COMBINATION OF A CELL THERAPY AND A GAMMA SECRETASE INHIBITOR

Cross-Reference to Related Applications

[0001] This application claims the benefit of priority to U.S. provisional patent application 62/582,308, entitled "COMBINATION OF A CELL THERAPY AND A GAMMA SECRETASE INHIBITOR" filed on November 6, 2017; U.S. provisional patent application 62/582,937, entitled "COMBINATION OF A CELL THERAPY AND A GAMMA SECRETASE INHIBITOR" filed on November 7, 2017; and U.S. provisional patent application 62/665,450, entitled "COMBINATION OF A CELL THERAPY AND A GAMMA SECRETASE INHIBITOR" filed on May 1, 2018, the contents of which are incorporated by reference in their entirety.

Incorporation by Reference of Sequence Listing

[0002] The present application is being filed along with a Sequence Listing in electronic format. The Sequence Listing is provided as a file entitled 735042014240SeqList.TXT, created November 6, 2018, which is 320,126 bytes in size. The information in the electronic format of the Sequence Listing is incorporated by reference in its entirety.

Field

[0003] The present disclosure relates in some aspects to methods, compositions and uses involving immunotherapies, such as adoptive cell therapy, e.g., T cell therapy, and a gamma secretase inhibitor. The provided methods, compositions and uses include those for combination therapies involving the administration or use of a gamma secretase inhibitor in conjunction with a T cell therapy, such as a genetically engineered T cell therapy involving cells engineered with a recombinant receptor, such as chimeric antigen receptor (CAR)-expressing T cells, such as anti-BCMA chimeric antigen receptor (CAR)-expressing T cells. Also provided are compositions, methods of administration to subjects, articles of manufacture and kits for use in the methods. In some aspects, features of the methods and cells provide for increased or improved activity, efficacy, persistence, expansion and/or proliferation of T cells for adoptive cell therapy or endogenous T cells recruited by immunotherapeutic agents.

Background

[0004] Various strategies are available for immunotherapy, for example administering engineered T cells for adoptive therapy. For example, strategies are available for engineering T cells expressing genetically engineered antigen receptors, such as CARs, and administering compositions containing such cells to subjects. Improved strategies are needed to improve efficacy of the cells, for example, improving the persistence, activity and/or proliferation of the cells upon administration to subjects. Provided are methods, compositions, kits, and systems that meet such needs.

Summary

[0005] Provided herein are combination therapies involving administration of an immunotherapy involving a cell therapy, such as a T cell therapy, and an inhibitor of gamma secretase. In some aspects, the provided methods of treating a subject having a disease or disorder involve administration of an immunotherapy or immunotherapeutic agent, such as a composition including cells for adoptive cell therapy, e.g., such as a T cell therapy (e.g. CAR-expressing T cells), and administration of a gamma secretase inhibitor. In some embodiments, the cell therapy, such as CAR-expressing T cells, comprises an antigen-binding domain that binds to a B Cell Maturation Antigen (BCMA), such as surface BCMA. In some embodiments, the method further comprises selecting a subject for treatment, wherein the subject has a low expression of surface BCMA on a cell (such as a tumor/cancer cell). In some embodiments, the cell therapy comprises a recombinant receptor, such as a receptor that does not bind to BCMA. In some embodiments, the gamma secretase inhibitor prohibits intramembrane cleavage of a receptor on a cell (such as a tumor/cancer cell), wherein the cell therapy specifically targets the receptor. In some embodiments, the combination therapy generally involves administration of a gamma secretase inhibitor and administration of a cell therapy, such as a composition including cells for adoptive cell therapy, e.g., such as a T cell therapy (e.g. CAR-expressing T cells), wherein the gamma secretase administration is subsequent to the administration of cell therapy. In some embodiments, the therapy modulates activity of a cell therapy.

[0006] Provided herein are methods of treatment that involve: (a) administering a cell therapy to a subject having a disease or disorder, said cell therapy comprising a dose of genetically engineered cells expressing a chimeric antigen receptor (CAR) comprising an

antigen-binding domain that specifically binds to surface B cell maturation antigen (BCMA); and (b) administering to the subject an inhibitor of gamma secretase, wherein binding of the CAR to surface BCMA, or a measure indicative of function or activity following exposure to cells expressing surface BCMA, is not reduced or blocked or is not substantially reduced or blocked in the presence of a soluble or shed form of the BCMA. In some embodiments, the binding of the CAR to surface BCMA, or a measure indicative of function or activity following exposure to cells expressing surface BCMA, is not reduced or blocked or is not substantially reduced or blocked in the presence of a soluble or shed form of the BCMA at a concentration or amount of the soluble or shed form of the BCMA corresponding to a concentration or amount present in serum or blood or plasma of the subject or of a multiple myeloma patient or on average in a patient population for the disease or disorder, or at a concentration or amount of the soluble or shed BCMA at which the binding or measure is reduced or blocked, or is substantially reduced or blocked, for cells expressing a reference anti-BCMA CAR in the same assay.

[0007] Provided here are methods of treatment that involve: administering a cell therapy to a subject having a disease or disorder, said cell therapy comprising a dose of genetically engineered cells expressing a chimeric antigen receptor comprising an antigen-binding domain that binds to surface B cell maturation antigen (BCMA), wherein binding of the CAR to surface BCMA or a measure indicative of function or activity, following exposure to cells expressing surface BCMA, is not reduced or blocked or is not substantially reduced or blocked in the presence of a soluble or shed form of the BCMA, optionally at a concentration or amount of the soluble or shed form of the BCMA corresponding to a concentration or amount present in serum or blood or plasma of the subject or of a multiple myeloma patient, or on average in a patient population for the disease or disorder, or at a concentration or amount of the soluble or shed BCMA at which the binding or measure is reduced or blocked, or is substantially reduced or blocked, for cells expressing a reference anti-BCMA CAR in the same assay, wherein, at the time of initiation of the administration of the cell therapy, the subject has been previously administered, and/or is undergoing treatment with, an inhibitor of gamma secretase.

[0008] Provided here are methods of treatment that involve: administering an inhibitor of gamma secretase to a subject having a disease or disorder, wherein, at the time of initiation of the administration of the inhibitor, the subject has been previously administered, and/or is undergoing treatment with, a cell therapy, said cell therapy comprising a dose of genetically

engineered cells expressing a chimeric antigen receptor comprising an antigen-binding domain that specifically binds surface B cell maturation antigen (BCMA), wherein binding of the CAR to surface BCMA or a measure indicative of function or activity, following exposure to cells expressing surface BCMA, is not reduced or blocked or is not substantially reduced or blocked in the presence of a soluble or shed form of the BCMA, optionally at a concentration or amount of the soluble or shed form of the BCMA corresponding to a concentration or amount present in serum or blood or plasma of the subject or of a multiple myeloma patient, or on average in a patient population for the disease or disorder, or at a concentration or amount of the soluble or shed BCMA at which the binding or measure is reduced or blocked, or is substantially reduced or blocked, for cells expressing a reference anti-BCMA CAR in the same assay.

[0009] Provided here are methods of treatment that involve: (a) selecting a subject having a cancer in which cells of the cancer in the subject (i) express surface CD138, surface CD38 or a surface plasma cell marker or are derived from plasma cells and (ii) comprise low expression of surface B cell maturation antigen (BCMA) and/or a level of expression of surface BCMA below a threshold level; (b) administering a cell therapy to the subject selected in (a), said cell therapy comprising a dose of genetically engineered cells expressing a chimeric antigen receptor comprising an antigen-binding domain that binds to B cell maturation antigen (BCMA); and (c) administering to the subject an inhibitor of gamma secretase.

[0010] Provided here are methods of treatment that involve: (a) administering an inhibitor of gamma secretase to a subject having a cancer, said subject selected as having plasma cells comprising low expression of surface B cell maturation antigen (BCMA) and/or a level of expression of surface BCMA below a threshold level; and (b) administering to the subject a cell therapy for treating the cancer, said cell therapy comprising a dose of genetically engineered cells expressing a chimeric antigen receptor comprising an antigen-binding domain that specifically binds to B cell maturation antigen (BCMA).

[0011] In some embodiments of any one of the methods provided herein, the threshold level of expression of surface BCMA is lower than the average or median expression or level of surface BCMA on plasma cells in a plurality of control subjects, optionally wherein the control subjects are a group of healthy or normal subjects.

[0012] In some embodiments of any one of the methods provided herein, the low expression of surface BCMA is present when less than or less than about 60%, less than or less than about

50%, less than or less than about 40%, less than or less than about 30%, less than or less than about 20% or less than or less than about 10% of the plasma cells, or cells with a plasma cell marker or phenotype or the cancer cells, in the subject express surface BCMA; or the threshold level of surface BCMA is less than or less than about 60%, less than or less than about 50%, less than or less than about 40%, less than or less than about 30%, less than or less than about 20% or less than or less than about 10% of the plasma cells, or cells with a plasma cell marker or phenotype or the cancer cells, in the subject that express surface BCMA.

[0013] In some embodiments of any one of the methods provided herein, the expression of surface BCMA is determined by flow cytometry and/or an immunoassay.

[0014] In some embodiments of any one of the embodiments provided herein, binding of the CAR to surface BCMA or a measure indicative of function or activity, following exposure to cells expressing surface BCMA, is not reduced or blocked or is not substantially reduced or blocked in the presence of a soluble or shed form of the BCMA, optionally at a concentration or amount of the soluble or shed form of the BCMA corresponding to a concentration or amount present in serum or blood or plasma of the subject or of a multiple myeloma patient, or on average in a patient population for the disease or disorder, or at a concentration or amount of the soluble or shed BCMA at which the binding or measure is reduced or blocked, or is substantially reduced or blocked, for cells expressing a reference anti-BCMA CAR in the same assay.

[0015] In some embodiments of any one of the methods provided herein, the antigen-binding domain does not bind soluble BCMA or binds to soluble BCMA with an affinity lower than the affinity of said antigen-binding domain binding to surface BCMA. In some embodiments, the affinity of the antigen-binding domain to the surface B cell maturation antigen (BCMA) is at least 1-fold, 2-fold, 3-fold, 4-fold, 5-fold, 6-fold, 7-fold, 8-fold, 9-fold, or 10-fold higher than the affinity to the soluble BCMA.

[0016] In some of any of the provided embodiments, the anti-BCMA CAR is an anti-BCMA CAR as described herein. Among a provided anti-BCMA CAR for use in the methods and uses herein is a CAR in which the antibody or antigen-binding fragment contains a V_H region comprising the sequence set forth in SEQ ID NO: 24 or an amino acid sequence having at least at or about 90%, at or about 91%, at or about 92%, at or about 93%, at or about 94%, at or about 95%, at or about 96%, at or about 97%, at or about 98%, or at or about 99% identity to SEQ ID NO:24; and contains a V_L region comprising the sequence set forth in SEQ ID NO:25 or an

amino acid sequence having at least at or about 90%, at or about 91%, at or about 92%, at or about 93%, at or about 94%, at or about 95%, at or about 96%, at or about 97%, at or about 98%, or at or about 99% identity to SEQ ID NO:25. In some embodiments, the antibody or antigen-binding fragment of the provided CAR contains a V_H region that has a CDRH1, a CDRH2 and a CDRH3 comprising the amino acid sequence of SEQ ID NOS: 173, 174 and 175, respectively and a V_L region that has a CDRL1, a CDRL2 and a CDRL3 comprising the amino acid sequence of SEQ ID NOS: 183, 184 and 185, respectively. In some embodiments, the antibody or antigen-binding fragment of the provided CAR contains a V_H region that has a CDRH1, a CDRH2 and a CDRH3 comprising the amino acid sequence of SEQ ID NOS: 176, 177 and 175, respectively and a V_L region that has a CDRL1, a CDRL2 and a CDRL3 comprising the amino acid sequence of SEQ ID NOS: 183, 184 and 185, respectively. In some embodiments, the antibody or antigen-binding fragment of the provided CAR contains a V_H region that has a CDRH1, a CDRH2 and a CDRH3 comprising the amino acid sequence of SEQ ID NOS: 178, 179 and 175, respectively and a V_L region that has a CDRL1, a CDRL2 and a CDRL3 comprising the amino acid sequence of SEQ ID NOS: 183, 184 and 185, respectively. In some embodiments, the antibody or antigen-binding fragment of the provided CAR contains a V_H region that has a CDRH1, a CDRH2 and a CDRH3 comprising the amino acid sequence of SEQ ID NOS: 180, 181 and 182, respectively and a V_L region that has a CDRL1, a CDRL2 and a CDRL3 comprising the amino acid sequence of SEQ ID NOS: 186, 187 and 185, respectively. In some embodiments, the V_H region comprises the sequence set forth in SEQ ID NO:24 and the V_L region comprises the sequence set forth in SEQ ID NO:25. In some embodiments, the antibody or antigen-binding fragment is a single-chain antibody fragment, such as an scFv. In some embodiments, the scFv comprises the sequence of amino acids set forth in SEQ ID NO:188 or a sequence of amino acids at least at or about 90%, at or about 91%, at or about 92%, at or about 93%, at or about 94%, at or about 95%, at or about 96%, at or about 97%, at or about 98%, or at or about 99% identity to SEQ ID NO:188. In some embodiments, the anti-BCMA CAR has the sequence of amino acids set forth in SEQ NO: 124 or a sequence of amino acids at least at or about 90%, at or about 91%, at or about 92%, at or about 93%, at or about 94%, at or about 95%, at or about 96%, at or about 97%, at or about 98%, or at or about 99% identity to SEQ ID NO:124. In some embodiments, the anti-BCMA CAR has the sequence of amino acids set forth in SEQ NO: 125 or a sequence of amino acids at least at or about 90%, at or about 91%,

at or about 92%, at or about 93%, at or about 94%, at or about 95%, at or about 96%, at or about 97%, at or about 98%, or at or about 99% identity to SEQ ID NO:125.

[0017] In some embodiments of any one of the methods provided herein, the gamma secretase inhibitor inhibits intramembrane cleavage of BCMA.

[0018] In some embodiments of any one of the methods provided herein, the gamma secretase inhibitor inhibits intramembrane cleavage of BCMA with a half-maximal inhibitory concentration (IC_{50}) of 0.01 nM to 10 nM, 0.01 nM to 5 nM, 0.01 nM to 1 nM, 0.01 nM to 0.5 nM, 0.01 nM to 0.35 nM, 0.01 nM to 0.25 nM, 0.01 nM to 0.1 nM, 0.01 nM to 0.05 nM, 0.05 nM to 10 nM, 0.05 nM to 5 nM, 0.05 nM to 1 nM, 0.05 nM to 0.5 nM, 0.05 nM to 0.35 nM, 0.05 nM to 0.25 nM, 0.05 nM to 0.1 nM, 0.1 nM to 10 nM, 0.1 nM to 5 nM, 0.1 nM to 1 nM, 0.1 nM, to 0.5 nM, 0.1 nM to 0.35 nM, 0.1 nM to 0.25 nM, 0.25 nM to 10 nM, 0.25 nM to 5 nM, 0.25 nM to 1 nM, 0.25 nM to 0.5 nM, 0.25 nM to 0.35 nM, 0.35 nM to 10 nM, 0.35 nM to 5 nM, 0.35 nM to 1 nM, 0.35 nM to 0.5 nM, 0.5 nM to 10 nM, 0.5 nM to 5 nM, 0.5 nM to 1 nM, 1 nM to 10 nM, 1 nM to 5 nM or 5 nM to 10 nM.

[0019] In some embodiments of any one of the methods provided herein, the gamma secretase inhibitor inhibits intramembrane cleavage of BCMA with a half-maximal inhibitory concentration (IC_{50}) of less than or less than about 10 nM, 5 nM, 1 nM, 0.5 nM, 0.35 nM, 0.25 nM, 0.1 nM, 0.05 nM or 0.01 nM or less.

[0020] Provided herein are methods of treatment that involve: (a) administering a cell therapy to a subject having a disease or disorder, said cell therapy comprising a dose of genetically engineered cells expressing a recombinant receptor, wherein the recombinant receptor does not specifically bind to surface B cell maturation antigen (BCMA); and (b) administering an inhibitor of gamma secretase.

[0021] Provided herein are methods of treatment that involve: administering a cell therapy to a subject having a disease or disorder, said cell therapy comprising a dose of genetically engineered cells expressing a recombinant receptor, wherein the recombinant receptor does not specifically bind to surface B cell maturation antigen (BCMA), wherein, at the time of initiation of the administration of the cell therapy, the subject has been previously administered, and/or is undergoing treatment with, an inhibitor of gamma secretase.

[0022] Provided herein are methods of treatment that involve: administering an inhibitor of gamma secretase to a subject having a disease or disorder, wherein, at the time of initiation of

the administration of the inhibitor, the subject has been previously administered, and/or is undergoing treatment with, a cell therapy, said cell therapy comprising a dose of genetically engineered cells expressing a recombinant receptor, wherein the recombinant receptor does not specifically bind to surface B cell maturation antigen (BCMA).

[0023] In some embodiments of any one of the methods provided herein, the recombinant receptor specifically binds to a target antigen associated with the disease or disorder.

[0024] In some embodiments of any one of the methods provided herein, the target antigen is selected from among, carbonic anhydrase 9 (CAIX), Her2/neu (receptor tyrosine kinase erbB2), L1-CAM, CD19, CD20, CD22, mesothelin, CEA, and hepatitis B surface antigen, anti-folate receptor, CD23, CD24, CD30, CD33, CD38, CD44, EGFR, epithelial glycoprotein 2 (EPG-2), epithelial glycoprotein 40 (EPG-40), EPHa2, erb-B2, erb-B3, erb-B4, erbB dimers, EGFR vIII, folate binding protein (FBP), FCRL5, FCRH5, fetal acetylcholine receptor, GD2, GD3, HMW-MAA, IL-22R-alpha, IL-13R-alpha2, kinase insert domain receptor (kdr), kappa light chain, Lewis Y, L1-cell adhesion molecule, (L1-CAM), Melanoma-associated antigen (MAGE)-A1, MAGE-A3, MAGE-A6, Preferentially expressed antigen of melanoma (PRAME), survivin, TAG72, B7-H6, IL-13 receptor alpha 2 (IL-13Ra2), CA9, GD3, HMW-MAA, CD171, G250/CAIX, HLA-AI MAGE Al, HLA-A2 NY-ESO-1, PSCA, folate receptor-a, CD44v6, CD44v7/8, avb6 integrin, 8H9, NCAM, VEGF receptors, 5T4, Foetal AchR, NKG2D ligands, CD44v6, dual antigen, a cancer-testes antigen, mesothelin, murine CMV, mucin 1 (MUC1), MUC16, PSCA, NKG2D, NY-ESO-1, MART-1, gp100, oncofetal antigen, ROR1, TAG72, VEGF-R2, carcinoembryonic antigen (CEA), Her2/neu, estrogen receptor, progesterone receptor, ephrinB2, CD123, c-Met, GD-2, O-acetylated GD2 (OGD2), CE7, Wilms Tumor 1 (WT-1), a cyclin, cyclin A2, CCL-1, CD138, a pathogen-specific antigen and an antigen associated with a universal tag.

[0025] Provided herein are methods of treatment that involve: (a) administering a cell therapy to a subject having a disease or disorder, said cell therapy comprising a dose of genetically engineered cells expressing a recombinant receptor; and (b) subsequent to the administration in (a) administering to the subject an inhibitor of gamma secretase.

[0026] Provided herein are methods of treatment that involve: administering an inhibitor of gamma secretase to a subject having a disease or disorder, wherein, at the time of initiation of the administration of the inhibitor, the subject has been previously administered, and/or is

undergoing treatment with, a cell therapy, said cell therapy comprising a dose of genetically engineered cells expressing recombinant receptor.

[0027] In some embodiments of any one of the methods provided herein, the recombinant receptor specifically binds to a target antigen associated with the disease or disorder.

[0028] In some embodiments of any one of the methods provided herein, the target antigen is selected from among, B cell maturation antigen (BCMA), carbonic anhydrase 9 (CAIX), Her2/neu (receptor tyrosine kinase erbB2), L1-CAM, CD19, CD20, CD22, mesothelin, CEA, and hepatitis B surface antigen, anti-folate receptor, CD23, CD24, CD30, CD33, CD38, CD44, EGFR, epithelial glycoprotein 2 (EPG-2), epithelial glycoprotein 40 (EPG-40), EPHa2, erb-B2, erb-B3, erb-B4, erbB dimers, EGFR vIII, folate binding protein (FBP), FCRL5, FCRH5, fetal acetylcholine receptor, GD2, GD3, HMW-MAA, IL-22R-alpha, IL-13R-alpha2, kinase insert domain receptor (kdr), kappa light chain, Lewis Y, L1-cell adhesion molecule, (L1-CAM), Melanoma-associated antigen (MAGE)-A1, MAGE-A3, MAGE-A6, Preferentially expressed antigen of melanoma (PRAME), survivin, TAG72, B7-H6, IL-13 receptor alpha 2 (IL-13Ra2), CA9, GD3, HMW-MAA, CD171, G250/CAIX, HLA-AI MAGE Al, HLA-A2 NY-ESO-1, PSCA, folate receptor-a, CD44v6, CD44v7/8, avb6 integrin, 8H9, NCAM, VEGF receptors, 5T4, Foetal AchR, NKG2D ligands, CD44v6, dual antigen, a cancer-testes antigen, mesothelin, murine CMV, mucin 1 (MUC1), MUC16, PSCA, NKG2D, NY-ESO-1, MART-1, gp100, oncofetal antigen, ROR1, TAG72, VEGF-R2, carcinoembryonic antigen (CEA), Her2/neu, estrogen receptor, progesterone receptor, ephrinB2, CD123, c-Met, GD-2, O-acetylated GD2 (OGD2), CE7, Wilms Tumor 1 (WT-1), a cyclin, cyclin A2, CCL-1, CD138, a pathogen-specific antigen and an antigen associated with a universal tag. In some embodiments, the target antigen is Muc1. In some embodiments of any one of the methods provided herein, the target antigen is BCMA. In some embodiments, wherein the BCMA is surface BCMA.

[0029] In some embodiments of any one of the methods provided herein, the gamma secretase inhibitor inhibits or reduces or is capable of inhibiting or reducing cleavage of one or more targets selected from BCMA, Notch 1, Notch 2, Notch 3, Notch 4, Muc1, Ephrin B2, Betaglycan (TGFBR3), CD43, CD44, CSF1R, CXCR1, CXCL16, Delta1, E-cadherein, N-cadherin, HLA-A2, IFNaR2, IL1R1, IL1R2, IL6R or ameloid precursor protein (APP). In some embodiments, the one or more targets the cleavage of which is reduced or inhibited comprises

Muc1. In some embodiments, the one or more targets the cleavage of which is reduced or inhibited comprises BCMA.

[0030] In some embodiments of any one of the methods provided herein, cleavage of one or more targets selected from BCMA, Notch 1, Notch 2, Notch 3, Notch 4, Muc1, Ephrin B2, Betaglycan (TGFBR3), CD43, CD44, CSF1R, CXCR1, CXCL16, Delta1, E-cadherein, N-cadherin, HLA-A2, IFNaR2, IL1R1, IL1R2, IL6R or ameloid precursor protein (APP) is inhibited or reduced or can be inhibited or reduced by the gamma secretase inhibitor. In some embodiments, the one or more targets the cleavage of which is reduced or inhibited comprises Muc1. In some embodiments, the one or more targets the cleavage of which is reduced or inhibited comprises BCMA.

[0031] In some embodiments of any one of the methods provided herein, the gamma secretase inhibitor inhibits the target with a half-maximal inhibitory concentration (IC_{50}) of 0.01 nM to 1 μM, 0.01 nM to 100 nM, 0.01 nM to 10 nM, 0.01 nM to 5 nM, 0.01 nM to 1 nM, 0.01 nM to 0.5 nM, 0.01 nM to 0.25 nM, 0.01 nM to 0.1 nM, 0.01 nM to 0.05 nM, 0.05 nM to 1 μM, 0.05 nM to 100 nM, 0.05 nM to 10 nM, 0.05 nM to 5 nM, 0.05 nM to 1 nM, 0.05 nM to 0.5 nM, 0.05 nM to 0.25 nM, 0.05 nM to 0.1 nM, 0.1 nM to 1 μM, 0.1 nM to 100 nM, 0.1 nM to 10 nM, 0.1 nM to 5 nM, 0.1 nM to 1 nM, 0.1 nM to 0.5 nM, 0.1 nM to 0.25 nM, 0.25 nM to 1 μM, 0.25 nM to 100 nM, 0.25 nM to 10 nM, 0.25 nM to 5 nM, 0.25 nM to 1 nM, 0.25 nM to 0.5 nM, 0.5 nM to 1 μM, 0.5 nM to 100 nM, 0.5 nM to 10 nM, 0.5 nM to 5 nM, 0.5 nM to 1 nM, 1 nM to 1 μM, 1 nM to 100 nM, 1 nM to 10 nM, 1 nM to 5 nM, 5 nM to 1 μM, 5 nM to 100 nM, 5 nM to 10 nM, 10 nM to 1 μM, 10 nM to 100 nM or 100 nM to 1 μM.

[0032] In some embodiments of any one of the methods provided herein, the gamma secretase inhibitor inhibits the target with a half-maximal inhibitory concentration (IC_{50}) of less than or less than about 1 μM, 100 nM, 10 nM, 5 nM, 1 nM, 0.5 nM, 0.35 nM, 0.25 nM, 0.1 nM, 0.05 nM or 0.01 nM or less.

[0033] In some embodiments of any one of the methods provided herein, the gamma secretase inhibitor is a peptide inhibitor or non-peptide inhibitor. In some embodiments, the gamma secretase inhibitor is a peptide inhibitor and the peptide inhibitor is selected from among peptide aldehydes derivatives, difluoroketones derivatives, hydroxyethylene dipeptide isotere derivatives, alpha-helical peptide derivatives and dipeptide analogs. In some embodiments, the

gamma secretase inhibitor is a non-peptide inhibitor and the non-peptide inhibitor is a benzodiazepines derivative or a sulfonamides derivative.

[0034] In some embodiments of any one of the methods provided herein, the gamma secretase inhibitor is a transition state inhibitor or non-transition state inhibitor.

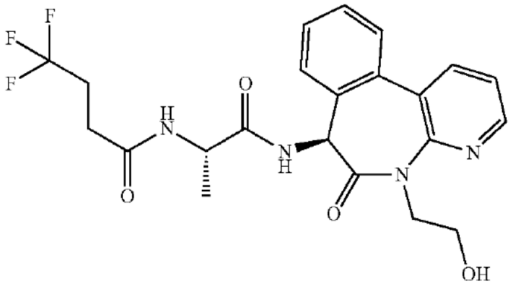
[0035] In some embodiments of any one of the methods provided herein, the gamma secretase inhibitor is a nonsteroidal anti-inflammatory drug.

