent Application Publication No. 20060121005, each of which is incorporated by reference in its entirety.

Generally, a population of immune effector cells e.g., T cells, may be expanded by contact with a surface having attached thereto an agent that stimulates a CD3/TCR complex associated signal and a ligand that stimulates a costimulatory molecule on the surface of the T cells. In particular, T cell populations may be stimulated as described herein, such as by contact with an anti-CD3 antibody, or antigen-binding fragment thereof, or an anti-CD2 antibody immobilized on a surface, or by contact with a protein kinase C activator (e.g., bryostatin) in conjunction with a calcium ionophore. For co-stimulation of an accessory molecule on the surface of the T cells, a ligand that binds the accessory molecule is used. For example, a population of T cells can be contacted with an anti-CD3 antibody and an anti-CD28 antibody, under conditions appropriate for stimulating proliferation of the T cells. To stimulate proliferation of either CD4+ T cells or CD8+ T cells, an anti-CD3 antibody and an anti-CD28 antibody can be used. Examples of an anti-CD28 antibody include 9.3, B-T3, XR-CD28 (Diaclone, Besançon, France) can be used as can other methods commonly known in the art (Berg et al., Transplant Proc. 30(8):3975-3977, 1998; Haanen et al., J. Exp. Med. 190(9):13191328, 1999; Garland et al., J. Immunol Meth. 227(1-2):53-63, 1999).

In certain aspects, the primary stimulatory signal and the costimulatory signal for the T cell may be provided by different protocols. For example, the agents providing each signal may be in solution or coupled to a surface. When coupled to a surface, the agents may be coupled to the same surface (i.e., in "cis" formation) or to separate surfaces (i.e., in "trans" formation). Alternatively, one agent may be coupled to a surface and the other agent in solution. In one aspect, the agent providing the costimulatory signal is bound to a cell surface and the agent providing the primary activation signal is in solution or coupled to a surface. In certain aspects, both agents can be in solution. In one aspect, the agents may be in soluble form, and then cross-linked to a surface, such as a cell expressing Fc receptors or an antibody or other binding agent which will bind to the agents. In this regard, see for example, U.S. Patent Application Publication Nos. 20040101519 and 20060034810 for artificial antigen presenting cells (aAPCs) that are contemplated for use in activating and expanding T cells in the present invention.

In one aspect, the two agents are immobilized on beads, either on the same bead, i.e., "cis," or to separate beads, i.e., "trans." By way of example, the agent providing the primary activation signal is an anti-CD3 antibody or an antigen-binding fragment thereof and the agent providing the costimulatory signal is an anti-CD28 antibody or antigen-binding fragment thereof; and both agents are co-immobilized to the same bead in equivalent molecular amounts. In one aspect, a 1:1 ratio of each antibody bound to the beads for CD4+ T cell expansion and T cell growth is used. In certain aspects of

the present invention, a ratio of anti CD3:CD28 antibodies bound to the beads is used such that an increase in T cell expansion is observed as compared to the expansion observed using a ratio of 1:1. In one particular aspect an increase of from about 1 to about 3 fold is observed as compared to the expansion observed using a ratio of 1:1. In one aspect, the ratio of CD3:CD28 antibody bound to the beads ranges from 100:1 to 1:100 and all integer values there between. In one aspect, more anti-CD28 antibody is bound to the particles than anti-CD3 antibody, i.e., the ratio of CD3:CD28 is less than one. In certain aspects, the ratio of anti CD28 antibody to anti CD3 antibody bound to the beads is greater than 2:1. In one particular aspect, a 1:100 CD3:CD28 ratio of antibody bound to beads is used. In one aspect, a 1:75 CD3:CD28 ratio of antibody bound to beads is used. In a further aspect, a 1:50 CD3:CD28 ratio of antibody bound to beads is used. In one aspect, a 1:30 CD3:CD28 ratio of antibody bound to beads is used. In one preferred aspect, a 1:10 CD3:CD28 ratio of antibody bound to beads is used. In one aspect, a 1:3 CD3:CD28 ratio of antibody bound to the beads is used. In yet one aspect, a 3:1 CD3:CD28 ratio of antibody bound to the beads is used.

Ratios of particles to cells from 1:500 to 500:1 and any integer values in between may be used to stimulate T cells or other target cells. As those of ordinary skill in the art can readily appreciate, the ratio of particles to cells may depend on particle size relative to the target cell. For example, small sized beads could only bind a few cells, while larger beads could bind many. In certain aspects the ratio of cells to particles ranges from 1:100 to 100:1 and any integer values in-between and in further aspects the ratio comprises 1:9 to 9:1 and any integer values in between, can also be used to stimulate T cells. The ratio of anti-CD3- and anti-CD28-coupled particles to T cells that result in T cell stimulation can vary as noted above, however certain preferred values include 1:100, 1:50, 1:40, 1:30, 1:20, 1:10, 1:9, 1:8, 1:7, 1:6, 1:5, 1:4, 1:3, 1:2, 1:1, 2:1, 3:1, 4:1, 5:1, 6:1, 7:1, 8:1, 9:1, 10:1, and 15:1 with one preferred ratio being at least 1:1 particles per T cell. In one aspect, a ratio of particles to cells of 1:1 or less is used. In one particular aspect, a preferred particle: cell ratio is 1:5. In further aspects, the ratio of particles to cells can be varied depending on the day of stimulation. For example, in one aspect, the ratio of particles to cells is from 1:1 to 10:1 on the first day and additional particles are added to the cells every day or every other day thereafter for up to 10 days, at final ratios of from 1:1 to 1:10 (based on cell counts on the day of addition). In one particular aspect, the ratio of particles to cells is 1:1 on the first day of stimulation and adjusted to 1:5 on the third and fifth days of stimulation. In one aspect, particles are added on a daily or every other day basis to a final ratio of 1:1 on the first day, and 1:5 on the third and fifth days of stimulation. In one aspect, the ratio of particles to cells is 2:1 on the first day of stimulation and adjusted to 1:10 on the third and fifth days of stimulation. In one aspect, particles are added on a

daily or every other day basis to a final ratio of 1:1 on the first day, and 1:10 on the third and fifth days of stimulation. One of skill in the art will appreciate that a variety of other ratios may be suitable for use in the present invention. In particular, ratios will vary depending on particle size and on cell size and type. In one aspect, the most typical ratios for use are in the neighborhood of 1:1, 2:1 and 3:1 on the first day.

In further aspects, the cells, such as T cells, are combined with agent-coated beads, the beads and the cells are subsequently separated, and then the cells are cultured. In an alternative aspect, prior to culture, the agent-coated beads and cells are not separated but are cultured together. In a further aspect, the beads and cells are first concentrated by application of a force, such as a magnetic force, resulting in increased ligation of cell surface markers, thereby inducing cell stimulation.

By way of example, cell surface proteins may be ligated by allowing paramagnetic beads to which anti-CD3 and anti-CD28 are attached (3x28 beads) to contact the T cells. In one aspect the cells (for example, 10^4 to 10^9 T cells) and beads (for example, DYNABEADS® M-450 CD3/CD28 T paramagnetic beads at a ratio of 1:1) are combined in a buffer, for example PBS (without divalent cations such as, calcium and magnesium). Again, those of ordinary skill in the art can readily appreciate any cell concentration may be used. For example, the target cell may be very rare in the sample and comprise only 0.01% of the sample or the entire sample (i.e., 100%) may comprise the target cell of interest. Accordingly, any cell number is within the context of the present invention. In certain aspects, it may be desirable to significantly decrease the volume in which particles and cells are mixed together (i.e., increase the concentration of cells), to ensure maximum contact of cells and particles. For example, in one aspect, a concentration of about 10 billion cells/ml, 9 billion/ml, 8 billion/ml, 7 billion/ml, 6 billion/ml, 5 billion/ml, or 2 billion cells/ml is used. In one aspect, greater than 100 million cells/ml is used. In a further aspect, a concentration of cells of 10, 15, 20, 25, 30, 35, 40, 45, or 50 million cells/ml is used. In yet one aspect, a concentration of cells from 75, 80, 85, 90, 95, or 100 million cells/ml is used. In further aspects, concentrations of 125 or 150 million cells/ml can be used. Using high concentrations can result in increased cell yield, cell activation, and cell expansion. Further, use of high cell concentrations allows more efficient capture of cells that may weakly express target antigens of interest, such as CD28-negative T cells. Such populations of cells may have therapeutic value and would be desirable to obtain in certain aspects. For example, using high concentration of cells allows more efficient selection of CD8+ T cells that normally have weaker CD28 expression.

In one embodiment, cells transduced with a nucleic acid encoding a CAR, e.g., a CAR described herein, are expanded, e.g., by a method described herein. In one embodiment, the cells are expanded in culture for a period of several hours (e.g., about 2, 3, 4, 5, 6, 7, 8, 9, 10, 15, 18, 21 hours) to about 14 days (e.g., 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13 or 14 days). In one embodiment, the cells are expanded for a period of 4 to 9 days. In one embodiment, the cells are expanded for a period of 8 days or less, e.g., 7, 6 or 5 days. In one embodiment, the cells are expanded in culture for 5 days, and the resulting cells are more potent than the same cells expanded in culture for 9 days under the same culture conditions. Potency can be defined, e.g., by various T cell functions, e.g. proliferation, target cell killing, cytokine production, activation, migration, or combinations thereof. In one embodiment, the cells are expanded for 5 days show at least a one, two, three or four fold increase in cells doublings upon antigen stimulation as compared to the same cells expanded in culture for 9 days under the same culture conditions. In one embodiment, the cells are expanded in culture for 5 days, and the resulting cells exhibit higher proinflammatory cytokine production, e.g., IFN-γ and/or GM-CSF levels, as compared to the same cells expanded in culture for 9 days under the same culture conditions. In one embodiment, the cells expanded for 5 days show at least a one, two, three, four, five, ten fold or more increase in pg/ml of proinflammatory cytokine production, e.g., IFN-γ and/or GM-CSF levels, as compared to the same cells expanded in culture for 9 days under the same culture conditions.

Several cycles of stimulation may also be desired such that culture time of T cells can be 60 days or more. Conditions appropriate for T cell culture include an appropriate media (e.g., Minimal Essential Media or RPMI Media 1640 or, X-vivo 15, (Lonza)) that may contain factors necessary for proliferation and viability, including serum (e.g., fetal bovine or human serum), interleukin-2 (IL-2), insulin, IFN-γ, IL-4, IL-7, GM-CSF, IL-10, IL-12, IL-15, TGFβ, and TNF-α or any other additives for the growth of cells known to the skilled artisan. Other additives for the growth of cells include, but are not limited to, surfactant, plasmanate, and reducing agents such as N-acetyl-cysteine and 2-mercaptoethanol. Media can include RPMI 1640, AIM-V, DMEM, MEM, α-MEM, F-12, X-Vivo 15, and X-Vivo 20, Optimizer, with added amino acids, sodium pyruvate, and vitamins, either serum-free or supplemented with an appropriate amount of serum (or plasma) or a defined set of hormones, and/or an amount of cytokine(s) sufficient for the growth and expansion of T cells. Antibiotics, e.g., penicillin and streptomycin, are included only in experimental cultures, not in cultures of cells that are to be infused into a subject. The target cells are maintained under conditions necessary to support growth, for example, an appropriate temperature (e.g., 37° C) and atmosphere (e.g., air plus 5% CO_2).

In one embodiment, the cells are expanded in an appropriate media (e.g., media described herein) that includes one or more interleukin that result in at least a 200-fold (e.g., 200-fold, 250-fold, 300-fold, 350-fold) increase in cells over a 14 day expansion period, e.g., as measured by a method described herein such as flow cytometry. In one embodiment, the cells are expanded in the presence of IL-15 and/or IL-7 (e.g., IL-15 and IL-7).

In embodiments, methods described herein, e.g., CAR-expressing cell manufacturing methods, comprise removing T regulatory cells, e.g., CD25+ T cells, from a cell population, e.g., using an anti-CD25 antibody, or fragment thereof, or a CD25-binding ligand, IL-2. Methods of removing T regulatory cells, e.g., CD25+ T cells, from a cell population are described herein. In embodiments, the methods, e.g., manufacturing methods, further comprise contacting a cell population (e.g., a cell population in which T regulatory cells, such as CD25+ T cells, have been depleted; or a cell population that has previously contacted an anti-CD25 antibody, fragment thereof, or CD25-binding ligand) with IL-15 and/or IL-7. For example, the cell population (e.g., that has previously contacted an anti-CD25 antibody, fragment thereof, or CD25-binding ligand) is expanded in the presence of IL-15 and/or IL-7.

In some embodiments a CAR-expressing cell described herein is contacted with a composition comprising a interleukin-15 (IL-15) polypeptide, a interleukin-15 receptor alpha (IL-15Ra) polypeptide, or a combination of both a IL-15 polypeptide and a IL-15Ra polypeptide e.g., hetIL-15, during the manufacturing of the CAR-expressing cell, e.g., ex vivo. In embodiments, a CAR-expressing cell described herein is contacted with a composition comprising a IL-15 polypeptide during the manufacturing of the CAR-expressing cell, e.g., ex vivo. In embodiments, a CAR-expressing cell described herein is contacted with a composition comprising a combination of both a IL-15 polypeptide and a IL-15 Ra polypeptide during the manufacturing of the CAR-expressing cell, e.g., ex vivo. In embodiments, a CAR-expressing cell described herein is contacted with a composition comprising hetIL-15 during the manufacturing of the CAR-expressing cell, e.g., ex vivo.

In one embodiment the CAR-expressing cell described herein is contacted with a composition comprising hetIL-15 during ex vivo expansion. In an embodiment, the CAR-expressing cell described herein is contacted with a composition comprising an IL-15 polypeptide during ex vivo expansion. In an embodiment, the CAR-expressing cell described herein is contacted with a composition comprising both an IL-15 polypeptide and an IL-15Ra polypeptide during ex vivo expansion. In one embodiment the contacting results in the survival and proliferation of a lymphocyte subpopulation, e.g., CD8+ T cells.

T cells that have been exposed to varied stimulation times may exhibit different characteristics. For example, typical blood or apheresed peripheral blood mononuclear cell products have a helper T cell population (TH, CD4+) that is greater than the cytotoxic or suppressor T cell population (TC, CD8+). Ex vivo expansion of T cells by stimulating CD3 and CD28 receptors produces a population of T cells that prior to about days 8-9 consists predominately of TH cells, while after about days 8-9, the population of T cells comprises an increasingly greater population of TC cells. Accordingly, depending on the purpose of treatment, infusing a subject with a T cell population comprising predominately of TH cells may be advantageous. Similarly, if an antigen-specific subset of TC cells has been isolated it may be beneficial to expand this subset to a greater degree.

Further, in addition to CD4 and CD8 markers, other phenotypic markers vary significantly, but in large part, reproducibly during the course of the cell expansion process. Thus, such reproducibility enables the ability to tailor an activated T cell product for specific purposes.

Once a CAR described herein is constructed, various assays can be used to evaluate the activity of the molecule, such as but not limited to, the ability to expand T cells following antigen stimulation, sustain T cell expansion in the absence of re-stimulation, and anti-cancer activities in appropriate in vitro and animal models. Assays to evaluate the effects of a CAR of the present invention are described in further detail below.

Western blot analysis of CAR expression in primary T cells can be used to detect the presence of monomers and dimers. See, *e.g.*, Milone *et al.*, Molecular Therapy 17(8): 1453-1464 (2009). Very briefly, T cells (1:1 mixture of $CD4^+$ and $CD8^+$ T cells) expressing the CARs are expanded *in vitro* for more than 10 days followed by lysis and SDS-PAGE under reducing conditions. CARs containing the full length TCR-ζ cytoplasmic domain and the endogenous TCR-ζ chain are detected by western blotting using an antibody to the TCR-ζ chain. The same T cell subsets are used for SDS-PAGE analysis under non-reducing conditions to permit evaluation of covalent dimer formation.

In vitro expansion of CAR^+ T cells following antigen stimulation can be measured by flow cytometry. For example, a mixture of $CD4^+$ and $CD8^+$ T cells are stimulated with αCD3/αCD28 aAPCs followed by transduction with lentiviral vectors expressing GFP under the control of the promoters to be analyzed. Exemplary promoters include the CMV IE gene, EF-1α, ubiquitin C, or phosphoglycerokinase (PGK) promoters. GFP fluorescence is evaluated on day 6 of culture in the $CD4^+$ and/or $CD8^+$ T cell subsets by flow cytometry. See, *e.g.*, Milone *et al.*, Molecular Therapy 17(8): 1453-1464 (2009). Alternatively, a mixture of $CD4^+$ and $CD8^+$ T cells are stimulated with αCD3/αCD28 coated magnetic

beads on day 0, and transduced with CAR on day 1 using a bicistronic lentiviral vector expressing CAR along with eGFP using a 2A ribosomal skipping sequence. Cultures are re-stimulated with either a cancer associated antigen as described herein$^+$ K562 cells (K562 expressing a cancer associated antigen as described herein), wild-type K562 cells (K562 wild type) or K562 cells expressing hCD32 and 4-1BBL in the presence of antiCD3 and anti-CD28 antibody (K562-BBL-3/28) following washing. Exogenous IL-2 is added to the cultures every other day at 100 IU/ml. GFP$^+$ T cells are enumerated by flow cytometry using bead-based counting. See, *e.g.*, Milone *et al.*, Molecular Therapy 17(8): 1453-1464 (2009).

Sustained CAR$^+$ T cell expansion in the absence of re-stimulation can also be measured. See, *e.g.*, Milone *et al.*, Molecular Therapy 17(8): 1453-1464 (2009). Briefly, mean T cell volume (fl) is measured on day 8 of culture using a Coulter Multisizer III particle counter, a Nexcelom Cellometer Vision or Millipore Scepter, following stimulation with αCD3/αCD28 coated magnetic beads on day 0, and transduction with the indicated CAR on day 1.

Animal models can also be used to measure a CART activity. For example, xenograft model using human a cancer associated antigen described herein-specific CAR$^+$ T cells to treat a primary human pre-B ALL in immunodeficient mice can be used. See, *e.g.*, Milone *et al.*, Molecular Therapy 17(8): 1453-1464 (2009). Very briefly, after establishment of ALL, mice are randomized as to treatment groups. Different numbers of a cancer associated antigen -specific CARengineered T cells are coinjected at a 1:1 ratio into NOD-SCID-$\gamma^{-/-}$ mice bearing B-ALL. The number of copies of a cancer associated antigen -specific CAR vector in spleen DNA from mice is evaluated at various times following T cell injection. Animals are assessed for leukemia at weekly intervals. Peripheral blood a cancer associate antigen as described herein$^+$ B-ALL blast cell counts are measured in mice that are injected with a cancer associated antigen described herein-ζ CAR$^+$ T cells or mock-transduced T cells. Survival curves for the groups are compared using the log-rank test. In addition, absolute peripheral blood CD4$^+$ and CD8$^+$ T cell counts 4 weeks following T cell injection in NOD-SCID-$\gamma^{-/-}$ mice can also be analyzed. Mice are injected with leukemic cells and 3 weeks later are injected with T cells engineered to express CAR by a bicistronic lentiviral vector that encodes the CAR linked to eGFP. T cells are normalized to 45–50% input GFP$^+$ T cells by mixing with mock-transduced cells prior to injection, and confirmed by flow cytometry. Animals are assessed for leukemia at 1-week intervals. Survival curves for the CAR$^+$ T cell groups are compared using the log-rank test.

Dose dependent CAR treatment response can be evaluated. See, *e.g.*, Milone *et al.*, Molecular Therapy 17(8): 1453-1464 (2009). For example, peripheral blood is obtained 35–70 days after

establishing leukemia in mice injected on day 21 with CAR T cells, an equivalent number of mock-transduced T cells, or no T cells. Mice from each group are randomly bled for determination of peripheral blood a cancer associate antigen as described herein$^+$ ALL blast counts and then killed on days 35 and 49. The remaining animals are evaluated on days 57 and 70.

Assessment of cell proliferation and cytokine production has been previously described, *e.g.*, at Milone *et al.*, Molecular Therapy 17(8): 1453-1464 (2009). Briefly, assessment of CAR-mediated proliferation is performed in microtiter plates by mixing washed T cells with K562 cells expressing a cancer associated antigen described herein (K19) or CD32 and CD137 (KT32-BBL) for a final T-cell:K562 ratio of 2:1. K562 cells are irradiated with gamma-radiation prior to use. Anti-CD3 (clone OKT3) and anti-CD28 (clone 9.3) monoclonal antibodies are added to cultures with KT32-BBL cells to serve as a positive control for stimulating T-cell proliferation since these signals support long-term CD8$^+$ T cell expansion *ex vivo*. T cells are enumerated in cultures using CountBright™ fluorescent beads (Invitrogen, Carlsbad, CA) and flow cytometry as described by the manufacturer. CAR$^+$ T cells are identified by GFP expression using T cells that are engineered with eGFP-2A linked CAR-expressing lentiviral vectors. For CAR+ T cells not expressing GFP, the CAR+ T cells are detected with biotinylated recombinant a cancer associate antigen as described herein protein and a secondary avidin-PE conjugate. CD4+ and CD8$^+$ expression on T cells are also simultaneously detected with specific monoclonal antibodies (BD Biosciences). Cytokine measurements are performed on supernatants collected 24 hours following re-stimulation using the human TH1/TH2 cytokine cytometric bead array kit (BD Biosciences, San Diego, CA) according the manufacturer's instructions. Fluorescence is assessed using a FACScalibur flow cytometer, and data is analyzed according to the manufacturer's instructions.

Cytotoxicity can be assessed by a standard 51Cr-release assay. See, *e.g.*, Milone *et al.*, Molecular Therapy 17(8): 1453-1464 (2009). Briefly, target cells (K562 lines and primary pro-B-ALL cells) are loaded with 51Cr (as NaCrO4, New England Nuclear, Boston, MA) at 37°C for 2 hours with frequent agitation, washed twice in complete RPMI and plated into microtiter plates. Effector T cells are mixed with target cells in the wells in complete RPMI at varying ratios of effector cell:target cell (E:T). Additional wells containing media only (spontaneous release, SR) or a 1% solution of triton-X 100 detergent (total release, TR) are also prepared. After 4 hours of incubation at 37°C, supernatant from each well is harvested. Released 51Cr is then measured using a gamma particle counter (Packard Instrument Co., Waltham, MA). Each condition is performed in at least triplicate, and the percentage of lysis is calculated

using the formula: % Lysis = (ER– SR) / (TR – SR), where ER represents the average 51Cr released for each experimental condition.

Imaging technologies can be used to evaluate specific trafficking and proliferation of CARs in tumor-bearing animal models. Such assays have been described, for example, in Barrett *et al*., Human Gene Therapy 22:1575-1586 (2011). Briefly, NOD/SCID/$\gamma c^{-/-}$ (NSG) mice are injected IV with Nalm-6 cells followed 7 days later with T cells 4 hour after electroporation with the CAR constructs. The T cells are stably transfected with a lentiviral construct to express firefly luciferase, and mice are imaged for bioluminescence. Alternatively, therapeutic efficacy and specificity of a single injection of CAR^+ T cells in Nalm-6 xenograft model can be measured as the following: NSG mice are injected with Nalm-6 transduced to stably express firefly luciferase, followed by a single tail-vein injection of T cells engineered to express a CAR of the present invention (e.g., transduced with nucleic acid encoding a CAR of the present invention) 7 days later. Animals are imaged at various time points post injection. For example, photon-density heat maps of firefly luciferasepositive leukemia in representative mice at day 5 (2 days before treatment) and day 8 (24 hr post CAR^+ PBLs) can be generated.

Other assays, including those described in the Example section herein as well as those that are known in the art can also be used to evaluate the CARs described herein.

Methods of treatment/Combination therapies

In another aspect, the present invention provides a method comprising administering the LSD1 inhibitors of the invention in combination with a population of immune effector cells, e.g., engineered to express a CAR molecule, e.g., as described herein, as a therapy. Typically, such administration will be in the form of cells (e.g., autologous or allogeneic host cells) expressing a CAR and separate administration of the LSD1 inhibitor. Alternatively, the LSD1 inhibitor may be administered in the same composition as the cell engineered to express a CAR molecule. Alternatively, the cell engineered to express a CAR molecule may also be engineered to express an LSD1 inhibitor. In one embodiment, the subject has a disorder described herein, e.g., the subject has cancer, e.g., the subject has a cancer which expresses a target antigen described herein. In one embodiment, the subject is a human.

Methods described herein that comprise administering an LSD1 inhibitor and a CAR-expressing cell described herein may be used in combination with other known agents and therapies.

Administered “in combination”, as used herein, means that two (or more) different treatments are delivered to the subject during the course of the subject's affliction with the disorder, e.g., the two or more treatments are delivered after the subject has been diagnosed with the disorder and before the

disorder has been cured or eliminated or treatment has ceased for other reasons. In some embodiments, the delivery of one treatment is still occurring when the delivery of the second begins, so that there is overlap in terms of administration. This is sometimes referred to herein as "simultaneous" or "concurrent delivery". In other embodiments, the delivery of one treatment ends before the delivery of the other treatment begins. In some embodiments of either case, the treatment is more effective because of combined administration. For example, the second treatment is more effective, e.g., an equivalent effect is seen with less of the second treatment, or the second treatment reduces symptoms to a greater extent, than would be seen if the second treatment were administered in the absence of the first treatment, or the analogous situation is seen with the first treatment. In some embodiments, delivery is such that the reduction in a symptom, or other parameter related to the disorder is greater than what would be observed with one treatment delivered in the absence of the other. The effect of the two treatments can be partially additive, wholly additive, or greater than additive. The delivery can be such that an effect of the first treatment delivered is still detectable when the second is delivered.

A CAR-expressing cell described herein, an LSD1 inhibitor, and the at least one additional therapeutic agent can be administered simultaneously, in the same or in separate compositions, or sequentially. The three agents can be administered in any order. For example, in sequential administration, the LSD1 inhibitor and the CAR-expressing cell described herein can be administered first, and the additional agent can be administered second, or the order of administration can be reversed.

The CAR therapy and/or other therapeutic agents, procedures or modalities can be administered during periods of active disorder, or during a period of remission or less active disease. The CAR therapy can be administered before another treatment, concurrently with the treatment, post-treatment, or during remission of the disorder.

In another aspect, the invention pertains to a method of treating a subject having a disease associated with expression of a cancer associated antigen as described herein comprising administering to the subject an effective amount of a cell comprising a CAR molecule, e.g., a CAR molecule described herein.

In one aspect, the invention provides methods for treating a disease associated with expression of a cancer associated antigen as described herein. The method comprises the administration of an LSD1 inhibitor and administrartion of a cell, e.g., a T cell, e.g., that expresses, or can express a CAR.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express an XCAR, wherein X represents a tumor marker (or cancer associated antigen; as used herein, the terms XCAR and CARX are used interchangeably, e.g., a CAR19 or CARCD19 is the same as a CD19 CAR) as described herein, and wherein said cancer cells express said X tumor marker (or cancer associated antigen).

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a CD19CAR, wherein the cancer cells express CD19. In one embodiment, the cancer to be treated is ALL (acute lymphoblastic leukemia), CLL (chronic lymphocytic leukemia), DLBCL (diffuse large B-cell lymphoma), MCL (Mantle cell lymphoma, or MM (multiple myeloma).

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express an EGFRvIIICAR, wherein the cancer cells express EGFRvIII. In one embodiment, the cancer to be treated is glioblastoma.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a mesothelin CAR, wherein the cancer cells express mesothelin. In one embodiment, the cancer to be treated is mesothelioma, pancreatic cancer, or ovarian cancer.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a CD123CAR, wherein the cancer cells express CD123. In one embodiment, the cancer to be treated is AML.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a CD22CAR, wherein the cancer cells express CD22. In one embodiment, the cancer to be treated is B cell malignancies.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are

engineered to express a CS-1CAR, wherein the cancer cells express CS-1. In one embodiment, the cancer to be treated is multiple myeloma.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a CLL-1CAR, wherein the cancer cells express CLL-1. In one embodiment, the cancer to be treated is AML.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a CD33CAR, wherein the cancer cells express CD33. In one embodiment, the cancer to be treated is AML.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a GD2CAR, wherein the cancer cells express GD2. In one embodiment, the cancer to be treated is neuroblastoma.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a BCMACAR, wherein the cancer cells express BCMA. In one embodiment, the cancer to be treated is multiple myeloma.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a TnCAR, wherein the cancer cells express Tn antigen. In one embodiment, the cancer to be treated is ovarian cancer.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a PSMACAR, wherein the cancer cells express PSMA. In one embodiment, the cancer to be treated is prostate cancer.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a ROR1CAR, wherein the cancer cells express ROR1. In one embodiment, the cancer to be treated is B cell malignancies.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a FLT3 CAR, wherein the cancer cells express FLT3. In one embodiment, the cancer to be treated is AML.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereo an LSD1 inhibitor and f immune effector cells (e.g., T cells, NK cells) that are engineered to express a TAG72CAR, wherein the cancer cells express TAG72. In one embodiment, the cancer to be treated is gastrointestinal cancer.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a CD38CAR, wherein the cancer cells express CD38. In one embodiment, the cancer to be treated is multiple myeloma.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a CD44v6CAR, wherein the cancer cells express CD44v6. In one embodiment, the cancer to be treated is cervical cancer, AML, or MM.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a CEACAR, wherein the cancer cells express CEA. In one embodiment, the cancer to be treated is pastrointestinal cancer, or pancreatic cancer.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a EPCAMCAR, wherein the cancer cells express EPCAM. In one embodiment, the cancer to be treated is gastrointestinal cancer.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a B7H3CAR, wherein the cancer cells express B7H3.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are

engineered to express a KITCAR, wherein the cancer cells express KIT. In one embodiment, the cancer to be treated is gastrointestinal cancer.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a IL-13Ra2CAR, wherein the cancer cells express IL-13Ra2. In one embodiment, the cancer to be treated is glioblastoma.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a CD30CAR, wherein the cancer cells express CD30. In one embodiment, the cancer to be treated is lymphomas, or leukemias.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a GD3CAR, wherein the cancer cells express GD3. In one embodiment, the cancer to be treated is melanoma.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a CD171CAR, wherein the cancer cells express CD171. In one embodiment, the cancer to be treated is neuroblastoma, ovarian cancer, melanoma, breast cancer, pancreatic cancer, colon cancers, or NSCLC (non-small cell lung cancer).

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a IL-11RaCAR, wherein the cancer cells express IL-11Ra. In one embodiment, the cancer to be treated is osteosarcoma.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a PSCACAR, wherein the cancer cells express PSCA. In one embodiment, the cancer to be treated is prostate cancer.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are

engineered to express a VEGFR2CAR, wherein the cancer cells express VEGFR2. In one embodiment, the cancer to be treated is a solid tumor.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a LewisYCAR, wherein the cancer cells express LewisY. In one embodiment, the cancer to be treated is ovarian cancer, or AML.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a CD24CAR, wherein the cancer cells express CD24. In one embodiment, the cancer to be treated is pancreatic cancer.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a PDGFR-betaCAR, wherein the cancer cells express PDGFR-beta. In one embodiment, the cancer to be treated is breast cancer, prostate cancer, GIST (gastrointestinal stromal tumor), CML, DFSP (dermatofibrosarcoma protuberans), or glioma.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a SSEA-4CAR, wherein the cancer cells express SSEA-4. In one embodiment, the cancer to be treated is glioblastoma, breast cancer, lung cancer, or stem cell cancer.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a CD20CAR, wherein the cancer cells express CD20. In one embodiment, the cancer to be treated is B cell malignancies.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a Folate receptor alphaCAR, wherein the cancer cells express folate receptor alpha. In one embodiment, the cancer to be treated is ovarian cancer, NSCLC, endometrial cancer, renal cancer, or other solid tumors.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are

engineered to express an ERBB2CAR, wherein the cancer cells express ERBB2 (Her2/neu). In one embodiment, the cancer to be treated is breast cancer, gastric cancer, colorectal cancer, lung cancer, or other solid tumors.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a MUC1CAR, wherein the cancer cells express MUC1. In one embodiment, the cancer to be treated is breast cancer, lung cancer, or other solid tumors.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a EGFRCAR, wherein the cancer cells express EGFR. In one embodiment, the cancer to be treated is glioblastoma, SCLC (small cell lung cancer), SCCHN (squamous cell carcinoma of the head and neck), NSCLC, or other solid tumors.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a NCAMCAR, wherein the cancer cells express NCAM. In one embodiment, the cancer to be treated is neuroblastoma, or other solid tumors.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a CAIXCAR, wherein the cancer cells express CAIX. In one embodiment, the cancer to be treated is renal cancer, CRC, cervical cancer, or other solid tumors.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express an EphA2CAR, wherein the cancer cells express EphA2. In one embodiment, the cancer to be treated is GBM.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a GD3CAR, wherein the cancer cells express GD3. In one embodiment, the cancer to be treated is melanoma.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a Fucosyl GM1CAR, wherein the cancer cells express Fucosyl GM

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a sLeCAR, wherein the cancer cells express sLe. In one embodiment, the cancer to be treated is NSCLC, or AML.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a GM3CAR, wherein the cancer cells express GM3.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a TGS5CAR, wherein the cancer cells express TGS5.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a HMWMAACAR, wherein the cancer cells express HMWMAA. In one embodiment, the cancer to be treated is melanoma, glioblastoma, or breast cancer.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express an o-acetyl-GD2CAR, wherein the cancer cells express o-acetyl-GD2. In one embodiment, the cancer to be treated is neuroblastoma, or melanoma.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a Folate receptor betaCAR, wherein the cancer cells express CD19. In one embodiment, the cancer to be treated is AML, or myeloma.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a TEM1/CD248CAR, wherein the cancer cells express TEM1/CD248. In one embodiment, the cancer to be treated is a solid tumor.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a TEM7RCAR, wherein the cancer cells express TEM7R. In one embodiment, the cancer to be treated is solid tumor.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a CLDN6CAR, wherein the cancer cells express CLDN6. In one embodiment, the cancer to be treated is ovarian cancer, lung cancer, or breast cancer.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a TSHRCAR, wherein the cancer cells express TSHR. In one embodiment, the cancer to be treated is thyroid cancer, or multiple myeloma.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a GPRC5DCAR, wherein the cancer cells express GPRC5D. In one embodiment, the cancer to be treated is multiple myeloma.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a CXORF61CAR, wherein the cancer cells express CXORF61.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a CD97CAR, wherein the cancer cells express CD97. In one embodiment, the cancer to be treated is B cell malignancies, gastric cancer, pancreatic cancer, esophageal cancer, glioblastoma, breast cancer, or colorectal cancer.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a CD179aCAR, wherein the cancer cells express CD179a. In one embodiment, the cancer to be treated is B cell malignancies.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are

engineered to express an ALK CAR, wherein the cancer cells express ALK. In one embodiment, the cancer to be treated is NSCLC, ALCL (anaplastic large cell lymphoma), IMT (inflammatory myofibroblastic tumor), or neuroblastoma.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a Plysialic acid CAR, wherein the cancer cells express Plysialic acid. In one embodiment, the cancer to be treated is small cell lung cancer.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a PLAC1CAR, wherein the cancer cells express PLAC1. In one embodiment, the cancer to be treated is HCC (hepatocellular carcinoma).