[0036] In some embodiments of any one of the methods provided herein, the gamma secretase inhibitor is selected from LY3039478, secretase inhibitor I (GSI I) Z-Leu-Leu-Norleucine; γ-secretase inhibitor II (GSI II); γ-secretase inhibitor III (GSI III), N-Benzyloxycarbonyl-Leu- leucinal, N-(2-Naphthoyl)-Val- phenylalaninal; γ-secretase inhibitor III (GSI IV); γ-secretase inhibitor III (GSI V), N- Benzyloxycarbonyl-Leu- phenylalaninal; γ-secretase inhibitor III (GSI VI), l-(S)-endo-N- (1,3,3)-Trimethylbicyclo[2.2.1]hept-2-yl)-4-fluorophenyl Sulfonamide; γ-secretase inhibitor III (GSI VII), Menthyloxycarbonyl-LL-CHO; γ-secretase inhibitor III (GSI IX), (DAPT), N- [N-(3,5- Difluorophenacetyl-L- alanyl)]-S-phenylglycine t- Butyl Ester; γ-secretase inhibitor X (GSI X), { 1 S-Benzyl-4R-[1-(1S-carbamoyl- 2- phenethylcarbamoyl)-lS-3- methylbutylcarb-amoyl]-2R- hydroxy-5-phenylpentyl}carbamic Acid tert-butyl Ester; γ- secretase inhibitor XI (GSI XI), 7-Amino-4-chloro-3- methoxyisocoumarin; γ-secretase inhibitor XII (GSI XII), Z-Ile-Leu-CHO; γ-secretase inhibitor XIII (GSI XIII), Z-Tyr-Ile-Leu- CHO; γ-secretase inhibitor XIV (GSI XIV), Z-Cys(t-Bu)-Ile- Leu-CHO; γ-secretase inhibitor XVI (GSI XVI), N-[N-3,5- Difluorophenacetyl]-L-alanyl-S- phenylglycine Methyl Ester; γ- secretase inhibitor XVII (GSI XVII); γ-secretase inhibitor XIX (GSI XIX), benzo[e][l,4]diazepin-3-yl)- butyramide; γ-secretase inhibitor XX (GSI XX), (S,S)- 2-[2-(3,5- Difluorophenyl)acetylamino]- N-(5-methyl-6-oxo-6,7- dihydro-5H-dibenzo[b,d]azepin-7- yl)propionamide; γ-secretase inhibitor XXI (GSI XXI), (S,S)-2-[2-(3,5-Difluorophenyl)- acetylamino]-N-(l-methyl-2- oxo-5-phenyl-2-,3-dihydro-lH-benzo[e][l,4]diazepin-3- yl)- propionamide; Gamma40 secretase inhibitor I, N-trans-3,5-Dimethoxycinnamoyl-Ile- leucinal; Gamma40 secretase inhibitor II, N-tert-Butyloxycarbonyl-Gly-Val-Valinal Isovaleryl-V V-Sta-A-Sta- OCH3; MK-0752; MRK-003 (Merck); semagacestat/LY450139; RO4929097; PF-03084,014; BMS-708163; MPC-7869 (γ- secretase modifier), YO-01027 (Dibenzazepine), Compound E ([(2S)-2-{[(3,5-Difluorophenyl)acetyl]amino}-N-[(3S)-l-methyl-2-oxo-5-phenyl-2,3-dihydro-lH-l,4-

benzodiazepin-3-yl]propanamide], LY411575, L-685,458, BMS-289948 (4-chloro-N-(2,5-difluorophenyl)-N-((lR)- {4-fluoro-2-[3-(lH-imidazol-l-yl)propyl]phenyl}ethyl)benzenesulfonamide hydrochloride) and BMS-299897 (4-[2-((lR)-l-{[(4-chlorophenyl)sulfonyl]-2,5-difluoroanilino}ethyl)-5- fluorophenyljbutanoic acid).

[0037] In some embodiments of any one of the methods provided herein, the gamma secretase inhibitor is a compound of the structure:

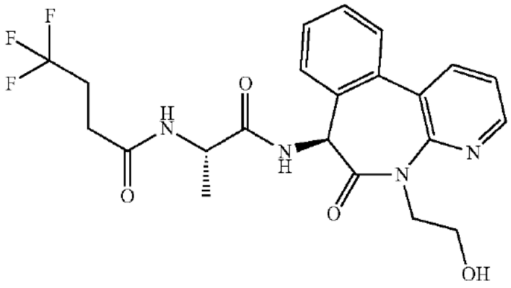
Compound 1



or a pharmaceutically acceptable salt of hydrate thereof.

[0038] Provided herein are methods of treatment that involve: (a) administering a cell therapy to a subject having a disease or disorder, said cell therapy comprising a dose of genetically engineered cells expressing a recombinant receptor; and (b) administering to the subject a compound of the structure:

Compound 1

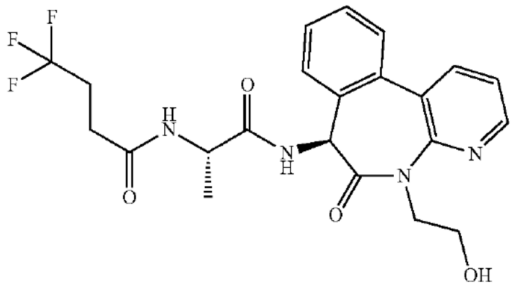


or a pharmaceutically acceptable salt of hydrate thereof.

[0038a] Disclosed herein is the use of a cell therapy and a compound in the manufacture of a medicament for combination ~~comprising a cell~~ therapy ~~and a compound, in the manufacture of a medicament for the treatment~~ of a B cell maturation antigen (BCMA)-expressing cancer in a subject, wherein (a) the cell therapy ~~is to be administered to the subject having the BCMA-expressing cancer said cell therapy emprising~~comprises a dose of T cells expressing a

recombinant receptor, wherein the recombinant receptor specifically binds to BCMA; and (b) the compound is ~~to be administered to the subject, wherein the compound is of the structure: or a stereoisomer thereof, or a pharmaceutically acceptable salt or hydrate of either of the foregoing~~

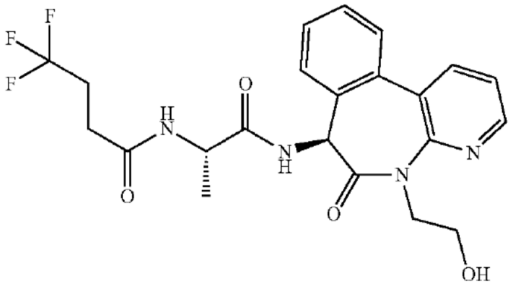
Compound 1



or a stereoisomer thereof, or a pharmaceutically acceptable salt or hydrate of either of the foregoing.

[0039] Provided herein are methods of treatment that involve: administering a cell therapy to a subject having a disease or disorder, said cell therapy comprising a dose of genetically engineered cells expressing a recombinant receptor, wherein, at the time of initiation of the administration of the cell therapy, the subject has been previously administered, and/or is undergoing treatment with, a compound of the structure:

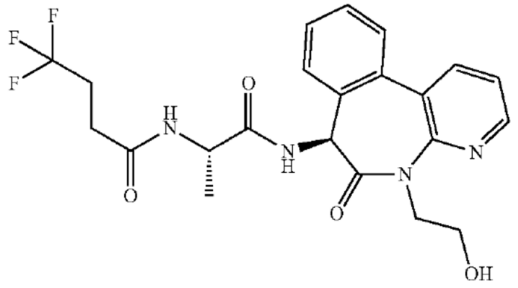
Compound 1



or a pharmaceutically acceptable salt of hydrate thereof.

[0039a] Disclosed herein is the use of a cell therapy for preparation of a medicament for treating a B cell maturation antigen (BCMA)-expressing cancer in a subject, wherein the cell therapy is to be administered to a subject having the BCMA-expresing cancer, said cell therapy comprising a dose of T cells expressing a recombinant receptor, wherein the recombinant receptor specifically binds to BCMA, and wherein, at the time of initiation of the administration of the cell therapy, the subject has been previously administered, and/or is undergoing treatment

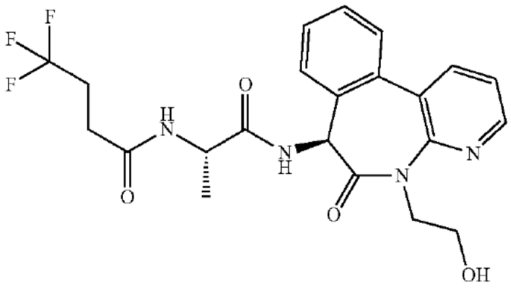
Compound 1



with, a compound having the structure: or a stereoisomer thereof, or a pharmaceutically acceptable salt or hydrate of either of the foregoing.

[0040] Provided herein are methods of treatment that involve: administering to a subject having a disease or disorder a compound of the structure:

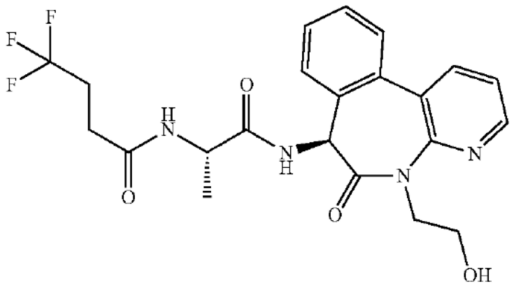
Compound 1



or a pharmaceutically acceptable salt of hydrate thereof, wherein, at the time of initiation of the administration, the subject has been previously administered, and/or is undergoing treatment with a cell therapy, said cell therapy comprising a dose of genetically engineered cells expressing a recombinant receptor.

[0040a] Disclosed herein is the use of a compound in the manufacture of a medicament for the treatment of a B cell maturation antigen (BCMA)-expressing cancer in a subject, wherein the compound is to be administered to the subject having the BCMA-expressing cancer, said

Compound 1



compound ~~having the structure~~is : or a

stereoisomer thereof, or a pharmaceutically acceptable salt or hydrate of either of the foregoing, wherein, at the time of initiation of the administration of the compound, the subject has been previously administered, and/or is undergoing treatment with, a cell therapy, said cell therapy comprising a dose ofT cells expressing a recombinant receptor, wherein the recombinant receptor specifically binds to BCMA.

[0041] In some embodiments of any one of the methods provided herein, the recombinant receptor specifically binds to a target antigen associated with the disease or disorder. In some embodiments, the target antigen is selected from among, B cell maturation antigen (BCMA), carbonic anhydrase 9 (CAIX), Her2/neu (receptor tyrosine kinase erbB2), L1-CAM, CD19, CD20, CD22, mesothelin, CEA, and hepatitis B surface antigen, anti-folate receptor, CD23, CD24, CD30, CD33, CD38, CD44, EGFR, epithelial glycoprotein 2 (EPG-2), epithelial glycoprotein 40 (EPG-40), EPHa2, erb-B2, erb-B3, erb-B4, erbB dimers, EGFR vIII, folate binding protein (FBP), FCRL5, FCRH5, fetal acetylcholine receptor, GD2, GD3, HMW-MAA, IL-22R-alpha, IL-13R-alpha2, kinase insert domain receptor (kdr), kappa light chain, Lewis Y, L1-cell adhesion molecule, (L1-CAM), Melanoma-associated antigen (MAGE)-A1, MAGE-A3, MAGE-A6, Preferentially expressed antigen of melanoma (PRAME), survivin, TAG72, B7-H6, IL-13 receptor alpha 2 (IL-13Ra2), CA9, GD3, HMW-MAA, CD171, G250/CAIX, HLA-AI MAGE Al, HLA-A2 NY-ESO-1, PSCA, folate receptor-a, CD44v6, CD44v7/8, avb6 integrin, 8H9, NCAM, VEGF receptors, 5T4, Foetal AchR, NKG2D ligands, CD44v6, dual antigen, a cancer-testes antigen, mesothelin, murine CMV, mucin 1 (MUC1), MUC16, PSCA, NKG2D, NY-ESO-1, MART-1, gp100, oncofetal antigen, ROR1, TAG72, VEGF-R2, carcinoembryonic antigen (CEA), Her2/neu, estrogen receptor, progesterone receptor, ephrinB2, CD123, c-Met, GD-2, O-acetylated GD2 (OGD2), CE7, Wilms Tumor 1 (WT-1), a cyclin, cyclin A2, CCL-1, CD138, a pathogen-specific antigen and an antigen associated with a universal tag. In some embodiments, the target antigen is Muc1. In some embodiments, the target antigen is BCMA. In some embodiments, the BCMA is surface BCMA. In some embodiments, the recombinant receptor does not bind soluble BCMA or binds to soluble BCMA with an affinity lower than the affinity of said recombinant receptor for binding to surface BCMA. In some embodiments, the affinity of the antigen-binding domain to the surface B cell maturation antigen (BCMA) is at least 1-fold, 2-fold, 3-fold, 4-fold, 5-fold, 6-fold, 7-fold, 8-fold, 9-fold, or 10-fold higher than the affinity to the soluble BCMA.

[0042] In some embodiments of any one of the methods provided herein, the subject comprises plasma cells, or cancer cells or myeloma cells or cells expressing plasma cell markers, expressing surface BCMA.

[0043] In some embodiments of any one of the methods provided herein, the subject is selected as having cells comprising low expression of surface B cell maturation antigen (BCMA); and/or selecting a subject for administration of the cell therapy and/or the inhibitor based on having plasma cells comprising low expression of surface BCMA.

[0044] In some embodiments of any one of the methods provided herein, the target of the gamma secretase inhibitor is the target antigen and the gamma secretase inhibitor inhibits cleavage of the target antigen.

[0045] In some embodiments of any one of the methods provided herein, administration of the inhibitor: decreases BCMA cleavage or shedding from cells, optionally plasma cells, by greater than or greater than about 5%, 10%, 20%, 30%, 40%, 50%, 60%, 70%, 80%, 90% or more compared to the level of BCMA cleavage or shedding from cells in the subject prior to administration of the inhibitor; decreases the level or amount of BCMA detected in the serum of a subject by greater than or greater than about 5%, 10%, 20%, 30%, 40%, 50%, 60%, 70%, 80%, 90% or more compared to the level or amount of BCMA in the serum of the subject prior to administration of the inhibitor; and/or increases expression of surface BCMA on cells, optionally plasma cells by greater than or greater than about 5%, 10%, 20%, 30%, 40%, 50%, 60%, 70%, 80%, 90% or more compared to the level of surface BCMA on the cells in the subject prior to administration of the inhibitor.

[0046] In some embodiments of any one of the methods provided herein, administration of the inhibitor: decreases cleavage or shedding of the target or the target antigen, optionally BCMA or Muc1, from cells, optionally plasma cells, by greater than or greater than about 5%, 10%, 20%, 30%, 40%, 50%, 60%, 70%, 80%, 90% or more compared to the level of cleavage or shedding of the target or target antigen from cells in the subject prior to administration of the inhibitor; decreases the level or amount of the target or target antigen, optionally BCMA or Muc1, detected in the serum of a subject by greater than or greater than about 5%, 10%, 20%, 30%, 40%, 50%, 60%, 70%, 80%, 90% or more compared to the level or amount of the target or target antigen in the serum of the subject prior to administration of the inhibitor; and/or increases expression of surface target or target antigen, optionally BCMA or Muc1, on cells, optionally

plasma cells, by greater than or greater than about 5%, 10%, 20%, 30%, 40%, 50%, 60%, 70%, 80%, 90% or more compared to the level of surface target or target antigen on the cells in the subject prior to administration of the inhibitor.

[0047] In some embodiments of any one of the methods provided herein, the disease or disorder is a cancer. In some embodiments, the cancer is a B cell malignancy. In some embodiments, the cancer is multiple myeloma, plasmacytoma, a cancer of plasma cell origin and/or a cancer of B cell origin.

[0048] In some embodiments of any one of the methods provided herein, the inhibitor is administered orally, subcutaneously or intravenously. In some embodiments, the inhibitor is administered orally.

[0049] In some embodiments of any one of the methods provided herein, the inhibitor is administered at least or is administered six times daily, five times daily, four times daily, three times daily, twice daily, once daily, every other day, three times a week, at least once a week, or only one time. In some embodiments, the inhibitor is administered three times a week.

[0050] In some embodiments of any one of the methods provided herein, the administration of the inhibitor is carried out in a treatment cycle that is at least or at least about or 14 days, at least or at least about or 21 days or at least or at least about or 28 days.

[0051] In some embodiments of any one of the methods provided herein, the inhibitor is administered, or each administration of the inhibitor is independently administered, in an amount of 0.5 mg to 500 mg, 0.5 mg to 250 mg, 0.5 mg to 100 mg, 0.5 mg to 50 mg, 0.5 mg to 25 mg, 0.5 mg to 10 mg, 0.5 mg to 5.0 mg, 0.5 mg to 2.5 mg, 0.5 mg to 1.0 mg, 1.0 mg to 500 mg, 1.0 mg to 250 mg, 1.0 mg to 100 mg, 1.0 mg to 50 mg, 1.0 mg to 25 mg, 1.0 mg to 10 mg, 1.0 mg to 5.0 mg, 1.0 mg to 2.5 mg, 2.5 mg to 500 mg, 2.5 mg to 250 mg, 2.5 mg to 100 mg, 2.5 mg to 50 mg, 2.5 mg to 25 mg, 2.5 mg to 10 mg, 2.5 mg to 5.0 mg, 5.0 mg to 500 mg, 5.0 mg to 250 mg, 5.0 mg to 100 mg, 5.0 mg to 50 mg, 5.0 mg to 25 mg, 5.0 mg to 10 mg, 10 mg to 500 mg, 10 mg to 250 mg, 10 mg to 100 mg, 10 mg to 50 mg, 10 mg to 25 mg, 25 mg to 500 mg, 25 mg to 250 mg, 25 mg to 100 mg, 25 mg to 50 mg, 50 mg to 500 mg, 50 mg to 250 mg, 50 mg to 100 mg, 100 mg to 500 mg, 100 mg to 250 mg or 250 mg to 500 mg. In some embodiments, the inhibitor is administered, or each administration of the inhibitor is independently administered, in an amount that is at least or at least about or is or is about 0.5 mg, 1.0 mg, 2.5 mg, 5.0 mg, 10.0 mg, 25 mg, 50 mg, 100 mg, 250 mg or 500 mg.

[0052] In some embodiments of any one of the methods provided herein, the recombinant receptor is a transgenic T cell receptor (TCR) or a functional non-T cell receptor. In some embodiments, the recombinant receptor is a chimeric receptor, which optionally is a chimeric antigen receptor (CAR). In some embodiments, the chimeric antigen receptor (CAR) comprises an extracellular antigen-recognition domain that specifically binds to the antigen and an intracellular signaling domain comprising an ITAM. In some embodiments, the intracellular signaling domain comprises and intracellular domain of a CD3-zeta (CD3ζ) chain. In some embodiments, the chimeric antigen receptor (CAR) further comprises a costimulatory signaling region. In some embodiments, the costimulatory signaling region comprises a signaling domain derived from CD28 or 4-1BB, optionally human CD28 or human 4-1BB. In some embodiments, the costimulatory signaling region is a domain derived from 4-1BB, optionally human 4-1BB.

[0053] In some embodiments of any one of the methods provided herein, the subject is a human.

[0054] In some embodiments of any one of the methods provided herein, the BCMA is human BCMA or the target antigen is a human antigen.

[0055] In some embodiments of any one of the methods provided herein, the genetically engineered cells comprise T cells or NK cells. In some embodiments, the cell therapy is a T cell therapy and the dose of genetically engineered cells comprises T cells. In some embodiments, the T cells are CD4+ and/or CD8+. In some embodiments, the T cells are primary T cells obtained from a subject.

[0056] In some embodiments of any one of the methods provided herein, the cell therapy comprises cells that are autologous to the subject.

[0057] In some embodiments of any one of the methods provided herein, the cell therapy comprises cells that are allogeneic to the subject.

[0058] In some embodiments of any one of the methods provided herein, the cell therapy comprises the administration of from or from about 1×10^5 to 5×10^8 total recombinant receptor-expressing cells, total T cells, or total peripheral blood mononuclear cells (PBMCs); the cell therapy comprises the administration of from or from about 1×10^5 to 1×10^8 total recombinant receptor-expressing cells, total T cells, or total PBMCs; from or from about 5×10^5 to 1×10^7 total recombinant receptor-expressing cells, total T cells, or total PBMCs; or from or

from about 1 x 10^6 to 1 x 10^7 total recombinant receptor-expressing cells, total T cells, or total PBMCs, each inclusive.

[0059] In some embodiments of any one of the methods provided herein, the cell therapy comprises the administration of no more than 5 x 10^8 total recombinant receptor-expressing cells, total T cells, or total peripheral blood mononuclear cells (PBMCs); no more than 2.5 x 10^8 total recombinant receptor-expressing cells, total T cells, or total PBMCs; no more than 1 x 10^8 total recombinant receptor-expressing cells, total T cells, or total PBMCs; no more than 1 x 10^7 total recombinant receptor-expressing cells, total T cells, or total PBMCs; no more than 0.5 x 10^7 total recombinant receptor-expressing cells, total T cells, or total PBMCs; no more than 1 x 10^6 total recombinant receptor-expressing cells, total T cells, or total PBMCs; no more than 0.5 x 10^6 total recombinant receptor-expressing cells, total T cells, or total PBMCs.

[0060] In some embodiments of any one of the methods provided herein, the initiation of administration of the inhibitor is prior to, concurrently with or subsequently to initiation of administration of the cell therapy. In some embodiments, the inhibitor is administered prior to initiation of administration of the cell therapy. In some embodiments, initiation of administration of the inhibitor is within, or within about, 1 hours, 2 hour, 6 hours, 12 hours, 24 hours, 48 hours, 72 hours, 96 hours or 1 week prior to the initiation of the administration of the cell therapy. In some embodiments, the inhibitor is administered subsequently to initiation of administration of the cell therapy. In some embodiments, initiation of administration of the inhibitor is within, or within about, 1 hours, 2 hour, 6 hours, 12 hours, 24 hours, 48 hours, 72 hours, 96 hours,1 week, 14 days, 21 days, 28 days or more after the initiation of the administration of the cell therapy. In some embodiments, the inhibitor is administered up to 7 days, 14 days, 21 days, 28 days or more after initiation of the administration of the cells.

[0061] In some embodiments of any one of the methods provided herein, the inhibitor is administered at a time in which: the number of cells of the cell therapy detectable in the blood from the subject is decreased compared to in the subject at a preceding time point after initiation of the administration of the cells; the number of cells of the cell therapy detectable in the blood is less than or less than about 1.5-fold, 2-fold, 3-fold, 4-fold, 5-fold, 10-fold, 50-fold or 100-fold or less the peak or maximum number of the cells of the cell therapy detectable in the blood of the subject after initiation of administration of the administration of the cells; and/or at a time after a peak or maximum level of the cells of the cell therapy are detectable in the blood of the

subject, the number of cells of or derived from the cells detectable in the blood from the subject is less than less than 10%, less than 5%, less than 1% or less than 0.1% of total peripheral blood mononuclear cells (PBMCs) in the blood of the subject.

[0062] In some embodiments of any one of the methods provided herein, the method further comprises administering a lymphodepleting chemotherapy prior to administration of the cell therapy and/or wherein the subject has received a lymphodepleting chemotherapy prior to administration of the cells. In some embodiments, the lymphodepleting chemotherapy comprises administering fludarabine and/or cyclophosphamide to the subject.

[0063] In some embodiments of any one of the methods provided herein, the method further comprises administering a steroid, optionally wherein the steroid is administered prior to, concurrently with and/or subsequently to initiation of administration of the inhibitor, optionally wherein the steroid is administered during the cycle of treatment with the inhibitor. In some embodiments, the steroid is or comprises dexamethasone.

[0064] In some embodiments of any one of the methods provided herein, the cell therapy exhibits increased or prolonged expansion and/or persistence in the subject as compared to a method in which the cell therapy is administered to the subject in the absence of the gamma secretase inhibitor.

[0065] In some embodiments of any one of the methods provided herein, the method thereby prevents, reduces or ameliorates one or more symptoms or outcomes of the disease or disorder.

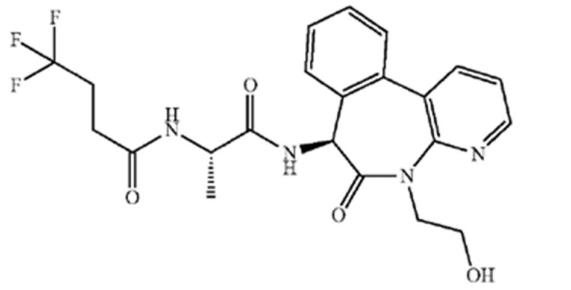
[0066] Provided herein are methods of treatment that involve: (a) administering to a subject having a disease or disorder, a therapeutic agent or therapy that targets or is specific for B cell maturation antigen (BCMA); and (b) administering a gamma secretase inhibitor to the subject.

[0067] In some embodiments of any one of the methods provided herein, the therapeutic agent or therapy is or comprises an antibody or antigen-binding fragment thereof, optionally a bispecific antibody. In some embodiments, therapeutic agent or therapy is a bispecific antibody that further targets or specifically binds to a T cell antigen, optionally CD2 or CD3. In some embodiments, the therapeutic agent or therapy is a bispecific antibody that further targets a second antigen, optionally wherein the second antigen is selected from CD19, CD20, CD22, CD33, CD38, CS1, ROR1, GPC3, CD123, IL-13R, CD138, c-Met, EGFRvIII, GD-2, NY-ESO-1, MAGE A3, and glycolipid F77.

[0068] In some embodiments of any one of the methods provided herein, the gamma secretase inhibitor is selected from LY3039478, secretase inhibitor I (GSI I) Z-Leu-Leu-Norleucine; γ-secretase inhibitor II (GSI II); γ- secretase inhibitor III (GSI III), N-Benzyloxycarbonyl-Leu- leucinal, N-(2-Naphthoyl)-Val- phenylalaninal; γ-secretase inhibitor III (GSI IV); γ-secretase inhibitor III (GSI V), N- Benzyloxycarbonyl-Leu- phenylalaninal; γ-secretase inhibitor III (GSI VI), l-(S)-endo-N- (1,3,3)- Trimethylbicyclo[2.2.1]hept-2-yl)-4-fluorophenyl Sulfonamide; γ-secretase inhibitor III (GSI VII), Menthyloxycarbonyl-LL-CHO; γ-secretase inhibitor III (GSI IX), (DAPT), N- [N-(3,5- Difluorophenacetyl-L- alanyl)]-S-phenylglycine t- Butyl Ester; γ-secretase inhibitor X (GSI X), { 1 S-Benzyl-4R-[1-(1S-carbamoyl- 2- phenethylcarbamoyl)-lS-3- methylbutylcarb-amoyl]-2R- hydroxy-5-phenylpentyl}carbamic Acid tert-butyl Ester; γ- secretase inhibitor XI (GSI XI), 7-Amino-4-chloro-3- methoxyisocoumarin; γ-secretase inhibitor XII (GSI XII), Z-Ile-Leu-CHO; γ-secretase inhibitor XIII (GSI XIII), Z-Tyr-Ile-Leu- CHO; γ-secretase inhibitor XIV (GSI XIV), Z-Cys(t-Bu)-Ile- Leu-CHO; γ-secretase inhibitor XVI (GSI XVI), N-[N-3,5- Difluorophenacetyl]-L-alanyl-S- phenylglycine Methyl Ester; γ- secretase inhibitor XVII (GSI XVII); γ-secretase inhibitor XIX (GSI XIX), benzo[e][l,4]diazepin-3-yl)- butyramide; γ-secretase inhibitor XX (GSI XX), (S,S)- 2-[2-(3,5- Difluorophenyl)acetylamino]- N-(5-methyl-6-oxo-6,7- dihydro-5H-dibenzo[b,d]azepin-7- yl)propionamide; γ-secretase inhibitor XXI (GSI XXI), (S,S)-2-[2-(3,5-Difluorophenyl)- acetylamino]-N-(l-methyl-2- oxo-5-phenyl-2-,3-dihydro-lH-benzo[e][l,4]diazepin-3- yl)- propionamide; Gamma40 secretase inhibitor I, N-trans-3,5-Dimethoxycinnamoyl-Ile- leucinal; Gamma40 secretase inhibitor II, N-tert-Butyloxycarbonyl-Gly-Val-Valinal Isovaleryl-V V-Sta-A-Sta- OCH3; MK-0752; MRK-003 (Merck); semagacestat/LY450139; RO4929097; PF-03084,014; BMS-708163; MPC-7869 (γ- secretase modifier), YO-01027 (Dibenzazepine), Compound E ([(2S)-2-{[(3,5-Difluorophenyl)acetyl]amino}-N-[(3S)-l-methyl-2-oxo-5-phenyl-2,3-dihydro-lH-l,4-benzodiazepin-3-yl]propanamide], LY411575, L-685,458, BMS-289948 (4-chloro-N-(2,5-difluorophenyl)-N-((lR)- {4-fluoro-2-[3-(lH-imidazol-l-yl)propyl]phenyl}ethyl)benzenesulfonamide hydrochloride) and BMS-299897 (4-[2-((lR)-l-{[(4-chlorophenyl)sulfonyl]-2,5-difluoroanilino}ethyl)-5- fluorophenyljbutanoic acid).