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a GloboHCAR, wherein the cancer cells express GloboH. In one embodiment, the cancer to be treated is ovarian cancer, gastric cancer, prostate cancer, lung cancer, breast cancer, or pancreatic cancer.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a NY-BR-1CAR, wherein the cancer cells express NY-BR-1. In one embodiment, the cancer to be treated is breast cancer.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a UPK2CAR, wherein the cancer cells express UPK2. In one embodiment, the cancer to be treated is bladder cancer.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a HAVCR1CAR, wherein the cancer cells express HAVCR1. In one embodiment, the cancer to be treated is renal cancer.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are

engineered to express a ADRB3CAR, wherein the cancer cells express ADRB3. In one embodiment, the cancer to be treated is Ewing sarcoma.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a PANX3CAR, wherein the cancer cells express PANX3. In one embodiment, the cancer to be treated is osteosarcoma.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a GPR20CAR, wherein the cancer cells express GPR20. In one embodiment, the cancer to be treated is GIST.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a LY6KCAR, wherein the cancer cells express LY6K. In one embodiment, the cancer to be treated is breast cancer, lung cancer, ovary caner, or cervix cancer.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a OR51E2CAR, wherein the cancer cells express OR51E2. In one embodiment, the cancer to be treated is prostate cancer.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a TARPCAR, wherein the cancer cells express TARP. In one embodiment, the cancer to be treated is prostate cancer.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a WT1CAR, wherein the cancer cells express WT1.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a NY-ESO-1CAR, wherein the cancer cells express NY-ESO-1.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a LAGE-1a CAR, wherein the cancer cells express LAGE-1a.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a MAGE-A1CAR, wherein the cancer cells express MAGE-A1. In one embodiment, the cancer to be treated is melanoma.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a MAGE A1CAR, wherein the cancer cells express MAGE A1.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a ETV6-AML CAR, wherein the cancer cells express ETV6-AML.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a sperm protein 17 CAR, wherein the cancer cells express sperm protein 17. In one embodiment, the cancer to be treated is ovarian cancer, HCC, or NSCLC.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a XAGE1CAR, wherein the cancer cells express XAGE1. In one embodiment, the cancer to be treated is Ewings, or rhabdo cancer.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a Tie 2 CAR, wherein the cancer cells express Tie 2. In one embodiment, the cancer to be treated is a solid tumor.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a MAD-CT-1CAR, wherein the cancer cells express MAD-CT-1. In one embodiment, the cancer to be treated is prostate cancer, or melanoma.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a MAD-CT-2CAR, wherein the cancer cells express MAD-CT-2. In one embodiment, the cancer to be treated is prostate cancer, melanoma.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a Fos-related antigen 1 CAR, wherein the cancer cells express Fos-related antigen 1. In one embodiment, the cancer to be treated is glioma, squamous cell cancer, or pancreatic cancer.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a p53CAR, wherein the cancer cells express p53.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a prostein CAR, wherein the cancer cells express prostein.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a survivin and telomerase CAR, wherein the cancer cells express survivin and telomerase.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a PCTA-1/Galectin 8 CAR, wherein the cancer cells express PCTA-1/Galectin 8.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a MelanA/MART1CAR, wherein the cancer cells express MelanA/MART1.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a Ras mutant CAR, wherein the cancer cells express Ras mutant.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a p53 mutant CAR, wherein the cancer cells express p53 mutant.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a hTERT CAR, wherein the cancer cells express hTERT.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a sarcoma translocation breakpoints CAR, wherein the cancer cells express sarcoma translocation breakpoints. In one embodiment, the cancer to be treated is sarcoma.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a ML-IAP CAR, wherein the cancer cells express ML-IAP. In one embodiment, the cancer to be treated is melanoma.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a ERGCAR, wherein the cancer cells express ERG (TMPRSS2 ETS fusion gene).

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a NA17CAR, wherein the cancer cells express NA17. In one embodiment, the cancer to be treated is melanoma.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a PAX3CAR, wherein the cancer cells express PAX3. In one embodiment, the cancer to be treated is alveolar rhabdomyosarcoma.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express an androgen receptor CAR, wherein the cancer cells express androgen receptor. In one embodiment, the cancer to be treated is metastatic prostate cancer.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a Cyclin B1CAR, wherein the cancer cells express Cyclin B1.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a MYCNCAR, wherein the cancer cells express MYCN.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a RhoC CAR, wherein the cancer cells express RhoC.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a TRP-2CAR, wherein the cancer cells express TRP-2. In one embodiment, the cancer to be treated is melanoma.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a CYP1B1CAR, wherein the cancer cells express CYP1B1. In one embodiment, the cancer to be treated is breast cancer, colon cancer, lung cancer, esophagus cancer, skin cancer, lymph node cancer, brain cancer, or testis cancer.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a BORIS CAR, wherein the cancer cells express BORIS. In one embodiment, the cancer to be treated is lung cancer.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a SART3CAR, wherein the cancer cells express SART3

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a PAX5CAR, wherein the cancer cells express PAX5.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a OY-TES1CAR, wherein the cancer cells express OY-TES1.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a LCK CAR, wherein the cancer cells express LCK.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a AKAP-4CAR, wherein the cancer cells express AKAP-4.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a SSX2CAR, wherein the cancer cells express SSX2.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a RAGE-1CAR, wherein the cancer cells express RAGE-1. In one embodiment, the cancer to be treated is RCC (renal cell cancer), or other solid tumors

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a human telomerase reverse transcriptaseCAR, wherein the cancer cells express human telomerase reverse transcriptase. In one embodiment, the cancer to be treated is solid tumors.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a RU1CAR, wherein the cancer cells express RU1.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a RU2CAR, wherein the cancer cells express RU2.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express an intestinal carboxyl esteraseCAR, wherein the cancer cells express intestinal carboxyl esterase. In one embodiment, the cancer to be treated is thyroid cancer, RCC, CRC (colorectal cancer), breast cancer, or other solid tumors.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a Prostase CAR, wherein the cancer cells express Prostase.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a PAPCAR, wherein the cancer cells express PAP.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express an IGF-I receptor CAR, wherein the cancer cells express IGF-I receptor.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a gp100 CAR, wherein the cancer cells express gp100.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a bcr-abl CAR, wherein the cancer cells express bcr-abl.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a tyrosinase CAR, wherein the cancer cells express tyrosinase.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof immune effector cells (e.g., T cells, NK cells) that are engineered to express a Fucosyl GM1CAR, wherein the cancer cells express Fucosyl GM1.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof an LSD1 inhibitor and immune effector cells (e.g., T cells, NK cells) that are engineered to express a mut hsp70-2CAR, wherein the cancer cells express mut hsp70-2. In one embodiment, the cancer to be treated is melanoma.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof immune effector cells (e.g., T cells, NK cells) that are engineered to express a CD79a CAR, wherein the cancer cells express CD79a.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof immune effector cells (e.g., T cells, NK cells) that are engineered to express a CD79b CAR, wherein the cancer cells express CD79b.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof immune effector cells (e.g., T cells, NK cells) that are engineered to express a CD72 CAR, wherein the cancer cells express CD72.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof immune effector cells (e.g., T cells, NK cells) that are engineered to express a LAIR1 CAR, wherein the cancer cells express LAIR1.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof immune effector cells (e.g., T cells, NK cells) that are engineered to express a FCAR CAR, wherein the cancer cells express FCAR.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof immune effector cells (e.g., T cells, NK cells) that are engineered to express a LILRA2 CAR, wherein the cancer cells express LILRA2.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof immune effector cells (e.g., T cells, NK cells) that are engineered to express a CD300LF CAR, wherein the cancer cells express CD300LF.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof immune effector cells (e.g., T cells, NK cells) that are engineered to express a CLEC12A CAR, wherein the cancer cells express CLEC12A.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof immune effector cells (e.g., T cells, NK cells) that are engineered to express a BST2 CAR, wherein the cancer cells express BST2.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof immune effector cells (e.g., T cells, NK cells) that are engineered to express an EMR2 CAR, wherein the cancer cells express EMR2.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof immune effector cells (e.g., T cells, NK cells) that are engineered to express a LY75 CAR, wherein the cancer cells express LY75.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof immune effector cells (e.g., T cells, NK cells) that are engineered to express a GPC3 CAR, wherein the cancer cells express GPC3.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof immune effector cells (e.g., T cells, NK cells) that are engineered to express a FCRL5 CAR, wherein the cancer cells express FCRL5.

In one aspect, the present invention provides methods of treating cancer by providing to the subject in need thereof immune effector cells (e.g., T cells, NK cells) that are engineered to express an IGLL1 CAR, wherein the cancer cells express IGLL1.

In one aspect, the present invention relates to treating a subject in vivo using an LSD1 inhibitor and a PD1 CAR such that growth of cancerous tumors is inhibited. A PD1 CAR may be used alone to inhibit the growth of cancerous tumors. Alternatively, PD1 CAR may be used in conjunction with other CARs, immunogenic agents, standard cancer treatments, or other antibodies. In one embodiment, the subject is treated with a PD1 CAR and an XCAR described herein. In an embodiment, a PD1 CAR is used in conjunction with another CAR, e.g., a CAR described herein, and a kinase inhibitor, e.g., a kinase inhibitor described herein.

In another aspect, a method of treating a subject, e.g., reducing or ameliorating, a hyperproliferative condition or disorder (e.g., a cancer), e.g., a solid tumor, a soft tissue tumor, a hematological cancer, or a metastatic lesion, in a subject, is provided. In embodiments the method comprises administration of an LSD1 inhibitor and a population of immune effector cells, e.g., T cells, engineered to express a CAR.

In yet another aspect, the invention features a method of treating a subject having a disease associated with expression of a tumor antigen (e.g., an antigen described herein), comprising administering to the subject an effective amount of a cell, e.g., an immune effector cell (*e.g.*, a population of immune effector cells) comprising a CAR molecule, wherein the CAR molecule comprises an antigen binding domain, a transmembrane domain, and an intracellular domain, said intracellular domain comprises a costimulatory domain and/or a primary signaling domain, wherein said antigen

binding domain binds to the tumor antigen associated with the disease, e.g. a tumor antigen as described herein.

In a related aspect, the invention features a method of treating a subject having a disease associated with expression of a tumor antigen. The method comprises administering to the subject an effective amount of a cell, e.g., an immune effector cell (*e.g.*, a population of immune effector cells) comprising a CAR molecule, in combination (in addition to the LSD1 inhibitor) with an agent that increases the efficacy of the immune cell, wherein:

the agent that increases the efficacy of the immune cell is chosen from one or more of:

(i) a protein phosphatase inhibitor;

(ii) a kinase inhibitor;

(iii) a cytokine;

(iv) an inhibitor of an immune inhibitory molecule; or

(v) an agent that decreases the level or activity of a T_{REG} cell.

In another aspect, the invention features a composition comprising an immune effector cell (*e.g.*, a population of immune effector cells) comprising a CAR molecule (e.g., a CAR molecule as described herein) for use in treating a subject having a disease associated with expression of a tumor antigen, e.g., a disorder as described herein.

In certain embodiments of any of the aforesaid methods or uses, the disease associated with a tumor antigen, e.g., a tumor antigen described herein, is selected from a proliferative disease such as a cancer or malignancy or a precancerous condition such as a myelodysplasia, a myelodysplastic syndrome or a preleukemia, or is a non-cancer related indication associated with expression of a tumor antigen described herein. In one embodiment, the disease is a cancer described herein, e.g., a cancer described herein as being associated with a target described herein. In one embodiment, the disease is a hematologic cancer. In one embodiment, the hematologic cancer is leukemia. In one embodiment, the cancer is selected from the group consisting of one or more acute leukemias including but not limited to B-cell acute lymphoid leukemia (“BALL”), T-cell acute lymphoid leukemia (“TALL”), acute lymphoid leukemia (ALL); one or more chronic leukemias including but not limited to chronic myelogenous leukemia (CML), chronic lymphocytic leukemia (CLL); additional hematologic cancers or hematologic conditions including, but not limited to B cell prolymphocytic leukemia, blastic plasmacytoid dendritic

cell neoplasm, Burkitt's lymphoma, diffuse large B cell lymphoma, follicular lymphoma, hairy cell leukemia, small cell- or a large cell-follicular lymphoma, malignant lymphoproliferative conditions, MALT lymphoma, mantle cell lymphoma, Marginal zone lymphoma, multiple myeloma, myelodysplasia and myelodysplastic syndrome, non-Hodgkin lymphoma, Hodgkin lymphoma, plasmablastic lymphoma, plasmacytoid dendritic cell neoplasm, Waldenstrom macroglobulinemia, and “preleukemia” which are a diverse collection of hematological conditions united by ineffective production (or dysplasia) of myeloid blood cells, and to disease associated with expression of a tumor antigen described herein include, but not limited to, atypical and/or non-classical cancers, malignancies, precancerous conditions or proliferative diseases expressing a tumor antigen as described herein; and any combination thereof. In another embodiment, the disease associated with a tumor antigen described herein is a solid tumor.

In certain embodiments, the methods or uses are carried out in combination with an agent that increases the efficacy of the immune effector cell, e.g., an agent as described herein.

In any of the aforesaid methods or uses, the disease associated with expression of the tumor antigen is selected from the group consisting of a proliferative disease, a precancerous condition, a cancer, and a non-cancer related indication associated with expression of the tumor antigen.

The cancer can be a hematologic cancer, e.g., a cancer chosen from one or more of chronic lymphocytic leukemia (CLL), acute leukemias, acute lymphoid leukemia (ALL), acute myeloid leukemia (AML), B-cell acute lymphoid leukemia (B-ALL), T-cell acute lymphoid leukemia (T-ALL), chronic myelogenous leukemia (CML), B cell prolymphocytic leukemia, blastic plasmacytoid dendritic cell neoplasm, Burkitt's lymphoma, diffuse large B cell lymphoma, follicular lymphoma, hairy cell leukemia, small cell- or a large cell-follicular lymphoma, malignant lymphoproliferative conditions, MALT lymphoma, mantle cell lymphoma, marginal zone lymphoma, multiple myeloma, myelodysplasia and myelodysplastic syndrome, non-Hodgkin’s lymphoma, Hodgkin’s lymphoma, plasmablastic lymphoma, plasmacytoid dendritic cell neoplasm, Waldenstrom macroglobulinemia, or pre-leukemia.

The cancer can also be chosen from colon cancer, rectal cancer, renal-cell carcinoma, liver cancer, non-small cell carcinoma of the lung, cancer of the small intestine, cancer of the esophagus, melanoma, bone cancer, pancreatic cancer, skin cancer, cancer of the head or neck, cutaneous or intraocular malignant melanoma, uterine cancer, ovarian cancer, rectal cancer, cancer of the anal region, stomach cancer, testicular cancer, uterine cancer, carcinoma of the fallopian tubes, carcinoma of the endometrium, carcinoma of the cervix, carcinoma of the vagina, carcinoma of the vulva, Hodgkin's Disease, non-Hodgkin's lymphoma, cancer of the endocrine system, cancer of the thyroid gland, cancer

of the parathyroid gland, cancer of the adrenal gland, sarcoma of soft tissue, cancer of the urethra, cancer of the penis, solid tumors of childhood, cancer of the bladder, cancer of the kidney or ureter, carcinoma of the renal pelvis, neoplasm of the central nervous system (CNS), primary CNS lymphoma, tumor angiogenesis, spinal axis tumor, brain stem glioma, pituitary adenoma, Kaposi's sarcoma, epidermoid cancer, squamous cell cancer, T-cell lymphoma, environmentally induced cancers, combinations of said cancers, and metastatic lesions of said cancers.

In certain embodiments of the methods or uses described herein, the CAR molecule is administered in combination with an agent that increases the efficacy of the immune effector cell (in addition to the LSD1 inhibitor), e.g., one or more of a protein phosphatase inhibitor, a kinase inhibitor, a cytokine, an inhibitor of an immune inhibitory molecule; or an agent that decreases the level or activity of a T_{REG} cell.

In certain embodiments of the methods or uses described herein, the protein phosphatase inhibitor is a SHP-1 inhibitor and/or an SHP-2 inhibitor.

In other embodiments of the methods or uses described herein, kinase inhibitor is chosen from one or more of a CDK4 inhibitor, a CDK4/6 inhibitor (e.g., palbociclib), a BTK inhibitor (e.g., ibrutinib or RN-486), an mTOR inhibitor (e.g., rapamycin or everolimus (RAD001)), an MNK inhibitor, or a dual P13K/mTOR inhibitor. In one embodiment, the BTK inhibitor does not reduce or inhibit the kinase activity of interleukin-2-inducible kinase (ITK).

In other embodiments of the methods or uses described herein, the agent that inhibits the immune inhibitory molecule comprises an antibody or antibody fragment, an inhibitory nucleic acid, a clustered regularly interspaced short palindromic repeats (CRISPR), a transcription-activator like effector nuclease (TALEN), or a zinc finger endonuclease (ZFN) that inhibits the expression of the inhibitory molecule.

In other embodiments of the methods or uses described herein, the agent that decreases the level or activity of the TREG cells is chosen from cyclophosphamide, anti-GITR antibody, CD25-depletion, or a combination thereof.

In certain embodiments of the methods or uses described herein, the immune inhibitory molecule is selected from the group consisting of PD1, PD-L1, CTLA-4, TIM-3, LAG-3, VISTA, BTLA, TIGIT, LAIR1, CD160, 2B4, TGF beta, CEACAM-1, CEACAM-3, and CEACAM-5.

In other embodiments, the agent that inhibits the inhibitory molecule comprises a first polypeptide comprising an inhibitory molecule or a fragment thereof and a second polypeptide that provides a positive signal to the cell, and wherein the first and second polypeptides are expressed on the CAR-containing immune cells, wherein (i) the first polypeptide comprises PD1, PD-L1, CTLA-4, TIM-3, LAG3, VISTA, BTLA, TIGIT, LAIR1, CD160, 2B4, TGF beta, CEACAM-1, CEACAM-3, and CEACAM-5 or a fragment thereof; and/or (ii) the second polypeptide comprises an intracellular signaling domain comprising a primary signaling domain and/or a costimulatory signaling domain. In one embodiment, the primary signaling domain comprises a functional domain of CD3 zeta; and/or the costimulatory signaling domain comprises a functional domain of a protein selected from 41BB, CD27 and CD28.

In other embodiments, cytokine is chosen from IL-7, IL-15 or IL-21, or combinations thereof.

In other embodiments, the immune effector cell comprising the CAR molecule and a second, e.g., any of the combination therapies disclosed herein (e.g., the agent that that increases the efficacy of the immune effector cell) are administered substantially simultaneously or sequentially.

In other embodiments, the immune cell comprising the CAR molecule is administered in combination with a molecule that targets GITR and/or modulates GITR function. In certain embodiments, the molecule targeting GITR and/or modulating GITR function is administered prior to the CAR-expressing cell or population of cells, or prior to apheresis.

In one embodiment, lymphocyte infusion, for example allogeneic lymphocyte infusion, is used in the treatment of the cancer, wherein the lymphocyte infusion comprises at least one CAR-expressing cell of the present invention. In one embodiment, autologous lymphocyte infusion is used in the treatment of the cancer, wherein the autologous lymphocyte infusion comprises at least one CAR-expressing cell described herein.

In one embodiment, the cell is a T cell and the T cell is diaglycerol kinase (DGK) deficient. In one embodiment, the cell is a T cell and the T cell is Ikaros deficient. In one embodiment, the cell is a T cell and the T cell is both DGK and Ikaros deficient.

In one embodiment, the method includes administering a cell expressing the CAR molecule, as described herein, in combination with an agent which enhances the activity of a CAR-expressing cell, wherein the agent is a cytokine, e.g., IL-7, IL-15, IL-18, IL-21, or a combination thereof. The cytokine can be delivered in combination with, e.g., simultaneously or shortly after, administration of the CAR-expressing cell. Alternatively, the cytokine can be delivered after a prolonged period of time after

administration of the CAR-expressing cell, e.g., after assessment of the subject's response to the CAR-expressing cell. In one embodiment the cytokine is administered to the subject simultaneously (e.g., administered on the same day) with or shortly after administration (e.g., administered 1 day, 2 days, 3 days, 4 days, 5 days, 6 days, or 7 days after administration) of the cell or population of cells of any of claims 61-80. In other embodiments, the cytokine is administered to the subject after a prolonged period of time (e.g., e.g., at least 2 weeks, 3 weeks, 4 weeks, 6 weeks, 8 weeks, 10 weeks, or more) after administration of the cell or population of cells of any of claims 61-80, or after assessment of the subject's response to the cell.

In other embodiments, the cells expressing a CAR molecule are administered in combination with an agent that ameliorates one or more side effects associated with administration of a cell expressing a CAR molecule. Side effects associated with the CAR-expressing cell can be chosen from cytokine release syndrome (CRS) or hemophagocytic lymphohistiocytosis (HLH).

In embodiments of any of the aforeseaid methods or uses, the cells expressing the CAR molecule are administered in combination with an agent that treats the disease associated with expression of the tumor antigen, e.g., any of the second or third therapies disclosed herein. Additional exemplary combinations include one or more of the following.

In another embodiment, the cell expressing the CAR molecule, e.g., as described herein, can be administered in combination with another agent, e.g., a kinase inhibitor and/or checkpoint inhibitor described herein. In an embodiment, a cell expressing the CAR molecule can further express another agent, e.g., an agent which enhances the activity of a CAR-expressing cell.

For example, in one embodiment, the agent that enhances the activity of a CAR-expressing cell can be an agent which inhibits an inhibitory molecule (e.g., an immune inhibitor molecule). Examples of inhibitory molecules include PD1, PD-L1, CTLA-4, TIM-3, CEACAM (e.g., CEACAM-1, CEACAM-3 and/or CEACAM-5), LAG-3, VISTA, BTLA, TIGIT, LAIR1, CD160, 2B4 and TGF beta.

In one embodiment, the agent that inhibits the inhibitory molecule is an inhibitory nucleic acid is a dsRNA, a siRNA, or a shRNA. In embodiments, the inhibitory nucleic acid is linked to the nucleic acid that encodes a component of the CAR molecule. For example, the inhibitory molecule can be expressed on the CAR-expressing cell.

In another embodiment, the agent which inhibits an inhibitory molecule, e.g., is a molecule described herein, e.g., an agent that comprises a first polypeptide, e.g., an inhibitory molecule,

associated with a second polypeptide that provides a positive signal to the cell, e.g., an intracellular signaling domain described herein. In one embodiment, the agent comprises a first polypeptide, e.g., of an inhibitory molecule such as PD-1, PD-L1, CTLA-4, TIM-3, CEACAM (e.g., CEACAM-1, CEACAM-3 and/or CEACAM-5), LAG-3, VISTA, BTLA, TIGIT, LAIR1, CD160, 2B4 or TGF beta, or a fragment of any of these (e.g., at least a portion of the extracellular domain of any of these), and a second polypeptide which is an intracellular signaling domain described herein (e.g., comprising a costimulatory domain (e.g., 41BB, CD27 or CD28, e.g., as described herein) and/or a primary signaling domain (e.g., a CD3 zeta signaling domain described herein). In one embodiment, the agent comprises a first polypeptide of PD1 or a fragment thereof (e.g., at least a portion of the extracellular domain of PD1), and a second polypeptide of an intracellular signaling domain described herein (e.g., a CD28 signaling domain described herein and/or a CD3 zeta signaling domain described herein).

In one embodiment, the CAR-expressing immune effector cell of the present invention, e.g., T cell or NK cell, is administered to a subject that has received a previous stem cell transplantation, e.g., autologous stem cell transplantation.

In one embodiment, the CAR–expressing immune effector cell of the present invention, e.g., T cell or NK cells, is administered to a subject that has received a previous dose of melphalan.

In one embodiment, the cell expressing a CAR molecule, e.g., a CAR molecule described herein, is administered in combination with an agent that increases the efficacy of a cell expressing a CAR molecule, e.g., an agent described herein.

In one embodiment, the cells expressing a CAR molecule, e.g., a CAR molecule described herein, are administered in combination with an LSD1 inhibitor. While not wishing to be bound by theory, it is believed that treatment with an LSD1 inhibitor is accompanied by a decrease in PD-1 positive T cells or an increase in PD-1 negative cells. PD-1 positive T cells, but not PD-1 negative T cells, can be exhausted by engagement with cells which express a PD-1 ligand, e.g., PD-L1 or PD-L2.

In an embodiment this approach can be used to optimize the performance of CAR cells described herein in the subject. While not wishing to be bound by theory, it is believed that, in an embodiment, the performance of endogenous, non-modified immune effector cells, e.g., T cells or NK cells, is improved. While not wishing to be bound by theory, it is believed that, in an embodiment, the performance of a target antigen CAR- expressing cell is improved. In other embodiments, cells, e.g., T cells or NK cells, which have, or will be engineered to express a CAR, can be treated ex vivo by contact with an amount of an LSD1 inhibitor that increases the number of PD1 negative immune effector cells,

e.g., T cells or increases the ratio of PD1 negative immune effector cells, e.g., T cells/ PD1 positive immune effector cells, e.g., T cells.

In an embodiment, administration of an LSD1 inhibitor is initiated prior to administration of an CAR expressing cell described herein, e.g., T cells or NK cells. In an embodiment, the CAR cells are administered after a sufficient time, or sufficient dosing, of an LSD1 inhibitor, such that the level of PD1 negative immune effector cells, e.g., T cells or NK cells, or the ratio of PD1 negative immune effector cells, e.g., T cells/ PD1 positive immune effector cells, e.g., T cells, has been, at least transiently, increased.

In an embodiment, the cell, e.g., T cell or NK cell, to be engineered to express a CAR, is harvested after a sufficient time, or after sufficient dosing of the LSD1 inhibitor, such that the level of PD1 negative immune effector cells, e.g., T cells, or the ratio of PD1 negative immune effector cells, e.g., T cells/ PD1 positive immune effector cells, e.g., T cells, in the subject or harvested from the subject has been, at least transiently, increased.

In one embodiment, the cell expressing a CAR molecule, e.g., a CAR molecule described herein, is administered in combination with an agent that ameliorates one or more side effect associated with administration of a cell expressing a CAR molecule, e.g., an agent described herein.

In one embodiment, the cell expressing a CAR molecule, e.g., a CAR molecule described herein, is administered in combination with an agent that treats the disease associated with a cancer associated antigen as described herein, e.g., an agent described herein.

In one embodiment, a cell expressing two or more CAR molecules, e.g., as described herein, is administered to a subject in need thereof to treat cancer. In one embodiment, a population of cells including a CAR expressing cell, e.g., as described herein, is administered to a subject in need thereof to treat cancer.

In one embodiment, the cell expressing a CAR molecule, e.g., a CAR molecule described herein, is administered at a dose and/or dosing schedule described herein.

In one embodiment, the CAR molecule is introduced into immune effector cells (e.g., T cells, NK cells), e.g., using *in vitro* transcription, and the subject (e.g., human) receives an initial administration of cells comprising a CAR molecule, and one or more subsequent administrations of cells comprising a CAR molecule, wherein the one or more subsequent administrations are administered less than 15 days, e.g., 14, 13, 12, 11, 10, 9, 8, 7, 6, 5, 4, 3, or 2 days after the previous administration. In one embodiment,

more than one administration of cells comprising a CAR molecule are administered to the subject (e.g., human) per week, e.g., 2, 3, or 4 administrations of cells comprising a CAR molecule are administered per week. In one embodiment, the subject (e.g., human subject) receives more than one administration of cells comprising a CAR molecule per week (e.g., 2, 3 or 4 administrations per week) (also referred to herein as a cycle), followed by a week of no administration of cells comprising a CAR molecule, and then one or more additional administration of cells comprising a CAR molecule (e.g., more than one administration of the cells comprising a CAR molecule per week) is administered to the subject. In another embodiment, the subject (e.g., human subject) receives more than one cycle of cells comprising a CAR molecule, and the time between each cycle is less than 10, 9, 8, 7, 6, 5, 4, or 3 days. In one embodiment, the cells comprising a CAR molecule are administered every other day for 3 administrations per week. In one embodiment, the cells comprising a CAR molecule are administered for at least two, three, four, five, six, seven, eight or more weeks.

In one embodiment, the cells expressing a CAR molecule, e.g., a CAR molecule described herein, are administered as a first line treatment for the disease, e.g., the cancer, e.g., the cancer described herein. In another embodiment, the cells expressing a CAR molecule, e.g., a CAR molecule described herein, are administered as a second, third, fourth line treatment for the disease, e.g., the cancer, e.g., the cancer described herein.

In one embodiment, a population of cells described herein is administered.

In another aspect, the invention pertains to the isolated nucleic acid molecule encoding a CAR of the invention, the isolated polypeptide molecule of a CAR of the invention, the vector comprising a CAR of the invention, and the cell comprising a CAR of the invention for use as a medicament.

In another aspect, the invention pertains to a the isolated nucleic acid molecule encoding a CAR of the invention, the isolated polypeptide molecule of a CAR of the invention, the vector comprising a CAR of the invention, and the cell comprising a CAR of the invention for use in the treatment of a disease expressing a cancer associated antigen as described herein.

In another aspect, the invention pertains to a cell expressing a CAR molecule described herein for use as a medicament in combination with a cytokine, e.g., IL-7, IL-15 and/or IL-21 as described herein. In another aspect, the invention pertains to a cytokine described herein for use as a medicament in combination with a cell expressing a CAR molecule described herein.

In another aspect, the invention pertains to a cell expressing a CAR molecule described herein for use as a medicament in combination with a kinase inhibitor and/or a checkpoint inhibitor as described herein. In another aspect, the invention pertains to a kinase inhibitor and/or a checkpoint inhibitor described herein for use as a medicament in combination with a cell expressing a CAR molecule described herein.

In another aspect, the invention pertains to a cell expressing a CAR molecule described herein for use in combination with a cytokine, e.g., IL-7, IL-15 and/or IL-21 as described herein, in the treatment of a disease expressing a tumor antigen targeted by the CAR. In another aspect, the invention pertains to a cytokine described herein for use in combination with a cell expressing a CAR molecule described herein, in the treatment of a disease expressing a tumor antigen targeted by the CAR.

In another aspect, the invention pertains to a cell expressing a CAR molecule described herein for use in combination with a kinase inhibitor and/or a checkpoint inhibitor as described herein, in the treatment of a disease expressing a tumor antigen targeted by the CAR. In another aspect, the invention pertains to a kinase inhibitor and/or a checkpoint inhibitor described herein for use in combination with a cell expressing a CAR molecule described herein, in the treatment of a disease expressing a tumor antigen targeted by the CAR.

In another aspect, the present invention provides a method comprising administering a CAR molecule, e.g., a CAR molecule described herein, or a cell comprising a nucleic acid encoding a CAR molecule, e.g., a CAR molecule described herein. In one embodiment, the subject has a disorder described herein, e.g., the subject has cancer, e.g., the subject has a cancer and has tumor-supporting cells which express a tumor-supporting antigen described herein. In one embodiment, the subject is a human.

In another aspect, the invention pertains to a method of treating a subject having a disease associated with expression of a tumor-supporting antigen as described herein comprising administering to the subject an effective amount of a cell comprising a CAR molecule, e.g., a CAR molecule described herein.

In yet another aspect, the invention features a method of treating a subject having a disease associated with expression of a tumor-supporting antigen, comprising administering to the subject an effective amount of a cell, e.g., an immune effector cell (*e.g.*, a population of immune effector cells) comprising a CAR molecule, wherein the CAR molecule comprises an antigen binding domain, a transmembrane domain, and an intracellular domain, said intracellular domain comprises a

costimulatory domain and/or a primary signaling domain, wherein said antigen binding domain binds to the tumor-supporting antigen associated with the disease, e.g. a tumor-supporting antigen as described herein.

In another aspect, the invention features a composition comprising an immune effector cell (e.g., a population of immune effector cells) comprising a CAR molecule (e.g., a CAR molecule as described herein) for use in treating a subject having a disease associated with expression of a tumor-supporting antigen, e.g., a disorder as described herein.

In any of the aforesaid methods or uses, the disease associated with expression of the tumor-supporting antigen is selected from the group consisting of a proliferative disease, a precancerous condition, a cancer, and a non-cancer related indication associated with expression of the tumor-supporting antigen. In an embodiment, the disease associated with a tumor-supporting antigen described herein is a solid tumor.

In one embodiment of the methods or uses described herein, the CAR molecule is administered in combination with another agent. In one embodiment, the agent can be a kinase inhibitor, e.g., a CDK4/6 inhibitor, a BTK inhibitor, an mTOR inhibitor, a MNK inhibitor, or a dual PI3K/mTOR inhibitor, and combinations thereof. In one embodiment, the kinase inhibitor is a CDK4 inhibitor, e.g., a CDK4 inhibitor described herein, e.g., a CD4/6 inhibitor, such as, e.g., 6-Acetyl-8-cyclopentyl-5-methyl-2-(5-piperazin-1-yl-pyridin-2-ylamino)-8*H*-pyrido[2,3-*d*]pyrimidin-7-one, hydrochloride (also referred to as palbociclib or PD0332991). In one embodiment, the kinase inhibitor is a BTK inhibitor, e.g., a BTK inhibitor described herein, such as, e.g., ibrutinib. In one embodiment, the kinase inhibitor is an mTOR inhibitor, e.g., an mTOR inhibitor described herein, such as, e.g., rapamycin, a rapamycin analog, OSI-027. The mTOR inhibitor can be, e.g., an mTORC1 inhibitor and/or an mTORC2 inhibitor, e.g., an mTORC1 inhibitor and/or mTORC2 inhibitor described herein. In one embodiment, the kinase inhibitor is a MNK inhibitor, e.g., a MNK inhibitor described herein, such as, e.g., 4-amino-5-(4-fluoroanilino)-pyrazolo [3,4-*d*] pyrimidine. The MNK inhibitor can be, e.g., a MNK1a, MNK1b, MNK2a and/or MNK2b inhibitor. The dual PI3K/mTOR inhibitor can be, e.g., PF-04695102.