[0069] In some embodiments of any one of the methods provided herein, the gamma secretase inhibitor is a compound of the structure:

Compound I



or a pharmaceutically acceptable salt of hydrate thereof.

[0070] Provided herein are combinations that include: (a) genetically engineered cells expressing a chimeric receptor comprising an antigen binding domain that binds to surface B cell maturation antigen (BCMA); (b) an inhibitor of gamma secretase, wherein binding of the CAR to surface BCMA or a measure indicative of function or activity, following exposure to cells expressing surface BCMA, is not reduced or blocked or is not substantially reduced or blocked in the presence of a soluble or shed form of the BCMA, optionally at a concentration or amount of the soluble or shed form of the BCMA corresponding to a concentration or amount present in serum or blood or plasma of the subject or of a multiple myeloma patient, or on average in a patient population for the disease or disorder, or at a concentration or amount of the soluble or shed BCMA at which the binding or measure is reduced or blocked, or is substantially reduced or blocked, for cells expressing a reference anti-BCMA CAR in the same assay.

[0071] In some embodiments of any one of the combinations provided herein, the gamma secretase inhibitor inhibits intramembrane cleavage of BCMA.

[0072] In some embodiments of any one of the combinations provided herein, the gamma secretase inhibitor inhibits intramembrane cleavage of BCMA with a half-maximal inhibitory concentration (IC_{50}) of 0.01 nM to 10 nM, 0.01 nM to 5 nM, 0.01 nM to 1 nM, 0.01 nM to 0.5 nM, 0.01 nM to 0.35 nM, 0.01 nM to 0.25 nM, 0.01 nM to 0.1 nM, 0.01 nM to 0.05 nM, 0.05 nM to 10 nM, 0.05 nM to 5 nM, 0.05 nM to 1 nM, 0.05 nM to 0.5 nM, 0.05 nM to 0.35 nM, 0.05 nM to 0.25 nM, 0.05 nM to 0.1 nM, 0.1 nM to 10 nM, 0.1 nM to 5 nM, 0.1 nM to 1 nM, 0.1 nM, to 0.5 nM, 0.1 nM to 0.35 nM, 0.1 nM to 0.25 nM, 0.25 nM to 10 nM, 0.25 nM to 5 nM, 0.25 nM to 1 nM, 0.25 nM to 0.5 nM, 0.25 nM to 0.35 nM, 0.35 nM to 10 nM, 0.35 nM to 5 nM, 0.35 nM to 1 nM, 0.35 nM to 0.5 nM, 0.5 nM to 10 nM, 0.5 nM to 5 nM, 0.5 nM to 1 nM, 1 nM to 10 nM, 1 nM to 5 nM or 5 nM to 10 nM. In some embodiments, the gamma secretase inhibitor

inhibits intramembrane cleavage of BCMA with a half-maximal inhibitory concentration (IC_{50}) of less than or less than about 10 nM, 5 nM, 1 nM., 0.5 nM, 0.35 nM, 0.25 nM, 0.1 nM, 0.05 nM or 0.01 nM or less.

[0073] Provided herein are combinations that include: (a) genetically engineered cells expressing a recombinant receptor, wherein the recombinant receptor does not specifically bind to surface B cell maturation antigen (BCMA); and (b) an inhibitor of gamma secretase.

[0074] Provided herein are combinations that include: (a) genetically engineered cells expressing a recombinant receptor, wherein the recombinant receptor does not specifically bind to surface B cell maturation antigen (BCMA); and (b) an inhibitor of gamma secretase, wherein binding of the CAR to surface BCMA or a measure indicative of function or activity, following exposure to cells expressing surface BCMA, is not reduced or blocked or is not substantially reduced or blocked in the presence of a soluble or shed form of the BCMA, optionally at a concentration or amount of the soluble or shed form of the BCMA corresponding to a concentration or amount present in serum or blood or plasma of the subject or of a multiple myeloma patient, or on average in a patient population for the disease or disorder, or at a concentration or amount of the soluble or shed BCMA at which the binding or measure is reduced or blocked, or is substantially reduced or blocked, for cells expressing a reference anti-BCMA CAR in the same assay.

[0075] In some embodiments of any one of the combinations provided herein, the recombinant receptor specifically binds to a target antigen associated with the disease or disorder. In some embodiments, the target antigen is selected from among, carbonic anhydrase 9 (CAIX), Her2/neu (receptor tyrosine kinase erbB2), L1-CAM, CD19, CD20, CD22, mesothelin, CEA, and hepatitis B surface antigen, anti-folate receptor, CD23, CD24, CD30, CD33, CD38, CD44, EGFR, epithelial glycoprotein 2 (EPG-2), epithelial glycoprotein 40 (EPG-40), EPHa2, erb-B2, erb-B3, erb-B4, erbB dimers, EGFR vIII, folate binding protein (FBP), FCRL5, FCRH5, fetal acetylcholine receptor, GD2, GD3, HMW-MAA, IL-22R-alpha, IL-13R-alpha2, kinase insert domain receptor (kdr), kappa light chain, Lewis Y, L1-cell adhesion molecule, (L1-CAM), Melanoma-associated antigen (MAGE)-A1, MAGE-A3, MAGE-A6, Preferentially expressed antigen of melanoma (PRAME), survivin, TAG72, B7-H6, IL-13 receptor alpha 2 (IL-13Ra2), CA9, GD3, HMW-MAA, CD171, G250/CAIX, HLA-AI MAGE Al, HLA-A2 NY-ESO-1, PSCA, folate receptor-a, CD44v6, CD44v7/8, avb6 integrin, 8H9, NCAM, VEGF receptors,

5T4, Foetal AchR, NKG2D ligands, CD44v6, dual antigen, a cancer-testes antigen, mesothelin, murine CMV, mucin 1 (MUC1), MUC16, PSCA, NKG2D, NY-ESO-1, MART-1, gp100, oncofetal antigen, ROR1, TAG72, VEGF-R2, carcinoembryonic antigen (CEA), Her2/neu, estrogen receptor, progesterone receptor, ephrinB2, CD123, c-Met, GD-2, O-acetylated GD2 (OGD2), CE7, Wilms Tumor 1 (WT-1), a cyclin, cyclin A2, CCL-1, CD138, a pathogen-specific antigen and an antigen associated with a universal tag. In some embodiments, the target antigen is Muc1.

[0076] In some embodiments of any one of the combinations provided herein, the gamma secretase inhibitor inhibits or reduces or is capable of inhibiting or reducing cleavage of one or more targets selected from BCMA, Notch 1, Notch 2, Notch 3, Notch 4, Muc1, Ephrin B2, Betaglycan (TGFBR3), CD43, CD44, CSF1R, CXCR1, CXCL16, Delta1, E-cadherein, N-cadherin, HLA-A2, IFNaR2, IL1R1, IL1R2, IL6R or ameloid precursor protein (APP). In some embodiments, the one or more targets the cleavage of which is reduced or inhibited comprises Muc1. In some embodiments, the one or more targets the cleavage of which is reduced or inhibited comprises BCMA.

[0077] In some embodiments of any one of the combinations provided herein, cleavage of one or more targets selected from BCMA, Notch 1, Notch 2, Notch 3, Notch 4, Muc1, Ephrin B2, Betaglycan (TGFBR3), CD43, CD44, CSF1R, CXCR1, CXCL16, Delta1, E-cadherein, N-cadherin, HLA-A2, IFNaR2, IL1R1, IL1R2, IL6R or ameloid precursor protein (APP) is inhibited or reduced or can be inhibited or reduced by the gamma secretase inhibitor. In some embodiments, the one or more targets the cleavage of which is inhibited or reduced comprises Muc1. In some embodiments, the one or more targets the cleavage of which is reduced or inhibited comprises BCMA.

[0078] In some embodiments of any one of the combinations provided herein, the gamma secretase inhibitor inhibits the target with a half-maximal inhibitory concentration (IC_{50}) of 0.01 nM to 1 μM, 0.01 nM to 100 nM, 0.01 nM to 10 nM, 0.01 nM to 5 nM, 0.01 nM to 1 nM, 0.01 nM to 0.5 nM, 0.01 nM to 0.25 nM, 0.01 nM to 0.1 nM, 0.01 nM to 0.05 nM, 0.05 nM to 1 μM, 0.05 nM to 100 nM, 0.05 nM to 10 nM, 0.05 nM to 5 nM, 0.05 nM to 1 nM, 0.05 nM to 0.5 nM, 0.05 nM to 0.25 nM, 0.05 nM to 0.1 nM, 0.1 nM to 1 μM, 0.1 nM to 100 nM, 0.1 nM to 10 nM, 0.1 nM to 5 nM, 0.1 nM to 1 nM, 0.1 nM to 0.5 nM, 0.1 nM to 0.25 nM, 0.25 nM to 1 μM, 0.25 nM to 100 nM, 0.25 nM to 10 nM, 0.25 nM to 5 nM, 0.25 nM to 1 nM, 0.25 nM to 0.5 nM,

0.5 nM to 1 μM, 0.5 nM to 100 nM, 0.5 nM to 10 nM, 0.5 nM to 5 nM, 0.5 nM to 1 nM, 1 nM to 1 μM, 1 nM to 100 nM, 1 nM to 10 nM, 1 nM to 5 nM, 5 nM to 1 μM, 5 nM to 100 nM, 5 nM to 10 nM, 10 nM to 1 μM, 10 nM to 100 nM or 100 nM to 1 μM. In some embodiments, the gamma secretase inhibitor inhibits the target with a half-maximal inhibitory concentration (IC_{50}) of less than or less than about 1 μM, 100 nM, 10 nM, 5 nM, 1 nM, 0.5 nM, 0.35 nM, 0.25 nM, 0.1 nM, 0.05 nM or 0.01 nM or less.

[0079] In some embodiments of any one of the combinations provided herein, the gamma secretase inhibitor is a peptide inhibitor or non-peptide inhibitor. In some embodiments, the gamma secretase inhibitor is a peptide inhibitor and the peptide inhibitor is selected from among peptide aldehydes derivatives, difluoroketones derivatives, hydroxyethylene dipeptide isotere derivatives, alpha-helical peptide derivatives and dipeptide analogs. In some embodiments, the gamma secretase inhibitor is a non-peptide inhibitor and the non-peptide inhibitor is a benzodiazepines derivative or a sulfonamides derivative.

[0080] In some embodiments of any one of the combinations provided herein, the gamma secretase inhibitor is a transition state inhibitor or non-transition state inhibitor.

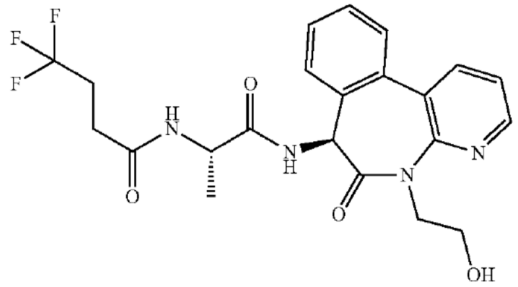
[0081] In some embodiments of any one of the combinations provided herein, the gamma secretase inhibitor is a nonsteroidal anti-inflammatory drug.

[0082] In some embodiments of any one of the combinations provided herein, the gamma secretase inhibitor is selected from LY3039478, secretase inhibitor I (GSI I) Z-Leu-Leu-Norleucine; γ-secretase inhibitor II (GSI II); γ- secretase inhibitor III (GSI III), N-Benzyloxycarbonyl-Leu- leucinal, N-(2-Naphthoyl)-Val- phenylalaninal; γ-secretase inhibitor III (GSI IV); γ-secretase inhibitor III (GSI V), N- Benzyloxycarbonyl-Leu- phenylalaninal; γ-secretase inhibitor III (GSI VI), l-(S)-endo-N- (1,3,3)- Trimethylbicyclo[2.2.1]hept-2-yl)-4-fluorophenyl Sulfonamide; γ-secretase inhibitor III (GSI VII), Menthyloxycarbonyl-LL-CHO; γ-secretase inhibitor III (GSI IX), (DAPT), N- [N-(3,5- Difluorophenacetyl-L- alanyl)]-S-phenylglycine t- Butyl Ester; γ-secretase inhibitor X (GSI X), { 1 S-Benzyl-4R-[1-(1S-carbamoyl- 2- phenethylcarbamoyl)-lS-3- methylbutylcarb-amoyl]-2R- hydroxy-5-phenylpentyl}carbamic Acid tert-butyl Ester; γ- secretase inhibitor XI (GSI XI), 7-Amino-4-chloro-3- methoxyisocoumarin; γ-secretase inhibitor XII (GSI XII), Z-Ile-Leu-CHO; γ-secretase inhibitor XIII (GSI XIII), Z-Tyr-Ile-Leu- CHO; γ-secretase inhibitor XIV (GSI XIV), Z-Cys(t-Bu)-Ile- Leu-CHO; γ-secretase inhibitor XVI (GSI XVI), N-[N-3,5- Difluorophenacetyl]-L-

alanyl-S- phenylglycine Methyl Ester; γ- secretase inhibitor XVII (GSI XVII); γ-secretase inhibitor XIX (GSI XIX), benzo[e][l,4]diazepin-3-yl)- butyramide; γ-secretase inhibitor XX (GSI XX), (S,S)- 2-[2-(3,5- Difluorophenyl)acetylamino]- N-(5-methyl-6-oxo-6,7- dihydro-5H-dibenzo[b,d]azepin-7- yl)propionamide; γ-secretase inhibitor XXI (GSI XXI), (S,S)-2-[2-(3,5-Difluorophenyl)- acetylamino]-N-(l-methyl-2- oxo-5-phenyl-2-,3-dihydro-lH-benzo[e][l,4]diazepin-3- yl)- propionamide; Gamma40 secretase inhibitor I, N-trans-3,5-Dimethoxycinnamoyl-Ile- leucinal; Gamma40 secretase inhibitor II, N-tert-Butyloxycarbonyl-Gly-Val-Valinal Isovaleryl-V V-Sta-A-Sta- OCH3; MK-0752; MRK-003 (Merck); semagacestat/LY450139; RO4929097; PF-03084,014; BMS-708163; MPC-7869 (γ- secretase modifier), YO-01027 (Dibenzazepine), Compound E ([(2S)-2-{[(3,5-Difluorophenyl)acetyl]amino}-N-[(3S)-l-methyl-2-oxo-5-phenyl-2,3-dihydro-lH-l,4-benzodiazepin-3-yl]propanamide], LY411575, L-685,458, BMS-289948 (4-chloro-N-(2,5-difluorophenyl)-N-((lR)- {4-fluoro-2-[3-(lH-imidazol-l-yl)propyl]phenyl}ethyl)benzenesulfonamide hydrochloride) and BMS-299897 (4-[2-((lR)-l-{[(4-chlorophenyl)sulfonyl]-2,5-difluoroanilino}ethyl)-5- fluorophenyljbutanoic acid).

[0083] In some embodiments of any one of the combinations provided herein, the gamma secretase inhibitor is a compound of the structure:

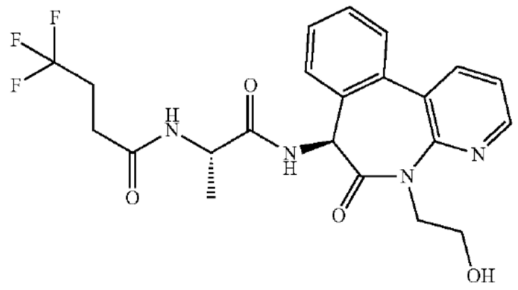
Compound 1



or a pharmaceutically acceptable salt of hydrate thereof.

[0084] Provided herein are combinations that include: (a) genetically engineered cells expressing a recombinant receptor; and (b) a compound of the structure:

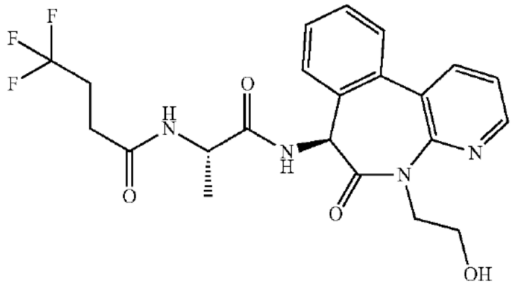
Compound 1



or a pharmaceutically acceptable salt of hydrate thereof.

[0084a] ~~Disclosed herein is~~ In one embodiment, there is provided a combination, comprising: (a) T cells expressing a recombinant receptor that specifically binds to B cell maturation antigen (BCMA); and (b) a compound having the structure:

Compound 1



or a stereoisomer thereof, or a pharmaceutically acceptable salt or hydrate of any of the foregoing.

[0085] In some embodiments of any one of the combinations provided herein, the recombinant receptor specifically binds to a target antigen associated with the disease or disorder. In some embodiments, the target antigen is selected from among, B cell maturation antigen (BCMA), carbonic anhydrase 9 (CAIX), Her2/neu (receptor tyrosine kinase erbB2), L1-CAM, CD19, CD20, CD22, mesothelin, CEA, and hepatitis B surface antigen, anti-folate receptor, CD23, CD24, CD30, CD33, CD38, CD44, EGFR, epithelial glycoprotein 2 (EPG-2), epithelial glycoprotein 40 (EPG-40), EPHa2, erb-B2, erb-B3, erb-B4, erbB dimers, EGFR vIII, folate binding protein (FBP), FCRL5, FCRH5, fetal acetylcholine receptor, GD2, GD3, HMW-MAA, IL-22R-alpha, IL-13R-alpha2, kinase insert domain receptor (kdr), kappa light chain, Lewis Y, L1-cell adhesion molecule, (L1-CAM), Melanoma-associated antigen (MAGE)-A1, MAGE-A3, MAGE-A6, Preferentially expressed antigen of melanoma (PRAME), survivin,

TAG72, B7-H6, IL-13 receptor alpha 2 (IL-13Ra2), CA9, GD3, HMW-MAA, CD171, G250/CAIX, HLA-AI MAGE Al, HLA-A2 NY-ESO-1, PSCA, folate receptor-a, CD44v6, CD44v7/8, avb6 integrin, 8H9, NCAM, VEGF receptors, 5T4, Foetal AchR, NKG2D ligands, CD44v6, dual antigen, a cancer-testes antigen, mesothelin, murine CMV, mucin 1 (MUC1), MUC16, PSCA, NKG2D, NY-ESO-1, MART-1, gp100, oncofetal antigen, ROR1, TAG72, VEGF-R2, carcinoembryonic antigen (CEA), Her2/neu, estrogen receptor, progesterone receptor, ephrinB2, CD123, c-Met, GD-2, O-acetylated GD2 (OGD2), CE7, Wilms Tumor 1 (WT-1), a cyclin, cyclin A2, CCL-1, CD138, a pathogen-specific antigen and an antigen associated with a universal tag. In some embodiments, the target antigen is Muc1.

[0086] In some embodiments of any one of the combinations provided herein, the target antigen is BCMA. In some embodiments, the BCMA is surface BCMA.In some embodiments, the binding of the CAR to surface BCMA or a measure indicative of function or activity, following exposure to cells expressing surface BCMA, is not reduced or blocked or is not substantially reduced or blocked in the presence of a soluble or shed form of the BCMA, optionally at a concentration or amount of the soluble or shed form of the BCMA corresponding to a concentration or amount present in serum or blood or plasma of the subject or of a multiple myeloma patient, or on average in a patient population for the disease or disorder, or at a concentration or amount of the soluble or shed BCMA at which the binding or measure is reduced or blocked, or is substantially reduced or blocked, for cells expressing a reference anti-BCMA CAR in the same assay. In some embodiments, the recombinant receptor does not bind soluble BCMA or binds to soluble BCMA with an affinity lower than the affinity of said recombinant receptor for binding to surface BCMA.

[0087] In some embodiments of any one of the combinations provided herein, the recombinant receptor is a transgenic T cell receptor (TCR) or a functional non-T cell receptor. In some embodiments, the recombinant receptor is a chimeric receptor, which optionally is a chimeric antigen receptor (CAR). In some embodiments, the chimeric antigen receptor (CAR) comprises an extracellular antigen-recognition domain that specifically binds to the antigen and an intracellular signaling domain comprising an ITAM. In some embodiments, the intracellular signaling domain comprises and intracellular domain of a CD3-zeta (CD3ζ) chain. In some embodiments, the chimeric antigen receptor (CAR) further comprises a costimulatory signaling region. In some embodiments, the costimulatory signaling region comprises a signaling domain

derived from CD28 or 4-1BB, optionally human CD28 or human 4-1BB. In some embodiments, the costimulatory signaling region is a domain derived from 4-1BB, optionally human 4-1BB.

[0088] In some embodiments of any one of the combinations provided herein, the BCMA is human BCMA or the target antigen is a human antigen.

[0089] In some embodiments of any one of the combinations provided herein, the genetically engineered cells comprise T cells or NK cells. In some embodiments, the genetically engineered cells comprise T cells. In some embodiments, the T cells are CD4+ and/or CD8+. In some embodiments, the T cells are primary T cells obtained from a subject.

[0090] In some embodiments of any one of the combinations provided herein, the genetically engineered cells are formulated as a pharmaceutical composition for administration to a subject, optionally wherein the cells are formulated for administration in one or more unit doses for treating a disease or condition.

[0091] In some embodiments of any one of the combinations provided herein, the gamma secretase inhibitor is formulated as a pharmaceutical composition for administration to a subject, optionally wherein the gamma secretase inhibitor is formulated for administration in one or more unit doses.

[0092] Provided herein are kits that contain: the combination provided hereinand instructions for administering the components of the combination to a subject having a disease or disorder.

[0092a] ~~Disclosed herein is~~In one embodiment, there is provided a kit comprising the combination provided herein and instructions for the immune cells that are to be administered, and/or for the compound, stereoisomer, pharmaceutically acceptable salt or hydrate that is to be administered to a subject having a disease or disorder.

[0093] Provided herein are kits that contain: (a) a reagent for detecting expression of B cell maturation antigen (BCMA) on the surface of a cell; (b) an inhibitor of gamma secretase, optionally formulated for administration in one or more unit doses; and (c) instructions for administering the inhibitor to a subject based on the results of use of the reagent for detecting BCMA on the surface of plasma cells in a subject having a cancer.

[0094] In some embodiments of any one of the kits provided herein, the instructions specify administering the inhibitor to the subject if the plasma cells comprise low expression of surface BCMA and/or a level of expression of surface BCMA below a threshold level. In some

embodiments, the threshold level of expression of surface BCMA is lower than the average or median expression or level of surface BCMA on plasma cells in a plurality of control subjects, optionally wherein the control subjects are a group of healthy or normal subjects. In some embodiments, the low expression of surface BCMA is present when less than or less than about 60%, less than or less than about 50%, less than or less than about 40%, less than or less than about 30%, less than or less than about 20% or less than or less than about 10% of the plasma cells, or cells with a plasma cell marker or phenotype, or the cancer cells in the subject express surface BCMA; or the threshold level of surface BCMA is less than or less than about 60%, less than or less than about 50%, less than or less than about 40%, less than or less than about 30%, less than or less than about 20% or less than or less than about 10% of the plasma cells, or cells with a plasma cell marker or phenotype or the cancer cells in the subject that express surface BCMA.

[0095] In some embodiments of any one of the kits provided herein, the kits further contain: genetically engineered cells expressing a recombinant receptor, optionally wherein the genetically engineered cells are formulated for administration of one or more unit doses to a subject having a disease or condition. In some embodiments, the recombinant receptor specifically binds to a target antigen associated with the disease or condition.

[0096] In some embodiments of any one of the kits provided herein, the instructions specify administering the gamma secretase inhibitor, or one or more unit doses thereof, to a subject having a disease or disorder prior to, concurrently with or after initiation of administration of a dose of the genetically engineered cells to the subject, optionally wherein the genetically engineered cells are formulated for administration of one or more unit doses to a subject having a disease or condition. In some embodiments, the instructions specify administering the gamma secretase inhibitor, or one or more unit doses thereof, to a subject having a disease or disorder prior to initiation of administration of a dose of the genetically engineered cells to the subject. In some embodiments, the instructions specify administering the inhibitor within, or within about, 1 hours, 2 hour, 6 hours, 12 hours, 24 hours, 48 hours, 72 hours, 96 hours or 1 week prior to the initiation of the administration of the cell therapy. In some embodiments, the instructions specify administering the gamma secretase inhibitor, or one or more unit doses thereof, to a subject having a disease or disorder after initiation of administration of a dose of the genetically engineered cells to the subject.

[0097] Provided herein are kits that contain: (a) a gamma secretase inhibitor, optionally wherein the gamma secretase inhibitor is formulated in one or more unit doses; and (b) instructions for administering the gamma secretase inhibitor to a subject after initiation of administration of a cell therapy to a subject, the cell therapy comprising a dose of genetically engineered cells expressing a recombinant receptor.

[0098] Provided herein are kits that contain: (a) a gamma secretase inhibitor, optionally wherein the gamma secretase inhibitor is formulated in one or more unit doses; and (b) instructions for administering the gamma secretase inhibitor to a subject after initiation of administration of a cell therapy to a subject, the cell therapy comprising a dose of genetically engineered cells expressing a recombinant receptor comprising a chimeric antigen receptor (CAR) comprising an antigen-binding domain that binds to surface B cell maturation antigen (BCMA), wherein binding of the CAR to surface BCMA or a measure indicative of function or activity, following exposure to cells expressing surface BCMA, is not reduced or blocked or is not substantially reduced or blocked in the presence of a soluble or shed form of the BCMA, optionally at a concentration or amount of the soluble or shed form of the BCMA corresponding to a concentration or amount present in serum or blood or plasma of the subject or of a multiple myeloma patient, or on average in a patient population for the disease or disorder, or at a concentration or amount of the soluble or shed BCMA at which the binding or measure is reduced or blocked, or is substantially reduced or blocked, for cells expressing a reference anti-BCMA CAR in the same assay.

[0099] In some embodiments of any one of the kits provided herein, initiation of administration of the inhibitor is within, or within about, 1 hours, 2 hour, 6 hours, 12 hours, 24 hours, 48 hours, 72 hours, 96 hours,1 week, 14 days, 21 days, 28 days or more after the initiation of the administration of the dose of genetically engineered cells.

[0100] In some embodiments of any one of the kits provided herein, the instructions specify the inhibitor is for administeration at a time in which: the number of cells of the cell therapy detectable in the blood from the subject is decreased compared to in the subject at a preceding time point after initiation of the administration of the cells; the number of cells of the cell therapy detectable in the blood is less than or less than about 1.5-fold, 2-fold, 3-fold, 4-fold, 5-fold, 10-fold, 50-fold or 100-fold or less the peak or maximum number of the cells of the cell therapy detectable in the blood of the subject after initiation of administration of the

administration of the cells; and/or at a time after a peak or maximum level of the cells of the cell therapy are detectable in the blood of the subject, the number of cells of or derived from the cells detectable in the blood from the subject is less than less than 10%, less than 5%, less than 1% or less than 0.1% of total peripheral blood mononuclear cells (PBMCs) in the blood of the subject.

[0101] In some embodiments of any one of the kits provided herein, the instructions specify the inhibitor is for administration up to 7 days, 14 days, 21 days, 28 days or more after initiation of the administration of the cells.

[0102] In some embodiments of any one of the kits provided herein, the recombinant receptor specifically binds to a target antigen associated with the disease or disorder. In some embodiments, the target antigen is selected from among, B cell maturation antigen (BCMA), carbonic anhydrase 9 (CAIX), Her2/neu (receptor tyrosine kinase erbB2), L1-CAM, CD19, CD20, CD22, mesothelin, CEA, and hepatitis B surface antigen, anti-folate receptor, CD23, CD24, CD30, CD33, CD38, CD44, EGFR, epithelial glycoprotein 2 (EPG-2), epithelial glycoprotein 40 (EPG-40), EPHa2, erb-B2, erb-B3, erb-B4, erbB dimers, EGFR vIII, folate binding protein (FBP), FCRL5, FCRH5, fetal acetylcholine receptor, GD2, GD3, HMW-MAA, IL-22R-alpha, IL-13R-alpha2, kinase insert domain receptor (kdr), kappa light chain, Lewis Y, L1-cell adhesion molecule, (L1-CAM), Melanoma-associated antigen (MAGE)-A1, MAGE-A3, MAGE-A6, Preferentially expressed antigen of melanoma (PRAME), survivin, TAG72, B7-H6, IL-13 receptor alpha 2 (IL-13Ra2), CA9, GD3, HMW-MAA, CD171, G250/CAIX, HLA-AI MAGE Al, HLA-A2 NY-ESO-1, PSCA, folate receptor-a, CD44v6, CD44v7/8, avb6 integrin, 8H9, NCAM, VEGF receptors, 5T4, Foetal AchR, NKG2D ligands, CD44v6, dual antigen, a cancer-testes antigen, mesothelin, murine CMV, mucin 1 (MUC1), MUC16, PSCA, NKG2D, NY-ESO-1, MART-1, gp100, oncofetal antigen, ROR1, TAG72, VEGF-R2, carcinoembryonic antigen (CEA), Her2/neu, estrogen receptor, progesterone receptor, ephrinB2, CD123, c-Met, GD-2, O-acetylated GD2 (OGD2), CE7, Wilms Tumor 1 (WT-1), a cyclin, cyclin A2, CCL-1, CD138, a pathogen-specific antigen and an antigen associated with a universal tag. In some embodiments, the target antigen is Muc1, optionally human Muc1. In some embodiments of any embodiment, the target antigen is a human antigen.

[0103] In some embodiments, the target antigen is BCMA, optionally human BCMA. In some embodiments, binding of the CAR to surface BCMA or a measure indicative of function or activity, following exposure to cells expressing surface BCMA, is not reduced or blocked or

is not substantially reduced or blocked in the presence of a soluble or shed form of the BCMA, optionally at a concentration or amount of the soluble or shed form of the BCMA corresponding to a concentration or amount present in serum or blood or plasma of the subject or of a multiple myeloma patient, or on average in a patient population for the disease or disorder, or at a concentration or amount of the soluble or shed BCMA at which the binding or measure is reduced or blocked, or is substantially reduced or blocked, for cells expressing a reference anti-BCMA CAR in the same assay. In some embodiments, the recombinant receptor does not bind soluble BCMA or binds to soluble BCMA with an affinity lower than the affinity of said recombinant receptor for binding to surface BCMA.

[0104] In some embodiments of any one of the kits provided herein, the recombinant receptor is a transgenic T cell receptor (TCR) or a functional non-T cell receptor. In some embodiments, the recombinant receptor is a chimeric receptor, which optionally is a chimeric antigen receptor (CAR). In some embodiments, the chimeric antigen receptor (CAR) comprises an extracellular antigen-recognition domain that specifically binds to the antigen and an intracellular signaling domain comprising an ITAM. In some embodiments, the intracellular signaling domain comprises and intracellular domain of a CD3-zeta (CD3ζ) chain. In some embodiments, the chimeric antigen receptor (CAR) further comprises a costimulatory signaling region. In some embodiments, the costimulatory signaling region comprises a signaling domain derived from CD28 or 4-1BB, optionally human CD28 or human 4-1BB. In some embodiments, the costimulatory signaling region is a domain derived from 4-1BB, optionally human 4-1BB.

[0105] In some embodiments of any one of the kits provided herein, the genetically engineered cells comprise T cells or NK cells. In some embodiments, the genetically engineered cells comprise T cells. In some embodiments, the T cells are CD4+ and/or CD8+. In some embodiments, the T cells are primary T cells obtained from a subject.

[0106] In some embodiments of any one of the kits provided herein, the gamma secretase inhibitor inhibits or reduces or is capable of inhibiting or reducing cleavage of one or more targets selected from BCMA, Notch 1, Notch 2, Notch 3, Notch 4, Muc1, Ephrin B2, Betaglycan (TGFBR3), CD43, CD44, CSF1R, CXCR1, CXCL16, Delta1, E-cadherein, N-cadherin, HLA-A2, IFNaR2, IL1R1, IL1R2, IL6R or ameloid precursor protein (APP). In some embodiments, the one or more targets the cleavage of which is reduced or inhibited comprises Muc1. In some

embodiments, the one or more targets the cleavage of which is reduced or inhibited comprises BCMA.

[0107] In some embodiments of any one of the kits provided herein, cleavage of one or more targets selected from BCMA, Notch 1, Notch 2, Notch 3, Notch 4, Muc1, Ephrin B2, Betaglycan (TGFBR3), CD43, CD44, CSF1R, CXCR1, CXCL16, Delta1, E-cadherein, N-cadherin, HLA-A2, IFNaR2, IL1R1, IL1R2, IL6R or ameloid precursor protein (APP) is inhibited or reduced or can be inhibited or reduced the gamma secretase inhibitor. In some embodiments, the one or more targets the cleavage of which is reduced or inhibited comprises Muc1. In some embodiments, the one or more targets the cleavage of which is reduced or inhibited comprises BCMA.

[0108] In some embodiments of any one of the kits provided herein, the gamma secretase inhibitor inhibits the target with a half-maximal inhibitory concentration (IC_{50}) of 0.01 nM to 1 μM, 0.01 nM to 100 nM, 0.01 nM to 10 nM, 0.01 nM to 5 nM, 0.01 nM to 1 nM, 0.01 nM to 0.5 nM, 0.01 nM to 0.25 nM, 0.01 nM to 0.1 nM, 0.01 nM to 0.05 nM, 0.05 nM to 1 μM, 0.05 nM to 100 nM, 0.05 nM to 10 nM, 0.05 nM to 5 nM, 0.05 nM to 1 nM, 0.05 nM to 0.5 nM, 0.05 nM to 0.25 nM, 0.05 nM to 0.1 nM, 0.1 nM to 1 μM, 0.1 nM to 100 nM, 0.1 nM to 10 nM, 0.1 nM to 5 nM, 0.1 nM to 1 nM, 0.1 nM to 0.5 nM, 0.1 nM to 0.25 nM, 0.25 nM to 1 μM, 0.25 nM to 100 nM, 0.25 nM to 10 nM, 0.25 nM to 5 nM, 0.25 nM to 1 nM, 0.25 nM to 0.5 nM, 0.5 nM to 1 μM, 0.5 nM to 100 nM, 0.5 nM to 10 nM, 0.5 nM to 5 nM, 0.5 nM to 1 nM, 1 nM to 1 μM, 1 nM to 100 nM, 1 nM to 10 nM, 1 nM to 5 nM, 5 nM to 1 μM, 5 nM to 100 nM, 5 nM to 10 nM, 10 nM to 1 μM, 10 nM to 100 nM or 100 nM to 1 μM.

[0109] In some embodiments of any one of the kits provided herein, the gamma secretase inhibitor inhibits the target with a half-maximal inhibitory concentration (IC_{50}) of less than or less than about 1 μM, 100 nM, 10 nM, 5 nM, 1 nM, 0.5 nM, 0.35 nM, 0.25 nM, 0.1 nM, 0.05 nM or 0.01 nM or less.

[0110] In some embodiments of any one of the kits provided herein, the gamma secretase inhibitor is a peptide inhibitor or non-peptide inhibitor. In some embodiments, the gamma secretase inhibitor is a peptide inhibitor and the peptide inhibitor is selected from among peptide aldehydes derivatives, difluoroketones derivatives, hydroxyethylene dipeptide isotere derivatives, alpha-helical peptide derivatives and dipeptide analogs. In some embodiments, the

gamma secretase inhibitor is a non-peptide inhibitor and the non-peptide inhibitor is a benzodiazepines derivative or a sulfonamides derivative.

[0111] In some embodiments of any one of the kits provided herein, the gamma secretase inhibitor is a transition state inhibitor or non-transition state inhibitor.

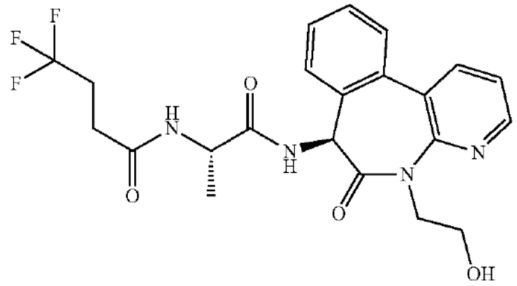
[0112] In some embodiments of any one of the kits provided herein, the gamma secretase inhibitor is a nonsteroidal anti-inflammatory drug.

[0113] In some embodiments of any one of the kits provided herein, the gamma secretase inhibitor is selected from LY3039478, secretase inhibitor I (GSI I) Z-Leu-Leu- Norleucine; γ-secretase inhibitor II (GSI II); γ- secretase inhibitor III (GSI III), N- Benzyloxycarbonyl-Leu-leucinal, N-(2-Naphthoyl)-Val- phenylalaninal; γ-secretase inhibitor III (GSI IV); γ-secretase inhibitor III (GSI V), N- Benzyloxycarbonyl-Leu- phenylalaninal; γ- secretase inhibitor III (GSI VI), l-(S)-endo-N- (1,3,3)- Trimethylbicyclo[2.2.1]hept-2-yl)-4- fluorophenyl Sulfonamide; γ-secretase inhibitor III (GSI VII), Menthyloxycarbonyl-LL-CHO; γ- secretase inhibitor III (GSI IX), (DAPT), N- [N-(3,5- Difluorophenacetyl-L- alanyl)]-S- phenylglycine t- Butyl Ester; γ-secretase inhibitor X (GSI X), { 1 S-Benzyl-4R-[1-(1S- carbamoyl- 2- phenethylcarbamoyl)-lS-3- methylbutylcarb-amoyl]-2R- hydroxy-5- phenylpentyl}carbamic Acid tert-butyl Ester; γ-secretase inhibitor XI (GSI XI), 7-Amino-4-chloro-3- methoxyisocoumarin; γ-secretase inhibitor XII (GSI XII), Z-Ile-Leu-CHO; γ-secretase inhibitor XIII (GSI XIII), Z-Tyr-Ile-Leu- CHO; γ-secretase inhibitor XIV (GSI XIV), Z-Cys(t-Bu)-Ile- Leu-CHO; γ-secretase inhibitor XVI (GSI XVI), N-[N-3,5- Difluorophenacetyl]-L- alanyl-S- phenylglycine Methyl Ester; γ- secretase inhibitor XVII (GSI XVII); γ-secretase inhibitor XIX (GSI XIX), benzo[e][l,4]diazepin-3-yl)-butyramide; γ-secretase inhibitor XX (GSI XX), (S,S)- 2-[2-(3,5- Difluorophenyl)acetylamino]-N-(5-methyl-6-oxo-6,7- dihydro-5H- dibenzo[b,d]azepin-7- yl)propionamide; γ-secretase inhibitor XXI (GSI XXI), (S,S)-2-[2-(3,5- Difluorophenyl)- acetylamino]-N-(l-methyl-2- oxo-5-phenyl-2-,3-dihydro-lH- benzo[e][l,4]diazepin-3- yl)- propionamide; Gamma40 secretase inhibitor I, N-trans-3,5- Dimethoxycinnamoyl-Ile- leucinal; Gamma40 secretase inhibitor II, N-tert-Butyloxycarbonyl- Gly-Val-Valinal Isovaleryl-V V-Sta-A-Sta- OCH3; MK-0752; MRK-003 (Merck); semagacestat/LY450139; RO4929097; PF-03084,014; BMS-708163; MPC-7869 (γ- secretase modifier), YO-01027 (Dibenzazepine), Compound E ([(2S)-2-{[(3,5-Difluorophenyl)acetyl]amino}-N-[(3S)-l-methyl-2-oxo-5-phenyl-2,3-dihydro-lH-l,4-benzodiazepin-3-yl]propanamide], LY411575, L-685,458, BMS-289948 (4-chloro-N-(2,5-

difluorophenyl)-N-((lR)- {4-fluoro-2-[3-(lH-imidazol-l-yl)propyl]phenyl}ethyl)benzenesulfonamide hydrochloride) and BMS-299897 (4-[2-((lR)-l-{[(4-chlorophenyl)sulfonyl]-2,5-difluoroanilino}ethyl)-5- fluorophenyljbutanoic acid).

[0114] In some embodiments of any one of the kits provided herein, the gamma secretase inhibitor is a compound of the structure:

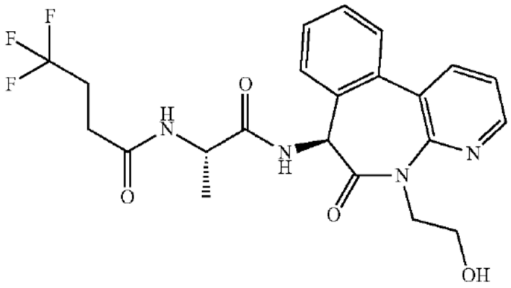
Compound 1



or a pharmaceutically acceptable salt of hydrate thereof.

[0114a] ~~Disclosed herein~~In one embodiment, there is provided a kit, comprising: (a) a compound having the structure:

Compound 1



or a stereoisomer thereof, or a pharmaceutically acceptable salt or hydrate of either of the foregoing; and (b) instructions for the compound that is to be administered to a subject after initiation of administration of a cell therapy to a subject, the cell therapy comprising a dose of immune cells expressing a recombinant receptor.

[0115] In some embodiments of any one of the kits provided herein, the instructions specify administering a dose of genetically engineered cells to a subject having a disease or disorder. In some embodiments, the disease or disorder is a cancer. In some embodiments, the cancer is a B cell malignancy. In some embodiments, the cancer is multiple myeloma, plasmacytoma, a cancer of plasma cell origin and/or a cancer of B cell origin.

[0116] In some embodiments of any one of the kits provided herein, the dose comprises from or from about 1 x 10^5 to 5 x 10^8 total recombinant receptor-expressing cells, total T cells, or total peripheral blood mononuclear cells (PBMCs); the cell therapy comprises the administration of from or from about 1 x 10^5 to 1 x 10^8 total recombinant receptor-expressing cells, total T cells, or total PBMCs; from or from about 5 x 10^5 to 1 x 10^7 total recombinant receptor-expressing cells, total T cells, or total PBMCs; or from or from about 1 x 10^6 to 1 x 10^7 total recombinant receptor-expressing cells, total T cells, or total PBMCs, each inclusive; optionally wherein the instructions specify the administration of one or of a plurality of unit doses comprising the dose of cells and/or a volume corresponding to such one or plurality of unit doses comprising the dose of cells.

[0117] In some embodiments of any one of the kits provided herein, the dose comprises no more than 5 x 10^8 total recombinant receptor-expressing cells, total T cells, or total peripheral blood mononuclear cells (PBMCs); no more than 2.5 x 10^8 total recombinant receptor-expressing cells, total T cells, or total PBMCs; no more than 1 x 10^8 total recombinant receptor-expressing cells, total T cells, or total PBMCs; no more than 1 x 10^7 total recombinant receptor-expressing cells, total T cells, or total PBMCs; no more than 0.5 x 10^7 total recombinant receptor-expressing cells, total T cells, or total PBMCs; no more than 1 x 10^6 total recombinant receptor-expressing cells, total T cells, or total PBMCs; no more than 0.5 x 10^6 total recombinant receptor-expressing cells, total T cells, or total PBMCs; optionally wherein the instructions specify the administration of one or of a plurality of unit doses comprising the dose of cells and/or a volume corresponding to such one or plurality of unit doses comprising the dose of cells.

[0118] In some embodiments of any one of the kits provided herein, the instructions specify administering the inhibitor orally, subcutaneously or intravenously. In some embodiments, the instructions specify administering the inhibitor orally.

[0119] In some embodiments of any one of the kits provided herein, the instructions specify the inhibitor is to be administered at least six times daily, five times daily, four times daily, three times daily, twice daily, once daily, every other day, three times a week, at least once a week, or only one time. In some embodiments, the instructions specify the inhibitor is administered three times a week.

[0120] In some embodiments of any one of the kits provided herein, the instructions specify the administration of the inhibitor is to be carried out in a treatment cycle that is at least or at least about or 14 days, at least or at least about or 21 days or at least or at least about or 28 days.

[0121] In some embodiments of any one of the kits provided herein, the instructions specify the inhibitor is administered, or each administration of the inhibitor is independently administered, in an amount of 0.5 mg to 500 mg, 0.5 mg to 250 mg, 0.5 mg to 100 mg, 0.5 mg to 50 mg, 0.5 mg to 25 mg, 0.5 mg to 10 mg, 0.5 mg to 5.0 mg, 0.5 mg to 2.5 mg, 0.5 mg to 1.0 mg, 1.0 mg to 500 mg, 1.0 mg to 250 mg, 1.0 mg to 100 mg, 1.0 mg to 50 mg, 1.0 mg to 25 mg, 1.0 mg to 10 mg, 1.0 mg to 5.0 mg, 1.0 mg to 2.5 mg, 2.5 mg to 500 mg, 2.5 mg to 250 mg, 2.5 mg to 100 mg, 2.5 mg to 50 mg, 2.5 mg to 25 mg, 2.5 mg to 10 mg, 2.5 mg to 5.0 mg, 5.0 mg to 500 mg, 5.0 mg to 250 mg, 5.0 mg to 100 mg, 5.0 mg to 50 mg, 5.0 mg to 25 mg, 5.0 mg to 10 mg, 10 mg to 500 mg, 10 mg to 250 mg, 10 mg to 100 mg, 10 mg to 50 mg, 10 mg to 25 mg, 25 mg to 500 mg, 25 mg to 250 mg, 25 mg to 100 mg, 25 mg to 50 mg, 50 mg to 500 mg, 50 mg to 250 mg, 50 mg to 100 mg, 100 mg to 500 mg, 100 mg to 250 mg or 250 mg to 500 mg.

[0122] In some embodiments of any one of the kits provided herein, the instructions specify the inhibitor is administered, or each administration of the inhibitor is independently administered, in an amount that is at least or at least about or is or is about 0.5 mg, 1.0 mg, 2.5 mg, 5.0 mg, 10.0 mg, 25 mg, 50 mg, 100 mg, 250 mg or 500 mg.

Brief Description of the Drawings

[0123] FIG. 1 shows expression level of surface B Cell Maturation Antigen (BCMA) on OPM2, MM1s or RPMI-8826 cells co-cultured with different concentrations of a representative gamma secretase inhibitor (LY3039478) after 24 hours of co-culture. The bar near the X-axis suggests the estimated C_{max} and C_{min} for the inhibitor.

[0124] FIG. 2A and FIG. 2B show percent lysis (% killing) of OPM2 cells (FIG. 2A) or RPMI-8226 cells (FIG. 2B) co-cultured with anti-BCMA CAR-expressing T cells in the presence or absence of a representative of gamma secretase inhibitor (LY3039478) after about 150 hours of co-culture. FIG. 2C shows percent lysis (% killing) of OPM2 cells co-cultured with anti-BCMA CAR-expressing T cells in the presence or absence of a representative gamma secretase inhibitor (LY3039478). The bar near the X-axis suggests the estimated C_{max} and C_{min} for the inhibitor.

[0125] FIGS. 3A-3D show the production of IFN-gamma and IL-2 in the supernatant of anti-BCMA CAR-expressing T cells co-cultured with OPM2 (FIGS. 3A and 3C) or RPMI-8226 cells (FIGS. 3B and 3D) in the presence or absence of a representative of gamma secretase inhibitor (LY3039478) after 24 hours of co-culture. FIG. 3E shows the production of IFN-gamma in the supernatant of anti-BCMA CAR-expressing T cells co-cultured with OPM2 cells and different concentrations of a representative gamma secretase inhibitor (LY3039478). The bar near the X-axis suggests the estimated C_{max} and C_{min} for the inhibitor.

[0126] FIGS. 4A and 4B show the cell number of anti-BCMA CAR-expressing T cells originally obtained from a healthy donor (FIG. 4A) and a donor with multiple myeloma (FIG. 4B) after multiple rounds of stimulation with MM1S cells over a period of 10-17 days in the absence of a representative of gamma secretase inhibitor (LY3039478) (vehicle control, square) or in the presence of the inhibitor with a concentration of 1μM (upward triangle) or 0.001μM (downward triangle).

[0127] FIGS. 5A and 5B show the production of IFN-gamma, IL2, and TNF-alpha in the supernatant of anti-BCMA CAR-expressing T cells originally obtained from a healthy donor (FIG. 5A) and a donor with multiple myeloma (FIG. 5B) after a second round of stimulation with MM1S cells at day 4 a in the absence of a representative of gamma secretase inhibitor (LY3039478) (left bar) or in the presence of the inhibitor with a concentration of 0.001μM (middle bar) or 0.001μM (right bar).

[0128] FIG. 6 shows plasma levels of gamma secretase inhibitor LY3039478 at different time points following oral administration of a single dose of LY3039478 (3 mg/kg).

[0129] FIG. 7 shows plasma levels of BCMA over time following oral administration of a single dose of LY3039478 (3 mg/kg) in a mouse xenograft model for human multiple myeloma (RPMI-8226).

[0130] FIG. 8 shows surface BCMA expression, as assessed by flow cytometry, in subcutaneous tumor cells derived from a mouse xenograft model for human multiple myeloma (RPMI-8226) over time following oral administration of a single dose of LY3039478.

[0131] FIGS. 9A-9B show results from the study described in Example 5, assessing the impact of different treatments in a mouse xenograft model for human multiple myeloma. FIG. 9A depicts tumor volume (mm^3) over days during the study. The dashed arrow below the graph indicates the time period during which LY3039478 was administered as described in Example 5.

FIG. 9B shows percent survival among animals treated over the course of 65 days following CAR T injection.

[0132] FIGS. 10A-10B show CD4+ CAR T+ and CD8+ CAR T+ peripheral blood counts over time (days) following CAR T injection (Days Post-CAR T injection) of animals of a human multiple myeloma xenograft mouse model treated under various conditions as described in Example 5. FIG. 10A depicts CD4+ CAR T+ cell counts per μL of blood over time. FIG. 10B depicts CD8+ CAR T+ cell counts per μL of blood over time GSI bar with arrows indicates the timing of LY3039478 (GSI) delivery if administered.

[0133] FIG. 11 shows CD4+ (top) and CD8+ (bottom) CAR+ T cell counts per 1 x 10^6 cells from satellite tumor digests taken on days 7 and 15 following low dose (1e6) CAR T injection in the study described in Example 5.

[0134] FIG. 12A shows serum levels of BCMA on days 7 and 15 following commencement of different treatment regimes as describe in Example 5.

[0135] FIG. 12B shows surface BCMA expression, as assessed by flow cytometry, from tumor digests taken on days 7 and 15 following anti-BCMA CAR T cell injection in mice from various treatment groups as described in Example 5.

[0136] FIG. 13A shows RPMI-8226, OPM2, and MM1.S cell viability over time when treated with chronically stimulated anti-BCMA CAR+ T cells and DMSO (DMSO control) or the gamma secretase inhibitor compound LY3039478 (1 μM GSI), at an effector to target ratio of 0.3:1, or when untreated (RPMI, OPM2, or MM1.S alone).

[0137] FIG. 13B shows the percentage of killing of RPMI-8226, OPM2, and MM1.S cells when treated with chronically stimulated anti-BCMA CAR+ T cells and DMSO (DMSO control) or LY3039478 (1 μM GSI), or when untreated (RPMI, OPM2, or MM1.S alone).

[0138] FIG. 13C shows supernatant IL-2 concentrations following 24 hour incubation of RPMI-8226, OPM2, and MM1.S cells with chronically stimulated anti-BCMA CAR+ T cells and LY3039478 (1 μM GSI) or DMSO (DMSO control) at effector to target ratios of 0.3:1 or 1:1.

[0139] FIG. 14 shows IFNg, IL-2, and TNFa concentrations in supernatant following 24 hour incubation of anti-BCMA CAR+ T cells with BCMA-containing beads in different concentrations of gamma secretase inhibitor LY3039478 or DMSO (veh.).

[0140] FIG. 15A shows BCMA antibody binding capacity (BCMA ABC) on the surface of cells in samples derived from multiple myeloma patients cells before (Before GSI) and after (After GSI) administration of three doses of the small molecule gamma secretase inhibitor (GSI) LY3039478 by mouth as described in Example 8.

[0141] FIG. 15B shows the percentage of plasma cells in patient samples determined to have measurable surface BCMA expression, before (Before GSI) and after (After GSI) administration of three doses of the small molecule gamma secretase inhibitor (GSI) LY3039478 by mouth, as described in Example 8.

Detailed Description

[0142] Provided herein are combination therapies involving administration of an immunotherapy, such as a cell therapy, e.g a T cell therapy, and an inhibitor of gamma secretase. In some aspects, the provided methods involve administration of an immunotherapy or immunotherapeutic agent, such as a composition including cells for adoptive cell therapy, e.g., such as a T cell therapy (e.g. CAR-expressing T cells) and administration of a gamma secretase inhibitor to a subject having a disease or disorder. In some embodiments, the gamma secretase is administered prior to, concurrently with or subsequent to initiation of administration of the immunotherapy, e.g. cell therapy. In some embodiments, the gamma secretase inhibitor inhibits or reduces intramembrane cleavage of a target of a gamma secretase, e.g. BCMA, on a cell (such as a tumor/cancer cell). In some aspects, the target is the same as the target antigen of the cell therapy. In some embodiments, the cell therapy, such as CAR-expressing T cells, comprises an antigen-binding domain that binds to a B Cell Maturation Antigen (BCMA), such as surface BCMA. In some embodiments, the methods further comprise selecting a subject for treatment, wherein the subject has a low expression of surface BCMA on a cell, such as a tumor/cancer cell, e.g. cells of the cancer in the subject that express CD138, surface CD38 or a surface plasma cell marker or are derived from plasma cells. In some embodiments, the cell therapy comprises a recombinant receptor, such as chimeric antigen receptor (CAR), that binds to an antigen other than BCMA. In some embodiments, the methods are carried out in connection with adoptive cell therapy.