In one embodiment of the methods or uses described herein, the kinase inhibitor is a CDK4 inhibitor selected from aloisine A; flavopiridol or HMR-1275, 2-(2-chlorophenyl)-5,7-dihydroxy-8-[(3S,4R)-3-hydroxy-1-methyl-4-piperidinyl]-4-chromenone; crizotinib (PF-02341066; 2-(2-Chlorophenyl)-5,7-dihydroxy-8-[(2*R*,3*S*)-2-(hydroxymethyl)-1-methyl-3-pyrrolidinyl]- 4*H*-1-benzopyran-4-one, hydrochloride (P276-00); 1-methyl-5-[[2-[5-(trifluoromethyl)-1*H*-imidazol-2-yl]-4-pyridinyl]oxy]-*N*-[4-

(trifluoromethyl)phenyl]-1*H*-benzimidazol-2-amine (RAF265); indisulam (E7070); roscovitine (CYC202); palbociclib (PD0332991); dinaciclib (SCH727965); N-[5-[[(5-*tert*-butyloxazol-2-yl)methyl]thio]thiazol-2-yl]piperidine-4-carboxamide (BMS 387032); 4-[[9-chloro-7-(2,6-difluorophenyl)-5*H*-pyrimido[5,4-*d*][2]benzazepin-2-yl]amino]-benzoic acid (MLN8054); 5-[3-(4,6-difluoro-1H-benzimidazol-2-yl)-1H-indazol-5-yl]-N-ethyl-4-methyl-3-pyridinemethanamine (AG-024322); 4-(2,6-dichlorobenzoylamino)-1H-pyrazole-3-carboxylic acid N-(piperidin-4-yl)amide (AT7519); 4-[2-methyl-1-(1-methylethyl)-1*H*-imidazol-5-yl]-*N*-[4-(methylsulfonyl)phenyl]- 2-pyrimidinamine (AZD5438); and XL281 (BMS908662).

In one embodiment of the methods or uses described herein, the kinase inhibitor is a CDK4 inhibitor, e.g., palbociclib (PD0332991), and the palbociclib is administered at a dose of about 50 mg, 60 mg, 70 mg, 75 mg, 80 mg, 90 mg, 100 mg, 105 mg, 110 mg, 115 mg, 120 mg, 125 mg, 130 mg, 135 mg (e.g., 75 mg, 100 mg or 125 mg) daily for a period of time, e.g., daily for 14-21 days of a 28 day cycle, or daily for 7-12 days of a 21 day cycle. In one embodiment, 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12 or more cycles of palbociclib are administered.

In one embodiment of the methods or uses described herein, the kinase inhibitor is a BTK inhibitor selected from ibrutinib (PCI-32765); GDC-0834; RN-486; CGI-560; CGI-1764; HM-71224; CC-292; ONO-4059; CNX-774; and LFM-A13. In one embodiment, the BTK inhibitor does not reduce or inhibit the kinase activity of interleukin-2-inducible kinase (ITK), and is selected from GDC-0834; RN-486; CGI-560; CGI-1764; HM-71224; CC-292; ONO-4059; CNX-774; and LFM-A13.

In one embodiment of the methods or uses described herein, the kinase inhibitor is a BTK inhibitor, e.g., ibrutinib (PCI-32765), and the ibrutinib is administered at a dose of about 250 mg, 300 mg, 350 mg, 400 mg, 420 mg, 440 mg, 460 mg, 480 mg, 500 mg, 520 mg, 540 mg, 560 mg, 580 mg, 600 mg (e.g., 250 mg, 420 mg or 560 mg) daily for a period of time, e.g., daily for 21 day cycle, or daily for 28 day cycle. In one embodiment, 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12 or more cycles of ibrutinib are administered.

In one embodiment of the methods or uses described herein, the kinase inhibitor is a BTK inhibitor that does not inhibit the kinase activity of ITK, e.g., RN-486, and RN-486 is administered at a dose of about 100 mg, 110 mg, 120 mg, 130 mg, 140 mg, 150 mg, 160 mg, 170 mg, 180 mg, 190 mg, 200 mg, 210 mg, 220 mg, 230 mg, 240 mg, 250 mg (e.g., 150 mg, 200 mg or 250 mg) daily for a period of time, e.g., daily a 28 day cycle. In one embodiment, 1, 2, 3, 4, 5, 6, 7, or more cycles of RN-486 are administered.

In one embodiment of the methods or uses described herein, the kinase inhibitor is an mTOR inhibitor selected from temsirolimus; ridaforolimus (1*R*,2*R*,4*S*)-4-[(2*R*)-2 [(1*R*,9*S*,12*S*,15*R*,16*E*,18*R*,19*R*,21*R*, 23*S*,24*E*,26*E*,28*Z*,30*S*,32*S*,35*R*)-1,18-dihydroxy-19,30-dimethoxy-15,17,21,23, 29,35-hexamethyl-2,3,10,14,20-pentaoxo-11,36-dioxa-4-azatricyclo[30.3.1.$0^{4,9}$] hexatriaconta-16,24,26,28-tetraen-12-yl]propyl]-2-methoxycyclohexyl dimethylphosphinate, also known as AP23573 and MK8669; everolimus (RAD001); rapamycin (AY22989); simapimod; (5-{2,4-bis[(3**S**)-3-methylmorpholin-4-yl]pyrido[2,3-**d**]pyrimidin-7-yl}-2-methoxyphenyl)methanol (AZD8055); 2-amino-8-[*trans*-4-(2-hydroxyethoxy)cyclohexyl]-6-(6-methoxy-3-pyridinyl)-4-methyl-pyrido[2,3-*d*]pyrimidin-7(8*H*)-one (PF04691502); and N^2-[1,4-dioxo-4-[[4-(4-oxo-8-phenyl-4*H*-1-benzopyran-2-yl)morpholinium-4-yl]methoxy]butyl]-L-arginylglycyl-L-α-aspartylL-serine-, inner salt (SF1126) (SEQ ID NO: 1937); and XL765.

In one embodiment of the methods or uses described herein, the kinase inhibitor is an mTOR inhibitor, e.g., rapamycin, and the rapamycin is administered at a dose of about 3 mg, 4 mg, 5 mg, 6 mg, 7 mg, 8 mg, 9 mg, 10 mg (e.g., 6 mg) daily for a period of time, e.g., daily for 21 day cycle, or daily for 28 day cycle. In one embodiment, 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12 or more cycles of rapamycin are administered. In one embodiment, the kinase inhibitor is an mTOR inhibitor, e.g., everolimus and the everolimus is administered at a dose of about 2 mg, 2.5 mg, 3 mg, 4 mg, 5 mg, 6 mg, 7 mg, 8 mg, 9 mg, 10 mg, 11 mg, 12 mg, 13 mg, 14 mg, 15 mg (e.g., 10 mg) daily for a period of time, e.g., daily for 28 day cycle. In one embodiment, 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12 or more cycles of everolimus are administered.

In one embodiment of the methods or uses described herein, the kinase inhibitor is an MNK inhibitor selected from CGP052088; 4-amino-3-(p-fluorophenylamino)-pyrazolo [3,4-*d*] pyrimidine (CGP57380); cercosporamide; ETC-1780445-2; and 4-amino-5-(4-fluoroanilino)-pyrazolo [3,4-*d*] pyrimidine.

In one embodiment of the methods or uses described herein, the kinase inhibitor is a dual phosphatidylinositol 3-kinase (PI3K) and mTOR inhibitor selected from 2-Amino-8-[trans-4-(2-hydroxyethoxy)cyclohexyl]-6-(6-methoxy-3-pyridinyl)-4-methyl-pyrido[2,3-d]pyrimidin-7(8H)-one (PF-04691502); N-[4-[[4-(Dimethylamino)-1-piperidinyl]carbonyl]phenyl]-N'-[4-(4,6-di-4-morpholinyl-1,3,5-triazin-2-yl)phenyl]urea (PF-05212384, PKI-587); 2-Methyl-2-{4-[3-methyl-2-oxo-8-(quinolin-3-yl)-2,3-dihydro-1*H*-imidazo[4,5-c]quinolin-1-yl]phenyl}propanenitrile (BEZ-235); apitolisib (GDC-0980, RG7422); 2,4-Difluoro-N-{2-(methyloxy)-5-[4-(4-pyridazinyl)-6-quinolinyl]-3-pyridinyl}benzenesulfonamide

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Claims

1. A method of treating a subject, comprising administering to said subject an LSD1 inhibitor and a population of immune effector cells engineered to express a CAR.

2. The method of claim 1, wherein:

a) the LSD1 inhibitor is administered before the subject is administered said population of immune effector cells;

b) the LSD1 inhibitor is administered concurrently with said population of immune effector cells;

c) the LSD1 inhibitor is administered after the subject is administered said population of immune effector cells; or

d) any combination of a), b) and/or c).

3. The method of claim 2, wherein the LSD1 inhibitor is administered before the subject is administered said population of immune effector cells, and wherein said administration of the LSD1 inhibitor is continued for a period of time after the administration of said population of immune effector cells.

4. The method of claim 3, wherein the administration of the LSD1 inhibitor is in an amount sufficient to increase an anti-tumor effect of said population of immune effector relative to an equivalent population of said immune effector cells administered in the absence of said LSD1 inhibitor.

5. A method of increasing the therapeutic efficacy of a population of immune effector cells engineered to express a CAR, e.g., a CART19 (e.g., CTL019), comprising a step of decreasing the activity or expression of LSD1 in said cells, transiently or permanently.

6. The method of claim 5, wherein the step of decreasing the activity or expression of LSD1 in said cells comprises contacting the cells with an LSD1 inhibitor.

7. The method of any of claims 1-4 or 6, wherein the administration or the contacting of the LSD1 inhibitor results in:

1) an increase in the proportion of naive T cells, e.g., T_{SCM} cells;

2) an increase in the number of naive T cells, e.g., T_{SCM} cells;

3) a decrease in the number of T_{EM} cells;

4) a decrease in the proportion of T_{EM} cells;

5) an increase in the proportion of CD45RA+CD62L+ T cells;

6) an increase in the number of CD45RA+CD62L+ T cells;

7) an increase in the proportion of CD45RA+CCR7+ T cells;

8) an increase in the number of CD45RA+CCR7+ T cells;

9) a decrease in the proportion of PD-1 positive immune effector cells;

10) an increase in the ratio of PD-1 negative immune effector cells / PD-1 positive immune effector cells;

11) a decrease in the proportion of PD-1+/Lag3+/Tim3+ immune effector cells;

12) an increase in the ratio of PD-1-/Lag3-/Tim3- immune effector cells to PD-1+/Lag3+/Tim3+ immune effector cells;

13) an increase in the proliferation of the immune effector cells;

14) an increase in the production of cytokines (e.g., IFNg and/or IL-2) from said population of immune effector cells; or

15) a combination of two or more of the above;

optionally as compared to cells not contacted with the LSD1 inhibitor.

8. A method of treating a subject, comprising:

a) administering an LSD1 inhibitor to a subject;

b) collecting a population of immune effector cells from the subject of step a), after said administration of the LSD1 inhibitor;

c) providing said population of immune effector cells *ex vivo*;

d) contacting said *ex vivo* population of immune effector cells with the LSD1 inhibitor, wherein the contacting with the LSD1 inhibitor causes one or more of the following to occur:

1) an increase in the proportion of naive T cells, e.g., T_{SCM} cells;

2) an increase in the number of naive T cells, e.g., T_{SCM} cells;

3) a decrease in the number of T_{EM} cells;

4) a decrease in the proportion of T_{EM} cells;

5) an increase in the proportion of CD45RA+CD62L+ T cells;

6) an increase in the number of CD45RA+CD62L+ T cells;

7) an increase in the proportion of CD45RA+CCR7+ T cells;

8) an increase in the number of CD45RA+CCR7+ T cells;

9) a decrease in the proportion of PD-1 positive immune effector cells;

10) an increase in the ratio of PD-1 negative immune effector cells / PD-1 positive immune effector cells;

11) a decrease in the proportion of PD-1+/Lag3+/Tim3+ immune effector cells;

12) an increase in the ratio of PD-1-/Lag3-/Tim3- immune effector cells to PD-1+/Lag3+/Tim3+ immune effector cells;

13) an increase in the proliferation of the immune effector cells;

14 an increase in the production of cytokines (e.g., IFNg and/or IL-2) from said population of immune effector cells; or

15) a combination of two or more of the above;

as compared to a non-contacted *ex vivo* population of immune effector cells; and

e) administering the population of immune effector cells to a subject.

9. The method of claim 8, wherein the step of e) further comprises administering the LSD1 inhibitor to the subject of step e).

10. The method of claim 8 or 9, further comprising the step of inserting nucleic acid that encodes a CAR into cells of the *ex vivo* population of immune effector cells.

11. A method of making a population of immune effector cells comprising:

a) contacting a population of immune effector cells with an LSD1 inhibitor;

thereby making a population of immune effector cells,

wherein the contacting with the LSD1 inhibitor causes one or more of the following to occur:

1) an increase in the proportion of naive T cells, e.g., T_{SCM} cells;

2) an increase in the number of naive T cells, e.g., T_{SCM} cells;

3) a decrease in the number of T_{EM} cells;

4) a decrease in the proportion of T_{EM} cells;

5) an increase in the proportion of CD45RA+CD62L+ T cells;

6) an increase in the number of CD45RA+CD62L+ T cells;

7) an increase in the proportion of CD45RA+CCR7+ T cells;

8) an increase in the number of CD45RA+CCR7+ T cells;

9) a decrease in the proportion of PD-1 positive immune effector cells;

10) an increase in the ratio of PD-1 negative immune effector cells / PD-1 positive immune effector cells;

11) a decrease in the proportion of PD-1+/Lag3+/Tim3+ immune effector cells;

12) an increase in the ratio of PD-1-/Lag3-/Tim3- immune effector cells to PD-1+/Lag3+/Tim3+ immune effector cells;

13) an increase in the proliferation of the immune effector cells;

14 an increase in the production of cytokines (e.g., IFNg and/or IL-2) from said population of immune effector cells; or

15) a combination of two or more of the above;

as compared to a non-contacted population of immune effector cells.

12. The method of claim 11, further comprising the step of b) inserting nucleic acid that encodes the CAR into cells of the population of immune effector cells.

13. The method of claim 12 wherein said contacting of step a) occurs

1) prior to;

2) concurrently with;

3) after; or

4) any combination of two or more of 1) to 3) above;

said inserting of step b).

14. The method of any of claims 11-13, wherein the contacting of step a), and optionally the inserting of step b), is *ex vivo*.

15. A population of immune effector cells, made by the method of any of claims 11-14.

16. A population of immune effector cells engineered to express a chimeric antigen receptor (CAR), wherein the CAR comprises an antigen-binding domain, a transmembrane domain, and an intracellular signaling domain, and wherein expression and/or function of LSD1 in said cell has been reduced or eliminated.

17. The population of immune effector cells of claim 16, comprising an LSD1 inhibitor.

18. The population of immune effector cells of claim 16, wherein the population of immune effector cells has been contacted with an LSD1 inhibitor.

19. The method or population of immune effector cells of any of the preceding claims, wherein the population of immune effector cells comprise, e.g., consist of, T cells or NK cells.

20. The method or population of immune effector cells of claim 19, wherein the population of immune effector cells comprise, e.g., consist of, T cells.

21. The method or population of immune effector cells of claim 20, wherein the T cells are CD8+ T cells, CD4+ T cells, or a combination thereof.

22. The method or population of immune effector cells of any of the preceding claims, wherein the immune effector cells are human cells.

23. The method of any of claims 1-7, 10, 12-14 or 18-22, or population of immune effector cells of any of claims 15-22, wherein the CAR comprises an antigen binding domain (which is optionally an antibody or antibody fragment, TCR or TCR fragment), a transmembrane domain, and an intracellular signaling domain (which is optionally an intracellular signaling domain comprising a costimulatory domain and/or a primary signaling domain).

24. The method or population of immune effector cells of claim 23, wherein the antigen-binding domain binds to a tumor antigen selected from a group consisting of: TSHR, CD19, CD123, CD22, CD30, CD171, CS-1, CLL-1, CD33, EGFRvIII , GD2, GD3, BCMA, Tn Ag, PSMA, ROR1, FLT3, FAP, TAG72, CD38, CD44v6, CEA, EPCAM, B7H3, KIT, IL-13Ra2, Mesothelin, IL-11Ra, PSCA, PRSS21, VEGFR2, LewisY, CD24, PDGFR-beta, SSEA-4, CD20, Folate receptor alpha, ERBB2 (Her2/neu), MUC1, EGFR, NCAM, Prostase, PAP, ELF2M, Ephrin B2, IGF-I receptor, CAIX, LMP2, gp100, bcr-abl, tyrosinase, EphA2, Fucosyl GM1, sLe, GM3, TGS5, HMWMAA, o-acetyl-GD2, Folate receptor beta, TEM1/CD248, TEM7R, CLDN6, GPRC5D, CXORF61, CD97, CD179a, ALK, Polysialic acid, PLAC1, GloboH, NY-BR-1, UPK2, HAVCR1, ADRB3, PANX3, GPR20, LY6K, OR51E2, TARP, WT1, NY-ESO-1, LAGE-1a, MAGE-A1, legumain, HPV E6,E7, MAGE A1, ETV6-AML, sperm protein 17, XAGE1, Tie 2, MAD-CT-1, MAD-CT-2, Fos-related antigen 1, p53, p53 mutant, prostein, survivin and telomerase, PCTA-1/Galectin 8, MelanA/MART1, Ras mutant, hTERT, sarcoma translocation breakpoints, ML-IAP, ERG (TMPRSS2 ETS fusion gene), NA17, PAX3, Androgen receptor, Cyclin B1, MYCN, RhoC, TRP-2, CYP1B1, BORIS, SART3, PAX5, OY-TES1, LCK, AKAP-4, SSX2, RAGE-1,

human telomerase reverse transcriptase, RU1, RU2, intestinal carboxyl esterase, mut hsp70-2, CD79a, CD79b, CD72, LAIR1, FCAR, LILRA2, CD300LF, CLEC12A, BST2, EMR2, LY75, GPC3, FCRL5, and IGLL1.

25. The method or population of immune effector cells of claim 24, wherein the tumor antigen is CD19 or BCMA.

26. The method or population of immune effector cells of claim 25, wherein:

the antigen-binding domain is an antibody or antibody fragment comprising:

(i) the amino acid sequence of a CD19 binding domain according to Tables 6-9, e.g., the amino acid sequence of CTL019 scFv domain according to Table 9 or an amino acid sequence according to SEQ ID NO: 957, or an amino acid sequence at least 95% identical thereto;

(ii) the amino acid sequence of a humanized CD19 binding domain according to Tables 6-9, e.g., the amino acid sequence of CAR2 scFv domain according to Table 9 or an amino acid sequence according to SEQ ID NO: 898, or an amino acid sequence at least 95% identical thereto; or

(iii) the amino acid sequence of a BCMA binding domain according to Tables 11A-11B, e.g., the amino acid sequence of 139109 scFv domain according to Table 11A or an amino acid sequence according to SEQ ID NO: 967, or an amino acid sequence at least 95% identical thereto; or

wherein the CAR comprises:

(i) the amino acid sequence of a CD19 CAR according to Tables 6-9, e.g., the amino acid sequence of CTL019 according to Table 9 or an amino acid sequence according to SEQ ID NO: 956 or an amino acid sequence at least 95% identical thereto;

(ii) the amino acid sequence of a humanized CD19 CAR according to Tables 6-9, e.g., the amino acid sequence of CAR2 according to Table 9 or an amino acid sequence according to SEQ ID NO: 902, or an amino acid sequence at least 95% identical thereto; or

(iii) the amino acid sequence of a BCMA CAR according to Tables 11A-11B, e.g., the amino acid sequence of 139109 CAR according to Table 11A or an amino acid sequence according to SEQ ID NO: 971, or an amino acid sequence at least 95% identical thereto.

27. The method or population of immune effector cells of any of claims 23-26, wherein the transmembrane domain comprises:

(i) an amino acid sequence having at least one, two or three modifications but not more than 20, 10 or 5 modifications of an amino acid sequence of SEQ ID NO: 12, or a sequence with 95-99% identity to an amino acid sequence of SEQ ID NO: 12; or

(ii) the sequence of SEQ ID NO: 12.

28. The method or population of immune effector cells of any of claim 23-27, wherein the antigen binding domain is connected to the transmembrane domain by a hinge region, wherein said hinge region comprises SEQ ID NO: 2 or SEQ ID NO: 6, or a sequence with 95-99% identity thereof.

29. The method or population of immune effector cells of any of claims 23-28, wherein the intracellular signaling domain comprises a primary signaling domain and/or a costimulatory signaling domain, wherein the primary signaling domain comprises a functional signaling domain of a protein chosen from CD3 zeta, CD3 gamma, CD3 delta, CD3 epsilon, common FcR gamma (FCER1G), FcR beta (Fc Epsilon R1b), CD79a, CD79b, Fcgamma RIIa, DAP10, or DAP12.

30. The method or population of immune effector cells of any of claims 23-29, wherein the primary signaling domain comprises:

(i) an amino acid sequence having at least one, two or three modifications but not more than 20, 10 or 5 modifications of an amino acid sequence of SEQ ID NO: 18 or SEQ ID NO: 20, or a sequence with 95-99% identity to an amino acid sequence of SEQ ID NO: 18 or SEQ ID NO: 20; or

(ii) the amino acid sequence of SEQ ID NO:18 or SEQ ID NO: 20.

31. The method or population of immune effector cells of any of claims 23-30, wherein the intracellular signaling domain comprises a costimulatory signaling domain, or a primary signaling domain and a costimulatory signaling domain, wherein the costimulatory signaling domain comprises a functional signaling domain of a protein selected from the group consisting of CD27, CD28, 4-1BB (CD137), OX40, CD30, CD40, PD-1, ICOS, lymphocyte function-associated antigen-1 (LFA-1), CD2, CD7, LIGHT, NKG2C, B7-H3, a ligand that specifically binds with CD83, CDS, ICAM-1, GITR, BAFFR, HVEM (LIGHTR), SLAMF7, NKp80 (KLRF1), CD160, CD19, CD4, CD8alpha, CD8beta, IL2R beta, IL2R gamma, IL7R alpha, ITGA4, VLA1, CD49a, ITGA4, IA4, CD49D, ITGA6, VLA-6, CD49f, ITGAD, CD11d, ITGAE, CD103, ITGAL, CD11a, LFA-1, ITGAM, CD11b, ITGAX, CD11c, ITGB1, CD29, ITGB2, CD18, LFA-1, ITGB7, TNFR2, TRANCE/RANKL, DNAM1 (CD226), SLAMF4 (CD244, 2B4), CD84, CD96 (Tactile), CEACAM1, CRTAM, Ly9 (CD229), CD160 (BY55), PSGL1, CD100 (SEMA4D), CD69, SLAMF6 (NTB-A, Ly108), SLAM (SLAMF1, CD150,

IPO-3), BLAME (SLAMF8), SELPLG (CD162), LTBR, LAT, GADS, SLP-76, PAG/Cbp, NKp44, NKp30, NKp46, and NKG2D.

32. The method or population of immune effector cells of any of claims 23-31, wherein the costimulatory signaling domain comprises an amino acid sequence having at least one, two or three modifications but not more than 20, 10 or 5 modifications of an amino acid sequence of SEQ ID NO:14 or SEQ ID NO: 16, or a sequence with 95-99% identity to an amino acid sequence of SEQ ID NO:14 or SEQ ID NO: 16.

33. The method or population of immune effector cells of any of claims 23-32, wherein the costimulatory signaling domain comprises a sequence of SEQ ID NO: 14 or SEQ ID NO: 16.

34. The method or population of immune effector cells of any of claims 23-33, wherein the intracellular domain comprises the sequence of SEQ ID NO: 14 or SEQ ID NO: 16, and the sequence of SEQ ID NO: 18 or SEQ ID NO: 20, wherein the sequences comprising the intracellular signaling domain are expressed in the same frame and as a single polypeptide chain.

35. The method or population of immune effector cells of any of claims 23-34, wherein the CAR comprises a leader sequence comprising, e.g., consisting of, SEQ ID NO: 2.

36. The method or population of immune effector cells of any of the preceding claims, wherein the LSD1 inhibitor is: (1) a gene editing system targeted to one or more sites of the LSD1 gene, or its corresponding regulatory elements; (2) a nucleic acid (e.g., an siRNA or shRNA, or antisense oligonucleotide) comprising sequence complementary to a target sequence of the LSD1 gene; (3) a protein (e.g., a dominant negative LSD1, e.g., catalytically inactive LSD1, or a dominant negative binding partner of LSD1); (4) a small molecule; (5) a nucleic acid encoding any of (1)-(3); or (6) any combination of (1) –(5).

37. The method or population of immune effector cells of claim 36, wherein the LSD1 inhibitor is an shRNA or siRNA comprising sequence complementary to a target sequence of the LSD1 gene listed in Table 1, e.g., selected from SEQ ID NO: [43] to SEQ ID NO: [82].

38. The method or population of immune effector cells of claim 36, wherein the LSD1 inhibitor is an shRNA encoded by nucleic acid comprising any sequence encoding an anti-LSD1 shRNA of Table 1, e.g., encoded by nucleic acid comprising a sequence selected from SEQ ID NO: [83] to SEQ ID NO: [122].

39. The method or population of immune effector cells of claim 36, wherein the LSD1 inhibitor is nucleic acid comprising any sequence encoding an anti-LSD1 shRNA of Table 1, e.g., a sequence selected from SEQ ID NO: [83] to SEQ ID NO: [122].

40. The method or population of immune effector cells of claim 39, wherein said nucleic acid is disposed on a vector.

41. The method or population of immune effector cells of claim 40, wherein the vector further comprises a U6 or H1 promoter operably linked to said nucleic acid.

42. The method or population of immune effector cells of any of claims 40-41, wherein the vector is a retroviral vector, a lentiviral vector, an adenoviral vector, an adeno-associated viral (AAV) vector, a herpes simplex virus (HSV) vector, a plasmid, a minicircle, a nanoplasmid, or an RNA vector.

43. The method or population of immune effector cells of any of claims 40-42, wherein the vector further comprises sequence encoding a CAR.

44. The method or population of immune effector cells of claim 36, wherein the LSD1 inhibitor is a genome editing system specific for a sequence of the LSD1 gene (KDM1A) or its regulatory elements selected from a CRISPR genome editing system, a zinc finger nuclease genome editing system, a TALEN genome editing system and a meganuclease genome editing system.

45. The method or population of immune effector cells of claim 44, wherein the LSD1 inhibitor is a CRISPR genome editing system comprising a gRNA molecule comprising a targeting domain complementary to a sequence of the LSD1 gene (KDM1A) or its regulatory elements, e.g., comprising any one of SEQ ID NO: [132] to [862].

46. The method or population of immune effector cells of claim 36, wherein the LSD1 inhibitor is a small molecule.

47. The method or population of immune effector cells of claim 46, wherein the small molecule is a reversible or irreversible LSD1 inhibitor.

48. The method or population of immune effector cells of claim 46 or 47, wherein the LSD1 inhibitor is:

a) GSK2699537;

b) rel- 2-[[(1R,2S)-2-[4-[(4-chlorophenyl)methoxy]phenyl]cyclopropyl]amino]-1-(4-methyl-1-piperazinyl)-ethanone;

c) (R)-4-(5-(pyrrolidin-3-ylmethoxy)-2-(p-tolyl)pyridin-3-yl)benzonitrile;

d) (1S,2R)-N-((2-methoxypyridin-3-yl)methyl)-2-phenylcyclopropan-1-amine;

e) N,N-dimethyl-1-((4-(4-(4-(piperidin-4-yl)phenyl)-1H-indazol-1-yl)phenyl)sulfonyl)piperidin-4-amine;

f) 5-(6-chloro-4'-(methylsulfonyl)-[1,1'-biphenyl]-3-yl)-2-(piperazin-1-yl)-1H-pyrrole-3-carbonitrile;

g) rel- N-[(1R,2S)-2-Phenylcyclopropyl]-4-Piperidinamine;

h) 2-(1R,2S)-2-(4-(Benzyloxy)phenyl)cyclopropylamino)-1-(4-methylpiperazin-1-yl)ethanone;

i) Trans-3-(3-amino-2-methylphenyl)-1-(4-hydroxycyclohexyl)-6-methyl-1H-indole-5-carbonitrile; or

j) a pharmaceutically acceptable salt of any of the foregoing.

49. The method or population of immune effector cells of any of claims 46-48, wherein the LSD1 inhibitor is conjugated to an antibody or antigen-binding fragment thereof.

50. The method or population of immune effector cells of claim 49, wherein the antibody or antigen-binding fragment thereof recognizes an antigen on the surface of a T cell.

51. The method or population of immune effector cells of clam 50, wherein the antigen on the surface of a T cell is CD3.

52. The method or population of immune effector cells of claim 36, wherein the LSD1 inhibitor is a protein, e.g., is a dominant negative binding partner of LSD1 (e.g., a histone deacetylase (HDAC) that interacts with LSD1 or other member of the Co-REST or AR co-activator complex), or nucleic acid encoding said dominant negative binding partner of LSD1.

53. The method or population of immune effector cells of claim 36, wherein the inhibitor of LSD1 is a protein, e.g., is a dominant negative (e.g., catalytically inactive) LSD1, or nucleic acid encoding said dominant negative LSD1.

54. A method of treating a subject in need thereof, comprising administering to said subject an effective amount of the population of immune effector cells of any of claims 15-18.

55. The method of claim 54, wherein the method further comprises administering to said subject an LSD1 inhibitor.

56. The method of claim 55, wherein the subject receives a pre-treatment of the LSD1 inhibitor, prior to the administration of the population of immune effector cells.

57. The method of any of claims 55-56, wherein the subject receives concurrent treatment with an LSD1 inhibitor and the population of immune effector cells.

58. The method of any of claims 55-57, wherein the subject receives treatment with an LSD1 inhibitor after administration of the population of immune effector cells.

59. The method of any of claims 1-15 or 19-58, wherein the subject has a disease associated with expression of a tumor antigen, e.g., a proliferative disease, a precancerous condition, a cancer, and a non-cancer related indication associated with expression of the tumor antigen.

60. The method of claim 59, wherein the cancer is a hematologic cancer chosen from one or more of chronic lymphocytic leukemia (CLL), acute leukemias, acute lymphoid leukemia (ALL), acute myeloid leukemia (AML), B-cell acute lymphoid leukemia (B-ALL), T-cell acute lymphoid leukemia (T-ALL), chronic myelogenous leukemia (CML), B cell prolymphocytic leukemia, blastic plasmacytoid dendritic cell neoplasm, Burkitt's lymphoma, diffuse large B cell lymphoma, follicular lymphoma, hairy cell leukemia, small cell- or a large cell-follicular lymphoma, malignant lymphoproliferative conditions, MALT lymphoma, mantle cell lymphoma, marginal zone lymphoma, multiple myeloma, myelodysplasia and myelodysplastic syndrome, non-Hodgkin’s lymphoma, Hodgkin’s lymphoma, plasmablastic lymphoma, plasmacytoid dendritic cell neoplasm, Waldenstrom macroglobulinemia, or pre-leukemia.

61. The use or method of claim 59, wherein the cancer is selected from the group consisting of colon cancer, rectal cancer, renal-cell carcinoma, liver cancer, non-small cell carcinoma of the lung, cancer of the small intestine, cancer of the esophagus, melanoma, bone cancer, pancreatic cancer, skin cancer, cancer of the head or neck, cutaneous or intraocular malignant melanoma, uterine cancer, ovarian cancer, rectal cancer, cancer of the anal region, stomach cancer, testicular cancer, uterine cancer, carcinoma of the fallopian tubes, carcinoma of the endometrium, carcinoma of the cervix, carcinoma of the vagina, carcinoma of the vulva, Hodgkin's Disease, non-Hodgkin's lymphoma, cancer of

the endocrine system, cancer of the thyroid gland, cancer of the parathyroid gland, cancer of the adrenal gland, sarcoma of soft tissue, cancer of the urethra, cancer of the penis, solid tumors of childhood, cancer of the bladder, cancer of the kidney or ureter, carcinoma of the renal pelvis, neoplasm of the central nervous system (CNS), primary CNS lymphoma, tumor angiogenesis, spinal axis tumor, brain stem glioma, pituitary adenoma, Kaposi's sarcoma, epidermoid cancer, squamous cell cancer, T-cell lymphoma, environmentally induced cancers, combinations of said cancers, and metastatic lesions of said cancers.

62. The method or population of immune effector cells of any of the preceding claims, wherein the subject is a human.

63. The method or population of immune effector cells of claim 62, wherein the human has a disease associated with expression of a tumor antigen, e.g., a cancer.

64. A compound selected from N,N-dimethyl-1-((4-(4-(4-(piperidin-4-yl)phenyl)-1H-indazol-1-yl)phenyl)sulfonyl)piperidin-4-amine and 5-(6-chloro-4'-(methylsulfonyl)biphenyl-3-yl)-2-(piperazin-1-yl)-1H-pyrrole-3-carbonitrile; or a pharmaceutically acceptable salt thereof.

65. A pharmaceutically acceptable salt of the compound of claim 64.

66. The compound of any of claims 64-65, for use in therapy.

67. The compound of any of claims 64-65, for use in the method of any of claims 1-14, 19-36, 46-51, or 54-63.

68. A composition for use in *ex vivo* manufacturing a population of immune effector cells, comprising an LSD1 inhibitor.

69. The composition of claim 68, wherein the LSD1 inhibitor is a small molecule.

70. The composition of claim 69, wherein the concentration of the small molecule ranges from about 0.001 nM to about 10 mM, e.g., from about 0.001 nM to about 100 nM.

71. The composition of claim 70, wherein the concentration of the small molecule ranges from about 0.1 uM to about 10 uM.