[0143] Cell therapies, such as T cell-based therapies, for example, adoptive T cell therapies (including those involving the administration of cells expressing chimeric receptors specific for

a disease or disorder of interest, such as chimeric antigen receptors (CARs) and/or other recombinant antigen receptors, as well as other adoptive immune cell and adoptive T cell therapies) can be effective in the treatment of cancer and other diseases and disorders. The engineered expression of recombinant receptors, such as chimeric antigen receptors (CARs), on the surface of T cells enables the redirection of T-cell specificity. In clinical studies, CAR-T cells, for example anti-CD19 CAR-T cells, have produced durable, complete responses in both leukemia and lymphoma patients (Porter et al. (2015) Sci Transl Med., 7:303ra139; Kochenderfer (2015) J. Clin. Oncol., 33: 540-9; Lee et al. (2015) Lancet, 385:517-28; Maude et al. (2014) N Engl J Med, 371:1507-17).

[0144] In certain contexts, available approaches to adoptive cell therapy may not always be entirely satisfactory. In some contexts, optimal efficacy can depend on the ability of the administered cells to recognize and bind to a target, e.g., target antigen, to traffic, localize to and successfully enter appropriate sites within the subject, tumors, and environments thereof. In some contexts, optimal efficacy can depend on the ability of the administered cells to become activated, expand, to exert various effector functions, including cytotoxic killing and secretion of various factors such as cytokines, to persist, including long-term, to differentiate, transition or engage in reprogramming into certain phenotypic states (such as long-lived memory, less-differentiated, and effector states), to avoid or reduce immunosuppressive conditions in the local microenvironment of a disease, to provide effective and robust recall responses following clearance and re-exposure to target ligand or antigen, and avoid or reduce exhaustion, anergy, peripheral tolerance, terminal differentiation, and/or differentiation into a suppressive state.

[0145] In some embodiments, the exposure and persistence of engineered cells is reduced or declines after administration to the subject. Yet, observations indicate that, in some cases, increased exposure of the subject to administered cells expressing the recombinant receptors (e.g., increased number of cells or duration over time) may improve efficacy and therapeutic outcomes in adoptive cell therapy. Preliminary analyses conducted following the administration of different CD19-targeting CAR-expressing T cells to subjects with various CD19-expressing cancers in multiple clinical trials revealed a correlation between greater and/or longer degree of exposure to the CAR-expressing cells and treatment outcomes. Such outcomes included patient survival and remission, even in individuals with severe or significant tumor burden.

[0146] In some aspects, the provided methods and uses provide for or achieve improved or more durable responses or efficacy as compared to certain alternative methods, such as in particular groups of subjects treated. In some embodiments, the methods are advantageous by virtue of administering T cell therapy, such as a composition including cells for adoptive cell therapy, e.g., such as a T cell therapy (e.g. CAR-expressing T cells), and an inhibitor of gamma secretase.

[0147] The provided methods are based on observations that gamma secretase inhibitor improves one or more functional activities of recombinant receptor-expressing cells specific to BCMA following exposure to BCMA-expressing cells in the presence of a gamma secretase inhibitor. Such functional activities include, for example, antigen-dependent cytolytic activity and/or ability to produce one or more cytokines. In some aspects, the improved activity may be due to increased expression or stabilized expression of surface BCMA on cells (e.g., tumor/cancer cells, for example, multiple myeloma cells). In some aspects, increasing or stabilizing the expression of a target antigen, e.g. BCMA, on target cells improves the outcome for patients who may be or are low for expression of the target antigen, e.g. BCMA, so that sufficient antigen is available for targeting by the cell therapy, e.g. T cell therapy. In some embodiments, the provided embodiments reduce the variability in treatment outcomes among a group of subjects with variable surface expression of the target antigen, e.g. BCMA. In some embodiments of the provided methods, a patient is selected for administration of the provided combination therapy of a gamma secretase inhibitor and an immunotherapy, such as cell therapy (e.g. CAR-T cells) if cells of a cancer in the subject, such as cells that express CD138, surface CD38 or a surface plasma cell marker or are derived from plasma cells, have low expression of surface B cell maturation antigen (BCMA) and/or a level of surface BCMA about a threshold level. Hence, in some aspects, the gamma secretase inhibitor is administered in an amount to stabilize surface expression of BCMA, such as by reducing or inhibiting intramembrane cleavage of BCMA.

[0148] In some embodiments, the provided methods offer advantages in subjects that do not have, or that are not selected based on, high levels of serum or plasma or tumor levels of soluble or shed BCMA. In some embodiments, the patient is not selected for having high serum or plasma or tumor levels of soluble or shed BCMA. In some embodiments, the provided combination therapy involves administration of a cell therapy, e.g. CAR-expressing T cells,

having an antigen-binding domain that binds to surface BCMA in which the antigen-binding domain binds to soluble BCMA with an affinity lower than the affinity of said antigen-binding domain binding to surface BCMA. In some contexts, the provided combination therapy involves administration of a cell therapy, such as recombinant receptor-expressing cells, e.g. CAR-expressing T cells, in which binding to surface BCMA by the recombinant receptor or CAR or a measure indicative of function or activity of the recombinant receptor-expressing cells (e.g. CAR-expressing cells), following exposure to cells expressing surface BCMA, is not reduced or blocked or is not substantially reduced or blocked in the presence of a soluble or shed form of the BCMA.

[0149] In some aspects, such results are achieved while enhancing or improving the activity, in some cases the intrinsic activity, of cells, e.g. T cells, of the cell therapy. In some aspects, this can be achieved in the presence of a gamma secretase inhibitor that exhibits activity for other targets beyond BCMA, such as activity for cleavage of a Notch. An example of such an inhibitor is LY3039478 or compound 1 herein, or stereoisomers, pharmaceutically acceptable salts or hydrates thereof. In some aspects, although notch signaling can regulate T cell differentiation and/or proliferation, observations herein demonstrate enhanced activities of T cells in the presence of an exemplary inhibitor known to target notch signaling, e.g. LY3039478. For example, as shown herein, in an exemplary serial stimulation assay assessing anti-BCMA CAR-T cell activities following repeated stimulation with BCMA-expressing cells, increased CAR+ T cell expansion and/or survival was observed in the presence of the exemplary gamma secretase inhibitor. Such results indicate that the provided combination therapy improves T cell function, including functions related to the expansion, proliferation and persistence of T cells.

[0150] The provided findings indicate that combination therapy of the gamma secretase inhibitor in methods involving cell therapy, such as involving administration of adoptive T cell therapy, achieves improved function of the T cell therapy. In some embodiments, combination of the cell therapy (e.g., administration of engineered T cells) with the gamma secretase inhibitor improves or enhances one or more functions and/or effects of the T cell therapy, including after encounter with antigen, such as persistence, expansion, cytotoxicity, and/or therapeutic outcomes, e.g., ability to kill or reduce the burden of tumor or other disease or target cell. In some aspects, the provided methods increase overall response and/or survival by or more than 1.5-fold, 2.0-fold, 3.0-fold, 4.0-fold, 5.0-fold, 10-fold or more compared to an alternative

treatment, such as compared to a monotherapy involving administration of the T cell therapy (e.g. CAR-T cell) or a gamma secretase inhibitor, or a pharmaceutically acceptable salt of hydrate thereof.

[0151] In some aspects, the provided methods can enhance, increase or potentiate T cell therapy, such as to overcome lack of persistence and/or exhaustion of T cells. In some embodiments, a subject having received administration of a T cell therapy, e.g. CAR-T cell, is monitored for the presence, absence or level of T cells of the therapy in the subject, such as in a biological sample of the subject, e.g. in the blood of the subject. In some embodiments, a gamma secretase inhibitor, or a pharmaceutically acceptable salt of hydrate thereof, is administered to a subject having received the T cell therapy (e.g. CAR-T cells) but in which such cells have weakly expanded and/or are at or below a threshold level in a sample of the subject, e.g. blood sample, at a time when strong or robust expansion of the CAR-T cells in the subject is typically observed in a plurality of subjects administered a T cell therapy (e.g. CAR-T), in some cases, this same T cell therapy (e.g. same CAR-T cells). In some of any such embodiments, the level of engineered, e.g., CAR+, cells in the sample is determined as the number of the cells, e.g., CAR+ cells, per microliter of the sample; in some embodiments, the peak level is the highest such measurement following, optionally over a specified period of time following, administration of the cells or cell therapy to the subject. In some aspects, a gamma secretase inhibitor is administered to a subject having previously received a cell therapy, if, at or about days 1-15 days after initiation of administration of the T cell therapyto the subject, less than 10 cells per μL, such as less than 5 cells per μL or less than 1 cells per μL of cells of the T cell therapy, e.g. CAR-T cells, or a CD8+ or CD3+ subset thereof, are detectable in the blood. In some embodiments, a gamma secretase inhibitor is administered to a subject having previously received a cell therapy, if, at or about day 12-15 cells after initiation of administration of the T cell therapy to the subject, e.g. CAR-T cells, less than 10 cells per μL, such as less than 5 cell pers μL or less than 1 cell per μL of such cells, or a CD8+ or CD3+ subset thereof, are detectable in the blood.

[0152] In certain aspects, the provided methods can enhance, increase or potentiate T cell therapy in subjects in which a peak response (e.g. presence of T cells and/or reduction in tumor burden) to the T cell therapy has been observed, but in which the response, e.g. presence of T cells and/or reduction in tumor burden, has become reduced or is no longer detectable. In some

aspects, a gamma secretase inhibitor is administered to a subject within a week, such as within 1, 2 or 3 days after: (i) peak or maximum level of the cells of the T cell therapy are detectable in the blood of the subject; (ii) the number of cells of the T cell therapy detectable in the blood, after having been detectable in the blood, is not detectable or is reduced, optionally reduced compared to a preceding time point after administration of the T cell therapy; (iii) the number of cells of the T cell therapy detectable in the blood is decreased by or more than 1.5-fold, 2.0-fold, 3.0-fold, 4.0-fold, 5.0-fold, 10-fold or more compared to the peak or maximum number cells of the T cell therapy detectable in the blood of the subject after initiation of administration of the T cell therapy; (iv) at a time after a peak or maximum level of the cells of the T cell therapy are detectable in the blood of the subject, the number of cells of or derived from the T cells detectable in the blood from the subject is less than 10%, less than 5%, less than 1% or less than 0.1% of total peripheral blood mononuclear cells (PBMCs) in the blood of the subject; (v) the subject exhibits disease progression and/or has relapsed following remission after treatment with the T cell therapy; and/or (iv) the subject exhibits increased tumor burden as compared to tumor burden at a time prior to or after administration of the T cells and prior to initiation of administration of the gamma secretase inhibitor, or a pharmaceutically acceptable salt of hydrate thereof.

[0153] In some embodiments, a gamma secretase inhibitor is administered to a subject having received a cell therapy, e.g. CAR-T cells, but who has relapsed following treatment with the T cell therapy, such as at a time in which the response, e.g. presence of T cells and/or reduction in tumor burden, has become reduced or is no longer detectable. In some aspects, relapse may occur due to BCMA antigen loss, which, in some cases, can be due to BCMA antigen downregulation/antigen escape. In some cases, it may be due to cleavage or shedding of BCMA antigen from the cells surface. In some aspects, antigen loss may lead to a reduction in cells presenting a target antigen (e.g., BCMA), thereby diminishing or reducing the activity and/or function of the CAR T cells and/or decreasing the number of CAR T cells, for example, in the blood. In some embodiments, administration of gamma secretase inhibitor can be administered at a time in which the subject has relapsed or is suspected or likely to relapse to cell therapy to reduce or prevent cleavage or shedding of the target antigen (e.g., BCMA) from the cell surface. In such embodiments, administering gamma secretase inhibitor can reinvigorate

or boost CAR T cells following relapse by preventing or reducing cleavage and/or shedding of the target antigen (e.g., BCMA).

[0154] In some embodiments, the provided methods involving combination therapy with a cell therapy and a gamma secretase inhibitor result in genetically engineered cells with increased persistence and/or better potency in a subject to which it is administered compared to administration of the cell therapy in the absence of the gamma secretase inhibitor. In some embodiments, the persistence is increased at least or about at least 1.5-fold, 2-fold, 3-fold, 4-fold, 5-fold, 6-fold, 7-fold, 8-fold, 9-fold, 10-fold, 20-fold, 30-fold, 50-fold, 60-fold, 70-fold, 80-fold, 90-fold, 100-fold or more. In some embodiments, the degree or extent of persistence of administered cells can be detected or quantified after administration to a subject. For example, in some aspects, quantitative PCR (qPCR) is used to assess the quantity of cells expressing the recombinant receptor (e.g., CAR-expressing cells) in the blood or serum or organ or tissue (e.g., disease site) of the subject. In some aspects, persistence is quantified as copies of DNA or plasmid encoding the receptor, e.g., CAR, per microgram of DNA, or as the number of receptor-expressing, e.g., CAR-expressing, cells per microliter of the sample, e.g., of blood or serum, or per total number of peripheral blood mononuclear cells (PBMCs) or white blood cells or T cells per microliter of the sample. In some embodiments, flow cytometric assays detecting cells expressing the receptor generally using antibodies specific for the receptors also can be performed. Cell-based assays may also be used to detect the number or percentage of functional cells, such as cells capable of binding to and/or neutralizing and/or inducing responses, e.g., cytotoxic responses, against cells of the disease or condition or expressing the antigen recognized by the receptor. In any of such embodiments, the extent or level of expression of another marker associated with the recombinant receptor (e.g. CAR-expressing cells) can be used to distinguish the administered cells from endogenous cells in a subject.

[0155] In aspects of the methods provided herein, combining cell therapy with a gamma secretase inhibitor that increases potentcy and/or promotes sustained CAR T cell activity allows administration of a lower dose of CAR T cells to achieve the same therapeutic effect as that seen with a higher dose of CAR T cells without a gamma secretase inhibitor. In some embodiments, the dose of CAR T cells administered when combined with gamma secretase inhibitor can be reduced by 10%, 20%, 30%, 40%, 50%, 60%, 70%, or more of a dose of CAR T cells administered without gamma secretase inhibitor.

[0156] In some embodiments, the methods can be used for treating a disease or condition, such as a cancer, including a B cell malignancy or hematological malignancy. In some aspects, the cancer is multiple myeloma, plasmacytoma, a cancer of plasma cell origin and/or a cancer of B cell origin. In some aspects, such diseases, conditions or malignancies include those in which responses, e.g. complete response (CR), to treatment with the cell therapy alone, such as a composition including cells for adoptive cell therapy, e.g., such as a T cell therapy (*e.g.* CAR-expressing T cells), is relatively low compared to treatment with other T cell therapies or treatment of other diseases or malignacies (e.g. a CR in less than or less than about 60%, less than about 50% or less than about 45% of the subjects so treated).

[0157] In some embodiments, the provided methods reduce or ameliorate a symptom or outcome or burden of the disease or condition to a degree that is greater than the combination of (i) the degree of reduction or amelioration effected by the administration of the gamma secretase inhibitor alone, optionally on average in a population of subjects having the disease or condition, and (ii) the degree of reduction or amelioration by the administration of the T cell therapy alone, optionally on average in a population of subjects having the disease or condition. In some embodiments, the method reduces or ameliorates such symptoms, outcomes or burdens of the disease, e.g. compared to on average in a population of subjects having the disease or condition, by greater than or greater than about 1.5-fold, 2.0-fold, 3.0-fold, 4.0-fold, 5.0-fold, 6.0-fold, 7.0-fold, 8.0-fold, 9.0-fold, 10.0 fold, 20.0-fold, 30.0-fold, 40.0-fold, 50.0-fold or more.

[0158] Also provided are methods for engineering, preparing, and producing the cells (*e.g.* CAR-expressing T cells), compositions containing the cells and/or gamma secretase inhibitor, and kits and devices containing and for using, producing and administering the cells and/or gamma secretase inhibitor, such as in accord with the provided combination therapy methods.

[0159] All publications, including patent documents, scientific articles and databases, referred to in this application are incorporated by reference in their entirety for all purposes to the same extent as if each individual publication were individually incorporated by reference. If a definition set forth herein is contrary to or otherwise inconsistent with a definition set forth in the patents, applications, published applications and other publications that are herein incorporated by reference, the definition set forth herein prevails over the definition that is incorporated herein by reference.

[0160] The section headings used herein are for organizational purposes only and are not to be construed as limiting the subject matter described.

I. COMBINATION THERAPY

[0161] Provided herein are methods for combination therapy for treating a disease or disorder, e.g. a cancer or proliferative disease, that include administering to a subject a combination therapy of 1) a gamma secretase inhibitor and 2) an immunotherapy, e.g. a cell therapy, e.g. T cell therapy, e.g., CAR-expressing cell (e.g., T cell). In some embodiments, the cell therapy is an adoptive immune cell therapy comprising T cells that specifically recognize and/or target an antigen associated with a disease or disorder, e.g. a cancer or proliferative disease. Also provided are combinations and articles of manufacture, such as kits, that contain a composition comprising the immunotherapy, e.g. T cell therapy, and/or a composition comprising the gamma secretase inhibitor, and uses of such compositions and combinations to treat or prevent diseases, conditions, and disorders, including cancers.

[0162] In some embodiments, such methods can include administration of the gamma secretase inhibitor prior to, simultaneously with, during, during the course of (including once and/or periodically during the course of), and/or subsequently to, the administration (e.g., initiation of administration) of the immunotherapy, e.g. cell therapy (*e.g.* CAR-expressing T cells). In some embodiments, the administrations can involve sequential or intermittent administrations of the gamma secretase inhibitor and cell therapy.

[0163] In some embodiments, the cell therapy is adoptive cell therapy. In some embodiments, the cell therapy is or comprises a tumor infiltrating lymphocytic (TIL) therapy, a transgenic TCR therapy or a recombinant-receptor expressing cell therapy (optionally T cell therapy), which optionally is a chimeric antigen receptor (CAR)-expressing cell therapy. In some embodiments, the therapy is a B cell targeted therapy. In some embodiments, the therapy targets B cell maturation antigen (BCMA). In some embodiments, the therapy targets CD19. In some embodiments, the therapy targets cell surface associated Mucin 1 (MUC1). In some embodiments, the cells and dosage regimens for administering the cells can include any as described in the following subsection A under "Administration of Cells." In some embodiments, the dosage regimens for administering the gamma secretase inhibitor can include any as described in the following subsection B under "Administration of gamma secretase inhibitor."

[0164] In some embodiments, the immunotherapy, e.g. cell therapy (*e.g.* CAR-expressing T cells), and gamma secretase inhibitor are provided as pharmaceutical compositions for administration to the subject. In some embodiments, the pharmaceutical compositions contain therapeutically effective amounts of one or both of the agents for combination therapy, *e.g.*, T cells for adoptive cell therapy and a gamma secretase inhibitor as described. In some embodiments, the agents are formulated for administration in separate pharmaceutical compositions. In some embodiments, any of the pharmaceutical compositions provided herein can be formulated in dosage forms appropriate for each route of administration.

[0165] In some embodiments, the combination therapy, which includes administering an immunotherapy, e.g. cell therapy, including engineered cells, such as CAR-T cell therapy, and the gamma secretase inhibitor is administered to a subject or patient having a disease or condition to be treated (*e.g.* cancer) or at risk for having the disease or condition (*e.g.* cancer). In some aspects, the methods treat, *e.g.*, ameliorate one or more symptom of, the disease or condition, such as by lessening tumor burden in a cancer expressing an antigen recognized by the immunotherapy or immunotherapeutic agent, e.g. recognized by an engineered T cell.

[0166] In some embodiments, the disease or condition that is treated can be any in which expression of an antigen is associated with and/or involved in the etiology of a disease condition or disorder, *e.g.* causes, exacerbates or otherwise is involved in such disease, condition, or disorder. Exemplary diseases and conditions can include diseases or conditions associated with malignancy or transformation of cells (*e.g.* cancer), autoimmune or inflammatory disease, or an infectious disease, *e.g.* caused by bacterial, viral or other pathogens. Exemplary antigens, which include antigens associated with various diseases and conditions that can be treated, include any of antigens described herein. In particular embodiments, the recombinant receptor expressed on engineered cells of a combination therapy, including a chimeric antigen receptor or transgenic TCR, specifically binds to an antigen associated with the disease or condition.

[0167] In some embodiments, the disease or condition is a cancer or proliferative disease that expresses BCMA. In some embodiments, the provided methods employ a recombinant receptor-expressing cell (*e.g.* CAR-T cell) that targets BCMA. In particular embodiments, the disease or condition is a multiple myeloma. In some cases, the multiple myeloma is a relapsed or refractory multiple myeloma.

[0168] Among the diseases to be treated is any disease or disorder associated with BCMA or any disease or disorder in which BCMA is specifically expressed and/or in which BCMA has been targeted for treatment (also referred to herein interchangeably as a "BCMA-associated disease or disorder"). Cancers associated with BCMA expression include hematologic malignancies such as multiple myeloma, Waldenstrom macroglobulinemia, as well as both Hodgkin's and non-Hodgkin's lymphomas. *See* Coquery *et al.*, *Crit Rev Immunol*., 2012, 32(4):287-305 for a review of BCMA. Since BCMA has been implicated in mediating tumor cell survival, it is a potential target for cancer therapy. Chimeric antigen receptors containing anti-BCMA antibodies, including mouse anti-human BCMA antibodies and human anti-human antibodies, and cells expressing such chimeric receptors have been previously described. *See* Carpenter *et al.*, *Clin Cancer Res*., 2013, 19(8):2048-2060, WO 2016/090320, WO2016090327, WO2010104949A2 and WO2017173256. Exemplary CARS containing anti-BCMA antibodies are described herein.

[0169] In some embodiments, prior to the initiation of administration of the engineered cells, such as anti-BCMA CAR, the subject has received one or more prior therapies. In some embodiments, the subject has received at least 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19 or 20 or more prior therapies. In some embodiments, the subject has received at least 3, 4, 5, 6, 7, 8, 9, 10 or more prior therapies. In some aspects, the subject has relapsed following, or has been treatment refractory to, one or more of, for example, each, individually, of the one or more prior therapies. In some aspects, the prior therapies include treatment with autologous stem cell transplant (ASCT); an immunomodulatory agent, e.g. an IMiD; a proteasome inhibitor; and an anti-CD38 antibody; unless the subject was not a candidate for or was contraindicated for one or more of the therapies. In some embodiments, the immunomodulatory agent, e.g. an IMiD, is selected from among thalidomide, lenalidomide or pomalidomide. In some embodiments, the proteasome inhibitor is selected from among bortezomib, carfilzomib or ixazomib. In some embodiments, the anti-CD38 antibody is or comprises daratumumab. In some embodiments, the subject must have undergone at least 2 consecutive cycles of treatment for each regimen unless progressive disease was the best response to the regimen. In some aspects, a subject for treatment with the method herein has a myeloma that has relapsed or is treatment refractory with greater than 10% CD138+ malignant plasma cells immunohistochemistry (IHC) on bone marrow core biopsy, either following ASCT

or, if the subject has not undergone ASCT, the subject is transplant ineligible, e.g. due to age comorbidity, patient choice, state of disease and/or discretion of physician, and has disease that has persisted after more than four cycles of induction therapy and/or that is refractory to or is not tolerant to therapy with a proteasome inhibitor and immunomodulatory drug, e.g. IMiD .

[0170] In some embodiments, the disease or condition is a tumor, such as a solid tumor, lymphoma, leukemia, blood tumor, metastatic tumor, or other cancer or tumor type.

[0171] In some embodiments, the antigen associated with the disease or disorder is selected from the group consisting of B cell maturation antigen (BCMA), RORl, Her2, Ll-CAM, CD19, CD20, CD22, mesothelin, CEA, and hepatitis B surface antigen, anti-folate receptor, CD23, CD24, CD30, CD33, CD38, CD44, EGFR, EGP-2, EGP-4, EPHa2, ErbB2, 3, or 4, erbB dimers, EGFR vIII, FBP, FCRL5, FCRH5, fetal acethycholine e receptor, GD2, GD3, HMW-MAA, IL-22R-alpha, IL-13R-alpha2, kdr, kappa light chain, Lewis Y, L1-cell adhesion molecule, (L1-CAM), Melanoma-associated antigen (MAGE)-A1, MAGE-A3, MAGE-A6, Preferentially expressed antigen of melanoma (PRAME), survivin, EGP2, EGP40, TAG72, B7-H6, IL-13 receptor a2 (IL-13Ra2), CA9, GD3, HMW-MAA, CD171, G250/CAIX, HLA-AI MAGE Al, HLA-A2 NY-ESO-1, PSCA, folate receptor-a, CD44v6, CD44v7/8, avb6 integrin, 8H9, NCAM, VEGF receptors, 5T4, Foetal AchR, NKG2D ligands, CD44v6, dual antigen, and an antigen associated with a universal tag, a cancer-testes antigen, mesothelin, MUC1, MUC16, PSCA, NKG2D Ligands, NY-ESO-1, MART-1, gp100, oncofetal antigen, ROR1, TAG72, VEGF-R2, carcinoembryonic antigen (CEA), prostate specific antigen, PSMA, Her2/neu, estrogen receptor, progesterone receptor, ephrinB2, CD123, c-Met, GD-2, O-acetylated GD2 (OGD2), CE7, Wilms Tumor 1 (WT-1), a cyclin, cyclin A2, CCL-1, CD138, and a pathogen-specific antigen. In some embodiments, the antigen is associated with or is a universal tag. In some embodiments, the antigen associated with the disease or disorder is selected from the group consisting of Notch 1, Notch 2, Notch 3, Notch 4, cell surface associated Mucin 1 (MUC1), Ephrin B2, Betaglycan (TGFBR3), CD43, CD44, CSF1R, CX3CR1, CXCL16, Delta1, E-cadherin, N-cadherin, HLA-A2, IFNaR2, IL1R1, IL1R2, IL6R, and amyloid precursor protein (APP).

[0172] In some embodiments, the disease or disorder is a B cell-related disorder. In some embodiments, the disease or disorder is one or more diseases or conditions from among glioblastoma, lymphomatoid granulomatosis, post-transplant lymphoproliferative disorder, an

immunoregulatory disorder, heavy-chain disease, primary or immunocyte-associated amyloidosis, or monoclonal gammopathy of undetermined significance.

[0173] In some embodiments, the disease or disorder is an autoimmune disease or disorder. Such autoimmune diseases or disorder include, but are not limited to, systemic lupus erythematosus (SLE), lupus nephritis, inflammatory bowel disease, rheumatoid arthritis (*e.g.*, juvenile rheumatoid arthristis), ANCA associated vasculitis, idiopathic thrombocytopenia purpura (ITP), thrombotic thrombocytopenia purpura (TTP), autoimmune thrombocytopenia, Chagas' disease, Grave's disease, Wegener's granulomatosis, poly-arteritis nodosa, Sjogren's syndrome, pemphigus vulgaris, scleroderma, multiple sclerosis, psoriasis, IgA nephropathy, IgM polyneuropathies, vasculitis, diabetes mellitus, Reynaud's syndrome, anti-phospholipid syndrome, Goodpasture's disease, Kawasaki disease, autoimmune hemolytic anemia, myasthenia gravis, or progressive glomerulonephritis.

[0174] In some embodiments, the disease or disorder is a B cell malignancy. In some embodiments, the cancer (*e.g.*, a BCMA-expressing cancer) is a lymphoma, a leukemia, or a plasma cell malignancy. Lymphomas contemplated herein include, but are not limited to, Burkitt lymphoma (*e.g.*, endemic Burkitt's lymphoma or sporadic Burkitt's lymphoma), non-Hodgkin's lymphoma (NHL), Hodgkin's lymphoma, Waldenstrom macroglobulinemia, follicular lymphoma, small non-cleaved cell lymphoma, mucosa-associated lymphatic tissue lymphoma (MALT), marginal zone lymphoma, splenic lymphoma, nodal monocytoid B cell lymphoma, immunoblastic lymphoma, large cell lymphoma, diffuse mixed cell lymphoma, pulmonary B cell angiocentric lymphoma, small lymphocytic lymphoma, primary mediastinal B cell lymphoma, lymphoplasmacytic lymphoma (LPL), or mantle cell lymphoma (MCL). Leukemias contemplated herein, include, but are not limited to, chronic lymphocytic leukemia (CLL), plasma cell leukemia or acute lymphocytic leukemia (ALL). Also contemplated herein are plasma cell malignancies including, but not limited to, multiple myeloma (*e.g.*, non-secretory multiple myeloma, smoldering multiple myeloma) or plasmacytoma. In some embodiments the disease or condition is multiple myeloma, such as relapsed and/or refractory multiple myeloma. Among the diseases, disorders or conditions that can be treated include, but are not limited to, neuroblastoma, renal cell carcinoma, colon cancer, colorectal cancer, breast cancer, epithelial squamous cell cancer, melanoma, myeloma (*e.g.*, multiple myeloma), stomach cancer, brain cancer, lung cancer, pancreatic cancer, cervical cancer, ovarian cancer, liver cancer, bladder

cancer, prostate cancer, testicular cancer, thyroid cancer, uterine cancer, adrenal cancer and head and neck cancer.

[0175] In some embodiments, the cancer or proliferative disease is a B cell malignancy or hematological malignancy. In some embodiments the cancer or proliferative disease is acute lymphoblastic leukemia (ALL), non-Hodgkin's lymphoma (NHL), or chronic lymphocytic leukemia (CLL). In some embodiments, the cancer is CLL. In some embodiments, the methods can be used to treat a myeloma, a lymphoma or a leukemia. In some embodiments, the methods can be used to treat a non-Hodgkin lymphoma (NHL), an acute lymphoblastic leukemia (ALL), a chronic lymphocytic leukemia (CLL), a diffuse large B-cell lymphoma (DLBCL), acute myeloid leukemia (AML), or a myeloma, *e.g.*, a multiple myeloma (MM). In some embodiments, the methods can be used to treat a MM or a DBCBL.

[0176] In some embodiments, the methods can be used to treat a non-hematologic cancer, such as a solid tumor. In some embodiments, the methods can be used to treat bladder, lung, brain, melanoma (*e.g.* small-cell lung, melanoma), breast, cervical, ovarian, colorectal, pancreatic, endometrial, esophageal, kidney, liver, prostate, skin, thyroid, or uterine cancers. In some embodiments, the cancer or proliferative disease is pancreatic cancer, bladder cancer, colorectal cancer, breast cancer, prostate cancer, renal cancer, hepatocellular cancer, lung cancer, ovarian cancer, cervical cancer, pancreatic cancer, rectal cancer, thyroid cancer, uterine cancer, gastric cancer, esophageal cancer, head and neck cancer, melanoma, neuroendocrine cancers, CNS cancers, brain tumors, bone cancer, or soft tissue sarcoma.

[0177] In some embodiments, the disease or condition is an infectious disease or condition, such as, but not limited to, viral, retroviral, bacterial, and protozoal infections, immunodeficiency, Cytomegalovirus (CMV), Epstein-Barr virus (EBV), adenovirus, BK polyomavirus. In some embodiments, the disease or condition is an autoimmune or inflammatory disease or condition, such as arthritis, *e.g.*, rheumatoid arthritis (RA), Type I diabetes, systemic lupus erythematosus (SLE), inflammatory bowel disease, psoriasis, scleroderma, autoimmune thyroid disease, Graves disease, Crohn's disease, multiple sclerosis, asthma, and/or a disease or condition associated with transplant.

[0178] For the prevention or treatment of disease, the appropriate dosage of gamma secretase inhibitor and/or immunotherapy, such as a cell therapy (*e.g.* CAR-expressing T cells), may depend on the type of disease to be treated, cells and/or recombinant receptors expressed on

the cells, the severity and course of the disease, route of administration, whether the gamma secretase inhibitor and/or the T cell therapy are administered for preventive or therapeutic purposes, previous therapy, frequency of administration, the subject's clinical history and response to the cells, and the discretion of the attending physician. The compositions and cells are in some embodiments suitably administered to the subject at one time or over a series of treatments. Exemplary dosage regimens and schedules for the provided combination therapy are described.

[0179] In some embodiments, the immunotherapy, such as cell therapy (e.g.CAR-T cell therapy) and the gamma secretase inhibitor are administered as part of a further combination treatment, which can be administered simultaneously with or sequentially to, in any order, another therapeutic intervention. In some contexts, the immunotherapy, such as cell therapy, e.g. engineered T cells, such as CAR-expressing T cells, are co-administered with another therapy sufficiently close in time such that the cell therapy enhances the effect of one or more additional therapeutic agents, or vice versa. In some embodiments, the cells are administered prior to the one or more additional therapeutic agents. In some embodiments, the cell therapy, e.g. engineered T cells, such as CAR-expressing T cells, are administered after the one or more additional therapeutic agents. In some embodiments, the combination therapy methods further include a lymphodepleting therapy, such as administration of a chemotherapeutic agent. In some embodiments, the combination therapy further comprises administering another therapeutic agent, such as an anti-cancer agent, a checkpoint inhibitor, or another immune modulating agent. In some embodiments, the combination therapy further comprises administering a steroid, such as dexamethasone. Uses include uses of the combination therapies in such methods and treatments, and uses of such compositions in the preparation of a medicament in order to carry out such combination therapy methods. In some embodiments, the methods and uses thereby treat the disease or condition or disorder, such as a cancer or proliferative disease, in the subject.

[0180] Prior to, during or following administration of the immunotherapy (*e.g.* T cell therapy, such as CAR-T cell therapy) and/or a gamma secretase inhibitor, the biological activity of the T cell therapy, *e.g.* the biological activity of the engineered cell populations, in some embodiments is measured, *e.g.*, by any of a number of known methods. Parameters to assess include the ability of the engineered cells to destroy target cells, persistence and other measures

of T cell activity, such as measured using any suitable method known in the art, such as assays described further below in Section III. In some embodiments, the biological activity of the cells, *e.g.*, T cells administered for the T cell based therapy, is measured by assaying cytotoxic cell killing, expression and/or secretion of one or more cytokines, proliferation or expansion, such as upon restimulation with antigen. In some aspects the biological activity is measured by assessing the disease burden and/or clinical outcome, such as reduction in tumor burden or load. In some embodiments, administration of one or both agents of the combination therapy and/or any repeated administration of the therapy, can be determined based on the results of the assays before, during, during the course of or after administration of one or both agents of the combination therapy.

[0181] In some embodiments, the combined effect of the gamma secretase inhibitor in combination with the cell therapy can be synergistic compared to treatments involving only the gamma secretase inhibitor or monotherapy with the cell therapy. For example, in some embodiments, the methods provided herein result in an increase or an improvement in a desired therapeutic effect, such as an increased or an improvement in the reduction or inhibition of one or more symptoms associated with cancer.

[0182] In some embodiments, the gamma secretase inhibitor increases the expansion or proliferation of the engineered T cells, such as CAR T-Cells. In some embodiments, the increase in expansion or proliferation is observed in vivo upon administration to a subject. In some embodiments, the increase in the number of engineered T cells, e.g. CAR-T cells, is increased by greater than or greater than about 1.2-fold, 1.5-fold, 2.0-fold, 3.0-fold, 4.0-fold, 5.0-fold, 6.0-fold, 7.0-fold, 8.0-fold, 9.0-fold, 10.0 fold or more. In some embodiments, the gamma secretase inhibitor prolongs and/or sustains the activity and/or function of engineered T cells, such as CAR T-Cells.

A. ADMINISTRATION OF AN IMMUNOTHERAPY

[0183] In some embodiments, an immunotherapy that binds to or targets an antigen on an immune cell and/or that is involved in a disease or disorder is administered in accord with the provided combination therapy methods. In particular embodiments, the immunotherapy binds to and/or recognizes an antigen that is expressed on or in a cell or tissue. In certain embodiments, the antigen is expressed on or in a cell or tissue. In particular embodiments, the antigen is expressed on the surface of a cell. In some embodiments, the cell is a B cell, a T cell and/or a

tumor or cancer cell. In particular embodiments, the antigen is expressed in or on a circulating cell. In some embodiments, the antigen is expressed on the surface of a circulating cell.

[0184] In particular embodiments, the immunotherapy binds to and/or recognizes at least one antigen associated with a disease. Among the diseases, conditions, and disorders that may be treated in human subjects with the immunotherapy are tumors, including solid tumors, hematologic malignancies, and melanomas, and including localized and metastatic tumors, infectious diseases, such as infection with a virus or other pathogen, e.g., HIV, HCV, HBV, CMV, HPV, and parasitic disease, and autoimmune and inflammatory diseases. In some embodiments, the disease or condition is a tumor, cancer, malignancy, neoplasm, or other proliferative disease or disorder. Such diseases include but are not limited to leukemia, lymphoma, e.g., chronic lymphocytic leukemia (CLL), ALL, non-Hodgkin's lymphoma, acute myeloid leukemia, multiple myeloma, refractory follicular lymphoma, mantle cell lymphoma, indolent B cell lymphoma, B cell malignancies, cancers of the colon, lung, liver, breast, prostate, ovarian, skin, melanoma, bone, and brain cancer, ovarian cancer, epithelial cancers, renal cell carcinoma, pancreatic adenocarcinoma, Hodgkin lymphoma, cervical carcinoma, colorectal cancer, glioblastoma, neuroblastoma, Ewing sarcoma, medulloblastoma, osteosarcoma, synovial sarcoma, and/or mesothelioma.

[0185] In some embodiments, the immunotherapy binds to at least one antigen that is a target for cleavage by a gamma secretase. In some embodiments, the antigen is BCMA or is mucin 1 (Muc1).

1. Therapeutic Agents, e.g. antibodies

[0186] In certain embodiments, the immunotherapy is a therapeutic agent, such as an antibody or antigen-binding fragment thereof. In some aspects, the antibody or antigen-binding fragment is a bispecific antibody.

[0187] In some aspects, the immunotherapy is a therapeutic agent that is or contains an immune system activator or stimulator. In certain embodiments, the immune system stimulator is an agent or therapy that activates at least one immune cell. In some embodiments, the immune cell is a T cell. In certain embodiments, the immune cell activator is IL-2, e.g., Proleukin; rhu-IFN-alpha-2a and/or rhu-IFN-alpha-2b, e.g., Pegasys, Roferon-A, Intron-A, and PEG intron; Anti-CD3 monoclonal antibody, e.g., Muromonab-CD3 and/or Orthoclone OKT 3; TGN-1412; and/or Blinatumomab, e.g., anti-CD3xCD3 BiTE.

[0188] In some embodiments, the immunotherapy is or contains a T cell-engaging therapy that is or comprises a binding molecule capable of binding to a surface molecule expressed on a T cell. In some embodiments, the surface molecule is an activating component of a T cell, such as a component of the T cell receptor complex. In some embodiments, the surface molecule is CD3 or is CD2. In some embodiments, the T cell-engaging therapy is or comprises an antibody or antigen-binding fragment. In some embodiments, the T cell-engaging therapy is a bispecific antibody containing at least one antigen-binding domain binding to an activating component of the T cell (*e.g.* a T cell surface molecule, *e.g.* CD3 or CD2) and at least one antigen-binding domain binding to a surface antigen on a target cell, such as a surface antigen on a tumor or cancer cell, for example any of the listed antigens as described herein, *e.g.* BCMA or Muc1. In some embodiments, the simultaneous or near simultaneous binding of such an antibody to both of its targets can result in a temporary interaction between the target cell and T cell, thereby resulting in activation, *e.g.* cytotoxic activity, of the T cell and subsequent lysis of the target cell.

[0189] Among such exemplary bispecific antibody T cell-engagers are bispecific T cell engager (BiTE) molecules, which contain tandem scFv molecules fused by a flexible linker (see e.g. Nagorsen and Bauerle, Exp Cell Res 317, 1255-1260 (2011); tandem scFv molecules fused to each other via, e.g. a flexible linker, and that further contain an Fc domain composed of a first and a second subunit capable of stable association (WO2013026837); diabodies and derivatives thereof, including tandem diabodies (Holliger et al, Prot Eng 9, 299-305 (1996); Kipriyanov et al, J Mol Biol 293, 41-66 (1999)); dual affinity retargeting (DART) molecules that can include the diabody format with a C-terminal disulfide bridge; or triomabs that include whole hybrid mouse/rat IgG molecules (Seimetz et al, Cancer Treat Rev 36, 458-467 (2010). In some embodiments, the T-cell engaging therapy is blinatumomab or AMG 330. Any of such T cell-engagers can be used in the provided methods, compositions or combinations.

[0190] In some embodiments, the therapeutic agent is a BCMA-specific binding capable of binding to BCMA and at least one additional antigen. In some embodiments, the at least one additional antigen CD19, CD20, CD22, CD33, CD38, CS1, ROR1, GPC3, CD123, IL-13R, CD138, c-Met, EGFRvIII, GD-2, NY-ESO-1, MAGE A3, or glycolipid F77. Examples of such bispecific antibodies are described in WO2017/025038.

[0191] The therapeutic agent, such as immunotherapy, can be administered by any suitable means, for example, by bolus infusion, by injection, e.g., intravenous or subcutaneous injections,

intraocular injection, periocular injection, subretinal injection, intravitreal injection, trans-septal injection, subscleral injection, intrachoroidal injection, intracameral injection, subconjectval injection, subconjuntival injection, sub-Tenon's injection, retrobulbar injection, peribulbar injection, or posterior juxtascleral delivery. In some embodiments, the therapeutic agent, such as immunotherapy, is administered by parenteral, intrapulmonary, and intranasal, and, if desired for local treatment, intralesional administration. Parenteral infusions include intramuscular, intravenous, intraarterial, intraperitoneal, intrathoracic, intracranial, or subcutaneous administration.

[0192] In certain embodiments, one or more doses of a T cell engaging therapy and/or an immune system stimulator are administered. In particular embodiments, between 0.001 μg – 5,000 μg of the T cell engaging therapy and/or an immune system stimulator are administered. In particular embodiments, between 0.001 μg to 1,000 μg, 0.001 μg to1 μg, 0.01 μg to1 μg, 0.1 μg to10 μg, 0.01 μg to1 μg, 0.1 μg to 5 μg, 0.1 μg to 50 μg, 1 μg to 100 μg, 10 μg to 100 μg, 50 μg to 500 μg, 100 μg to 1,000 μg, 1,000 μg to 2,000 μg, or 2,000 μg to 5,000 μg of the T cell engaging therapy is administered. In some embodiments, the dose of the T cell engaging therapy is or includes between 0.01 μg/kg and 100 mg/kg, between 0.1 μg/kg and 10 μg/kg, between 10 μg/kg and 50 μg/kg, between 50 μg/kg and 100 μg/kg, between 0.1 mg/kg and 1 mg/kg, between 1 mg/kg and 10 mg/kg, between 10 mg/kg and 100 mg/kg, between 100 mg/kg and 500 mg/kg, between 200 mg/kg and 300 mg/kg, between 100 mg/kg and 250 mg/kg, between 200 mg/kg and 400 mg/kg, between 250 mg/kg and 500 mg/kg, between 250 mg/kg and 750 mg/kg, between 50 mg/kg and 750 mg/kg, between 1 mg/kg and 10 mg/kg, or between 100 mg/kg and 1,000 mg/kg (amount of the lymphodepleting agent over body weight). In some embodiments, the dose of the T cell engaging therapy is or is about 0.1 μg/kg, 0.5 μg/kg, 1 μg/kg, 5 μg/kg, 10 μg/kg, 20 μg/kg, 30 μg/kg, 40 μg/kg, 50 μg/kg, 60 μg/kg, 70 μg/kg, 80 μg/kg, 90 μg/kg, 0.1 mg/kg, 0.5 mg/kg, 1 mg/kg, 2.5 mg/kg, 5 mg/kg, 10 mg/kg, 15 mg/kg, 20 mg/kg, 25 mg/kg, 30 mg/kg, 35 mg/kg, 40 mg/kg, 45 mg/kg, 50 mg/kg, 55 mg/kg, 60 mg/kg, 65 mg/kg, 70 mg/kg, 75 mg/kg, 80 mg/kg, 85 mg/kg, 90 mg/kg, 95 mg/kg, 100 mg/kg, 200 mg/kg, 300 mg/kg, 400 mg/kg, 500 mg/kg, 600 mg/kg, 700 mg/kg, 800 mg/kg, 900 mg/kg, or 1,000 mg/kg,. In particular embodiments, the T cell engaging therapy is administered orally, intravenously, intraperitoneally, transdermally, intrathecally, intramuscularly, intranasally, transmucosally, subcutaneously, or rectally.

[0193] In some embodiments, the dose of CAR T cells administered can be reduced by 10%, 20%, 30%, 40%, 50%, 60%, 70%, or more of a dose of CAR T cells when administered in combination with a gamma secretase inhibitor. In some embodiments, the dose of CAR T cells administered when combined with gamma secretase inhibitor can be reduced by 10%, 20%, 30%, 40%, 50%, 60%, 70%, or more of a dose of CAR T cells administered without gamma secretase inhibitor.

2. Cell therapy

[0194] In some embodiments of the methods, compositions, combinations, kits and uses provided herein, the combination therapy includes administering to a subject an immune cell therapy, such as a T cell therapy (*e.g.* CAR-expressing T cells). Administration of such therapies can be initiated prior to, subsequent to, simultaneously with administration of the gamma secretase inhibitor as described. Exemplary cell therapies and engineered cells for use in the provided methods, compositions, combinations, kits and uses provided herein are described in Section II.

[0195] In some embodiments, the cell-based therapy is or comprises administration of cells, such as immune cells, for example T cell or NK cells, that target a molecule expressed on the surface of a lesion, such as a tumor or a cancer. In some embodiments, the immune cells express a T cell receptor (TCR) or other antigen-binding receptor. In some embodiments, the immune cells express a recombinant receptor, such as a transgenic TCR or a chimeric antigen receptor (CAR). In some embodiments, the cells are autologous to the subject. In some embodiments, the cells are allogeneic to the subject.

[0196] In some aspects, the cell therapy is or comprises a tumor infiltrating lymphocytic (TIL) therapy, a transgenic TCR therapy or a T cell therapy comprising genetically engineered cells, such as a recombinant-receptor expressing cell therapy. In some embodiments, the recombinant receptor specifically binds to a ligand, such as one associated with a disease or condition, *e.g.* associated with or expressed on a cell of a tumor or cancer. In some embodiments, the cell therapy includes administering T cells engineered to express a chimeric antigen receptor (CAR).

[0197] In some embodiments, the provided cells express and/or are engineered to express receptors, such as recombinant receptors, including those containing ligand-binding domains or binding fragments thereof, and T cell receptors (TCRs) and components thereof, and/or

functional non-TCR antigen receptors, such as chimeric antigen receptors (CARs). In some embodiments, the recombinant receptor contains an extracellular ligand-binding domain that specifically binds to an antigen. In some embodiments, the recombinant receptor is a CAR that contains an extracellular antigen-recognition domain that specifically binds to an antigen. In some embodiments, the ligand, such as an antigen, is a protein expressed on the surface of cells. In some embodiments, the CAR is a TCR-like CAR and the antigen is a processed peptide antigen, such as a peptide antigen of an intracellular protein, which, like a TCR, is recognized on the cell surface in the context of a major histocompatibility complex (MHC) molecule.

[0198] Among the engineered cells, including engineered cells containing recombinant receptors, are described furtherherein. Exemplary recombinant receptors, including CARs and recombinant TCRs, as well as methods for engineering and introducing the receptors into cells, include those described, for example, in international patent application publication numbers WO200014257, WO2013126726, WO2012/129514, WO2014031687, WO2013/166321, WO2013/071154, WO2013/123061 U.S. patent application publication numbers US2002131960, US2013287748, US20130149337, U.S. Patent Nos.: 6,451,995, 7,446,190, 8,252,592, 8,339,645, 8,398,282, 7,446,179, 6,410,319, 7,070,995, 7,265,209, 7,354,762, 7,446,191, 8,324,353, and 8,479,118, and European patent application number EP2537416,and/or those described by Sadelain *et al.*, Cancer Discov. 2013 April; 3(4): 388–398; Davila *et al.* (2013) *PLoS ONE* 8(4): e61338; Turtle *et al.*, *Curr. Opin. Immunol.*, 2012 October; 24(5): 633-39; Wu *et al.*, *Cancer*, 2012 March 18(2): 160-75. In some aspects, the genetically engineered antigen receptors include a CAR as described in U.S. Patent No.: 7,446,190, and those described in International Patent Application Publication No.: WO/2014055668 A1.

[0199] In some embodiments, the combination therapy includes administration to a subject cells, e.g. T cells, expressing a recombinant receptor that specifically recognize and/or target an antigen associated with the cancer and/or present on a universal tag. In some embodiments, the antigen recognized or targeted by the T cells is B cell maturation antigen (BCMA), ROR1,carbonic anhydrase 9 (CAIX), Her2/neu (receptor tyrosine kinase erbB2), L1-CAM, CD19, CD20, CD22, mesothelin, CEA, and hepatitis B surface antigen, anti-folate receptor, CD23, CD24, CD30, CD33, CD38, CD44, EGFR, epithelial glycoprotein 2 (EPG-2), epithelial glycoprotein 40 (EPG-40), EPHa2, erb-B2, erb-B3, erb-B4, erbB dimers, EGFR vIII, folate

binding protein (FBP), FCRL5, FCRH5, fetal acetylcholine receptor, GD2, GD3, HMW-MAA, IL-22R-alpha, IL-13R-alpha2, kinase insert domain receptor (kdr), kappa light chain, Lewis Y, L1-cell adhesion molecule, (L1-CAM), Melanoma-associated antigen (MAGE)-A1, MAGE-A3, MAGE-A6, Preferentially expressed antigen of melanoma (PRAME), survivin, TAG72, B7-H6, IL-13 receptor alpha 2 (IL-13Ra2), CA9, GD3, HMW-MAA, CD171, G250/CAIX, HLA-AI MAGE Al, HLA-A2 NY-ESO-1, PSCA, folate receptor-a, CD44v6, CD44v7/8, avb6 integrin, 8H9, NCAM, VEGF receptors, 5T4, Foetal AchR, NKG2D ligands, CD44v6, dual antigen, a cancer-testes antigen, mesothelin, murine CMV, mucin 1 (MUC1), MUC16, PSCA, NKG2D, NY-ESO-1, MART-1, gp100, oncofetal antigen, ROR1, TAG72, VEGF-R2, carcinoembryonic antigen (CEA), Her2/neu, estrogen receptor, progesterone receptor, ephrinB2, CD123, c-Met, GD-2, O-acetylated GD2 (OGD2), CE7, Wilms Tumor 1 (WT-1), a cyclin, cyclin A2, CCL-1, CD138, optionally a human antigen of any of the foregoing; a pathogen-specific antigen. In some embodiments, the antigen recognized and/or targeted by T cells is selected from the group consisting of Notch 1, Notch 2, Notch 3, Notch 4, cell surface associated Mucin 1 (MUC1), Ephrin B2, Betaglycan (TGFBR3), CD43, CD44, CSF1R, CX3CR1, CXCL16, Delta1, E-cadherin, N-cadherin, HLA-A2, IFNaR2, IL1R1, IL1R2, IL6R, and amyloid precursor protein (APP).

[0200] In some embodiments the antigen recognized and/targeted by T cells is B Cell Maturation Antigen (BCMA). Exemplary antigen-binding domains, and CARs containing such antigen-binding domains, that target or specifically bind BCMA are known, see e.g. WO 2016/090320, WO2016090327, WO2010104949A2 and WO2017173256. In some embodiments, the antigen binding domain is an scFv that contains a VH and a VL derived from an antibody or an antibody fragment specific to BCMA. In some embodiments, the antibody or antibody fragment that binds BCMA is or contains a VH and a VL from an antibody or antibody fragment set forth in International Patent Applications, Publication Number WO 2016/090327 and WO 2016/090320.

[0201] Methods for administration of engineered cells for adoptive cell therapy are known and may be used in connection with the provided methods and compositions. For example, adoptive T cell therapy methods are described, *e.g.*, in US Patent Application Publication No. 2003/0170238 to Gruenberg et al; US Patent No. 4,690,915 to Rosenberg; Rosenberg (2011) *Nat Rev Clin Oncol.* 8(10):577-85). See, *e.g.*, Themeli *et al.*, (2013) *Nat Biotechnol.* 31(10): 928-

933; Tsukahara *et al.*, (2013) *Biochem Biophys Res Commun* 438(1): 84-9; Davila *et al.*, (2013) *PLoS ONE* 8(4): e61338.

[0202] In some embodiments, the cell therapy, *e.g.*, adoptive T cell therapy, is carried out by autologous transfer, in which the cells are isolated and/or otherwise prepared from the subject who is to receive the cell therapy, or from a sample derived from such a subject. Thus, in some aspects, the cells are derived from a subject, *e.g.*, patient, in need of a treatment and the cells, following isolation and processing are administered to the same subject.

[0203] In some embodiments, the cell therapy, *e.g.*, adoptive T cell therapy, is carried out by allogeneic transfer, in which the cells are isolated and/or otherwise prepared from a subject other than a subject who is to receive or who ultimately receives the cell therapy, *e.g.*, a first subject. In such embodiments, the cells then are administered to a different subject, *e.g.*, a second subject, of the same species. In some embodiments, the first and second subjects are genetically identical. In some embodiments, the first and second subjects are genetically similar. In some embodiments, the second subject expresses the same HLA class or supertype as the first subject.

[0204] The cells can be administered by any suitable means. The cells are administered in a dosing regimen to achieve a therapeutic effect, such as a reduction in tumor burden. Dosing and administration may depend in part on the schedule of administration of the gamma secretase inhibitor, which can be administered prior to, subsequent to and/or simultaneously with initiation of administration of the T cell therapy. Various dosing schedules of the T cell therapy include but are not limited to single or multiple administrations over various time-points, bolus administration, and pulse infusion.

a. Compositions and formulations

[0205] In some embodiments, the dose of cells of the cell therapy, such a T cell therapy comprising cells engineered with a recombinant antigen receptor, *e.g.* CAR or TCR, is provided as a composition or formulation, such as a pharmaceutical composition or formulation. Such compositions can be used in accord with the provided methods, such as in the prevention or treatment of diseases, conditions, and disorders.