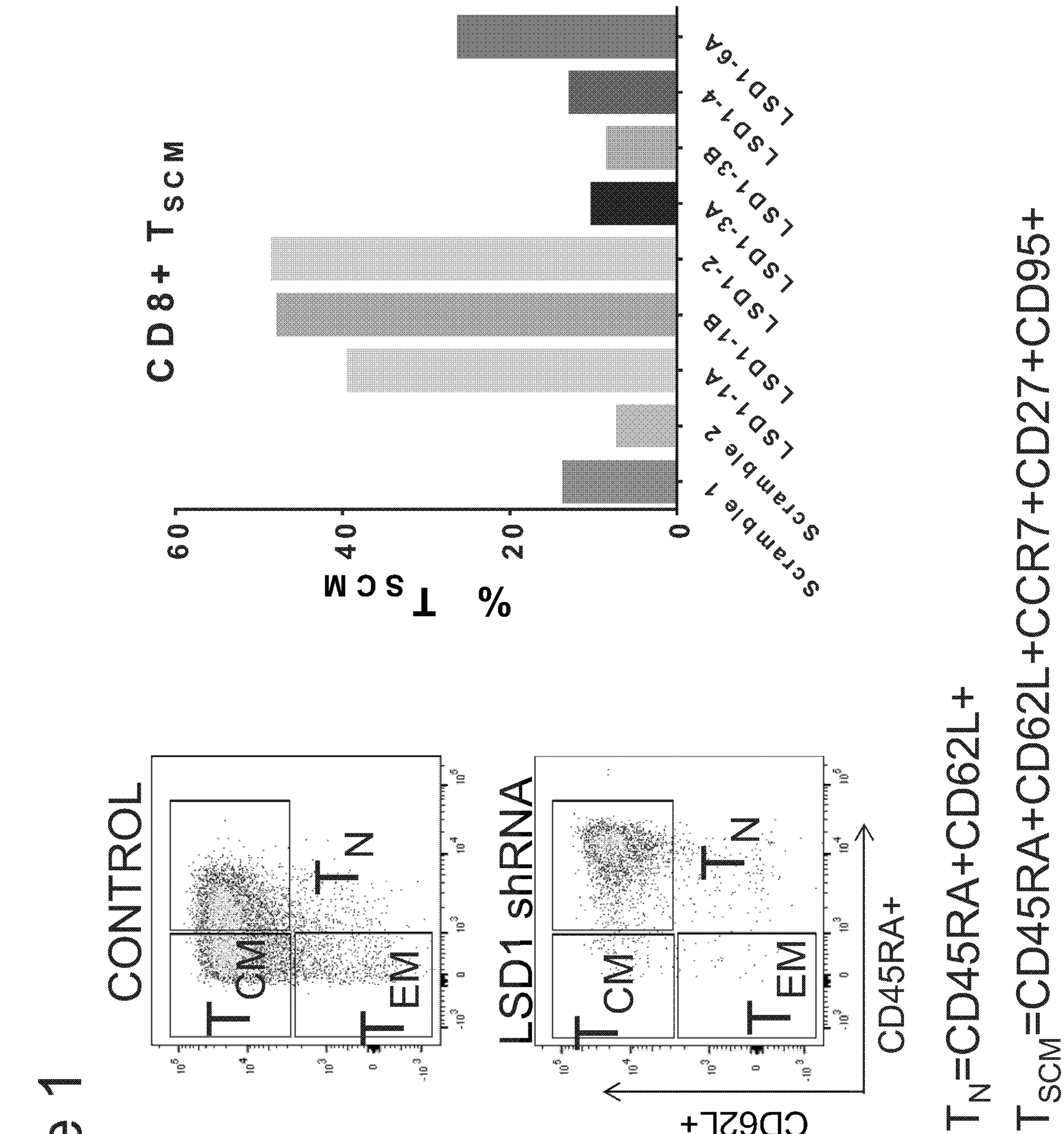
72. The composition of any of claims 68-71, wherein the population of immune effector cells comprises cells engineered to express a CAR.

73. An LSD1 inhibitor, for use in treating a subject, wherein said subject has received, is receiving, or is about to receive therapy comprising an immune effector cell, e.g., an immune effector cell engineered to express a CAR.

74. An LSD1 inhibitor, for use in the manufacture of an immune effector cell, e.g., an immune effector cell engineered to express a CAR.

75. A method of manufacturing an immune effector cells comprising introducing into said cells nucleic acid encoding a CAR, wherein the nucleic acid integrates into the genome of said cell within the LSD1 gene, such that LSD1 expression and/or function is reduced.

Figure 1


CONTROL
LSD1 shRNA
T_{CM}
T_{EM}
T_N
CD45RA+
CD62L+
T_N=CD45RA+CD62L+
T_{SCM}=CD45RA+CD62L+CCR7+CD27+CD95+
CD8+ T_{SCM}
% T_{SCM}
0
20
40
60
Scramble 1
Scramble 2
LSD1-1A
LSD1-1B
LSD1-2
LSD1-3A
LSD1-3B
LSD1-4
LSD1-6

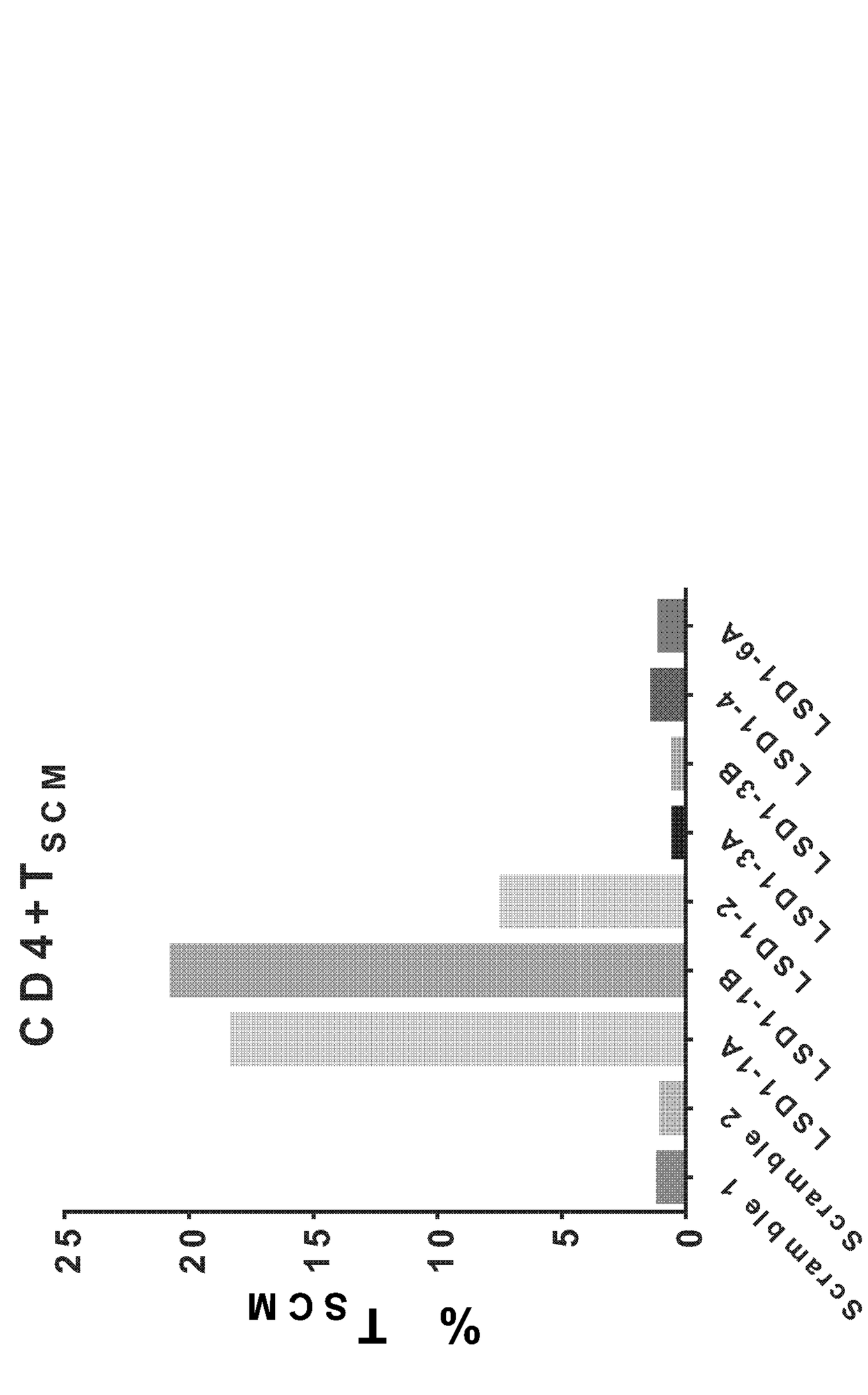

CD4+TSCM
%TSCM
0
5
10
15
20
25
Scramble 1
Scramble 2
LSD1-1A
LSD1-1B
LSD1-2
LSD1-3A
LSD1-3B
LSD1-4
LSD1-6A

Figure 2

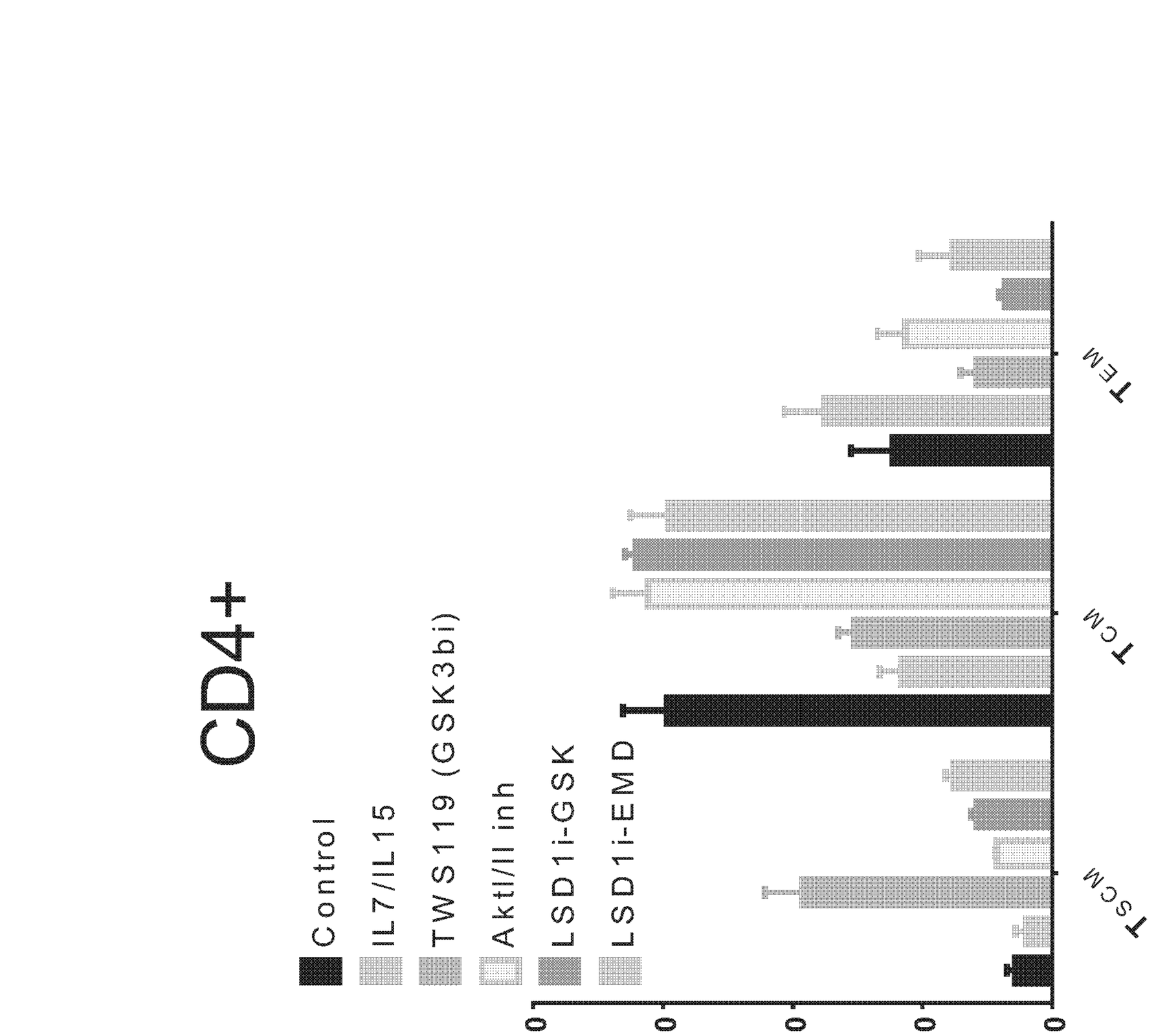

CD4+
Control
IL7/IL15
TWS119 (GSK3bi)
Aktl/II inh
LSD1i-GSK
LSD1i-EMD
% Cell Type
0
20
40
60
80
TSCM
TCM
TEM

Figure 3A

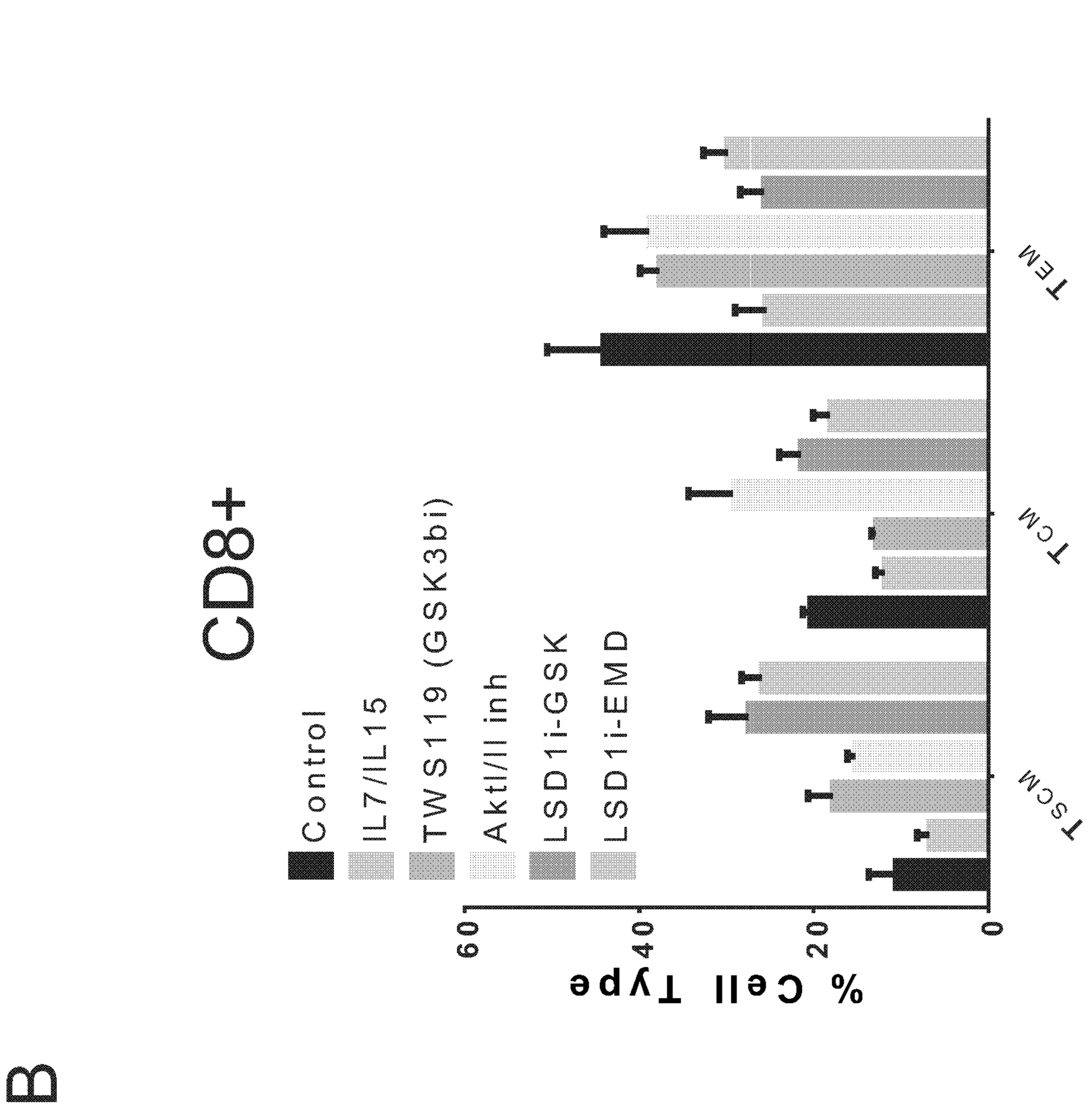

B
CD8+
Control
IL7/IL15
TWS119 (GSK3bi)
Akti/II inh
LSD1i-GSK
LSD1i-EMD
% Cell Type
0
20
40
60
TSCM
TCM
TEM

Figure 3

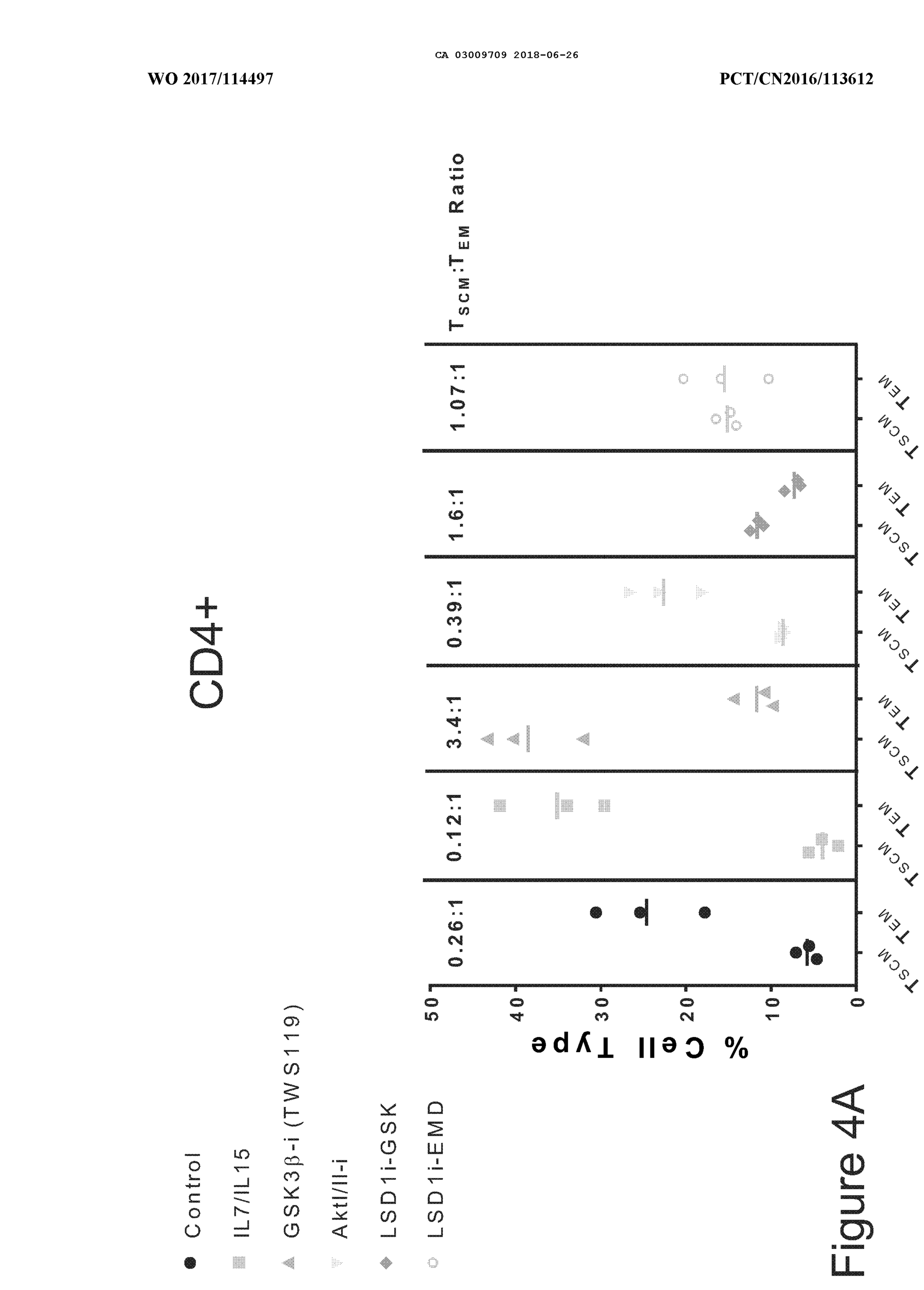

Tscm:Tem Ratio
CD4+
% Cell Type
0
10
20
30
40
50
0.26:1
0.12:1
3.4:1
0.39:1
1.6:1
1.07:1
Tscm
Tem
Control
IL7/IL15
GSK3β-i (TW
AktI/II-i
LSD1i-GSK
LSD1i-EMD
(S119)

Figure 4A

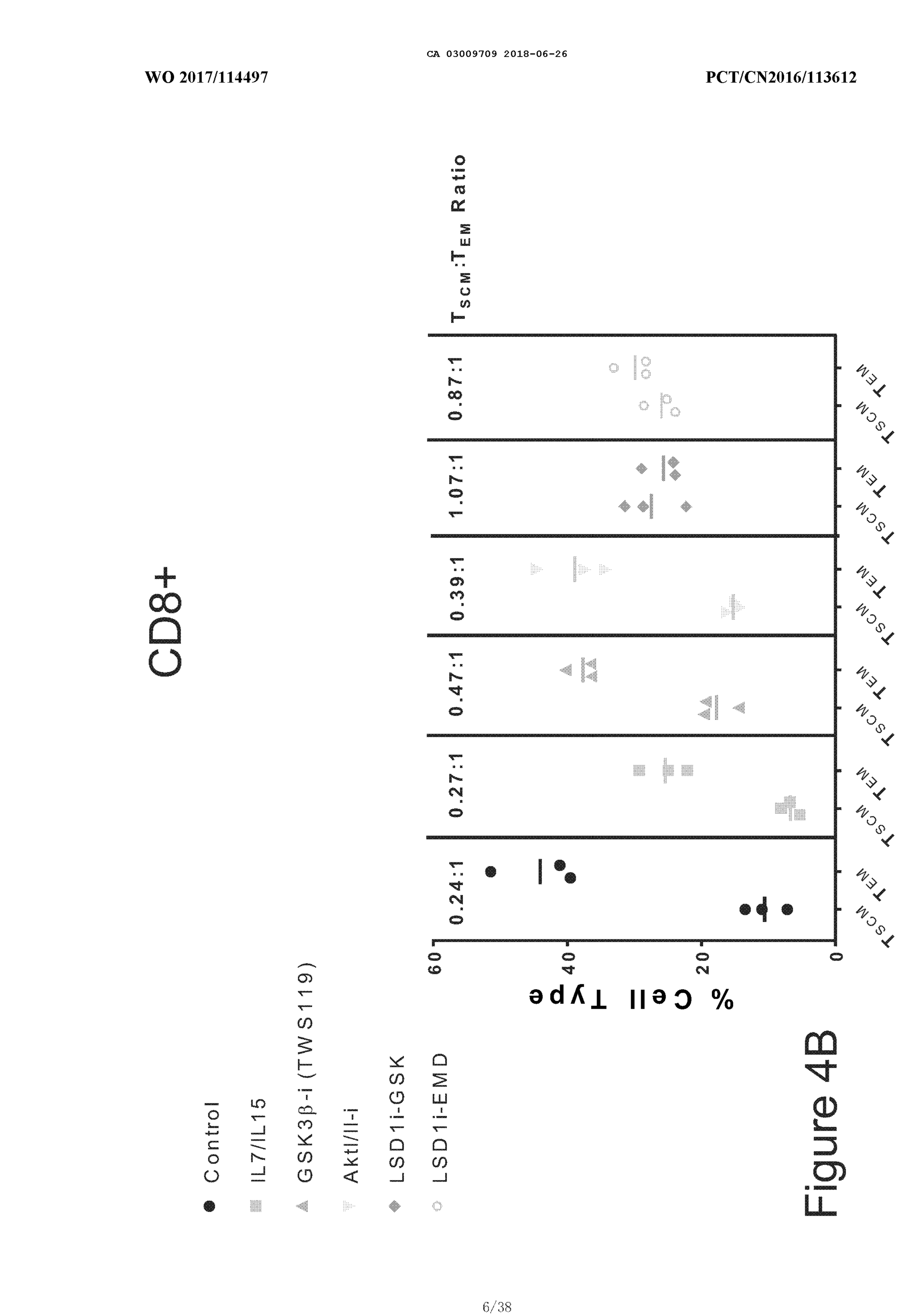

CD8+
TSCM:TEM Ratio
0.87:1
1.07:1
0.39:1
0.47:1
0.27:1
0.24:1
TSCM TEM
% Cell Type
60
40
20
0
Control
IL7/IL15
GSK3β-i (TWS119)
Akti/ii-i
LSD1i-GSK
LSD1i-EMD

Figure 4B

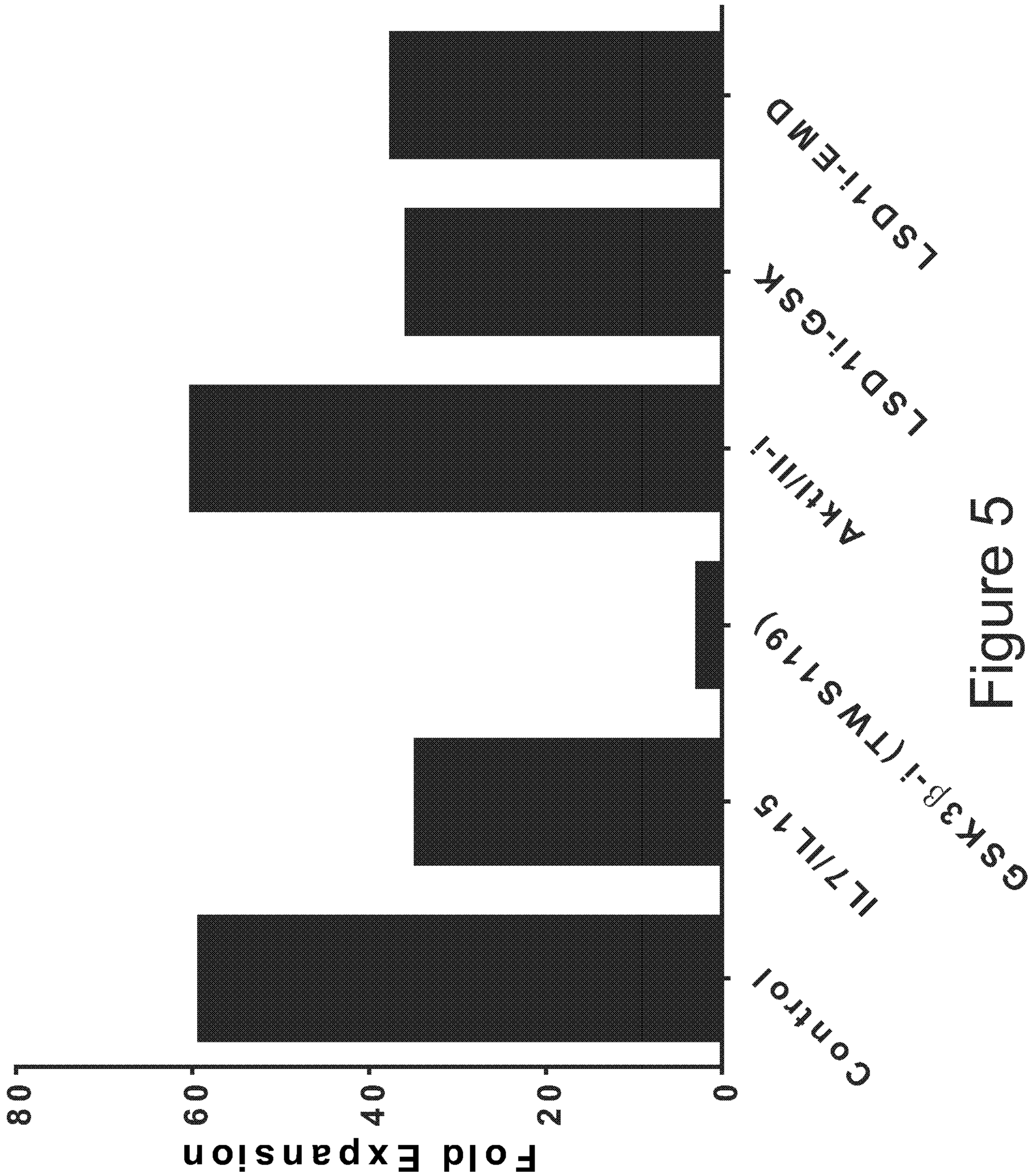

Fold Expansion
0
2
4
6
8
Control
IL7/IL-15
GSK3β-i (TWS119)
AKTi/MK-i
LSD1i-GSK
LSD1i-EMD

Figure 5

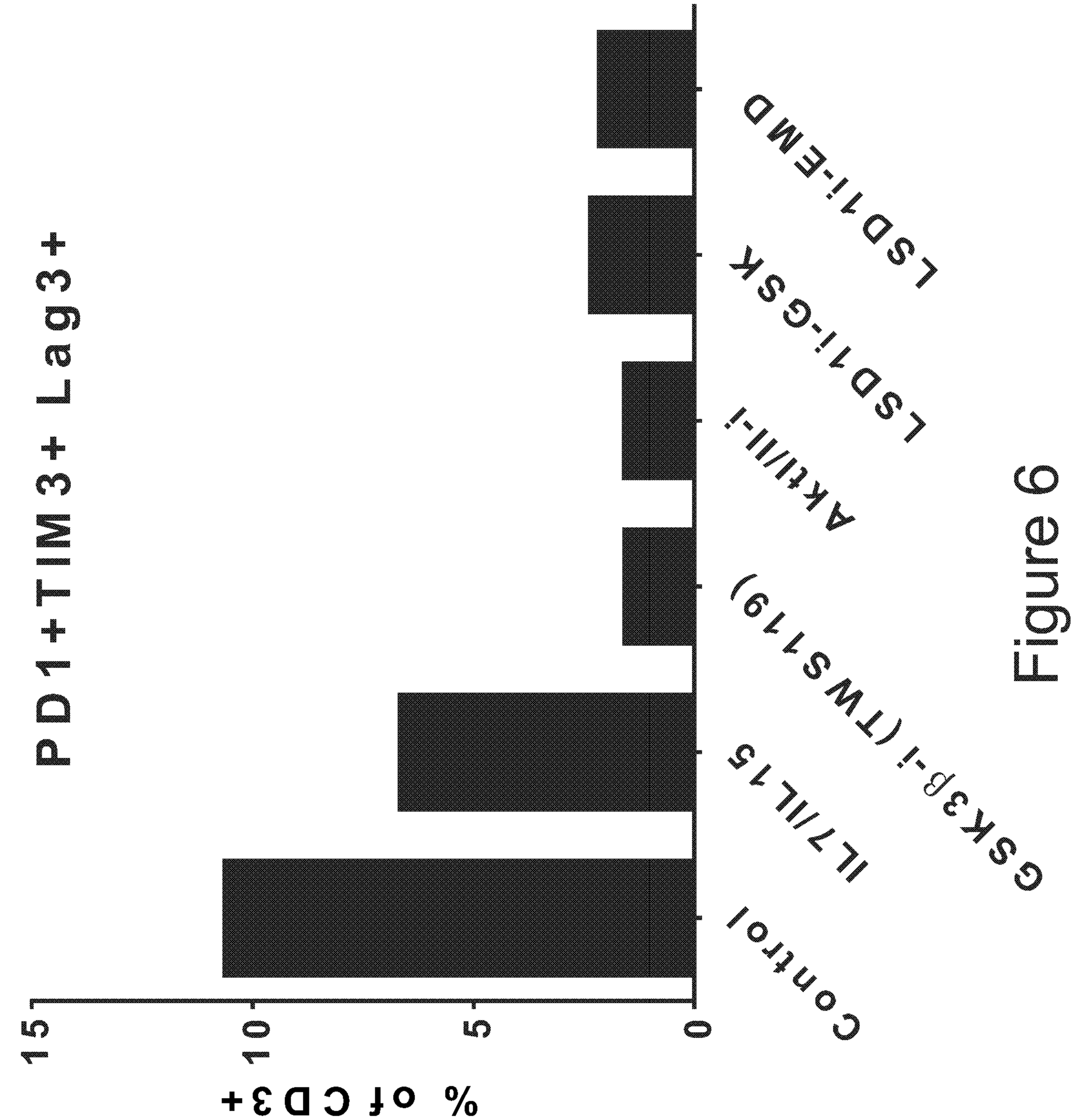

PD1+TIM3+Lag3+
% of CD3+
0
5
0
5
Control
IL7/IL15
GSK3β-i (TWS119)
AKTi/II-i
LSD1i-GSK
LSD1i-EMD

Figure 9

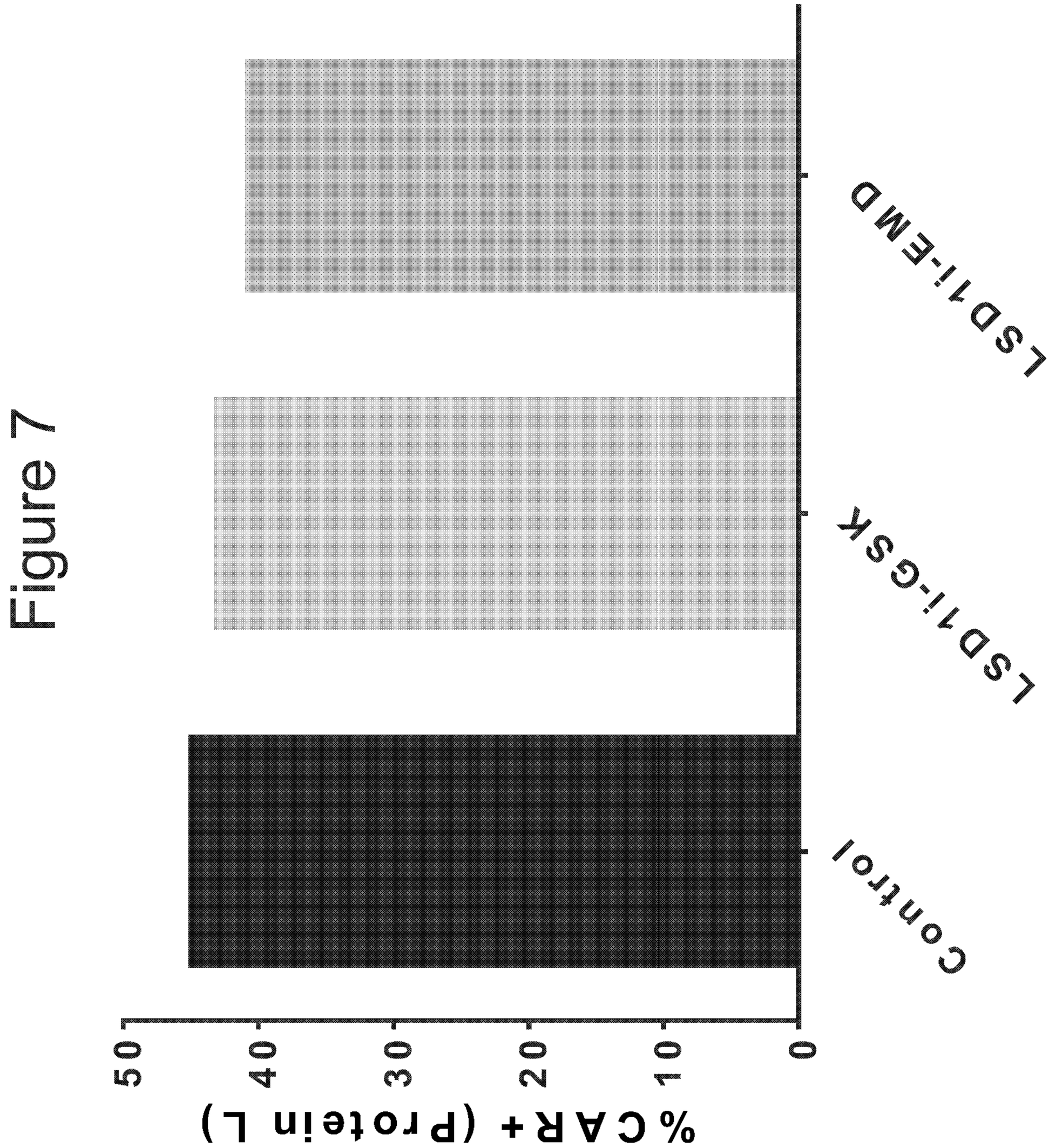


Figure 7

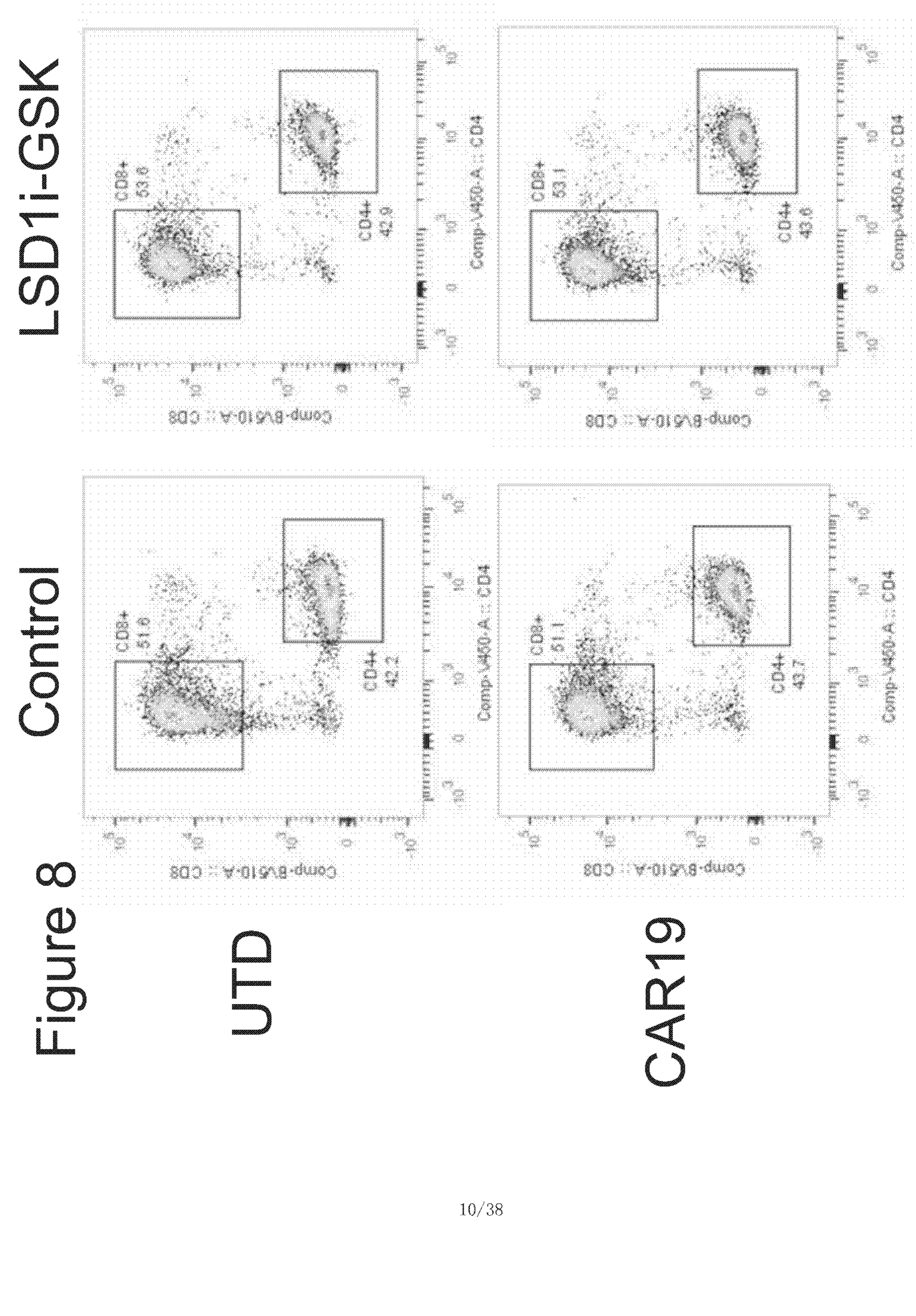

UTD
CAR19
Control
LSD1i-GSK
CD8+
51.0
CD4+
42.2
CD8+
61.1
CD4+
49.7
CD8+
63.6
CD4+
42.9
CD8+
63.1
CD4+
49.6
Comp-B\510-A :: CD8
Comp-V450-A :: CD4

Figure 8

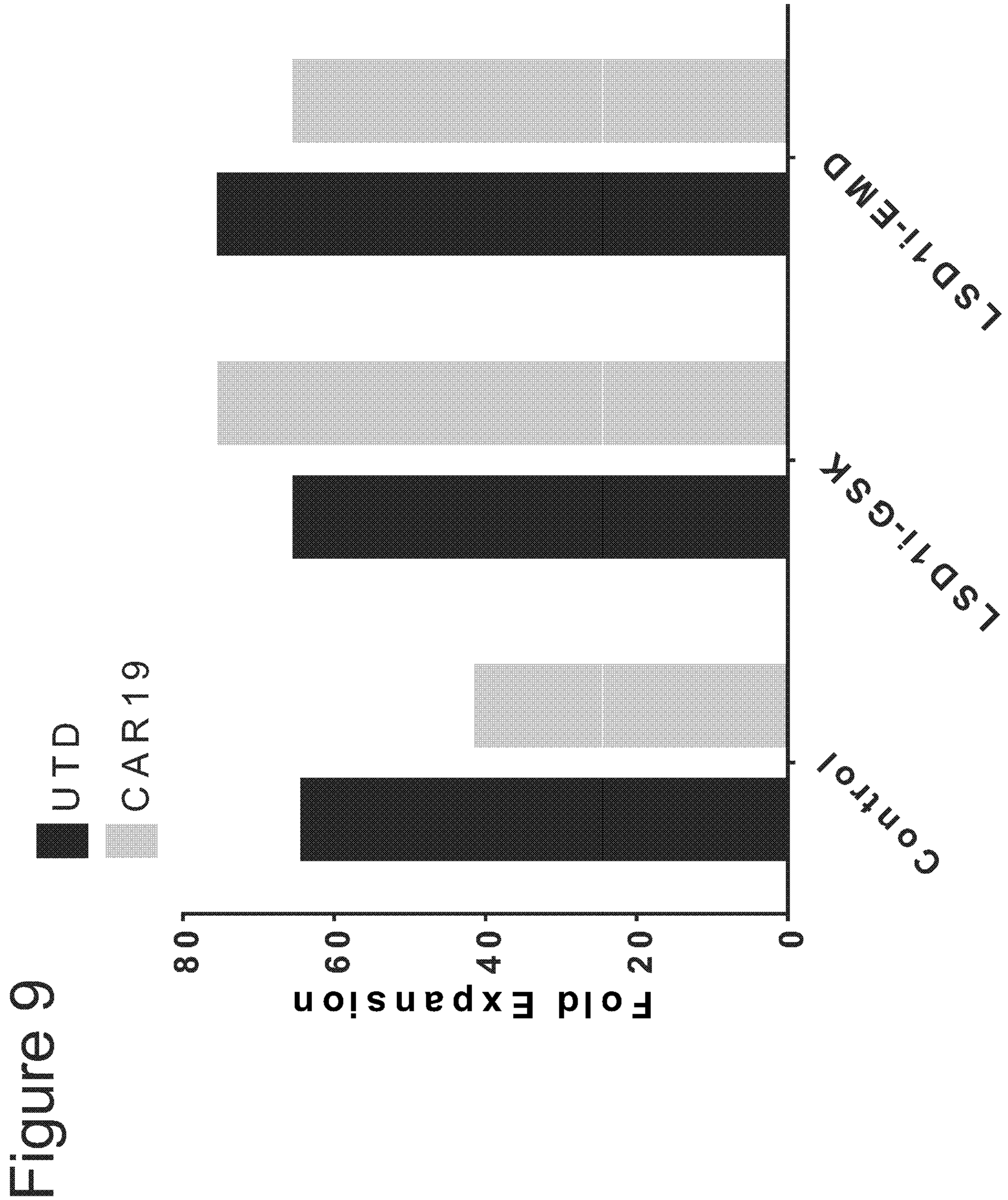

UTD
CAR19
Fold Expansion
0
20
40
60
80
Control
LSD1i-GSK
LSD1i-EMD

Figure 9

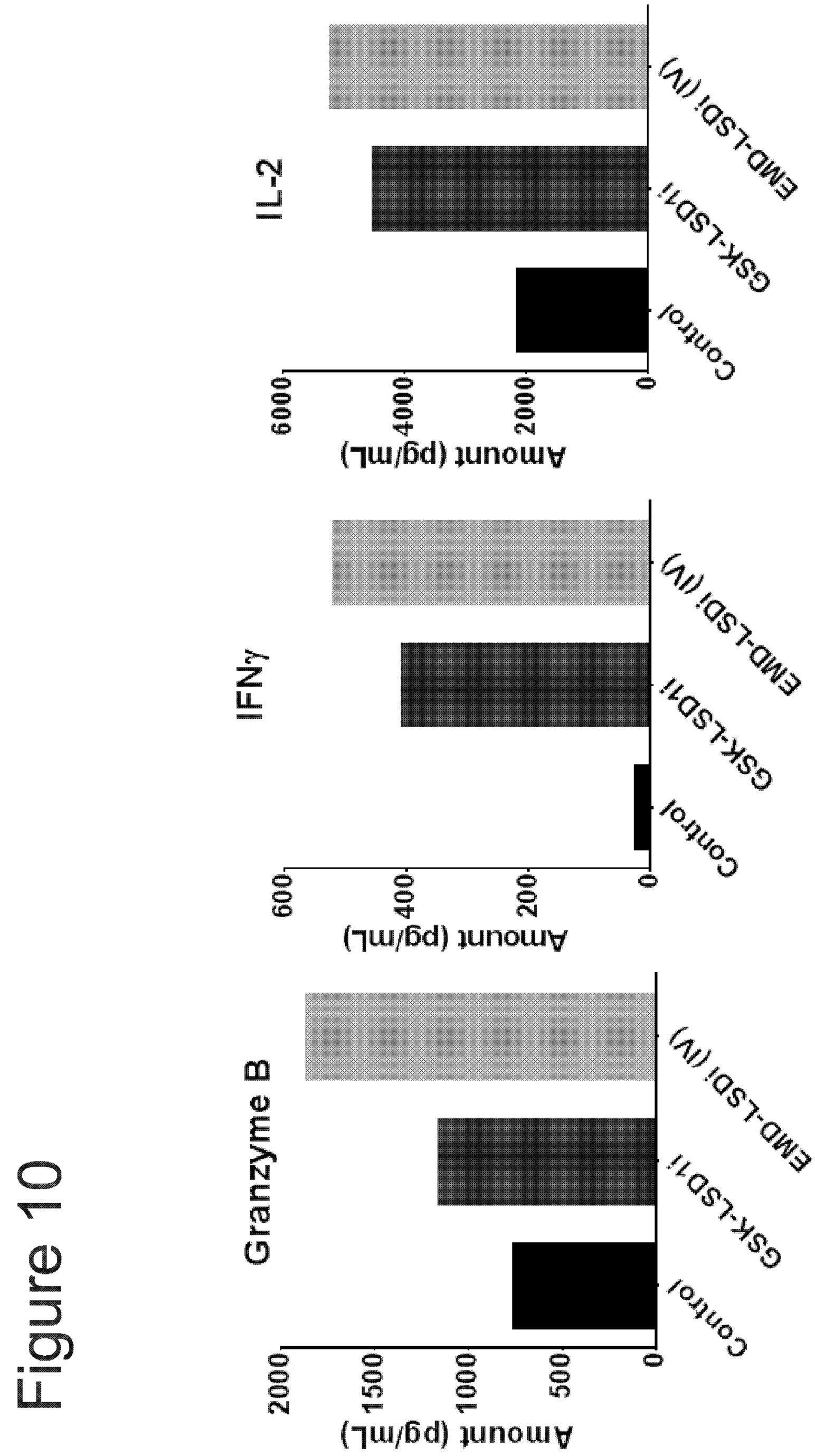

Granzyme B
Amount (pg/mL)
2000
1500
1000
500
0
Control
GSK-LSD1i
EMD-LSD1 (IV)
IFNγ
Amount (pg/mL)
600
400
200
0
Control
GSK-LSD1i
EMD-LSD1 (IV)
IL-2
Amount (pg/mL)
6000
4000
2000
0
Control
GSK-LSD1i
EMD-LSD1 (IV)

Figure 10

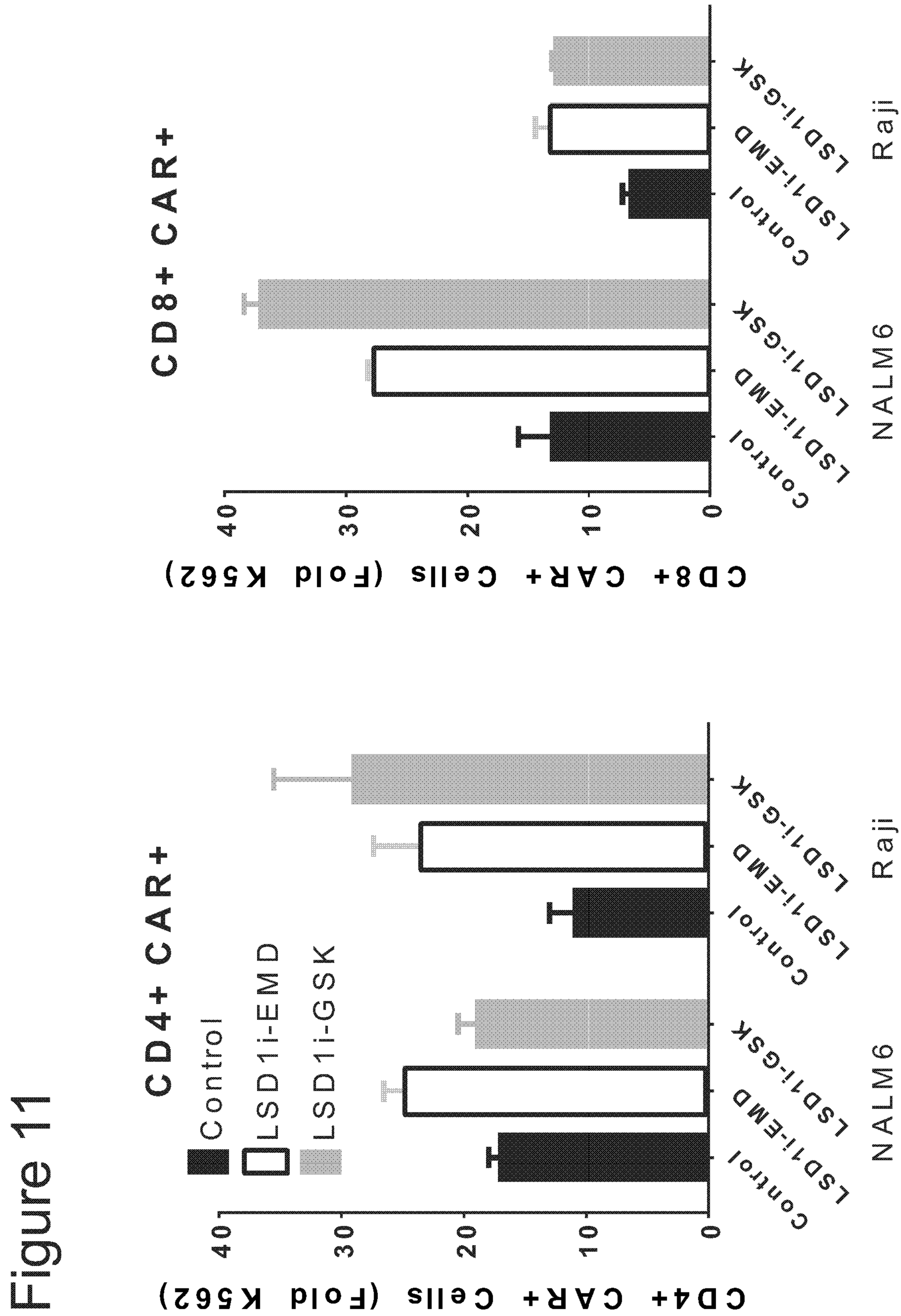

CD8+ CAR+ Cells (Fold K562)
40
30
20
10
0
Control
LSD1i-EMD
LSD1i-GSK
NALM6
Raji
CD4+ CAR+ Cells (Fold K562)
Control
LSD1i-EMD
LSD1i-GSK

Figure 11

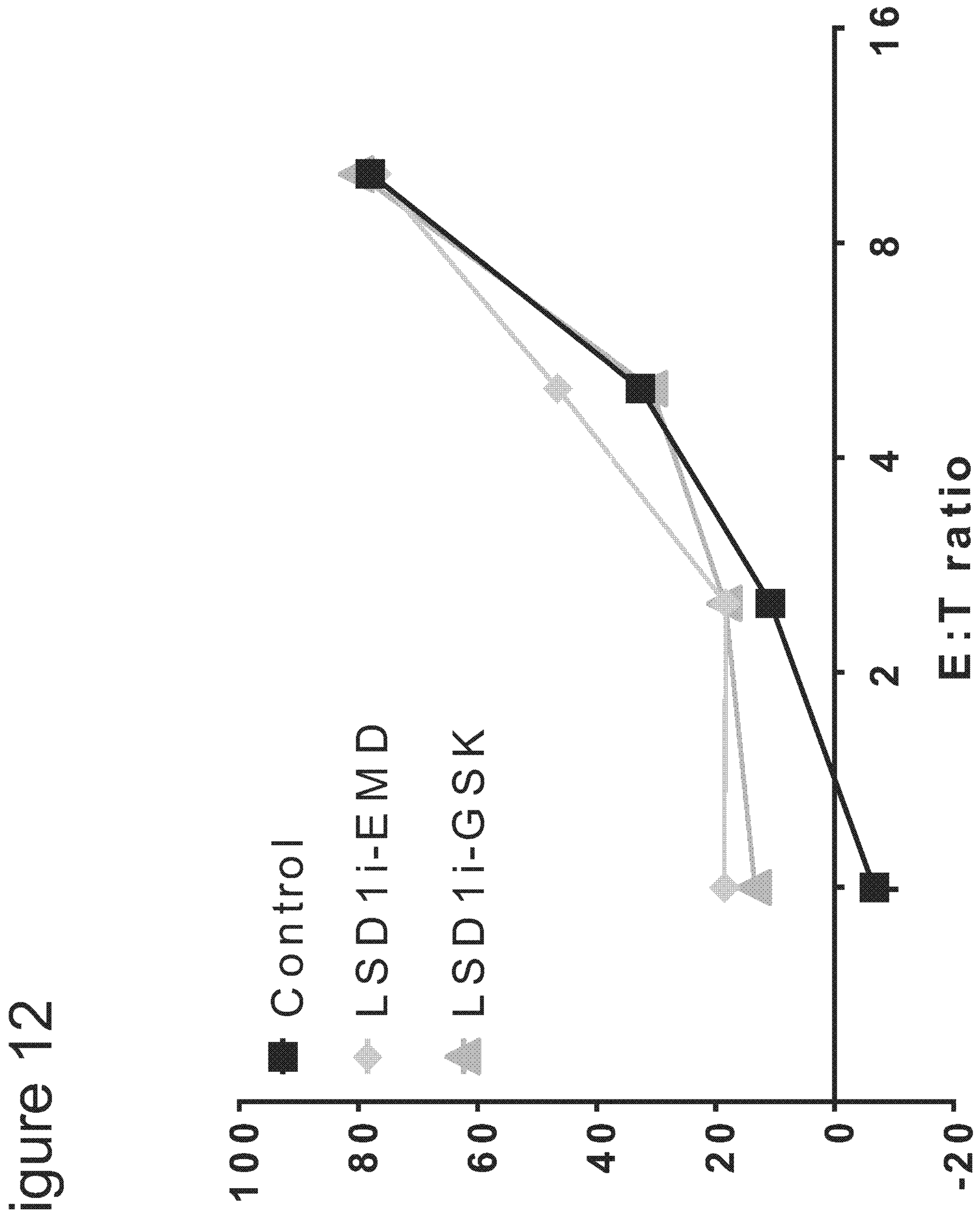

Control
LSD1i-EMD
LSD1i-GSK
100
80
60
40
20
0
-20
% Killing
E:T ratio
2
4
8
16

Figure 12

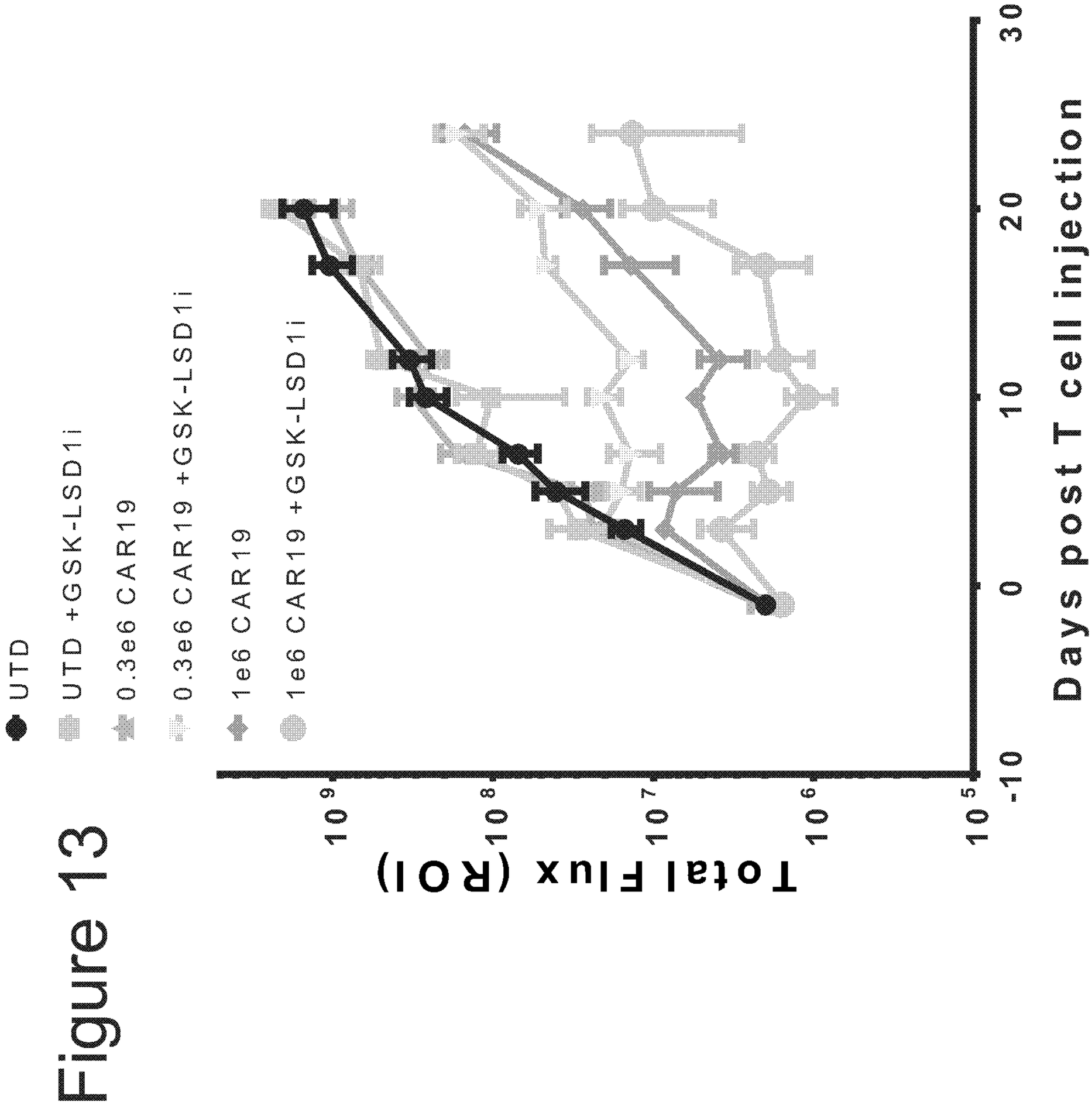

UTD
UTD +GSK-LSD1i
0.3e6 CAR19
0.3e6 CAR19 +GSK-LSD1i
1e6 CAR19
1e6 CAR19 +GSK-LSD1i
Total Flux (ROI)
10^9
10^8
10^7
10^6
10^5
-10
0
10
20
30
Days post T cell injection

Figure 13

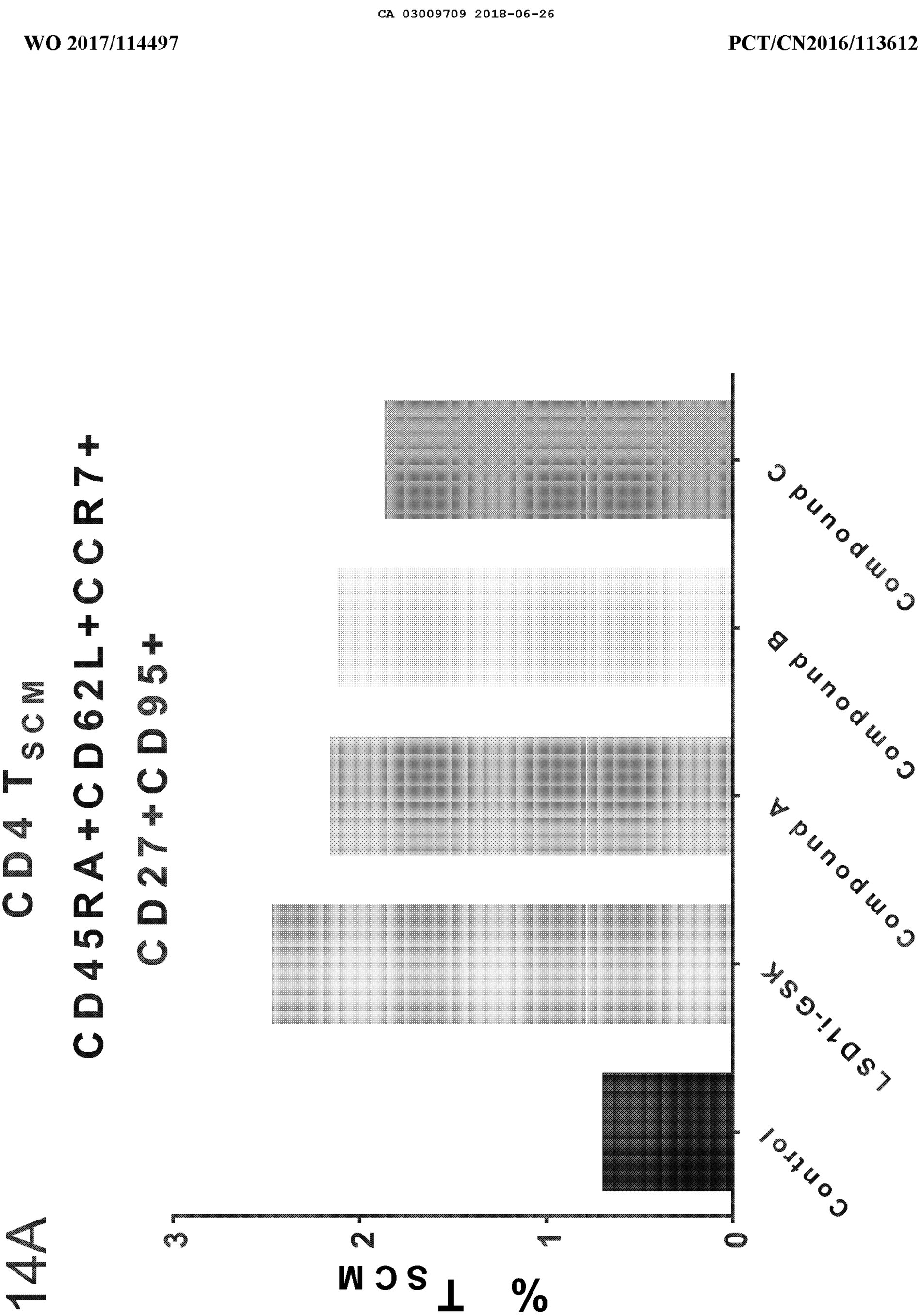

A
CD4 TSCM
CD45RA+CD62L+CCR7+
CD27+CD95+
% TSCM
3
2
1
0
Control
LSD1i-GSK
Compound A
Compound B
Compound C

Figure 14

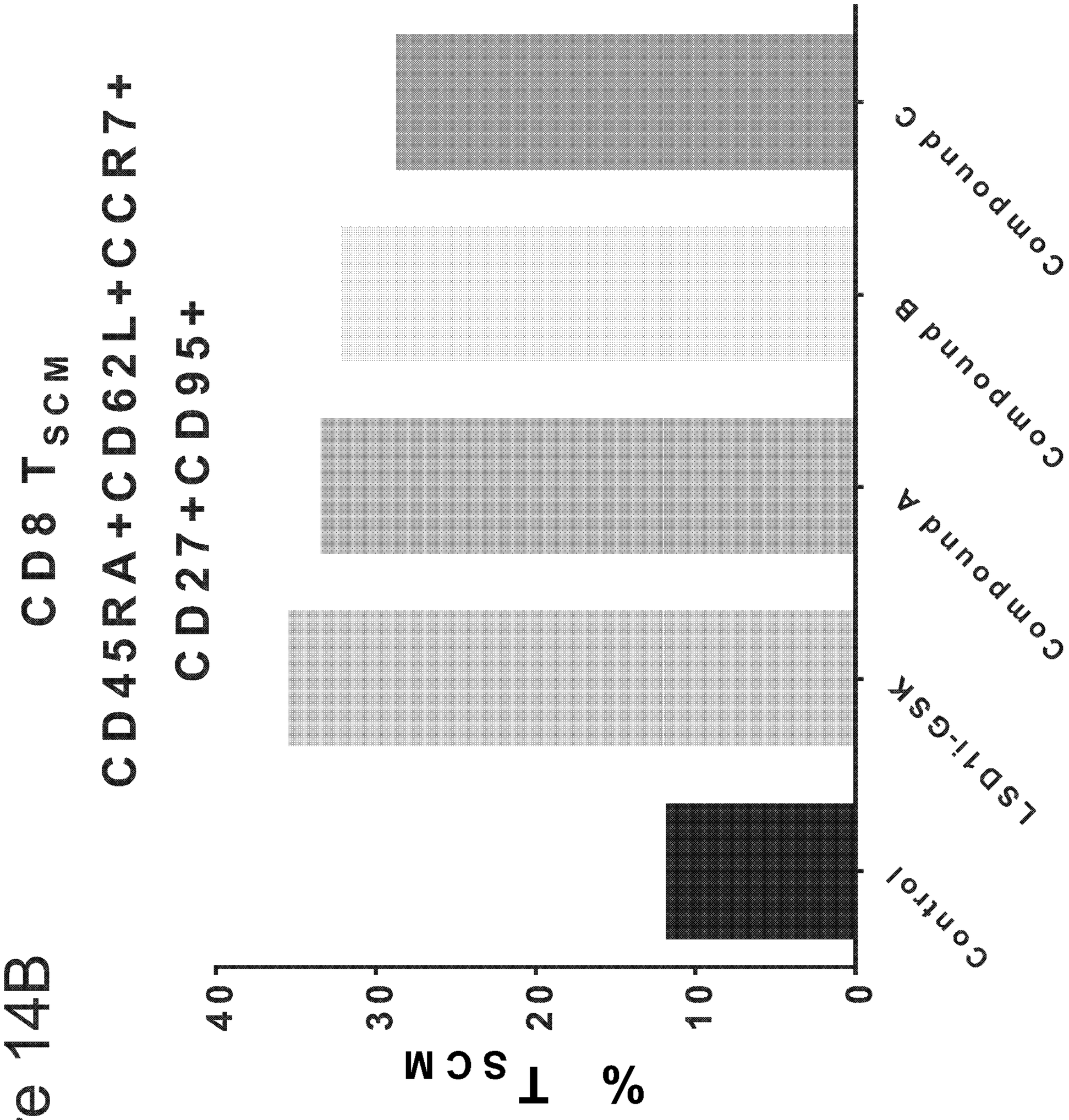

CD8 TSCM
CD45RA+CD62L+CCR7+
CD27+CD95+
40
30
20
10
0
TSCM %
Control
LSD1i-GSK
Compound A
Compound B
Compound C