[0206] In some embodiments, the cell therapy, such as engineered T cells (*e.g.* CAR T cells), are formulated with a pharmaceutically acceptable carrier. In some aspects, the choice of carrier is determined in part by the particular cell or agent and/or by the method of administration. Accordingly, there are a variety of suitable formulations. For example, the

pharmaceutical composition can contain preservatives. Suitable preservatives may include, for example, methylparaben, propylparaben, sodium benzoate, and benzalkonium chloride. In some aspects, a mixture of two or more preservatives is used. The preservative or mixtures thereof are typically present in an amount of about 0.0001% to about 2% by weight of the total composition. Carriers are described, *e.g.*, by Remington's Pharmaceutical Sciences 16th edition, Osol, A. Ed. (1980). Pharmaceutically acceptable carriers are generally nontoxic to recipients at the dosages and concentrations employed, and include, but are not limited to: buffers such as phosphate, citrate, and other organic acids; antioxidants including ascorbic acid and methionine; preservatives (such as octadecyldimethylbenzyl ammonium chloride; hexamethonium chloride; benzalkonium chloride; benzethonium chloride; phenol, butyl or benzyl alcohol; alkyl parabens such as methyl or propyl paraben; catechol; resorcinol; cyclohexanol; 3-pentanol; and m-cresol); low molecular weight (less than about 10 residues) polypeptides; proteins, such as serum albumin, gelatin, or immunoglobulins; hydrophilic polymers such as polyvinylpyrrolidone; amino acids such as glycine, glutamine, asparagine, histidine, arginine, or lysine; monosaccharides, disaccharides, and other carbohydrates including glucose, mannose, or dextrins; chelating agents such as EDTA; sugars such as sucrose, mannitol, trehalose or sorbitol; salt-forming counter-ions such as sodium; metal complexes (*e.g.* Zn-protein complexes); and/or non-ionic surfactants such as polyethylene glycol (PEG).

[0207] Buffering agents in some aspects are included in the compositions. Suitable buffering agents include, for example, citric acid, sodium citrate, phosphoric acid, potassium phosphate, and various other acids and salts. In some aspects, a mixture of two or more buffering agents is used. The buffering agent or mixtures thereof are typically present in an amount of about 0.001% to about 4% by weight of the total composition. Methods for preparing administrable pharmaceutical compositions are known. Exemplary methods are described in more detail in, for example, Remington: The Science and Practice of Pharmacy, Lippincott Williams & Wilkins; 21st ed. (May 1, 2005).

[0208] The formulations can include aqueous solutions. The formulation or composition may also contain more than one active ingredient useful for the particular indication, disease, or condition being prevented or treated with the cells or agents, where the respective activities do not adversely affect one another. Such active ingredients are suitably present in combination in amounts that are effective for the purpose intended. Thus, in some embodiments, the

pharmaceutical composition further includes other pharmaceutically active agents or drugs, such as chemotherapeutic agents, *e.g.*, asparaginase, busulfan, carboplatin, cisplatin, daunorubicin, doxorubicin, fluorouracil, gemcitabine, hydroxyurea, methotrexate, paclitaxel, rituximab, vinblastine, vincristine, etc.

[0209] The pharmaceutical composition in some embodiments contains cells in amounts effective to treat or prevent the disease or condition, such as a therapeutically effective or prophylactically effective amount. Therapeutic or prophylactic efficacy in some embodiments is monitored by periodic assessment of treated subjects. For repeated administrations over several days or longer, depending on the condition, the treatment is repeated until a desired suppression of disease symptoms occurs. However, other dosage regimens may be useful and can be determined. The desired dosage can be delivered by a single bolus administration of the composition, by multiple bolus administrations of the composition, or by continuous infusion administration of the composition.

[0210] The cells may be administered using standard administration techniques, formulations, and/or devices. Provided are formulations and devices, such as syringes and vials, for storage and administration of the compositions. With respect to cells, administration can be autologous or heterologous. For example, immunoresponsive cells or progenitors can be obtained from one subject, and administered to the same subject or a different, compatible subject. Peripheral blood derived immunoresponsive cells or their progeny (e.g., in vivo, ex vivo or in vitro derived) can be administered via localized injection, including catheter administration, systemic injection, localized injection, intravenous injection, or parenteral administration. When administering a therapeutic composition (e.g., a pharmaceutical composition containing a genetically modified immunoresponsive cell), it will generally be formulated in a unit dosage injectable form (solution, suspension, emulsion).

[0211] Formulations include those for oral, intravenous, intraperitoneal, subcutaneous, pulmonary, transdermal, intramuscular, intranasal, buccal, sublingual, or suppository administration. In some embodiments, the agent or cell populations are administered parenterally. The term “parenteral,” as used herein, includes intravenous, intramuscular, subcutaneous, rectal, vaginal, and intraperitoneal administration. In some embodiments, the agent or cell populations are administered to a subject using peripheral systemic delivery by intravenous, intraperitoneal, or subcutaneous injection.

[0212] Compositions in some embodiments are provided as sterile liquid preparations, e.g., isotonic aqueous solutions, suspensions, emulsions, dispersions, or viscous compositions, which may in some aspects be buffered to a selected pH. Liquid preparations are normally easier to prepare than gels, other viscous compositions, and solid compositions. Additionally, liquid compositions are somewhat more convenient to administer, especially by injection. Viscous compositions, on the other hand, can be formulated within the appropriate viscosity range to provide longer contact periods with specific tissues. Liquid or viscous compositions can comprise carriers, which can be a solvent or dispersing medium containing, for example, water, saline, phosphate buffered saline, polyoi (for example, glycerol, propylene glycol, liquid polyethylene glycol) and suitable mixtures thereof.

[0213] Sterile injectable solutions can be prepared by incorporating the cells in a solvent, such as in admixture with a suitable carrier, diluent, or excipient such as sterile water, physiological saline, glucose, dextrose, or the like. The compositions can also be lyophilized. The compositions can contain auxiliary substances such as wetting, dispersing, or emulsifying agents (e.g., methylcellulose), pH buffering agents, gelling or viscosity enhancing additives, preservatives, flavoring agents, colors, and the like, depending upon the route of administration and the preparation desired. Standard texts may in some aspects be consulted to prepare suitable preparations.

[0214] Various additives which enhance the stability and sterility of the compositions, including antimicrobial preservatives, antioxidants, chelating agents, and buffers, can be added. Prevention of the action of microorganisms can be ensured by various antibacterial and antifungal agents, for example, parabens, chlorobutanol, phenol, sorbic acid, and the like. Prolonged absorption of the injectable pharmaceutical form can be brought about by the use of agents delaying absorption, for example, aluminum monostearate and gelatin.

[0215] The formulations to be used for in vivo administration are generally sterile. Sterility may be readily accomplished, e.g., by filtration through sterile filtration membranes.

[0216] For the prevention or treatment of disease, the appropriate dosage may depend on the type of disease to be treated, the type of agent or agents, the type of cells or recombinant receptors, the severity and course of the disease, whether the agent or cells are administered for preventive or therapeutic purposes, previous therapy, the subject's clinical history and response to the agent or the cells, and the discretion of the attending physician. The compositions are in

some embodiments suitably administered to the subject at one time or over a series of treatments.

[0217] In some cases, the cell therapy is administered as a single pharmaceutical composition comprising the cells. In some embodiments, a given dose is administered by a single bolus administration of the cells or agent. In some embodiments, it is administered by multiple bolus administrations of the cells or agent, for example, over a period of no more than 3 days, or by continuous infusion administration of the cells or agent.

b. Dosage Schedule and Administration

[0218] In some embodiments, a dose of cells is administered to subjects in accord with the provided combination therapy methods. In some embodiments, the size or timing of the doses is determined as a function of the particular disease or condition in the subject. It is within the level of a skilled artisan to empirically determine the size or timing of the doses for a particular disease in view of the provided description.

[0219] In certain embodiments, the cells, or individual populations of sub-types of cells, are administered to the subject at a range of about 0.1 million to about 100 billion cells and/or that amount of cells per kilogram of body weight of the subject, such as, *e.g.*, 0.1 million to about 50 billion cells (*e.g.*, about 5 million cells, about 25 million cells, about 500 million cells, about 1 billion cells, about 5 billion cells, about 20 billion cells, about 30 billion cells, about 40 billion cells, or a range defined by any two of the foregoing values), 1 million to about 50 billion cells (*e.g.*, about 5 million cells, about 25 million cells, about 500 million cells, about 1 billion cells, about 5 billion cells, about 20 billion cells, about 30 billion cells, about 40 billion cells, or a range defined by any two of the foregoing values), such as about 10 million to about 100 billion cells (*e.g.*, about 20 million cells, about 30 million cells, about 40 million cells, about 60 million cells, about 70 million cells, about 80 million cells, about 90 million cells, about 10 billion cells, about 25 billion cells, about 50 billion cells, about 75 billion cells, about 90 billion cells, or a range defined by any two of the foregoing values), and in some cases about 100 million cells to about 50 billion cells (*e.g.*, about 120 million cells, about 250 million cells, about 350 million cells, about 450 million cells, about 650 million cells, about 800 million cells, about 900 million cells, about 3 billion cells, about 30 billion cells, about 45 billion cells) or any value in between these ranges and/or per kilogram of body weight of the subject. Dosages may vary depending on attributes particular to the disease or disorder and/or patient and/or other treatments. In some

embodiments, such values refer to numbers of recombinant receptor-expressing cells; in other embodiments, they refer to number of T cells or PBMCs or total cells administered.

[0220] In some embodiments, for example, where the subject is a human, the cells are administered at a dose of total recombinant receptor (e.g., CAR)-expressing cells, T cells, or peripheral blood mononuclear cells (PBMCs) that is in the range of about 1×10^6 to about 1.2×10^9 such cells, such as at or about 1×10^7, at or about 5×10^7, at or about 1×10^8, at or about 2×10^8, at or about 2.5×10^8, at or about 3×10^8, at or about 3.5×10^8, at or about 4×10^8, at or about 4.5×10^8 at or about 5×10^8 total such cells, at or about 6×10^8, at or about 7×10^8, at or about 8×10^8, at or about 9×10^8 or at or about 1×10^9 of such cells, or the range between any two of the foregoing values. In some embodiments, for example, where the subject is a human, the dose includes fewer than about or about 5×10^8 total recombinant receptor (e.g., CAR)-expressing cells, T cells, or peripheral blood mononuclear cells (PBMCs). In some embodiments, for example, where the subject is a human, the dose includes fewer than about or about 3.0×10^8 total recombinant receptor (e.g., CAR)-expressing cells, T cells, or peripheral blood mononuclear cells (PBMCs). In some embodiments, for example, where the subject is a human, the dose includes fewer than about or about 1.5×10^8 total recombinant receptor (e.g., CAR)-expressing cells, T cells, or peripheral blood mononuclear cells (PBMCs). In some embodiments, for example, where the subject is a human, the dose includes fewer than about 1×10^8 total recombinant receptor (e.g., CAR)-expressing cells, T cells, or peripheral blood mononuclear cells (PBMCs). In some embodiments, for example, where the subject is a human, the dose includes fewer than about 5×10^7 total recombinant receptor (e.g., CAR)-expressing cells, T cells, or peripheral blood mononuclear cells (PBMCs).

[0221] In some embodiments, the dose of total cells administered comprises from or from about 1×10^5 to 8×10^8 total CAR-expressing T cells, 1×10^5 to 5×10^8 total CAR-expressing T cells, 1×10^5 to 2.5×10^8 total CAR-expressing T cells, 1×10^5 to 1×10^8 total CAR-expressing T cells, 1×10^5 to 5×10^7 total CAR-expressing T cells, 1×10^5 to 2.5×10^7 total CAR-expressing T cells, 1×10^5 to 1×10^7 total CAR-expressing T cells, 1×10^5 to 5×10^6 total CAR-expressing T cells, 1×10^5 to 2.5×10^6 total CAR-expressing T cells, 1×10^5 to 1×10^6 total CAR-expressing T cells, 1×10^6 to 8×10^8 total CAR-expressing T cells, 1×10^6 to 5×10^8 total CAR-expressing T cells, 1×10^6 to 2.5×10^8 total CAR-expressing T cells, 1×10^6 to 1×10^8 total CAR-expressing T cells, 1×10^6 to 5×10^7 total CAR-expressing T cells, 1×10^6 to $2.5 \times$

10^7 total CAR-expressing T cells, 1 x 10^6 to 1 x 10^7 total CAR-expressing T cells, 1 x 10^6 to 5 x 10^6 total CAR-expressing T cells, 1 x 10^6 to 2.5 x 10^6 total CAR-expressing T cells, 2.5 x 10^6 to 8 x 10^8 total CAR-expressing T cells, 2.5 x 10^6 to 5 x 10^8 total CAR-expressing T cells, 2.5 x 10^6 to 2.5 x 10^8 total CAR-expressing T cells, 2.5 x 10^6 to 1 x 10^8 total CAR-expressing T cells, 2.5 x 10^6 to 5 x 10^7 total CAR-expressing T cells, 2.5 x 10^6 to 2.5 x 10^7 total CAR-expressing T cells, 2.5 x 10^6 to 1 x 10^7 total CAR-expressing T cells, 2.5 x 10^6 to 5 x 10^6 total CAR-expressing T cells, 5 x 10^6 to 8 x 10^8 total CAR-expressing T cells, 5 x 10^6 to 5 x 10^8 total CAR-expressing T cells, 5 x 10^6 to 2.5 x 10^8 total CAR-expressing T cells, 5 x 10^6 to 1 x 10^8 total CAR-expressing T cells, 5 x 10^6 to 5 x 10^7 total CAR-expressing T cells, 5 x 10^6 to 2.5 x 10^7 total CAR-expressing T cells, 5 x 10^6 to 1 x 10^7 total CAR-expressing T cells, 1 x 10^7 to 8 x 10^8 total CAR-expressing T cells, 1 x 10^7 to 5 x 10^8 total CAR-expressing T cells, 1 x 10^7 to 2.5 x 10^8 total CAR-expressing T cells, 1 x 10^7 to 1 x 10^8 total CAR-expressing T cells, 1 x 10^7 to 5 x 10^7 total CAR-expressing T cells, 1 x 10^7 to 2.5 x 10^7 total CAR-expressing T cells, 2.5 x 10^7 to 8 x 10^8 total CAR-expressing T cells, 2.5 x 10^7 to 5 x 10^8 total CAR-expressing T cells, 2.5 x 10^7 to 2.5 x 10^8 total CAR-expressing T cells, 2.5 x 10^7 to 1 x 10^8 total CAR-expressing T cells, 2.5 x 10^7 to 5 x 10^7 total CAR-expressing T cells, 5 x 10^7 to 8 x 10^8 total CAR-expressing T cells, 5 x 10^7 to 5 x 10^8 total CAR-expressing T cells, 5 x 10^7 to 2.5 x 10^8 total CAR-expressing T cells, 5 x 10^7 to 1 x 10^8 total CAR-expressing T cells, 1 x 10^8 to 8 x 10^8 total CAR-expressing T cells, 1 x 10^8 to 5 x 10^8 total CAR-expressing T cells, 1 x 10^8 to 2.5 x 10^8 total CAR-expressing T cells, 2.5 x 10^8 to 8 x 10^8 total CAR-expressing T cells, 2.5 x 10^8 to 5 x 10^8 total CAR-expressing T cells, or 5 x 10^8 total CAR-expressing T cells to 8 x 10^8 total CAR-expressing T cells, each inclusive.

[0222] In some embodiments, the dose of genetically engineered cells comprises from or from about 50 x 10^6 to 800 x 10^6 total CAR-expressing T cells, from or from about 50 x 10^6 to 450 x 10^6 total CAR-expressing T cells, from or from about 50 x 10^6 to 300 x 10^6 total CAR-expressing T cells, from or from about 50 x 10^6 to 150 x 10^6 total CAR-expressing T cells, from or from about 150 x 10^6 to 800 x 10^6 total CAR-expressing T cells, from or from about 150 x 10^6 to 450 x 10^6 total CAR-expressing T cells, from or from about 150 x 10^6 to 300 x 10^6 total CAR-expressing T cells, 300 x 10^6 to 800 x 10^6 total CAR-expressing T cells, from or from about 300 x 10^6 to 450 x 10^6 total CAR-expressing T cells, or 450 x 10^6 to 800 x 10^6 total CAR-expressing T cells. In some embodiments, the dose of genetically engineered cells is or is about

50 x 10^6 CAR-expressing T cells, is or is about 150 x 10^6 CAR-expressing T cells, cells, is or is about 450 x 10^6 CAR-expressing T cells, or is or is about 800 x 10^6 CAR-expressing T cells.

[0223] In some of any of the embodiments, the number is with reference to the total number of CD3+, total number of CD8+ or total number of CD4+ and CD8+ T cells, in some recombinant receptor-expressing (e.g. CAR+) cells of such cells.

[0224] In some embodiments, the cell therapy comprises administration of a dose comprising a number of cells from or from about 1 x 10^5 to 1 x 10^8 total recombinant receptor-expressing cells, total T cells, or total peripheral blood mononuclear cells (PBMCs); from or from about 5 x 10^5 to 1 x 10^7 total recombinant receptor-expressing cells, total T cells, or totalPBMCs or from or from about 1 x 10^6 to 1 x 10^7 total recombinant receptor-expressing cells, total T cells, or total PBMCs), each inclusive. In some embodiments, the cell therapy comprises administration of a dose of cells comprising a number of cells at least or about at least 1 x 10^5 total recombinant receptor-expressing cells, total T cells, or total PBMCs, such at least or at least 1 x 10^6, at least or about at least 1 x 10^7, at least or about at least 1 x 10^8 of such cells. In some embodiments, the number is with reference to the total number of CD3+ or CD8+, in some cases also recombinant receptor-expressing (e.g. CAR+) cells. In some embodiments, the cell therapy comprises administration of a dose comprising a number of cells from or from about 1 x 10^5 to 1 x 10^8 CD3+ or CD8+ total T cells or CD3+ or CD8+ recombinant receptor-expressing cells, from or from about 5 x 10^5 to 1 x 10^7 CD3+ or CD8+ total T cells or CD3+ or CD8+ recombinant receptor-expressing cells, or from or from about 1 x 10^6 to 1 x 10^7 CD3+ or CD8+ total T cells or CD3+ or CD8+recombinant receptor-expressing cells, each inclusive. In some embodiments, the cell therapy comprises administration of a dose comprising a number of cells from or from about 1 x 10^5 to 1 x 10^8 total CD3+/CAR+ or CD8+/CAR+ cells, from or from about 5 x 10^5 to 1 x 10^7 total CD3+/CAR+ or CD8+/CAR+ cells, or from or from about 1 x 10^6 to 1 x 10^7 total CD3+/CAR+ or CD8+/CAR+ cells, each inclusive.

[0225] In some embodiments, the T cells of the dose include CD4+ T cells, CD8+ T cells or CD4+ and CD8+ T cells.

[0226] In some embodiments, for example, where the subject is human, the CD8+ T cells of the dose, including in a dose including CD4+ and CD8+ T cells, includes between about 1 x 10^6 and 5 x 10^8 total recombinant receptor (e.g., CAR)-expressing CD8+cells, e.g., in the range of about 5 x 10^6 to 1 x 10^8 such cells, such cells 1 x 10^7, 2.5 x 10^7, 5 x 10^7, 7.5 x 10^7, 1 x 10^8, or 5 x

10^8 total such cells, or the range between any two of the foregoing values. In some embodiments, the patient is administered multiple doses, and each of the doses or the total dose can be within any of the foregoing values. In some embodiments, the dose of cells comprises the administration of from or from about 1 x 10^7 to 0.75 x 10^8 total recombinant receptor-expressing CD8+ T cells, 1 x 10^7 to 2.5 x 10^7 total recombinant receptor-expressing CD8+ T cells, from or from about 1 x 10^7 to 0.75 x 10^8 total recombinant receptor-expressing CD8+ T cells, each inclusive. In some embodiments, the dose of cells comprises the administration of or about 1 x 10^7, 2.5 x 10^7, 5 x 10^7 7.5 x 10^7, 1 x 10^8, or 5 x 10^8 total recombinant receptor-expressing CD8+ T cells.

[0227] In some embodiments, the cell therapy comprises administration of a dose comprising a number of cells that is at least or at least about or is or is about 0.1 x 10^6 cells/kg body weight of the subject, 0.2 x 10^6 cells/kg, 0.3 x 10^6 cells/kg, 0.4 x 10^6 cells/kg, 0.5 x 10^6 cells/kg, 1 x 10^6 cell/kg, 2.0 x 10^6 cells/kg, 3 x 10^6 cells/kg or 5 x 10^6 cells/kg.

[0228] In some embodiments, the cell therapy comprises administration of a dose comprising a number of cells between or between about 0.1 x 10^6 cells/kg body weight of the subject and 1.0 x 10^7 cells/kg, between or between about 0.5 x 10^6 cells/kg and 5 x 10^6 cells/kg, between or between about 0.5 x 10^6 cells/kg and 3 x 10^6 cells/kg, between or between about 0.5 x 10^6 cells/kg and 2 x 10^6 cells/kg, between or between about 0.5 x 10^6 cells/kg and 1 x 10^6 cell/kg, between or between about 1.0 x 10^6 cells/kg body weight of the subject and 5 x 10^6 cells/kg, between or between about 1.0 x 10^6 cells/kg and 3 x 10^6 cells/kg, between or between about 1.0 x 10^6 cells/kg and 2 x 10^6 cells/kg, between or between about 2.0 x 10^6 cells/kg body weight of the subject and 5 x 10^6 cells/kg, between or between about 2.0 x 10^6 cells/kg and 3 x 10^6 cells/kg, or between or between about 3.0 x 10^6 cells/kg body weight of the subject and 5 x 10^6 cells/kg, each inclusive.

[0229] In some embodiments, the dose of cells comprises between at or about 2 x 10^5 of the cells/kg and at or about 2 x 10^6 of the cells/kg, such as between at or about 4 x 10^5 of the cells/kg and at or about 1 x 10^6 of the cells/kg or between at or about 6 x 10^5 of the cells/kg and at or about 8 x 10^5 of the cells/kg. In some embodiments, the dose of cells comprises no more than 2 x 10^5 of the cells (*e.g.* antigen-expressing, such as CAR-expressing cells) per kilogram body weight of the subject (cells/kg), such as no more than at or about 3 x 10^5 cells/kg, no more than at or about 4 x 10^5 cells/kg, no more than at or about 5 x 10^5 cells/kg, no more than at or

about 6 x 10^5 cells/kg, no more than at or about 7 x 10^5 cells/kg, no more than at or about 8 x 10^5 cells/kg, nor more than at or about 9 x 10^5 cells/kg, no more than at or about 1 x 10^6 cells/kg, or no more than at or about 2 x 10^6 cells/kg. In some embodiments, the dose of cells comprises at least or at least about or at or about 2 x 10^5 of the cells (*e.g.* antigen-expressing, such as CAR-expressing cells) per kilogram body weight of the subject (cells/kg), such as at least or at least about or at or about 3 x 10^5 cells/kg, at least or at least about or at or about 4 x 10^5 cells/kg, at least or at least about or at or about 5 x 10^5 cells/kg, at least or at least about or at or about 6 x 10^5 cells/kg, at least or at least about or at or about 7 x 10^5 cells/kg, at least or at least about or at or about 8 x 10^5 cells/kg, at least or at least about or at or about 9 x 10^5 cells/kg, at least or at least about or at or about 1 x 10^6 cells/kg, or at least or at least about or at or about 2 x 10^6 cells/kg.

[0230] In some embodiments, the dose of cells, e.g., recombinant receptor-expressing T cells, is administered to the subject as a single dose or is administered only one time within a period of two weeks, one month, three months, six months, 1 year or more.

[0231] In some aspects, the pharmaceutical compositions and formulations are provided as unit dose form compositions including the number of cells for administration in a given dose or fraction thereof.

[0232] In some embodiments, cells of the dose may be administered by administration of a plurality of compositions or solutions, such as a first and a second, optionally more, each containing some cells of the dose. In some aspects, the plurality of compositions, each containing a different population and/or sub-types of cells, are administered separately or independently, optionally within a certain period of time. For example, the populations or sub-types of cells can include $CD8^+$ and $CD4^+$ T cells, respectively, and/or CD8+- and CD4+-enriched populations, respectively, e.g., CD4+ and/or CD8+ T cells each individually including cells genetically engineered to express the recombinant receptor. In some embodiments, the administration of the dose comprises administration of a first composition comprising a dose of CD8+ T cells or a dose of CD4+ T cells and administration of a second composition comprising the other of the dose of CD4+ T cells and the CD8+ T cells.

[0233] In some embodiments, the administration of the composition or dose, e.g., administration of the plurality of cell compositions, involves administration of the cell compositions separately. In some aspects, the separate administrations are carried out

simultaneously, or sequentially, in any order. In some embodiments, the dose comprises a first composition and a second composition, and the first composition and second composition are administered 0 to 12 hours apart, 0 to 6 hours apart or 0 to 2 hours apart. In some embodiments, the initiation of administration of the first composition and the initiation of administration of the second composition are carried out no more than 2 hours, no more than 1 hour, or no more than 30 minutes apart, no more than 15 minutes, no more than 10 minutes or no more than 5 minutes apart. In some embodiments, the initiation and/or completion of administration of the first composition and the completion and/or initiation of administration of the second composition are carried out no more than 2 hours, no more than 1 hour, or no more than 30 minutes apart, no more than 15 minutes, no more than 10 minutes or no more than 5 minutes apart.

[0234] In some composition, the first composition, e.g., first composition of the dose, comprises CD4+ T cells. In some composition, the first composition, e.g., first composition of the dose, comprises CD8+ T cells. In some embodiments, the first composition is administered prior to the second composition.

[0235] In some embodiments, the dose or composition of cells includes a defined or target ratio of CD4+ cells expressing a recombinant receptor to CD8+ cells expressing a recombinant receptor and/or of CD4+ cells to CD8+ cells, which ratio optionally is approximately 1:1 or is between approximately 1:3 and approximately 3:1, such as approximately 1:1. In some aspects, the administration of a composition or dose with the target or desired ratio of different cell populations (such as CD4+:CD8+ ratio or CAR+CD4+:CAR+CD8+ ratio, e.g., 1:1) involves the administration of a cell composition containing one of the populations and then administration of a separate cell composition comprising the other of the populations, where the administration is at or approximately at the target or desired ratio. In some aspects, administration of a dose or composition of cells at a defined ratio leads to improved expansion, persistence and/or antitumor activity of the T cell therapy.

[0236] In the context of adoptive cell therapy, administration of a given "dose" of cells encompasses administration of the given amount or number of cells as a single composition and/or single uninterrupted administration, *e.g.*, as a single injection or continuous infusion, and also encompasses administration of the given amount or number of cells as a split dose, provided in multiple individual compositions or infusions, over a specified period of time, which is no more than 3 days. Thus, in some contexts, the dose is a single or continuous administration

of the specified number of cells, given or initiated at a single point in time. In some contexts, however, the dose is administered in multiple injections or infusions over a period of no more than three days, such as once a day for three days or for two days or by multiple infusions over a single day period.

[0237] Thus, in some aspects, the cells of the dose are administered in a single pharmaceutical composition. In some embodiments, the cells of the dose are administered in a plurality of compositions, collectively containing the cells of the dose.

[0238] The term "split dose" refers to a dose that is split so that it is administered over more than one day. This type of dosing is encompassed by the present methods and is considered to be a single dose. In some embodiments, the cells of a split dose are administered in a plurality of compositions, collectively comprising the cells of the dose, over a period of no more than three days.