Figure 4B

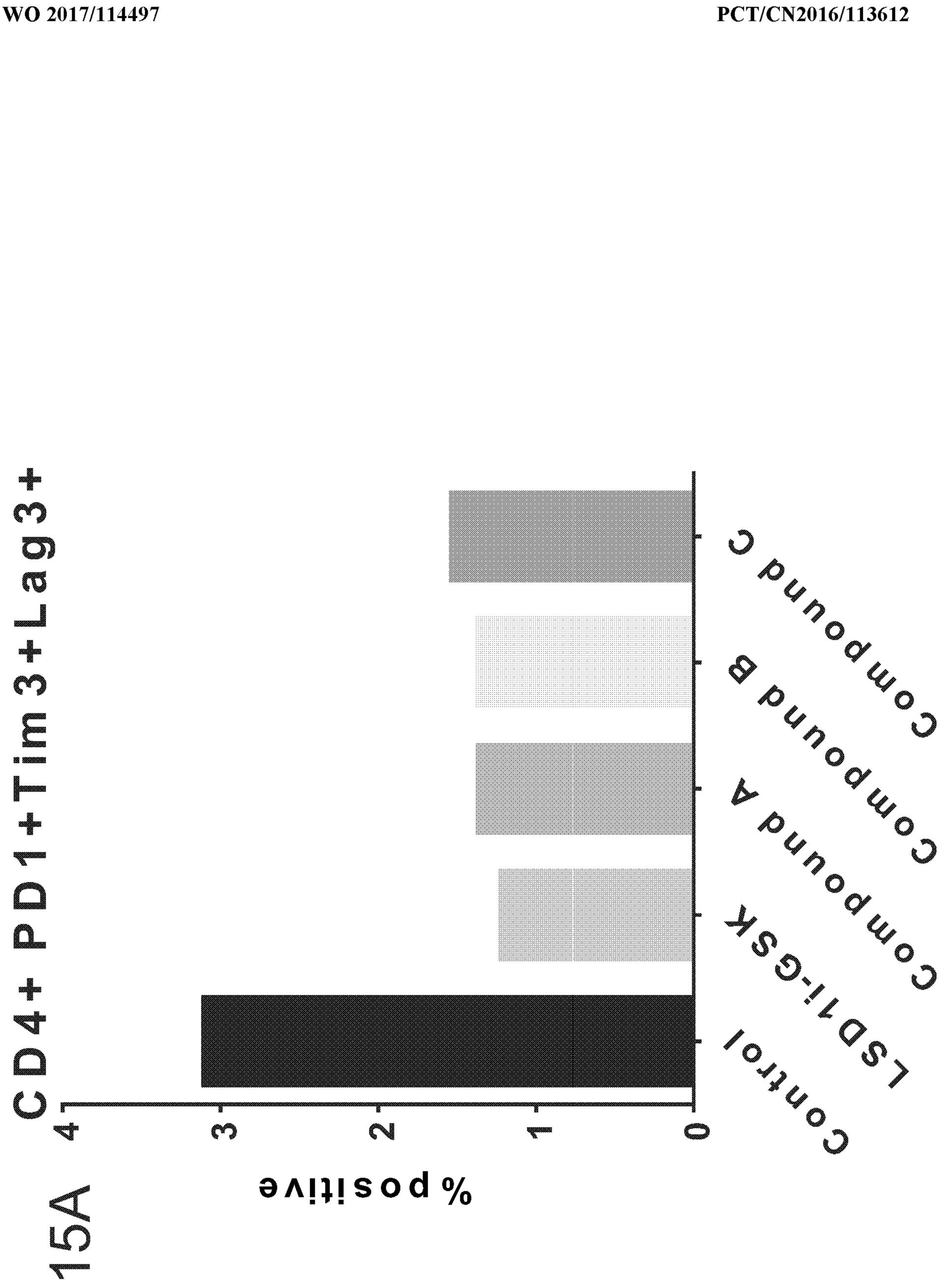

CD4+ PD1+ Tim3+ Lag3+
% positive
0
1
2
3
4
Control
LSD1i-GSK
Compound A
Compound B
Compound C

Figure 15A

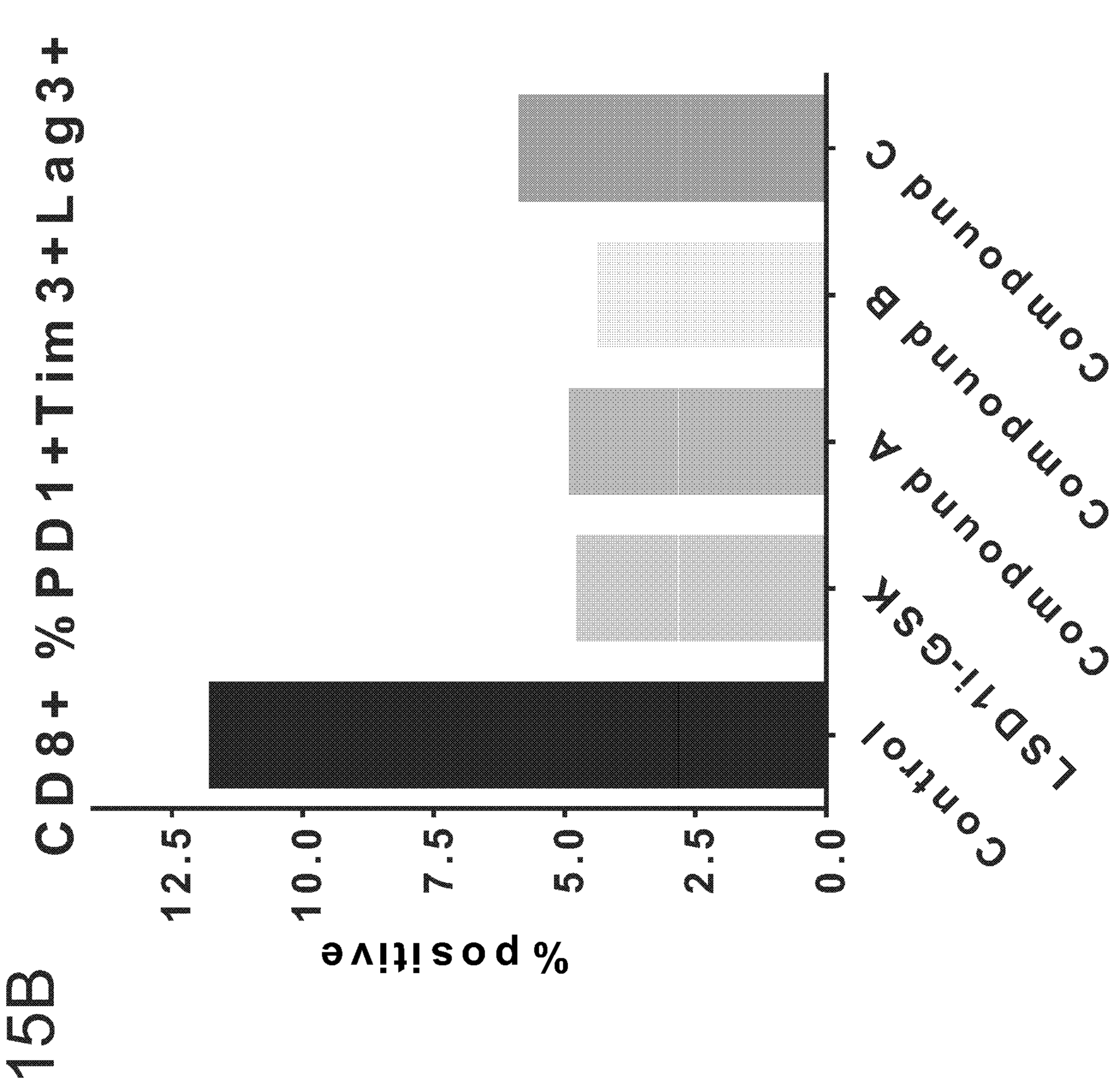

CD8+ %PD1+Tim3+Lag3+
% positive
12.5
10.0
7.5
5.0
2.5
0.0
Control
LSD1i-GSK
Compound A
Compound B
Compound C

Figure 15B

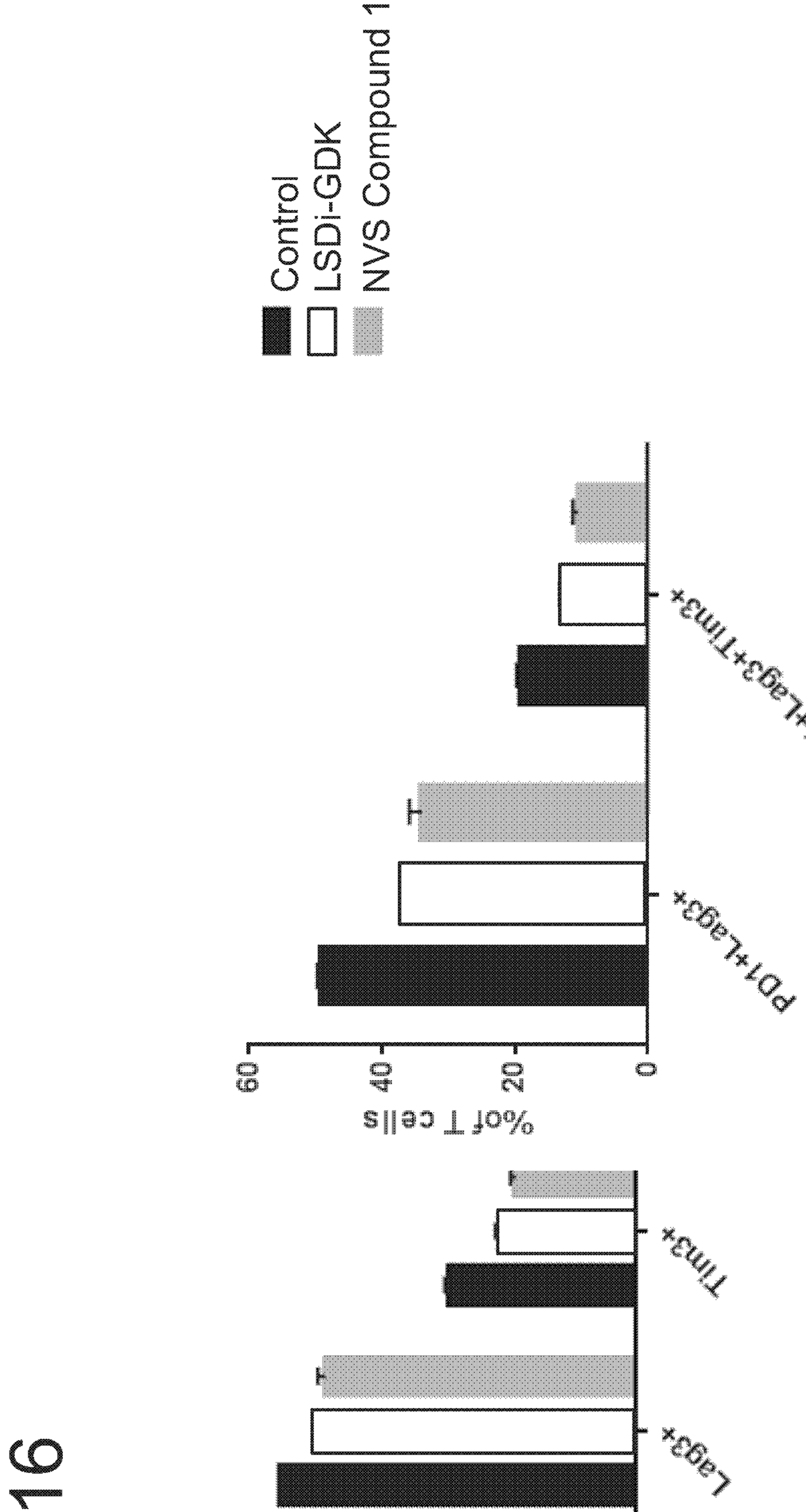

Control
LSDi-GDK
NVS Compound 1
PD1+Lag3+Tim3+
Lag3+
Tim3+
PD1+
%of T cells
0
20
40
60
80

Figure 19

Figure 17

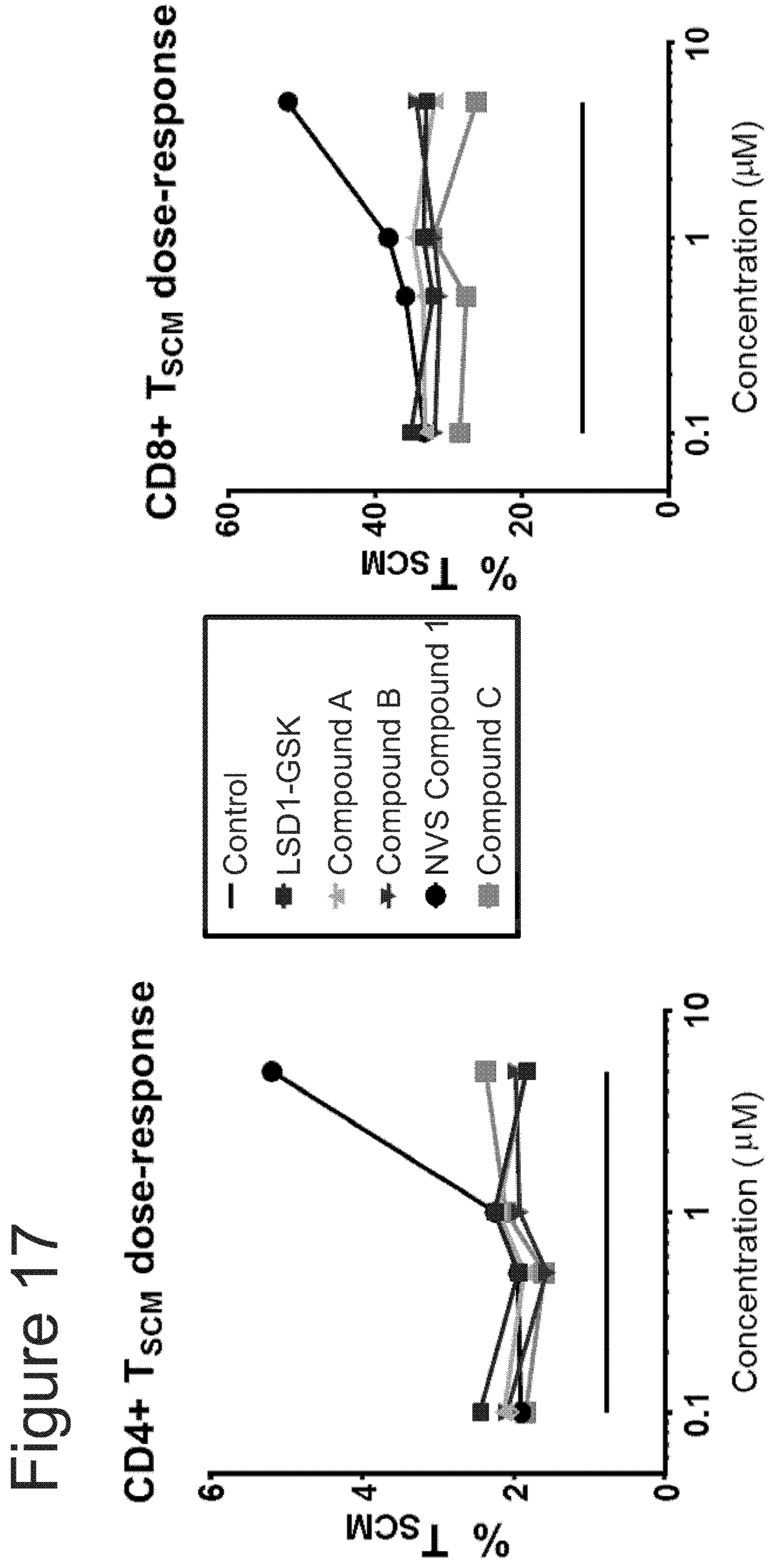

CD4+ TSCM % dose-response
%TSCM
60
40
20
0
0.1
1
10
Concentration (μM)
Control
LSD1-GSK
Compound A
Compound B
Compound C
CD8+ TSCM % dose-response
%TSCM
6
4
2
0
0.1
1
10
Concentration (μM)

Figure 18

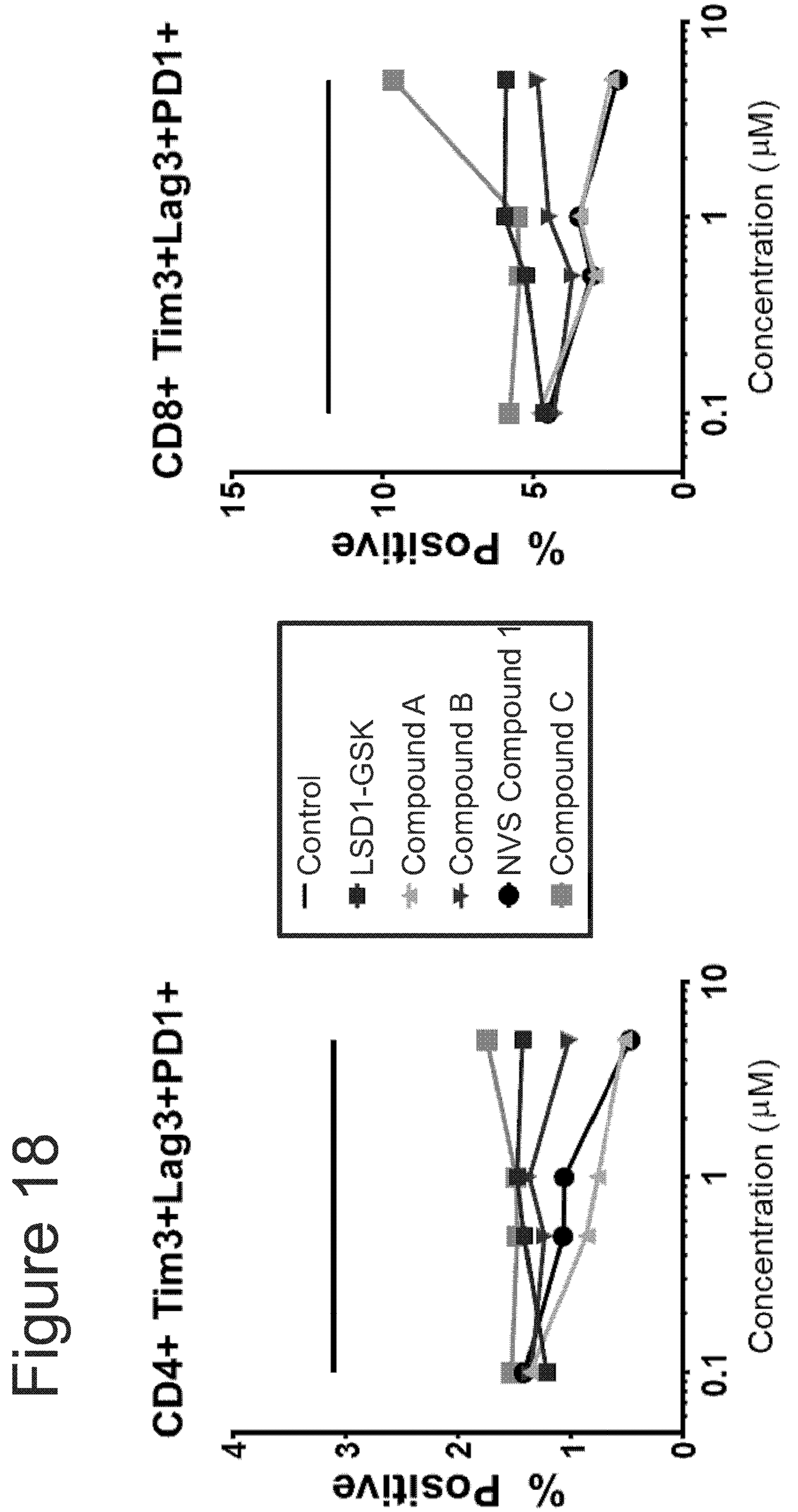

CD4+ Tim3+Lag3+PD1+
% Positive
0
1
2
3
4
0.1
1
10
Concentration (μM)
Control
LSD1-GSK
Compound A
Compound B
Compound C
NVS Compound 1
CD8+ Tim3+Lag3+PD1+
% Positive
0
5
10
15
0.1
1
10
Concentration (μM)

Figure 19

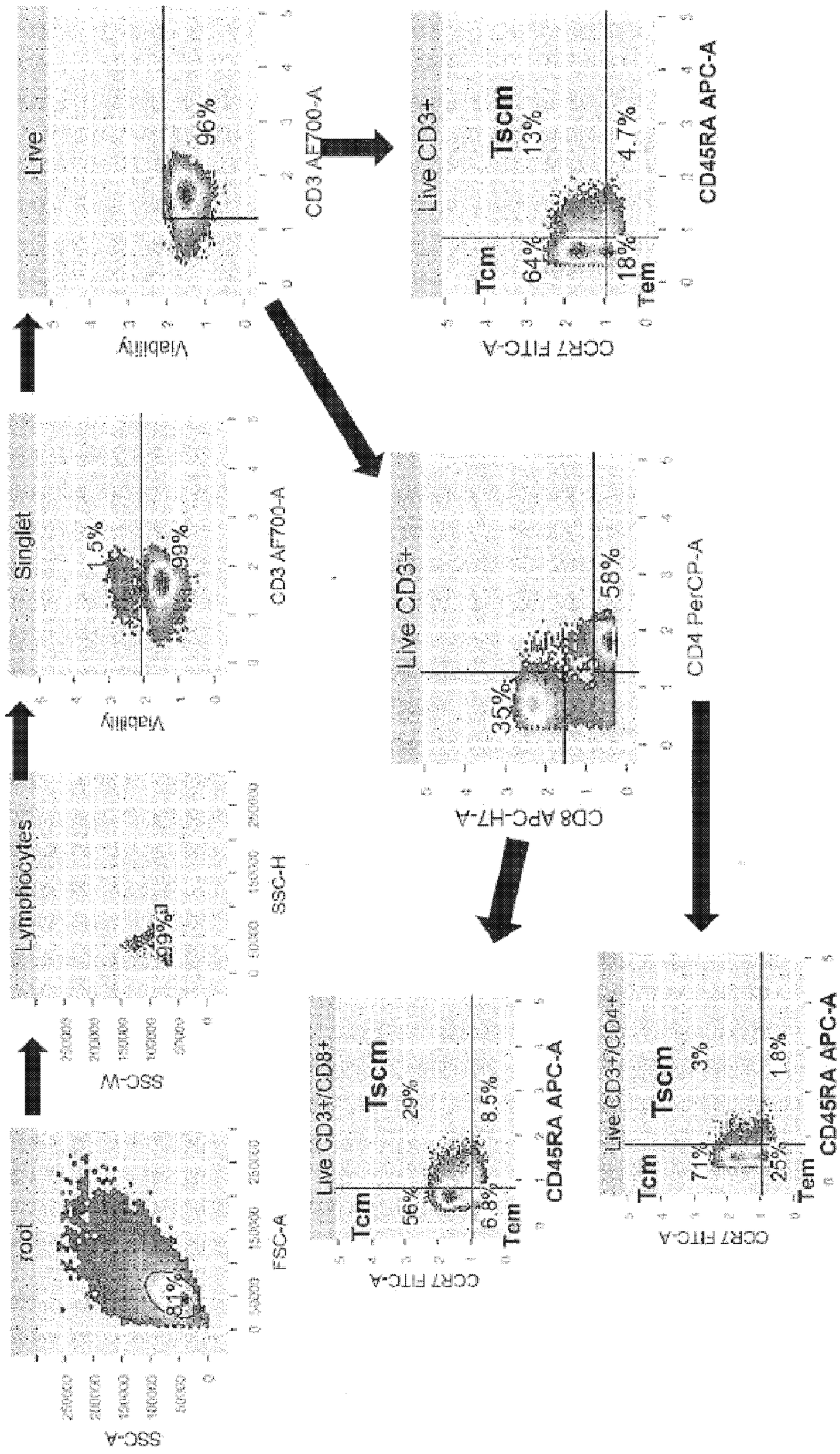

Live
96%
CD3 AF700-A
Viability
Live CD3+
Tscm
13%
Tcm
64%
18%
4.7%
Tem
CD45RA APC-A
CCR7 FITC-A
Singlet
1.5%
99%
CD3 AF700-A
Viability
Live CD3+
35%
58%
CD4 PerCP-A
CD8 APC-H7-A
Lymphocytes
99%
SSC-W
SSC-H
Tscm
29%
8.5%
Tcm
56%
6.8%
Tem
CD45RA APC-A
CCR7 FITC-A
Live CD3+/CD4+
Tscm
3%
71%
25%
1.8%
Tcm
Tem
CD45RA APC-A
CCR7 FITC-A
Root
81%
SSC-A
FSC-A

Figure 20A

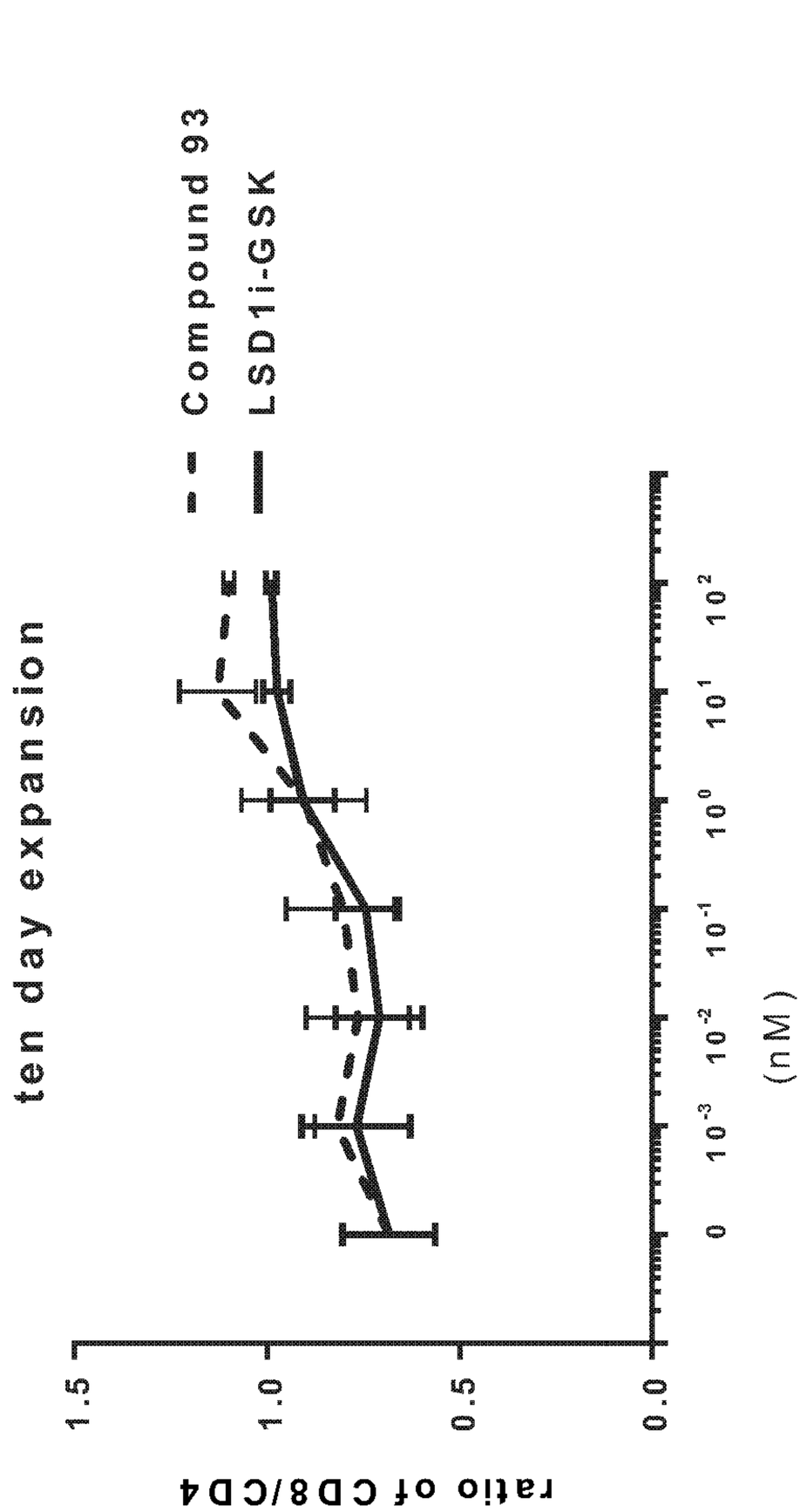
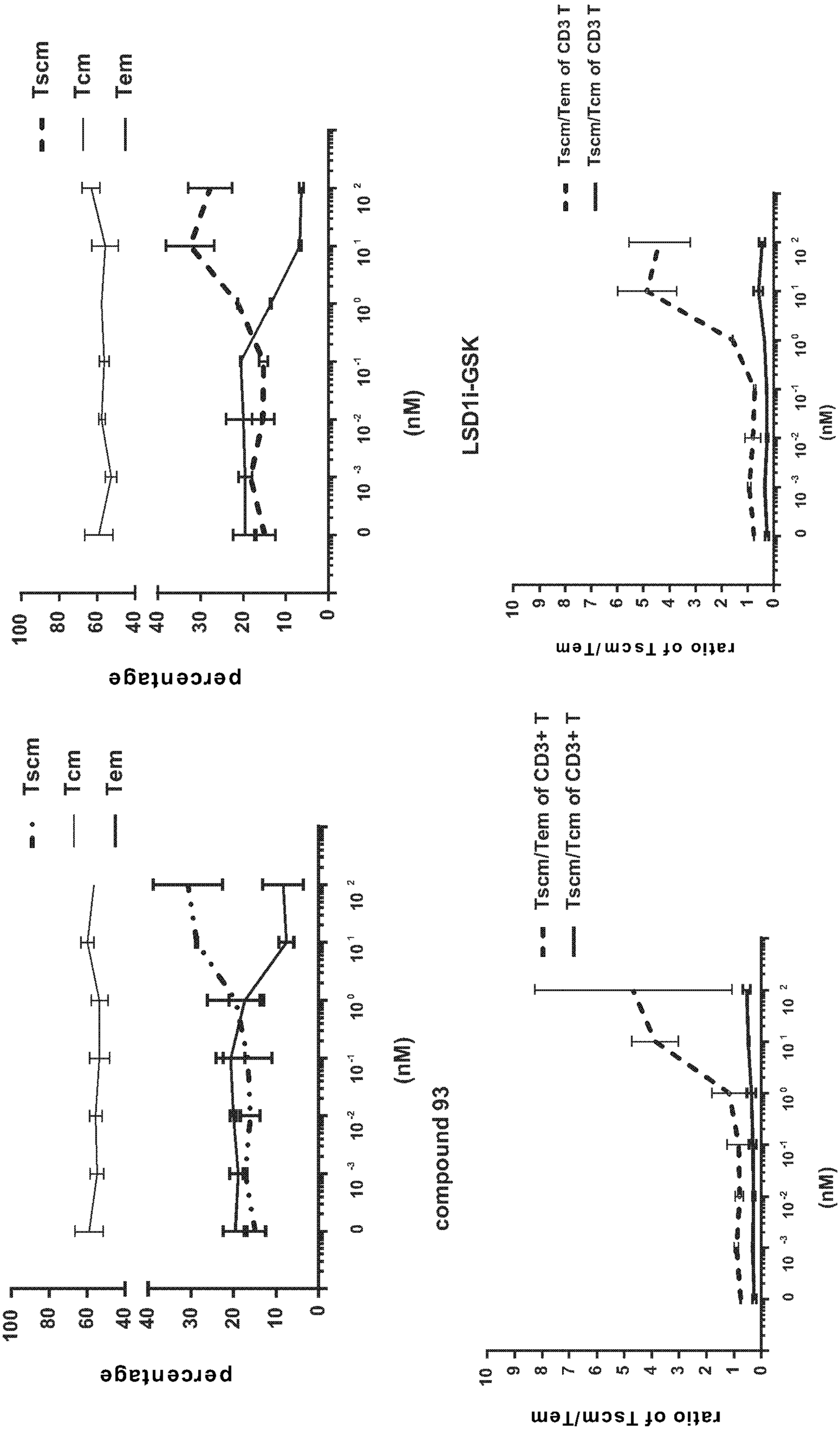

ten day expansion
ratio of CD8/CD4
1.5
1.0
0.5
0.0
0
10^{-3}
10^{-2}
10^{-1}
10^{0}
10^{1}
10^{2}
(nM)
Compound 93
LSD1i-GSK

Figure 20B


LSD1i-GSK (CD3+ T)
Tscm
Tcm
Tem
percentage
(nM)
LSD1i-GSK
Tscm/Tem of CD3 T
Tscm/Tcm of CD3 T
ratio of Tscm/Tem
compound 93 (CD3+ T)
compound 93

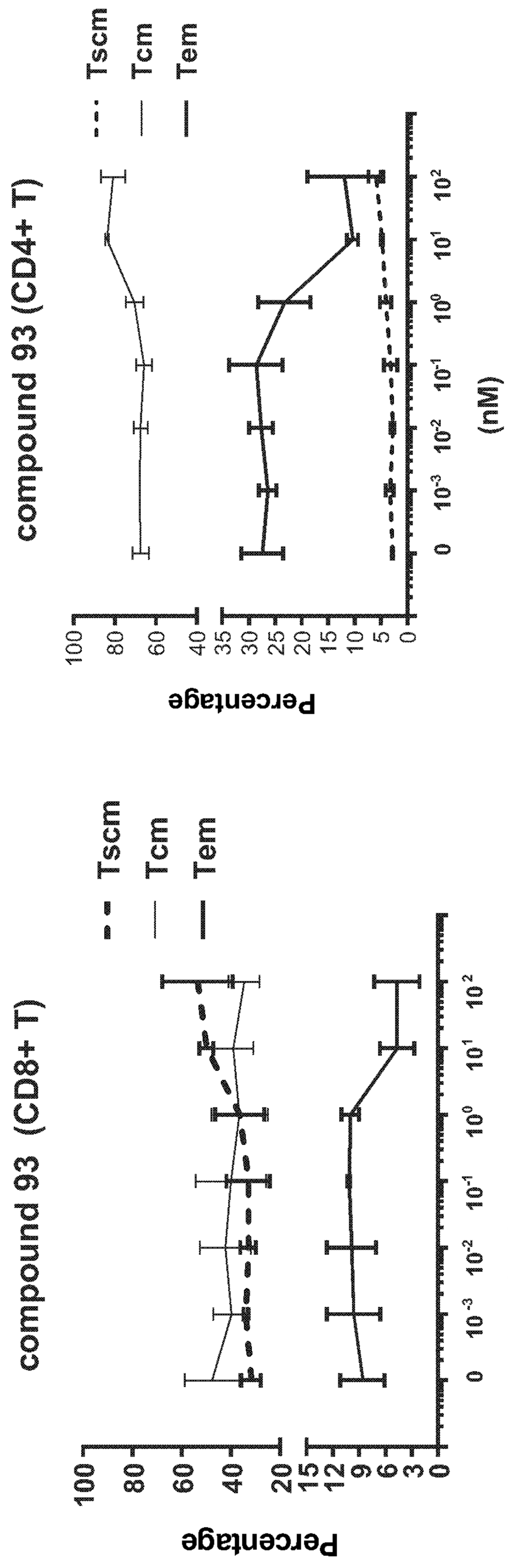
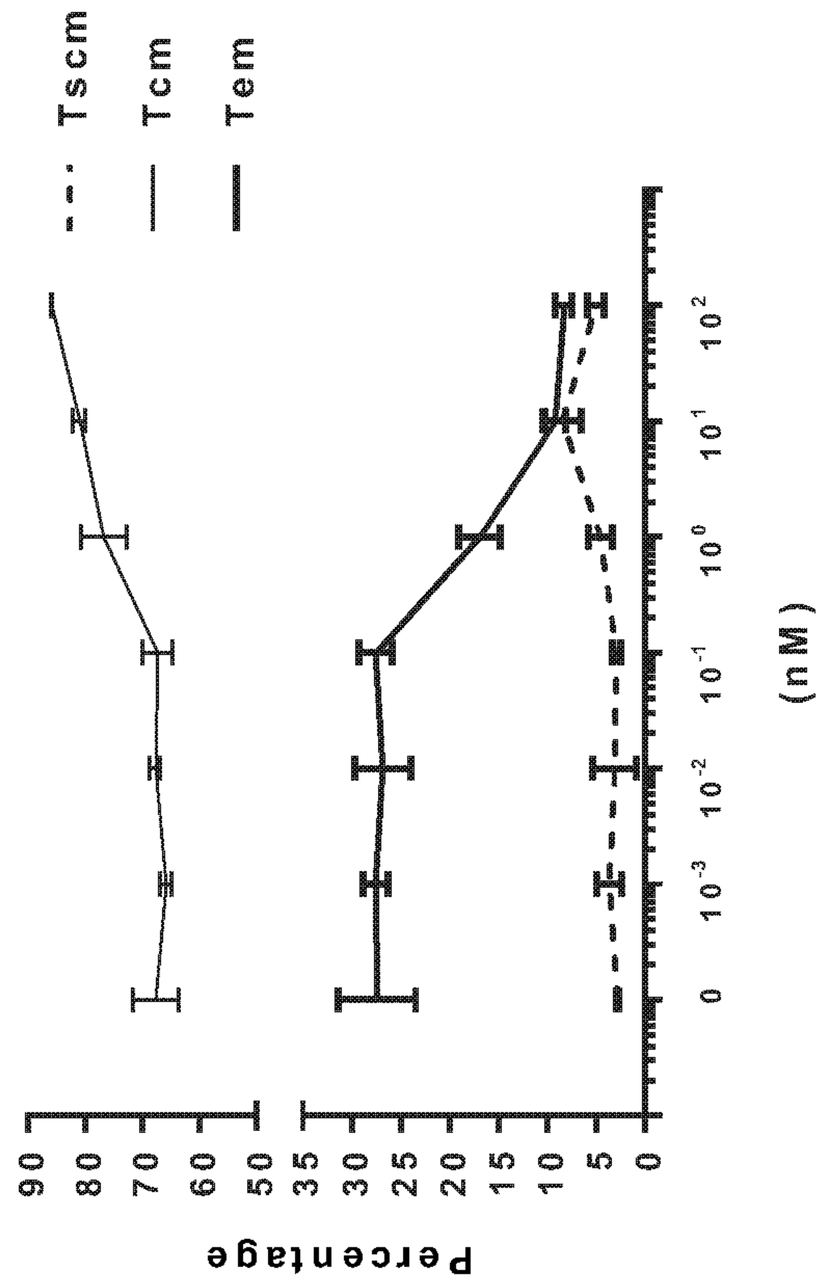
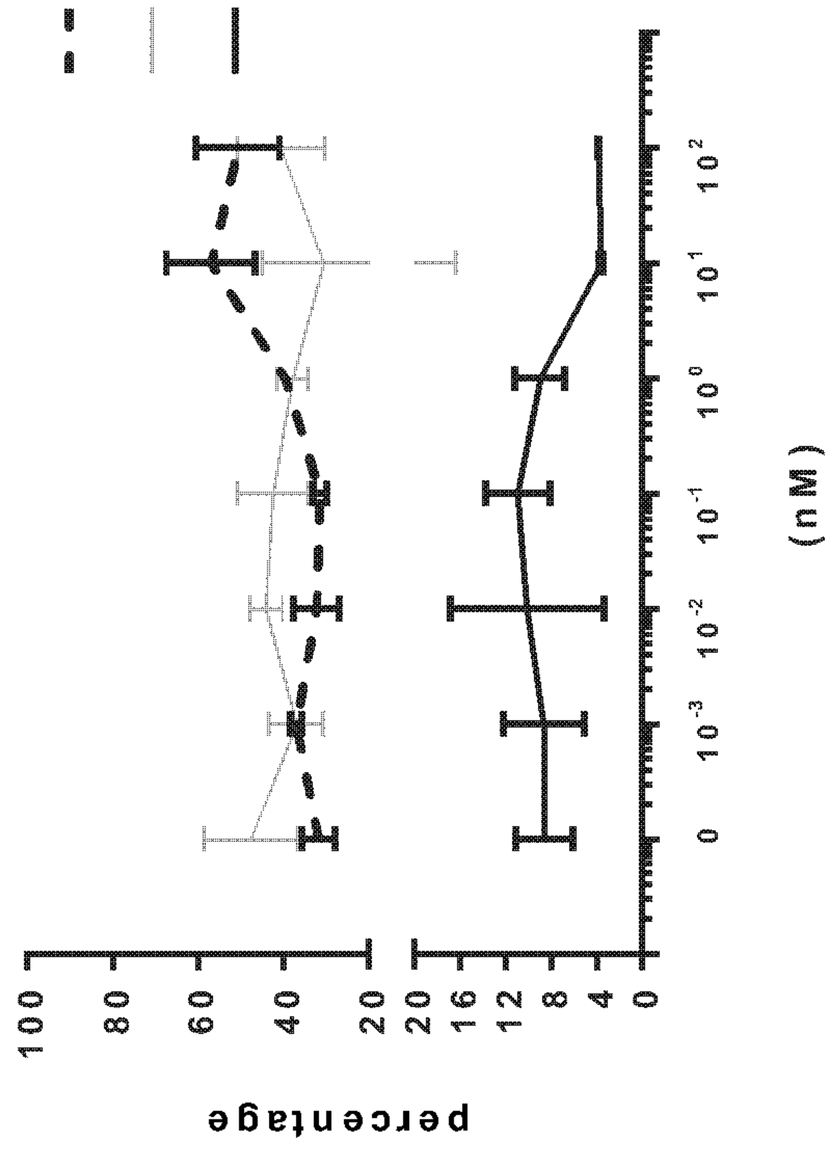

compound 93 (CD4+ T)
Tscm
Tcm
Tem
Percentage
100
80
60
40
35
30
25
20
15
10
5
0
0
10^{-3}
10^{-2}
10^{-1}
10^{0}
10^{1}
10^{2}
(nM)
compound 93 (CD8+ T)
Tscm
Tcm
Tem
Percentage
100
80
60
40
20
15
12
9
6
3
0
0
10^{-3}
10^{-2}
10^{-1}
10^{0}
10^{1}
10^{2}
(nM)
compound 93
Tscm/Tcm of CD8 T
Tscm/Tcm of CD4 T
ratio of Tscm/Tcm
2.5
2.0
1.5
1.0
0.5
0.0
0
10^{-3}
10^{-2}
10^{-1}
10^{0}
10^{1}
10^{2}
compound 93
Tscm/Tem of CD8 T
Tscm/Tem of CD4 T
ratio of Tscm/Tem
30
25
20
15
10
5
0
0
10^{-3}
10^{-2}
10^{-1}
10^{0}
10^{1}
10^{2}

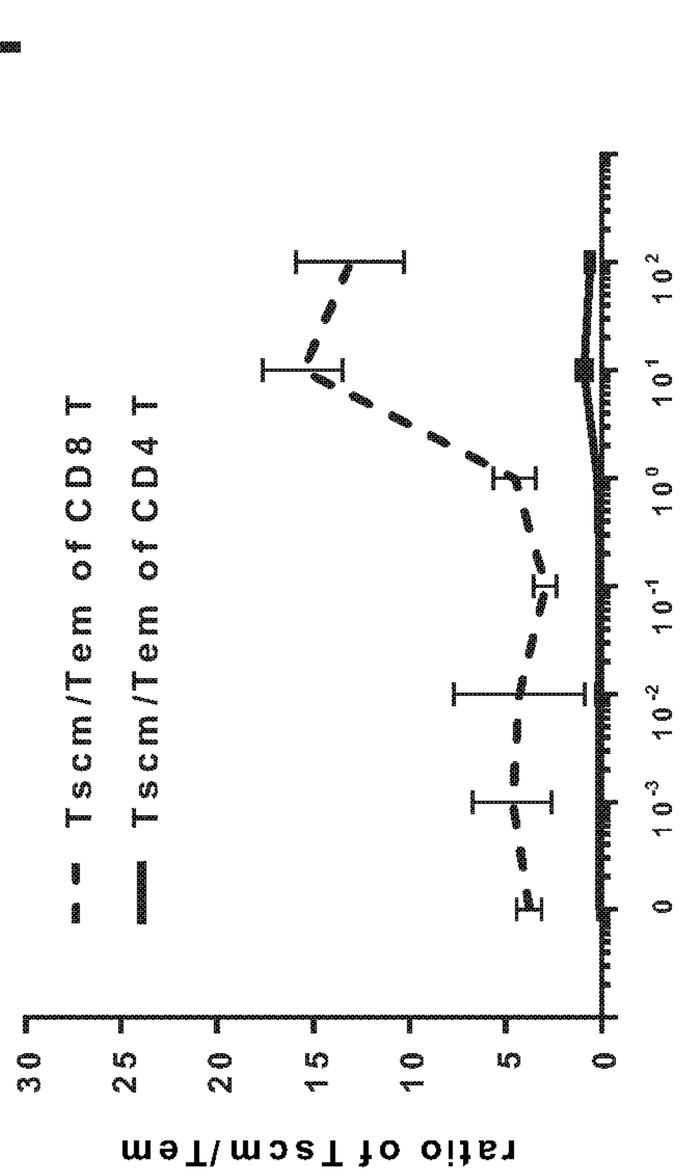
Figure 21

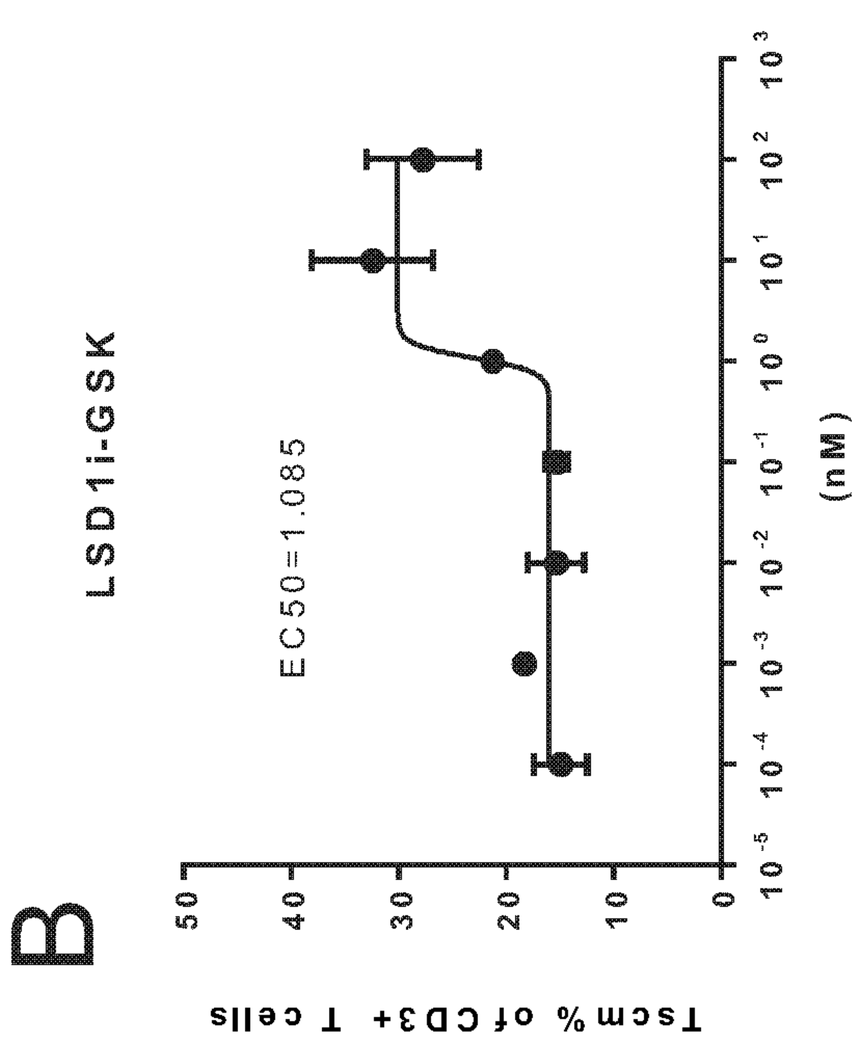

LSD1i-GSK (CD4+ T)
Tscm
Tcm
Tem
Percentage
(nM)

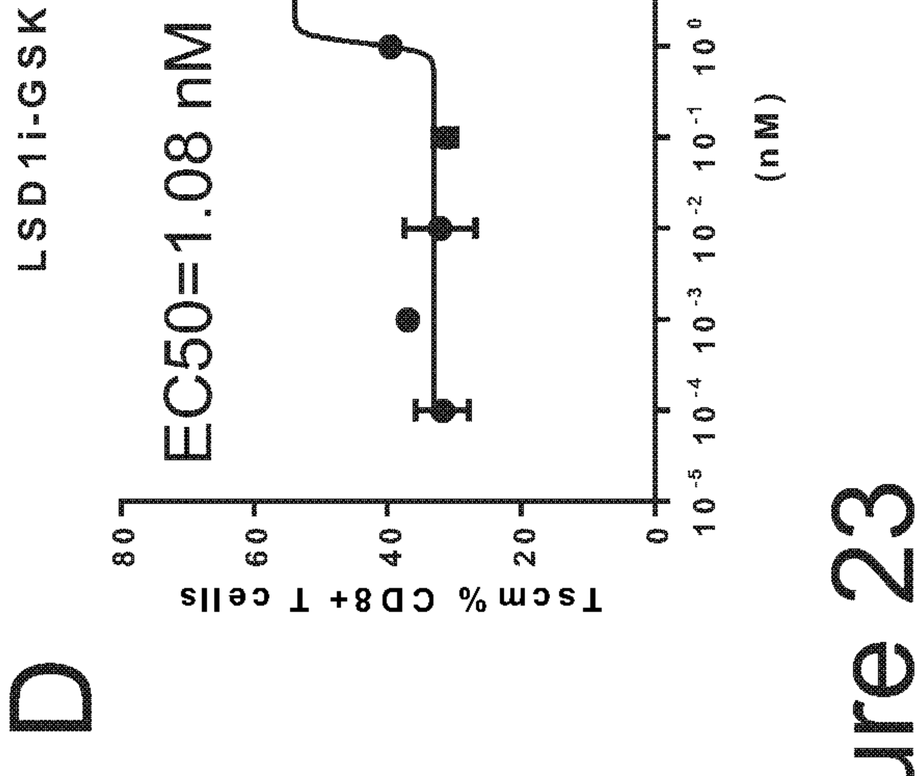

LSD1i-GSK
Tscm/Tcm of CD8 T
Tscm/Tcm of CD4 T
ratio of Tscm/Tcm
(nM)

Figure 22


LSD1i-GSK (CD8+ T)
Tscm
Tcm
Tem
percentage
(nM)


LSD1i-GSK
Tscm/Tem of CD8 T
Tscm/Tem of CD4 T
ratio of Tscm/Tem
(nM)


A
Tscm% of CD3+ T cells
50
40
30
20
10
0
10-5 10-4 10-3 10-2 10-1 100 101 102 103
(nM)
Compounf 93 (NVS compound 1)
EC50=2.59 nM

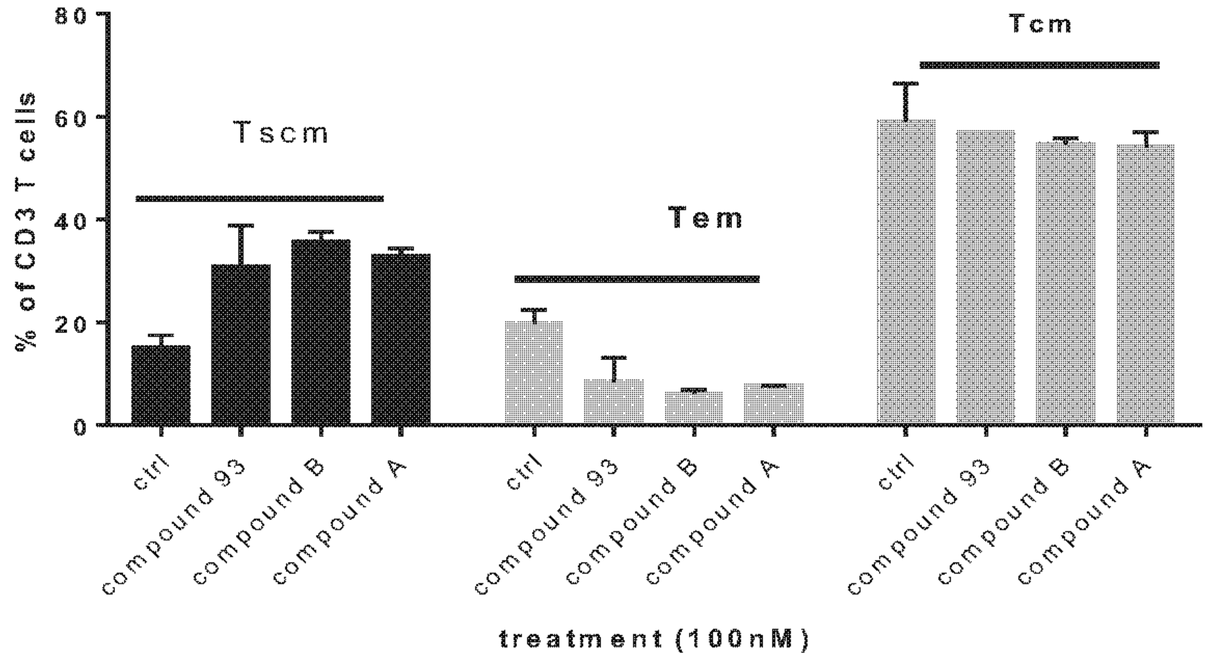

C
Tscm% of CD8+ T cells
80
60
40
20
0
10-5 10-4 10-3 10-2 10-1 100 101 102 103
(nM)
Compounf 93 (NVS compound 1)
EC50=3.25nM

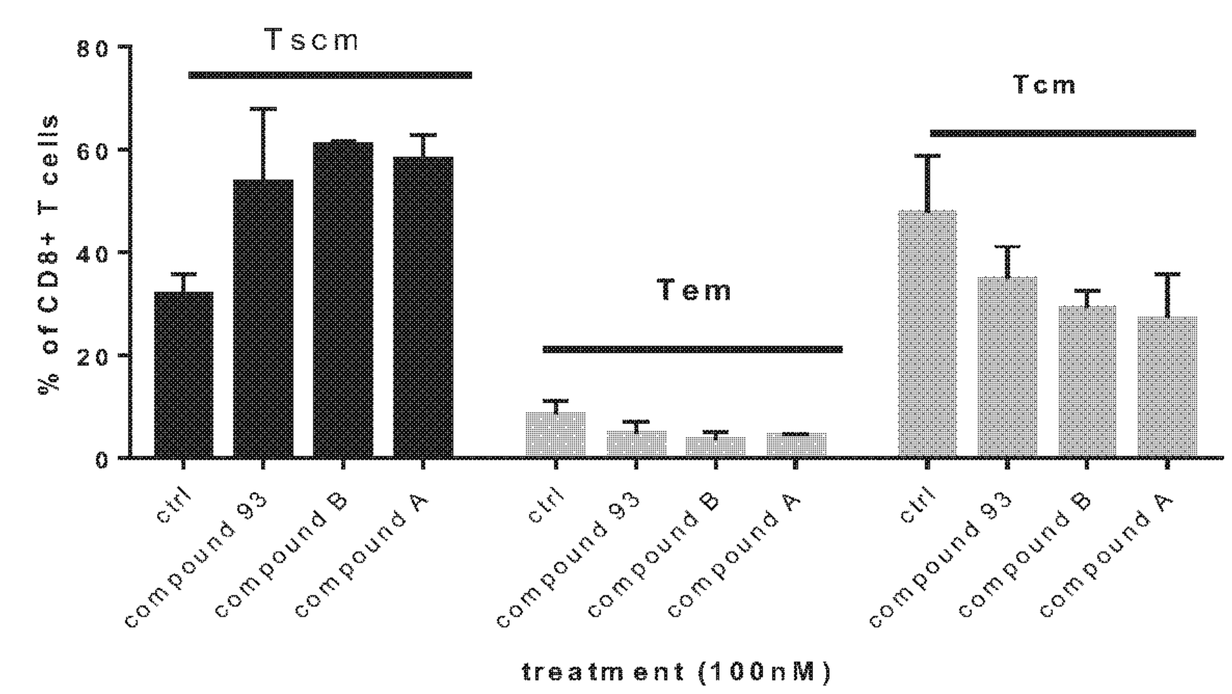
B
Tscm% of CD3+ T cells
50
40
30
20
10
0
10-5 10-4 10-3 10-2 10-1 100 101 102 103
(nM)
LSD1i-GSK
EC50=1.085

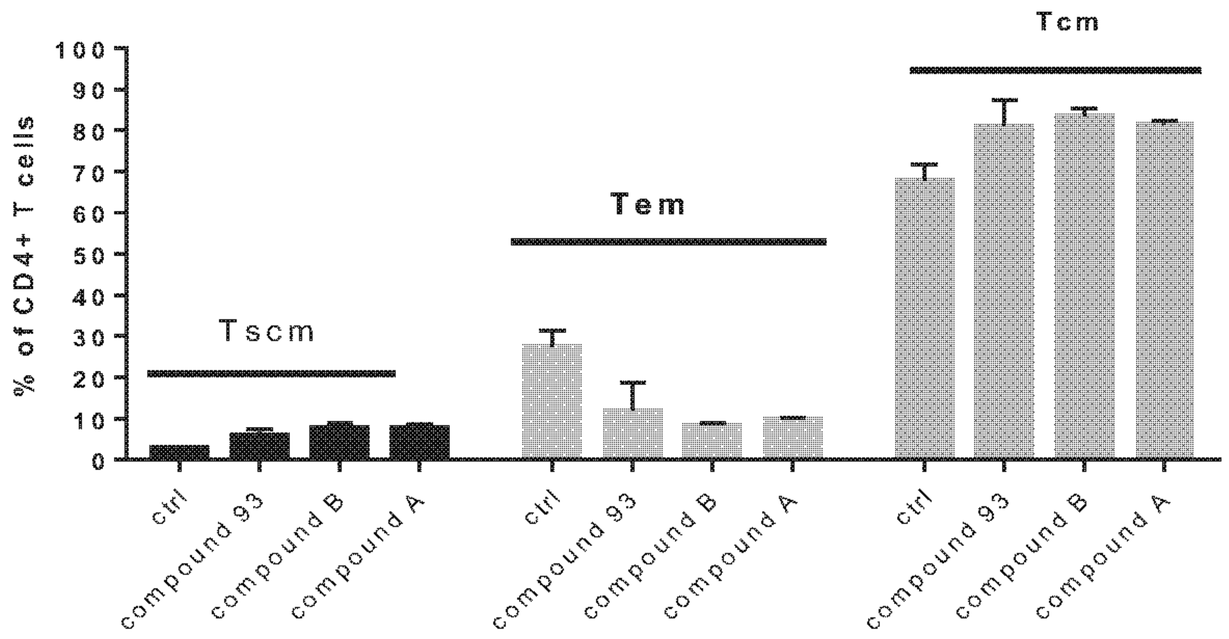
D
Tscm% CD8+ T cells
80
60
40
20
0
10-5 10-4 10-3 10-2 10-1 100 101 102 103
(nM)
LSD1i-GSK
EC50=1.08 nM

Figure 23

Figure 24


% of CD3 T cells
Tscm
Tem
Tcm
ctrl
compound 93
compound B
compound A
treatment (100nM)


% of CD8+ T cells
Tscm
Tem
Tcm
ctrl
compound 93
compound B
compound A
treatment (100nM)


% of CD4+ T cells
Tscm
Tem
Tcm
ctrl
compound 93
compound B
compound A
treatment (100nM)

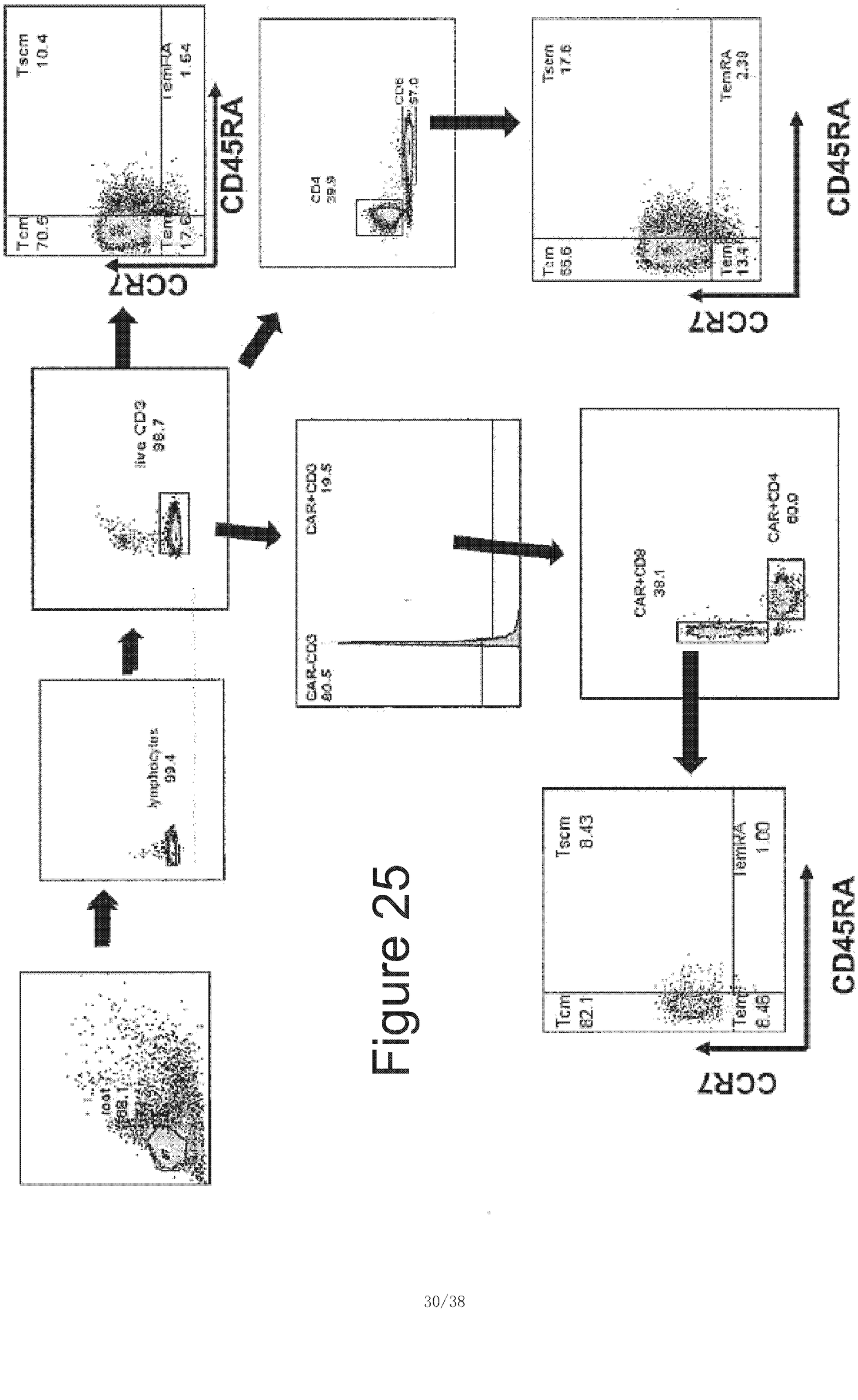
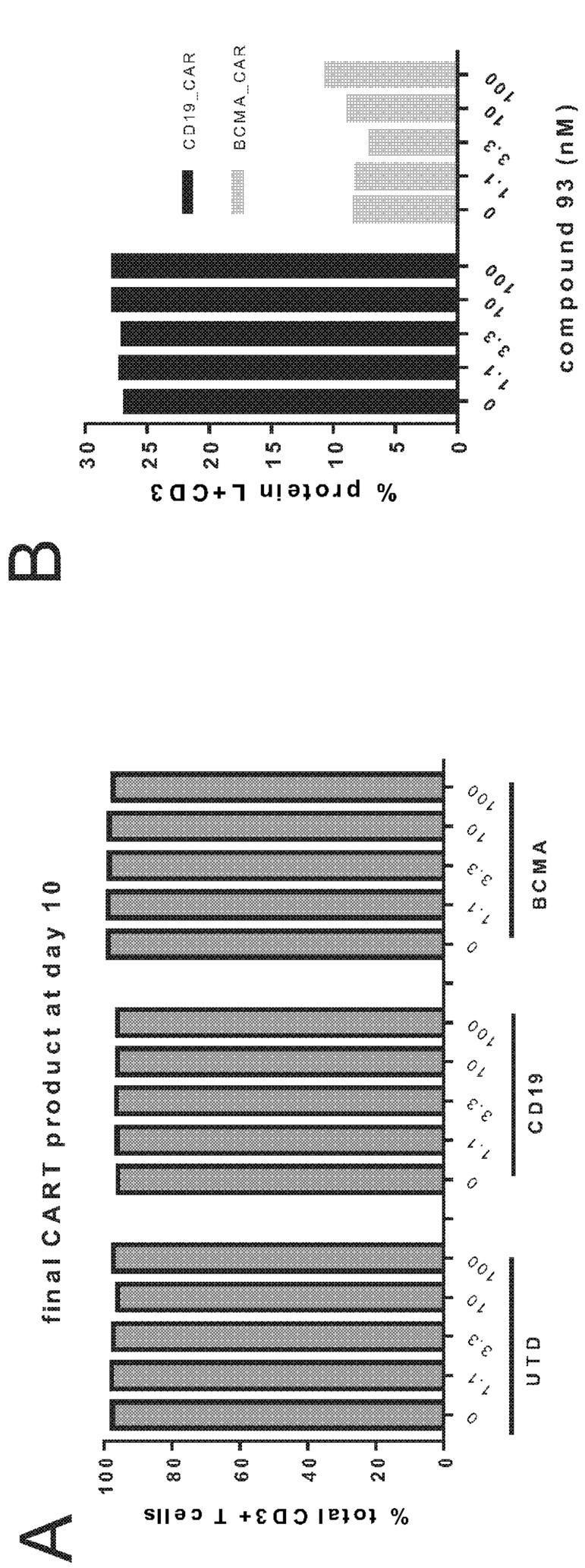
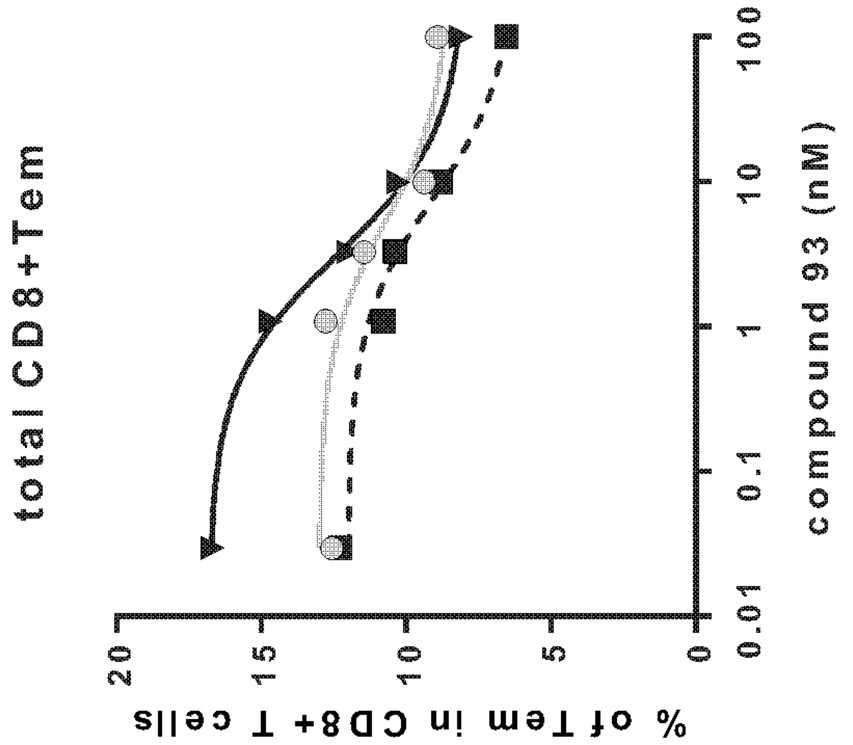
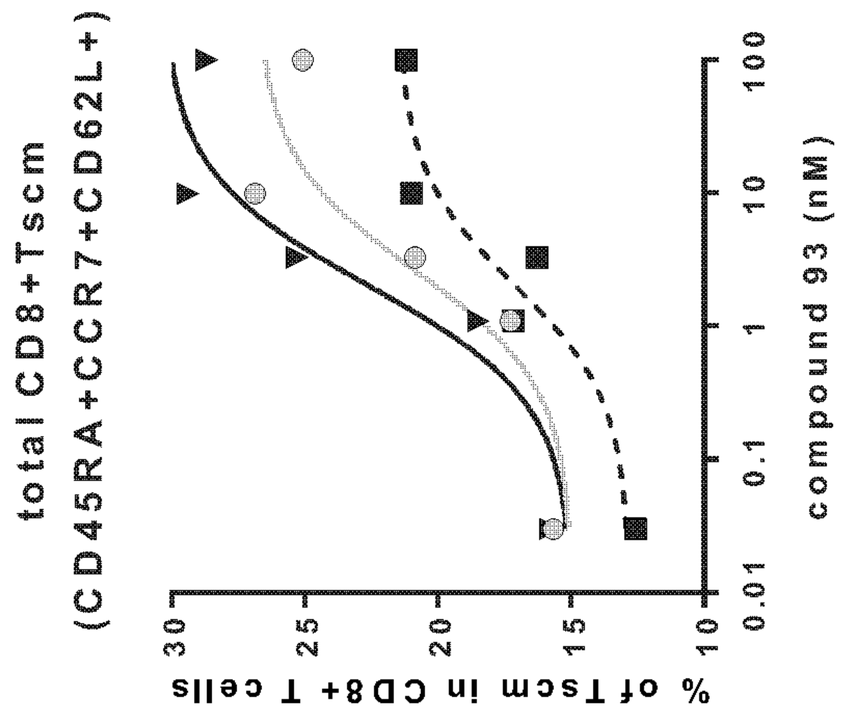

Tscm
10.4
TemRA
1.54
CD45RA
CCR7
Tcm
70.5
Tem
17.6
CD8
57.0
CD4
39.9
Tscm
17.6
TemRA
2.39
CD45RA
CCR7
Tcm
65.6
Tem
13.4
live CD3
98.7
CAR+CD3
19.5
CAR-CD3
80.5
CAR+CD4
60.9
CAR+CD8
38.1
lymphocytes
99.4
Tscm
8.43
TemRA
1.80
CD45RA
CCR7
Tcm
82.1
Tem
8.46
total
98.1

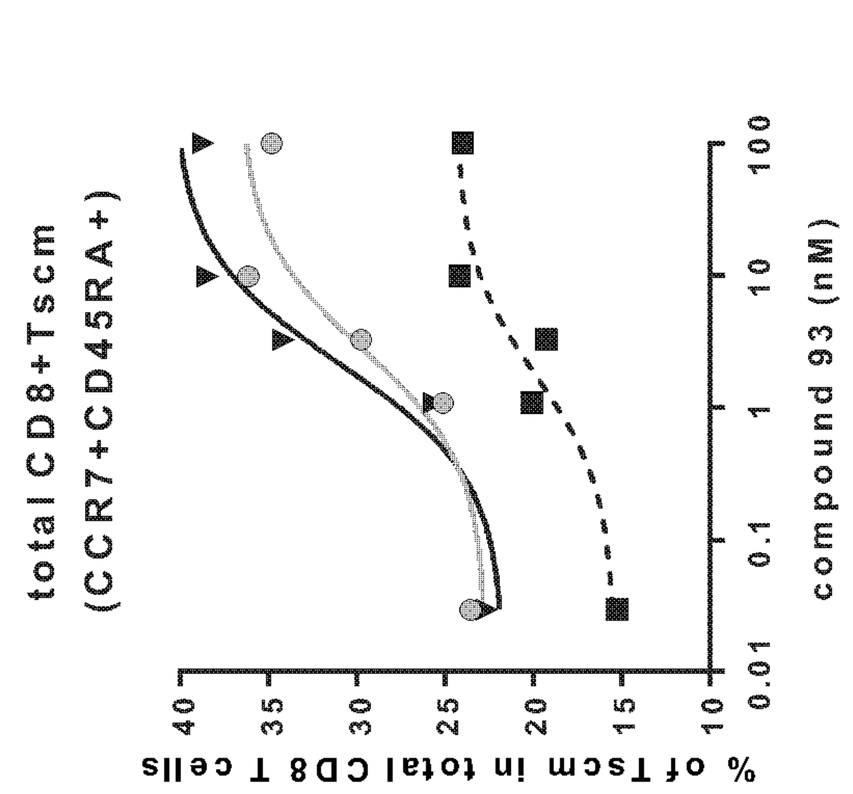
Figure 25

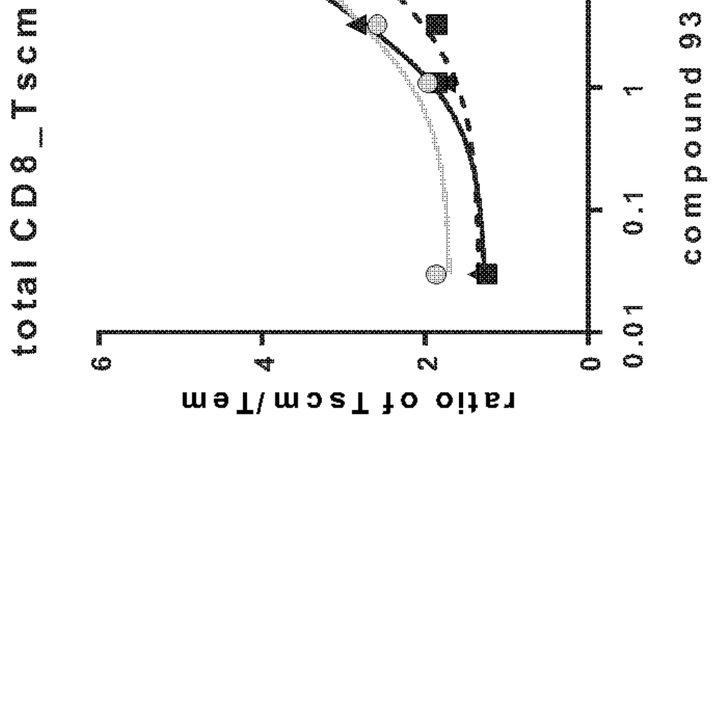
Figure 26


A
% total CD3+ T cells
100
80
60
40
20
0
final CART product at day 10
UTD
CD19
BCMA
0
1.1
3.3
10
100
treatment with compound 93 (nM)
B
% protein L+CD3
30
25
20
15
10
5
0
CD19_CAR
BCMA_CAR
compound 93 (nM)
C
CAR+ CD4 vs CD8
CD4_CAR+
CD8_CAR+
Percentage of protein L+
80
60
40
20
0
CD19
BCMA
compound 93 (nM)


% of Tem in CD8+ T cells
compound 93 (nM)
total CD8+Tem
UTD
CD19
BCMA


% of Tscm in CD8+ T cells
compound 93 (nM)
total CD8+Tscm
(CD45RA+CCR7+CD62L+)


% of Tscm in total CD8 T cells
compound 93 (nM)
total CD8+Tscm
(CCR7+CD45RA+)


ratio of Tscm/Tem
compound 93 (nM)
total CD8_Tscm vs Tem

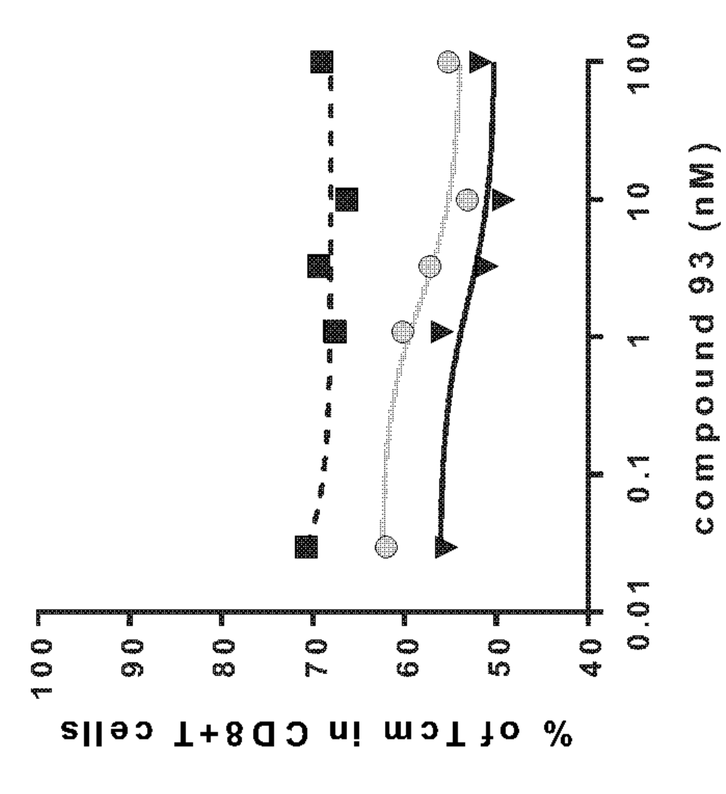

% of Tcm in CD8+T cells
compound 93 (nM)
total CD8+Tcm

Figure 27

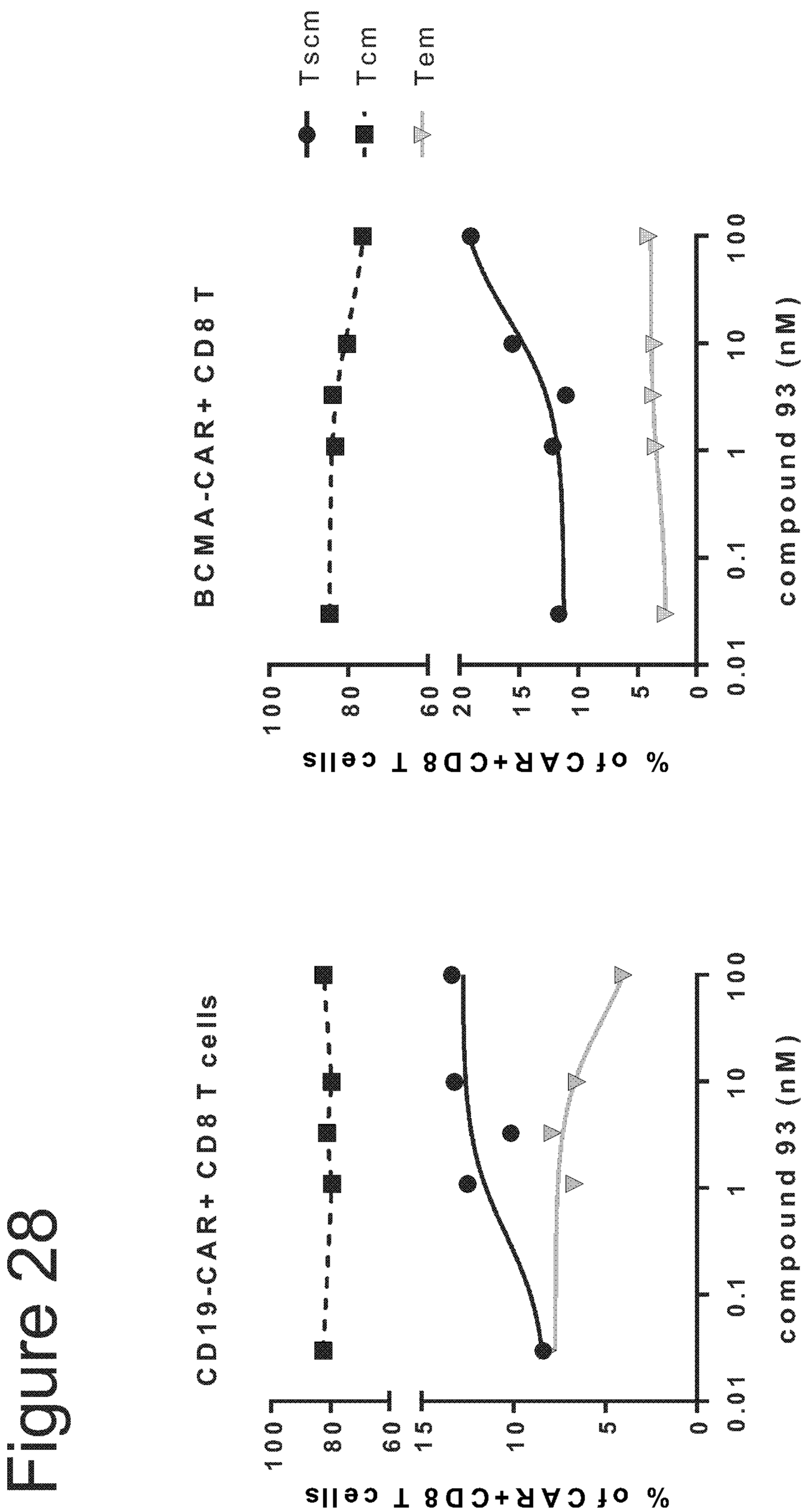

Tscm
Tcm
Tem
BCMA-CAR+ CD8 T
% of CAR+CD8 T cells
100
80
60
20
15
10
5
0
0.01
0.1
1
10
compound 93 (nM)
CD19-CAR+ CD8 T cells
% of CAR+CD8 T cells
100
80
60
15
10
5
0
0.01
0.1
1
10
100
compound 93 (nM)

Figure 28

Figure 29

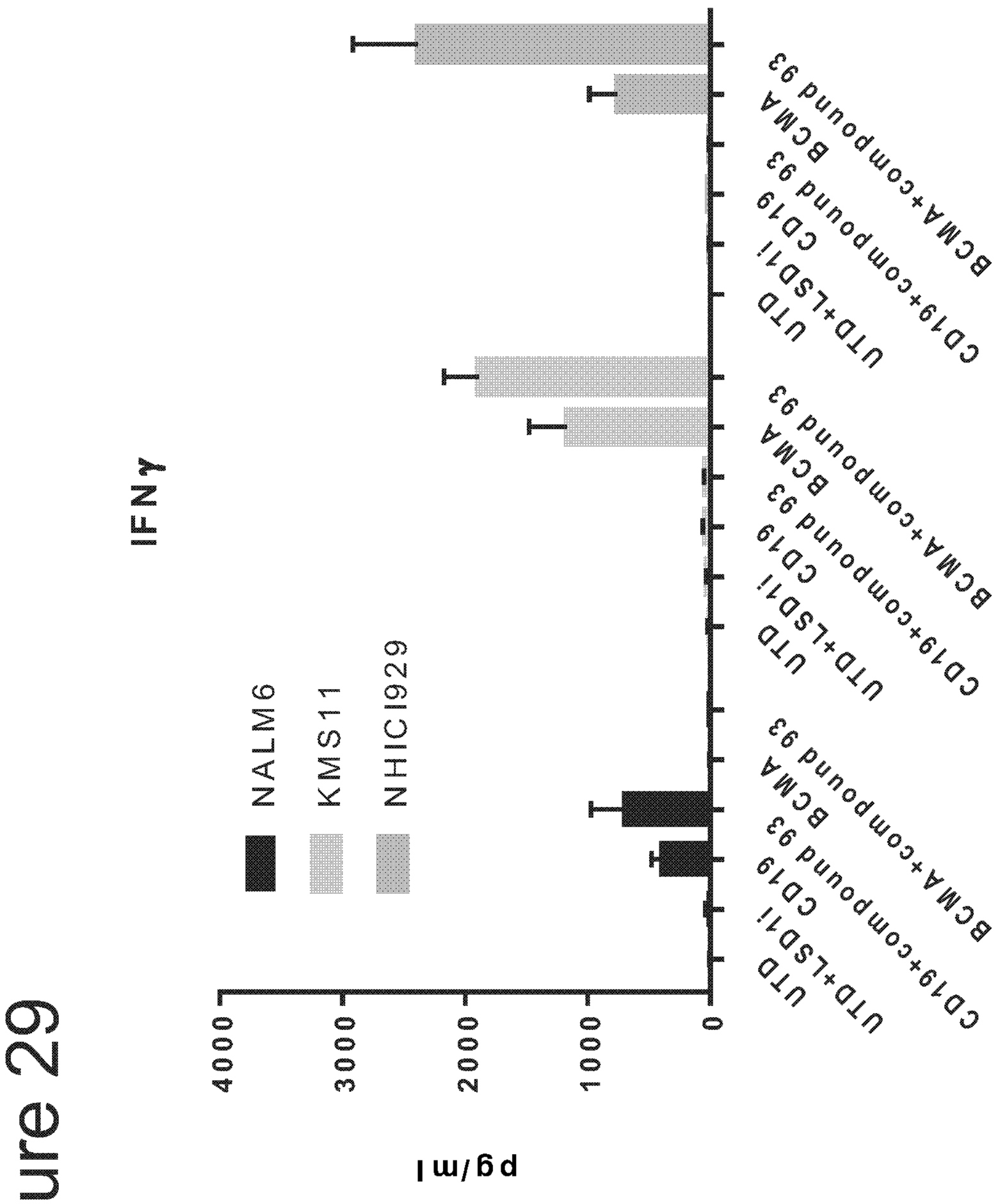

IFNγ
pg/ml
0
1000
2000
3000
4000
NALM6
KMS11
NHICI929
UTD
UTD+LSD1i
CD19+compound 93
BCMA+compound 93

Figure 30

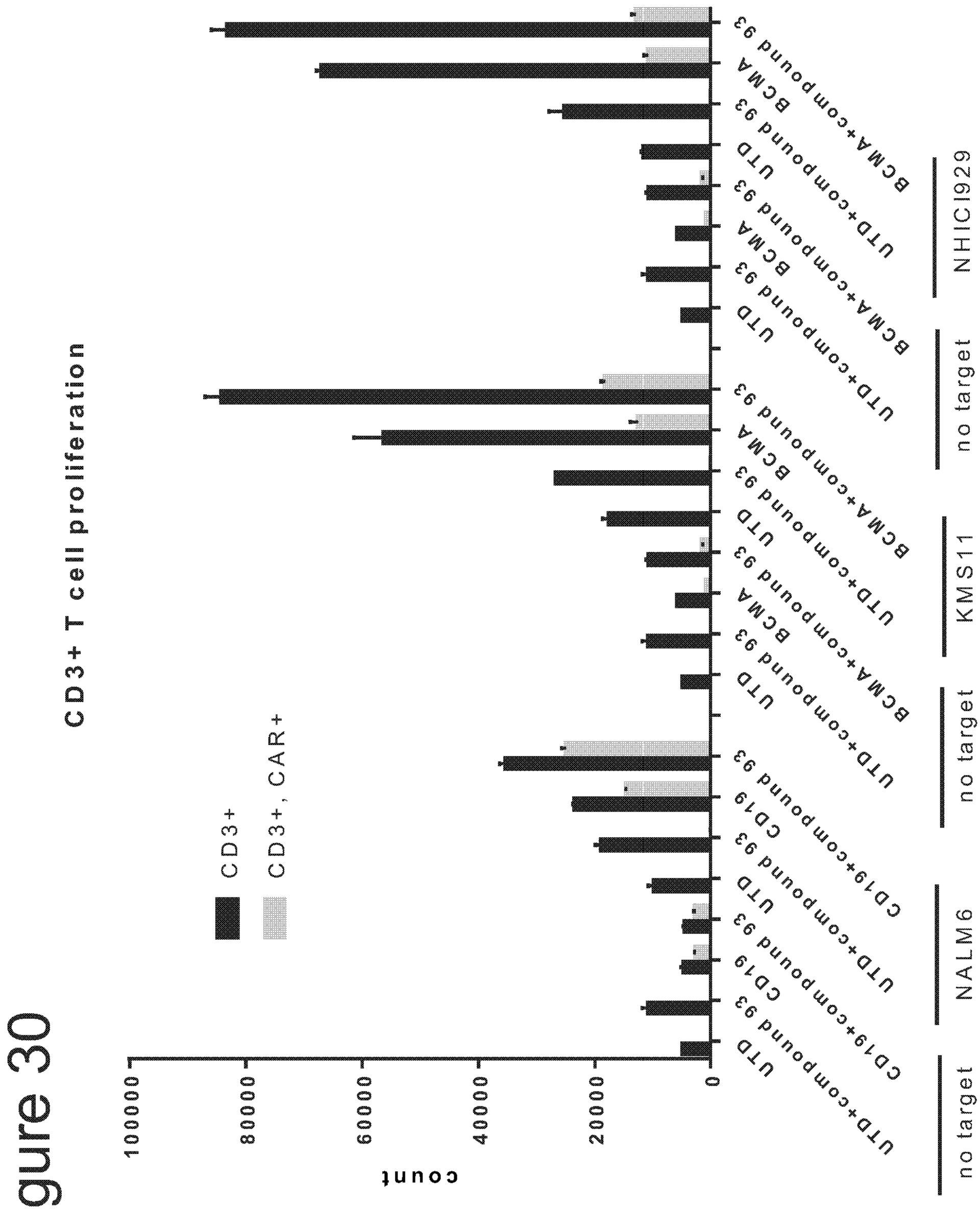

CD3+ T cell proliferation
CD3+
CD3+, CAR+
count
0
20000
40000
60000
80000
100000
UTD
UTD+compound 93
BCMA
BCMA+compound 93
UTD
UTD+compound 93
BCMA
BCMA+compound 93
UTD
UTD+compound 93
CD19
CD19+compound 93
UTD
UTD+compound 93
CD19
CD19+compound 93
UTD
UTD+compound 93
BCMA
BCMA+compound 93
UTD
UTD+compound 93
BCMA
BCMA+compound 93
no target
NALM6
no target
KMS11
no target
NHICI929

Figure 31

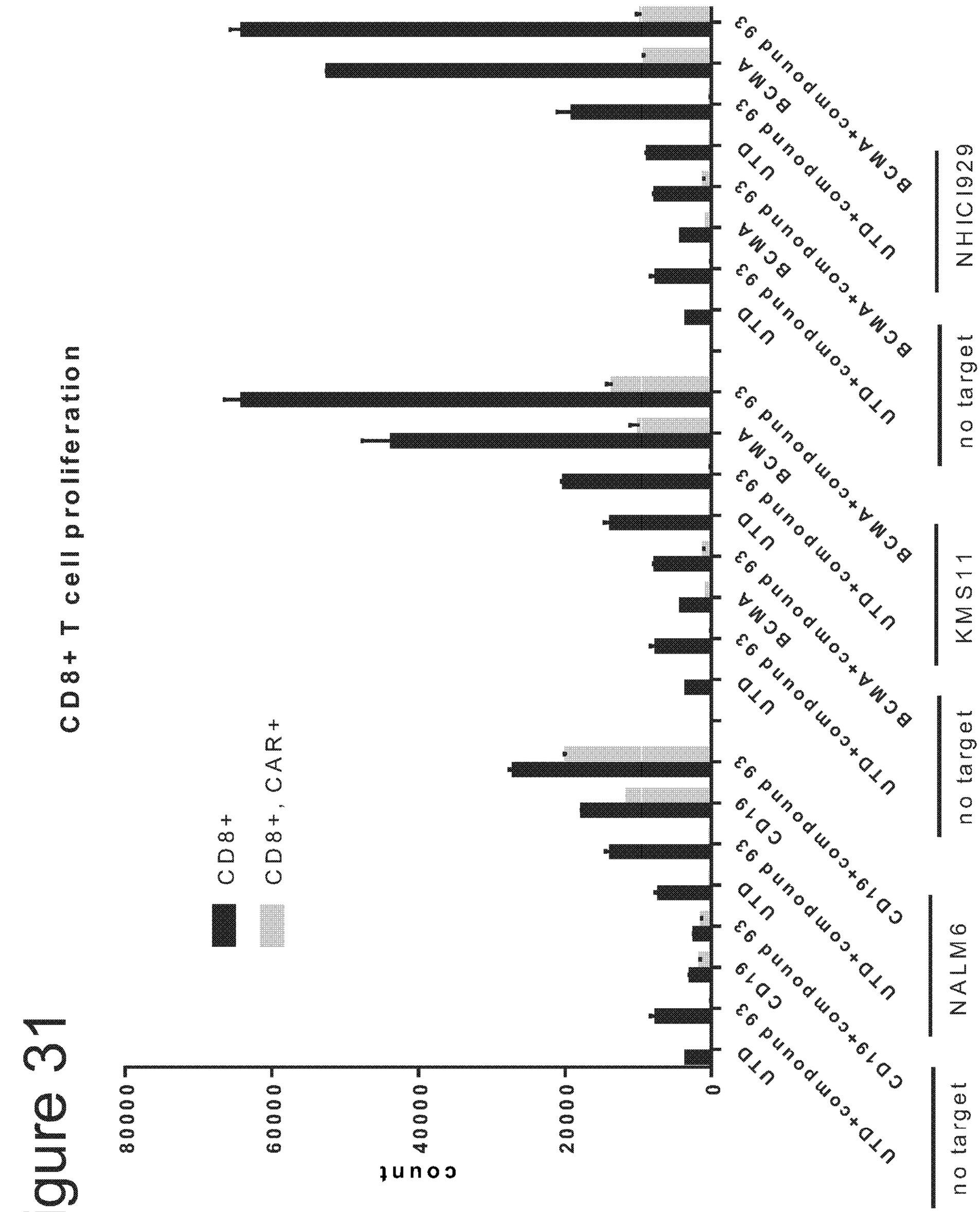
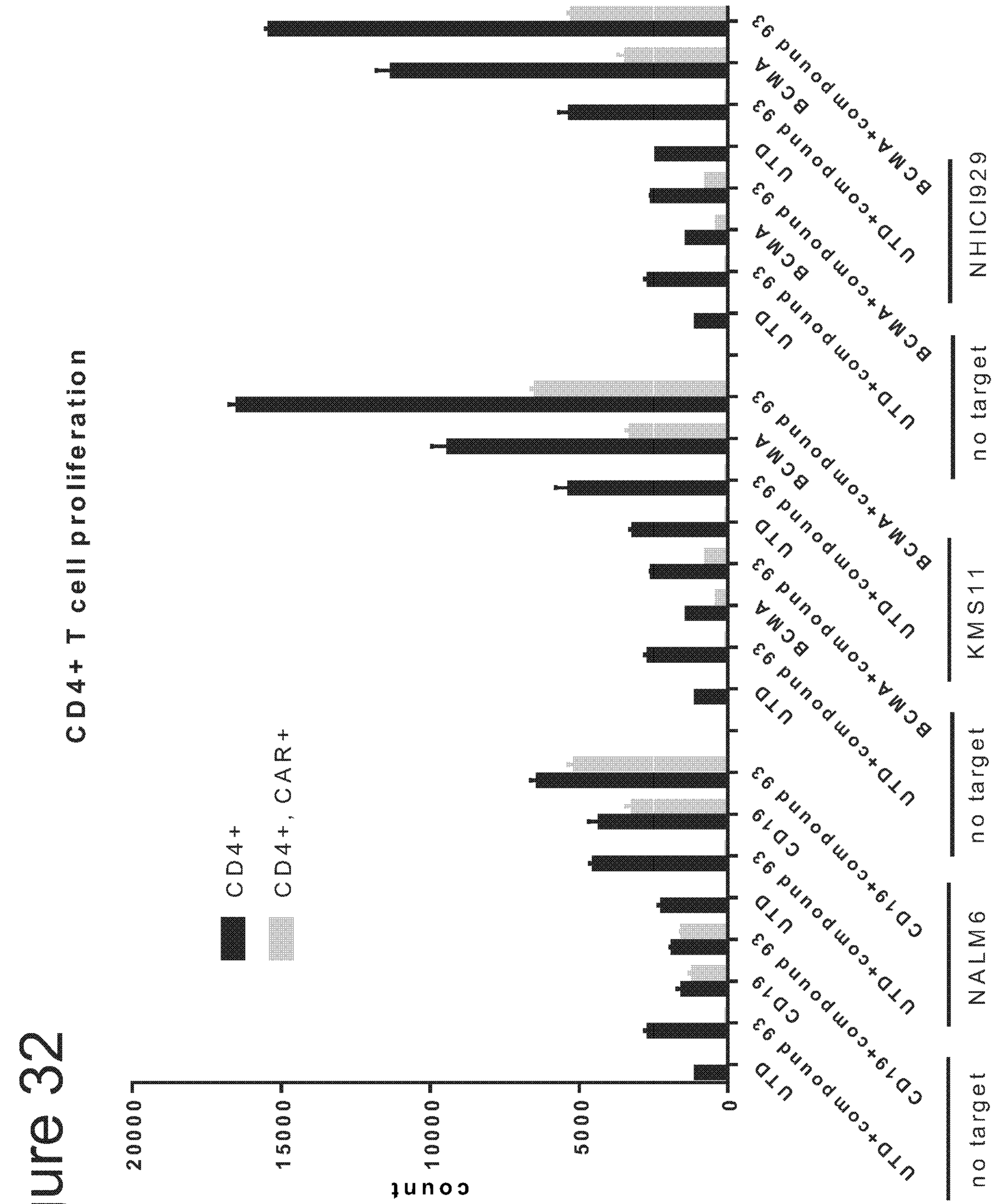

CD8+ T cell proliferation
CD8+
CD8+, CAR+
count
8000
6000
4000
2000
0
UTD
UTD+compound 93
BCMA
BCMA+compound 93
UTD
UTD+compound 93
CD19
CD19+compound 93
UTD
UTD+compound 93
BCMA
BCMA+compound 93
UTD
UTD+compound 93
BCMA
BCMA+compound 93
UTD
UTD+compound 93
CD19
CD19+compound 93
UTD
UTD+compound 93
CD19
CD19+compound 93
UTD
UTD+compound 93
BCMA
BCMA+compound 93
UTD
UTD+compound 93
BCMA
BCMA+compound 93
no target
NALM6
no target
KMS11
no target
NHICI929

Figure 32


CD4+ T cell proliferation
count
20000
15000
10000
5000
0
CD4+
CD4+, CAR+
UTD
UTD+compound 93
BCMA+compound 93
BCMA
CD19+compound 93
CD19
no target
NALM6
KMS11
NHICI929

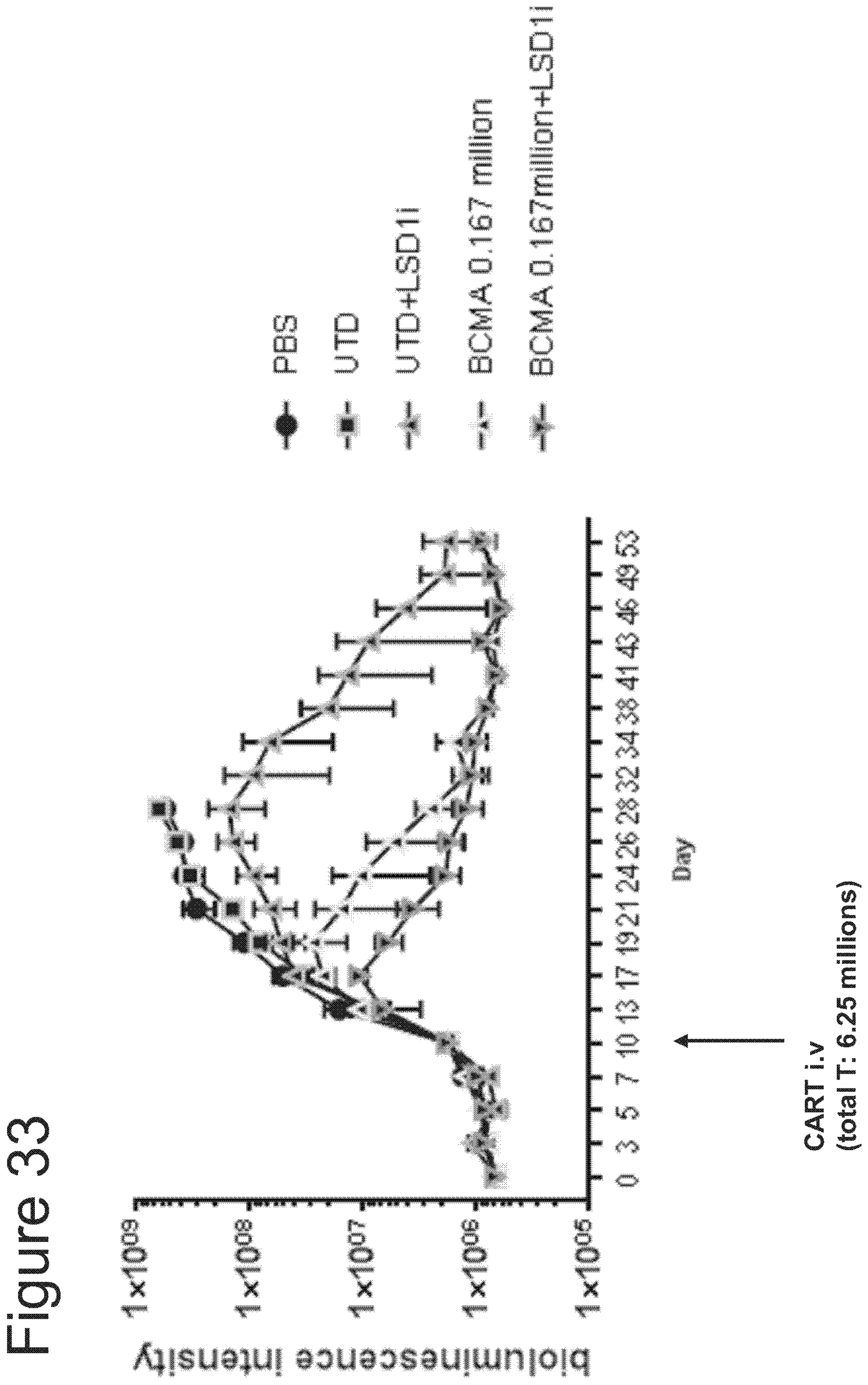

PBS
UTD
UTD+LSD1i
BCMA 0.1 million
BCMA 0.1 million+LSD1i
biolumineseence intensity
1x10^9
1x10^8
1x10^7
1x10^6
1x10^5
0 3 5 7 10 13 17 19 21 24 26 28 32 34 38 41 43 46 49 53
Day
CART i.v
(total T: 6.25 millions)

Figure 33