[0239] Thus, the dose of cells may be administered as a split dose. For example, in some embodiments, the dose may be administered to the subject over 2 days or over 3 days. Exemplary methods for split dosing include administering 25% of the dose on the first day and administering the remaining 75% of the dose on the second day. In other embodiments, 33% of the dose may be administered on the first day and the remaining 67% administered on the second day. In some aspects, 10% of the dose is administered on the first day, 30% of the dose is administered on the second day, and 60% of the dose is administered on the third day. In some embodiments, the split dose is not spread over more than 3 days.

[0240] In some embodiments, the subject receives multiple doses, e.g., two or more doses or multiple consecutive doses, of the cells. In some embodiments, two doses are administered to a subject. In some embodiments, the subject receives the consecutive dose, e.g., second dose, is administered approximately 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20 or 21 days after the first dose. In some embodiments, multiple consecutive doses are administered following the first dose, such that an additional dose or doses are administered following administration of the consecutive dose. In some aspects, the number of cells administered to the subject in the additional dose is the same as or similar to the first dose and/or consecutive dose. In some embodiments, the additional dose or doses are larger than prior doses.

[0241] In some aspects, the size of the first and/or consecutive dose is determined based on one or more criteria such as response of the subject to prior treatment, e.g. chemotherapy,

disease burden in the subject, such as tumor load, bulk, size, or degree, extent, or type of metastasis, stage, and/or likelihood or incidence of the subject developing toxic outcomes, e.g., CRS, macrophage activation syndrome, tumor lysis syndrome, neurotoxicity, and/or a host immune response against the cells and/or recombinant receptors being administered.

[0242] In some aspects, the time between the administration of the first dose and the administration of the consecutive dose is about 9 to about 35 days, about 14 to about 28 days, or 15 to 27 days. In some embodiments, the administration of the consecutive dose is at a time point more than about 14 days after and less than about 28 days after the administration of the first dose. In some aspects, the time between the first and consecutive dose is about 21 days. In some embodiments, an additional dose or doses, e.g. consecutive doses, are administered following administration of the consecutive dose. In some aspects, the additional consecutive dose or doses are administered at least about 14 and less than about 28 days following administration of a prior dose. In some embodiments, the additional dose is administered less than about 14 days following the prior dose, for example, 4, 5, 6, 7, 8, 9, 10, 11, 12, or 13 days after the prior dose. In some embodiments, no dose is administered less than about 14 days following the prior dose and/or no dose is administered more than about 28 days after the prior dose.

[0243] In some embodiments, the dose of cells, e.g., recombinant receptor-expressing cells, comprises two doses (e.g., a double dose), comprising a first dose of the T cells and a consecutive dose of the T cells, wherein one or both of the first dose and the second dose comprises administration of the split dose of T cells.

[0244] In some embodiments, the dose of cells is generally large enough to be effective in reducing disease burden.

[0245] In some embodiments, the cells are administered at a desired dosage, which in some aspects includes a desired dose or number of cells or cell type(s) and/or a desired ratio of cell types. Thus, the dosage of cells in some embodiments is based on a total number of cells (or number per kg body weight) and a desired ratio of the individual populations or sub-types, such as the CD4+ to CD8+ ratio. In some embodiments, the dosage of cells is based on a desired total number (or number per kg of body weight) of cells in the individual populations or of individual cell types. In some embodiments, the dosage is based on a combination of such

features, such as a desired number of total cells, desired ratio, and desired total number of cells in the individual populations.

[0246] In some embodiments, the populations or sub-types of cells, such as $CD8^+$ and $CD4^+$ T cells, are administered at or within a tolerated difference of a desired dose of total cells, such as a desired dose of T cells. In some aspects, the desired dose is a desired number of cells or a desired number of cells per unit of body weight of the subject to whom the cells are administered, *e.g.*, cells/kg. In some aspects, the desired dose is at or above a minimum number of cells or minimum number of cells per unit of body weight. In some aspects, among the total cells administered at the desired dose, the individual populations or sub-types are present at or near a desired output ratio (such as $CD4^+$ to $CD8^+$ ratio), *e.g.*, within a certain tolerated difference or error of such a ratio.

[0247] In some embodiments, the cells are administered at or within a tolerated difference of a desired dose of one or more of the individual populations or sub-types of cells, such as a desired dose of CD4+ cells and/or a desired dose of CD8+ cells. In some aspects, the desired dose is a desired number of cells of the sub-type or population, or a desired number of such cells per unit of body weight of the subject to whom the cells are administered, *e.g.*, cells/kg. In some aspects, the desired dose is at or above a minimum number of cells of the population or sub-type, or minimum number of cells of the population or sub-type per unit of body weight.

[0248] Thus, in some embodiments, the dosage is based on a desired fixed dose of total cells and a desired ratio, and/or based on a desired fixed dose of one or more, *e.g.*, each, of the individual sub-types or sub-populations. Thus, in some embodiments, the dosage is based on a desired fixed or minimum dose of T cells and a desired ratio of $CD4^+$ to $CD8^+$ cells, and/or is based on a desired fixed or minimum dose of $CD4^+$ and/or $CD8^+$ cells.

[0249] In some embodiments, the cells are administered at or within a tolerated range of a desired output ratio of multiple cell populations or sub-types, such as CD4+ and CD8+ cells or sub-types. In some aspects, the desired ratio can be a specific ratio or can be a range of ratios. For example, in some embodiments, the desired ratio (*e.g.*, ratio of $CD4^+$ to $CD8^+$ cells) is between at or about 5:1 and at or about 5:1 (or greater than about 1:5 and less than about 5:1), or between at or about 1:3 and at or about 3:1 (or greater than about 1:3 and less than about 3:1), such as between at or about 2:1 and at or about 1:5 (or greater than about 1:5 and less than about 2:1, such as at or about 5:1, 4.5:1, 4:1, 3.5:1, 3:1, 2.5:1, 2:1, 1.9:1, 1.8:1, 1.7:1, 1.6:1, 1.5:1,

1.4:1, 1.3:1, 1.2:1, 1.1:1, 1:1, 1:1.1, 1:1.2, 1:1.3, 1:1.4, 1:1.5, 1:1.6, 1:1.7, 1:1.8, 1:1.9: 1:2, 1:2.5, 1:3, 1:3.5, 1:4, 1:4.5, or 1:5. In some aspects, the tolerated difference is within about 1%, about 2%, about 3%, about 4% about 5%, about 10%, about 15%, about 20%, about 25%, about 30%, about 35%, about 40%, about 45%, about 50% of the desired ratio, including any value in between these ranges.

[0250] The cells can be administered by any suitable means. The cells are administered in a dosing regimen to achieve a therapeutic effect, such as a reduction in tumor burden. Dosing and administration may depend in part on the schedule of administration of the immunomodulatory compound, which can be administered prior to, subsequent to and/or simultaneously with initiation of administration of the T cell therapy. Various dosing schedules of the T cell therapy include but are not limited to single or multiple administrations over various time-points, bolus administration, and pulse infusion. In certain embodiments, the engineered T cells express a recombinant receptor. In certain embodiments, the engineered T cells express a CAR.

[0251] In particular embodiments, the numbers and/or concentrations of cells refer to the number of recombinant receptor (*e.g.*, CAR)-expressing cells. In other embodiments, the numbers and/or concentrations of cells refer to the number or concentration of all cells, T cells, or peripheral blood mononuclear cells (PBMCs) administered.

[0252] In some aspects, the size of the dose is determined based on one or more criteria such as response of the subject to prior treatment, *e.g.* chemotherapy, disease burden in the subject, such as tumor load, bulk, size, or degree, extent, or type of metastasis, stage, and/or likelihood or incidence of the subject developing toxic outcomes, *e.g.*, CRS, macrophage activation syndrome, tumor lysis syndrome, neurotoxicity, and/or a host immune response against the cells and/or recombinant receptors being administered.

[0253] In some embodiments, administration of the gamma secretase inhibitor in combination with the cells is able to increase, such as substantially or significantly increase, the expansion or proliferation of the cells, and thus a lower dose of cells can be administered to the subject. In some cases, the provided methods allow a lower dose of such cells to be administered, to achieve the same or better efficacy of treatment as the dose in a method in which the cell therapy is administered without administering the gamma secretase inhibitor, such as at least 1.5-fold, 2-fold, 3-fold, 4-fold, 5-fold or 10-fold less than the dose in a method in which the cell therapy is administered without administering the gamma secretase inhibitor.

[0254] In some embodiments, for example, the dose contains between or between about 5.0 x 10^6 and 2.25 x 10^7, 5.0 x 10^6 and 2.0 x 10^7, 5.0 x 10^6 and 1.5 x 10^7, 5.0 x 10^6 and 1.0 x 10^7, 5.0 x 10^6 and 7.5 x 10^6, 7.5 x 10^6 and 2.25 x 10^7, 7.5 x 10^6 and 2.0 x 10^7, 7.5 x 10^6 and 1.5 x 10^7, 7.5 x 10^6 and 1.0 x 10^7, 1.0 x 10^7 and 2.25 x 10^7, 1.0 x 10^7 and 2.0 x 10^7, 1.0 x 10^7 and 1.5 x 10^7, 1.5 x 10^7 and 2.25 x 10^7, 1.5 x 10^7 and 2.0 x 10^7, 2.0 x 10^7 and 2.25 x 10^7. In some embodiments, the dose of cells contains a number of cells, that is between at least or at least about 5 x 10^6, 6 x 10^6, 7 x 10^6, 8 x 10^6, 9 x 10^6, 10 x 10^6 and about 15 x 10^6 recombinant-receptor expressing cells, such as recombinant-receptor expressing cells that are CD8+. In some embodiments, such dose, such as such target number of cells refers to the total recombinant-receptor expressing cells in the administered composition.

[0255] In some embodiments, for example, the lower dose contains less than about 5 x 10^6 cells, recombinant receptor (e.g. CAR)-expressing cells, T cells, and/or PBMCs per kilogram body weight of the subject, such as less than about 4.5 x 10^6, 4 x 10^6, 3.5 x 10^6, 3 x 10^6, 2.5 x 10^6, 2 x 10^6, 1.5 x 10^6, 1 x 10^6, 5 x 10^5, 2.5 x 10^5, or 1 x 10^5 such cells per kilogram body weight of the subject. In some embodiments, the lower dose contains less than about 1 x 10^5, 2 x 10^5, 5 x 10^5, or 1 x 10^6 of such cells per kilogram body weight of the subject, or a value within the range between any two of the foregoing values. In some embodiments, such values refer to numbers of recombinant receptor-expressing cells; in other embodiments, they refer to the number of T cells or PBMCs or total cells administered.

[0256] In some embodiments, one or more subsequent dose of cells can be administered to the subject. In some embodiments, the subsequent dose of cells is administered greater than or greater than about 7 days, 14 days, 21 days, 28 days or 35 days after initiation of administration of the first dose of cells. The subsequent dose of cells can be more than, approximately the same as, or less than the first dose. In some embodiments, administration of the T cell therapy, such as administration of the first and/or second dose of cells, can be repeated.

[0257] In some embodiments, initiation of administration of the cell therapy, *e.g.* the dose of cells or a first dose of a split dose of cells, is administered before (prior to), concurrently with or after (subsequently or subsequent to) the administration of the gamma secretase inhibitor, or a pharmaceutically acceptable salt of hydrate thereof.

[0258] In some embodiments, the dose of cells, or the subsequent dose of cells, is administered concurrently with initiating administration of the gamma secretase inhitibor in

accord with the combination therapy methods. In some embodiments, the dose of cells, or the subsequent dose of cells, is administered on the same day as initiating administration of the gamma secretase inhibitor in accord with the combination therapy methods. In some embodiments, the dose of cells, or the subsequent dose of cells, is administered within 1 day, within 2 days, within 3 days, within 4 days, within 5 days, within 6 days, or within 7 days of initiating administration of the gamma secretase inhibitor in accord with the combination therapy methods.

[0259] In some embodiments, the dose of cells, or the subsequent dose of cells, is administered prior to starting or initiating administration of the gamma secretase inhibitor in accord with the provided combination therapy. In some embodiments, the dose of cells is administered at least or at least about 1 hour, at least or at least about 2 hours, at least or at least about 3 hours, at least or at least about 6 hours, at least or at least about 12 hours, at least or at least about 1 day, at least or at least about 2 days, at least or at least about 3 days, at least or about at least 4 days, at least or at least about 5 days, at least or about at least 6 days, at least or at least about 7 days, at least or about at least 12 days, at least or at least about 14 days, at least or about at least 15 days, at least or at least about 21 days, at least or at least about 28 days, at least or about at least 30 days, at least or at least about 35 days, at least or at least about 42 days, at least or about at least 60 days or at least or about at least 90 days prior to administering the gamma secretase inhibitor in accord with the provided combination therapy.

[0260] In some embodiments, the administration of the gamma secretase inhibitor in accord with the provided combination therapy is at a time in which the prior administration of the immunotherapy (*e.g.,* T cell therapy, such as CAR-T cell therapy) is associated with, or is likely to be associated with, a decreased functionality of the T cells compared to the functionality of the T cells at a time just prior to initiation of the immunotherapy (*e.g.,* T cell therapy, such as CAR-T cell therapy) or at a preceding time point after initiation of the T cell therapy. In some embodiments, the method involves, subsequent to administering the dose of cells of the T cell therapy, *e.g.*, adoptive T cell therapy, but prior to administering the gamma secretase inhibitor, assessing a sample from the subject for one or more functions of T cells, such as expansion or persistence of the cells, e.g. as determined by level or amount in the blood, or other phenotypes or desired outcomes as described herein, *e.g.*, such as those described in Section III. Various

parameters for determining or assessing the regimen of the combination therapy are described in Section III.

B. ADMINISTRATION OF GAMMA SECRETASE INHIBITOR

[0261] In some embodiments of the methods, compositions, combinations, kits and uses provided herein, the combination therapy can be administered in one or more compositions, *e.g.*, a pharmaceutical composition containing an inhibitor of gamma secretase, and/or the cell therapy, *e.g.*, T cell therapy.

[0262] In some embodiments, the gamma secretase inhibitor inhibits or reduces the intramembrane cleavage of one or more receptors on a cell, for example, a tumor/cancer cell. In some embodiments, the IC_{50} of the inhibition of the intramembrane cleavage of the cell surface receptor on the cancer/tumor cell is less than about 100μM, 50μM, 25μM, 10μM, 1μM, 0.75μM, 0.5 μM, 0.25 μM, 0.1 μM, 75nM, 50nM, 25nM, or 10nM. In some embodiments, he gamma secretase inhibitor inhibits intramembrane cleavage of the cell surface receptor on the cancer/tumor cell with a half-maximal inhibitory concentration (IC_{50}) of 0.01 nM to 10 nM, 0.01 nM to 5 nM, 0.01 nM to 1 nM, 0.01 nM, to 0.5 nM, 0.01 nM to 0.35 nM, 0.01 nM to 0.25 nM, 0.01 nM to 1.0 nM, 0.01 nM to 0.05 nM, 0.05 nM to 10 nM, 0.05 nM to 5 nM, 0.05 nM to 1 nM, 0.05 nM, to 0.5 nM, 0.05 nM to 0.35 nM, 0.05 nM to 0.25 nM, 0.05 nM to 1.0 nM, 1.0 nM to 10 nM, 1.0 nM nM to 5 nM, 1.0 nM nM to 1 nM, 1.0 nM nM, to 0.5 nM, 1.0 nM nM to 0.35 nM, 1.0 nM nM to 0.25 nM, 0.25 nM to 10 nM, 0.25 nM to 5 nM, 0.25 nM to 1 nM, 0.25 nM, to 0.5 nM, 0.25 nM to 0.35 nM, 0.35 nM to 10 nM, 0.35 nM to 5 nM, 0.35 nM to 1 nM, 0.35 nM, to 0.5 nM, 0.5 nM to 10 nM, 0.5 nM to 5 nM, 0.5 nM to 1 nM, 1 nM to 10 nM, 1 nM to 5 nM or 5 nM to 10 nM. In some embodiments, the IC_{50} of the inhibition of the intramembrane cleavage of the cell surface receptor on the cancer/tumor cell is about 10nM-25nM, 25nM-50nM, 50nM-75nM, 75nM-0.1μM, 0.1μM-0.25μM, 0.25μM-0.5μM, 0.5μM-0.75μM, 0.75μM-1μM, 1μM-10μM**,** 10μM-25μM, 25μM-50μM, 50μM-75μM, or 75μM-100μM. In some embodiments, the receptor(s) is/are selected from the group consisting of B cell maturation antigen (BCMA), Notch 1, Notch 2, Notch 3, Notch 4, cell surface associated Mucin 1 (MUC1), Ephrin B2, Betaglycan (TGFBR3), CD43, CD44, CSF1R, CX3CR1, CXCL16, Delta1, E-cadherin, N-cadherin, HLA-A2, IFNaR2, IL1R1, IL1R2, IL6R, and amyloid precursor protein (APP). In some embodiments, the receptor is cell surface associated Mucin 1 (MUC1).

[0263] In some embodiments, the an inhibitor of gamma secretase inhibits or reduces the intramembrane cleavage of one or more receptors described above on a cell, for example, a tumor/cancer cell, and wherein the administration of the gamma secretase increases the level of the receptor expression on the cell. In some embodiments, the receptor is surface B cell maturation antigen (BCMA). In some embodiments, the gamma secretase inhibitor prohibits intramembrane cleavage of surface B cell maturation antigen (BCMA), and wherein the IC_{50} of the inhibition of the cleavage of surface BCMA is less than about100μM, 50μM, 25μM, 10μM, 1μM, 0.75μM, 0.5 μM, 0.25 μM, 0.1 μM, 75nM, 50nM, 25nM, or 10nM. In some embodiments, the IC_{50} of the inhibition of the intramembrane cleavage of the cell surface receptor on the cancer/tumor cell is about 10nM-25nM, 25nM-50nM, 50nM-75nM, 75nM-0.1μM, 0.1μM-0.25μM, 0.25μM-0.5μM, 0.5μM-0.75μM, 0.75μM-1μM, 1μM-10μM**,** 10μM-25μM, 25μM-50μM, 50μM-75μM, or 75μM-100μM. In some embodiments, the IC_{50} of the inhibition of the cleavage of surface BCMA is less than 0.5nM. In some embodiments, the IC_{50} of the inhibition of the cleavage of surface BCMA is about 0.01nM to about 0.5nM, or 0.01nM to about 0.5nM, or about 0.01nM to about 0.35nM.

[0264] In some embodiments, the inhibitor of gamma secretase has a low brain penetration. In some embodiments, only 0.1%, 0.25%, 0.5%, 0.75%, 1%, 2.5%, 5%, 7.5%, 10%, 20%, 30%, 40%, 50%, 60%, 70%, 80% or 90% of the gamma secretase inhibitor can penetrate the brain after administration.

[0265] In some embodiments, the inhibitor of gamma secretase selectively inhibits or reduces intramembrane cleavage of part of the substrates of gamma secretase. In some embodiments, the gamma secretase inhibitor selectively inhibits or reduces the intramembrane cleavage of surface B cell maturation antigen (BCMA).

[0266] In some embodiments, the gamma secretase inhibitor is a peptide inhibitor or non-peptide inhibitor. In some embodiments, the gamma secretase inhibitor is a peptide inhibitor from among peptide aldehydes derivatives, difluoroketones derivatives, hydroxyethylene dipeptide isotere derivatives, alpha-helical peptide derivatives and dipeptide analogs. In some embodiments, the gamma secretase inhibitor is a non-peptide inhibitor selected from benzodiazepines derivatives and sulfonamides derivatives. In some embodiments, the gamma secretase is a transition state inhibitor, for example, LY-411575-I or LY685,458. In some embodiments, the gamma secretase inhibitor is a non-transition state inhibitor such as DAPT,

RO4929097. In some embodiments, the gamma secretase inhibitor is a gamma secretase modulator. In some embodiments, gamma secretase modulator is a nonsteroidal anti-inflammatory drugs-type modulator.

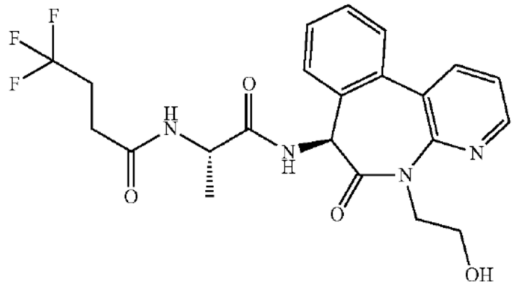
[0267] In some embodiments, the gamma secretase inhibitor is LY3039478, secretase inhibitor I (GSI I) Z-Leu-Leu- Norleucine; γ-secretase inhibitor II (GSI II); γ- secretase inhibitor III (GSI III), N- Benzyloxycarbonyl-Leu- leucinal, N-(2-Naphthoyl)-Val- phenylalaninal; γ-secretase inhibitor III (GSI IV); γ-secretase inhibitor III (GSI V), N- Benzyloxycarbonyl-Leu-phenylalaninal; γ- secretase inhibitor III (GSI VI), l-(S)-endo-N- (1,3,3)-Trimethylbicyclo[2.2.1]hept-2-yl)-4- fluorophenyl Sulfonamide; γ-secretase inhibitor III (GSI VII), Menthyloxycarbonyl-LL-CHO; γ- secretase inhibitor III (GSI IX), (DAPT), N- [N-(3,5-Difluorophenacetyl-L- alanyl)]-S- phenylglycine t- Butyl Ester; γ-secretase inhibitor X (GSI X), { 1 S-Benzyl-4R-[1-(1S- carbamoyl- 2- phenethylcarbamoyl)-lS-3- methylbutylcarb-amoyl]-2R-hydroxy-5- phenylpentyl}carbamic Acid tert-butyl Ester; γ- secretase inhibitor XI (GSI XI), 7-Amino-4-chloro-3- methoxyisocoumarin; γ-secretase inhibitor XII (GSI XII), Z-Ile-Leu-CHO; γ-secretase inhibitor XIII (GSI XIII), Z-Tyr-Ile-Leu- CHO; γ-secretase inhibitor XIV (GSI XIV), Z-Cys(t-Bu)-Ile- Leu-CHO; γ-secretase inhibitor XVI (GSI XVI), N-[N-3,5-Difluorophenacetyl]-L- alanyl-S- phenylglycine Methyl Ester; γ- secretase inhibitor XVII (GSI XVII); γ-secretase inhibitor XIX (GSI XIX), benzo[e][l,4]diazepin-3-yl)- butyramide; γ-secretase inhibitor XX (GSI XX), (S,S)- 2-[2-(3,5- Difluorophenyl)acetylamino]- N-(5-methyl-6-oxo-6,7- dihydro-5H- dibenzo[b,d]azepin-7- yl)propionamide; γ-secretase inhibitor XXI (GSI XXI), (S,S)-2-[2-(3,5- Difluorophenyl)- acetylamino]-N-(l-methyl-2- oxo-5-phenyl-2-,3-dihydro-lH- benzo[e][l,4]diazepin-3- yl)- propionamide; Gamma40 secretase inhibitor I, N-trans-3,5- Dimethoxycinnamoyl-Ile- leucinal; Gamma40 secretase inhibitor II, N-tert-Butyloxycarbonyl- Gly-Val-Valinal Isovaleryl-V V-Sta-A-Sta- OCH3; MK-0752 (Merck); MRK-003 (Merck); semagacestat/LY450139 (Eli Lilly); RO4929097; PF-03084,014; BMS-708163; MPC-7869 (γ- secretase modifier), YO-01027 (Dibenzazepine), Compound E ([(2S)-2-{[(3,5- Difluorophenyl)acetyl]amino}-N-[(3S)-l-methyl-2-oxo-5-phenyl-2,3-dihydro-lH-l,4-benzodiazepin-3-yl]propanamide], available from Alexis Biochemicals), LY411575 (Eli Lilly and Co.), L-685,458 (Sigma-Aldrich), BMS-289948 (4-chloro-N-(2,5-difluorophenyl)-N-((lR)-{4-fluoro-2-[3-(lH-imidazol-l-yl)propyl]phenyl}ethyl)benzenesulfonamide hydrochloride) and

BMS-299897 (4-[2-((lR)-l-{[(4-chlorophenyl)sulfonyl]-2,5-difluoroanilino}ethyl)-5-fluorophenyljbutanoic acid) (Bristol Myers Squibb).

[0268] In some embodiments, the gamma secretase inhibitor is LY3039478 or is a stereoisomer thereof or is a pharmaceutically acceptable salt or hydrate of the foregoing.

[0269] In some embodiments, the gamma secretase inhibitor is a compound of the structure:

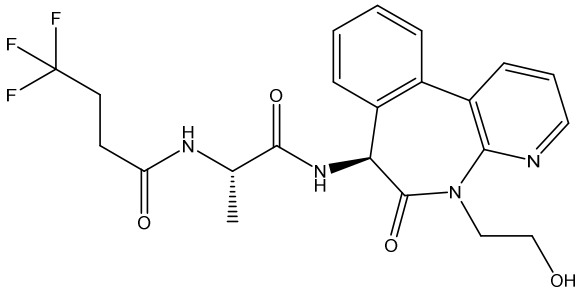
Compound 1



or a stereoisomer thereof or a pharmaceutically acceptable salt or hydrate of the foregoing.

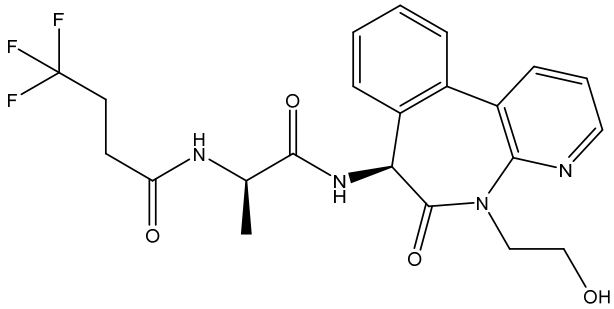
[0270] In some embodiments, Compound 1 is a single stereoisomer as depicted. In some cases, there are two chiral centers giving rise to four stereoisomers. Thus, in some embodiments, reference to Compound 1 also includes racemic mixtures including Compound 1. Specific stereoisomers can be prepared by stereospecific synthesis using enantiomerically pure or enriched starting materials. The specific stereoisomers of either starting materials, intermediates, or racemic mixtures including Compound 1 can be resolved by techniques well known in the art, such as those found in Stereochemistry of Organic Compounds, E. I. Eliel and S. H. Wilen (Wiley 1994) and Enantiomers, Racemates, and Resolutions, J., Jacques, A. Collet, and S. H. Wilen (Wiley 1991), including chromatography on chiral stationary phases, enzymatic resolutions, or fractional crystallization or chromatography of diastereomers formed for that purpose, such as diastereomeric salts.

[0271] In some embodiments, the gamma secretase inhibitor is a compound of the structure:



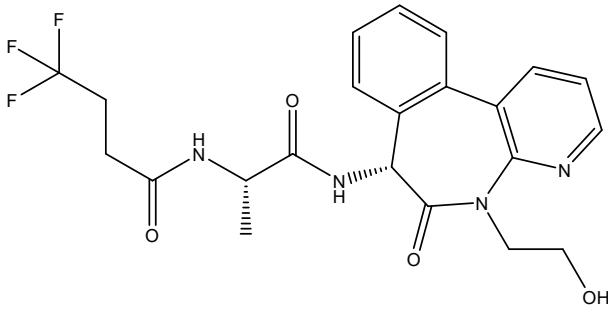
or a pharmaceutically acceptable salt or hydrate thereof.

[0272] In some embodiments, the gamma secretase inhibitor is a compound of the structure:



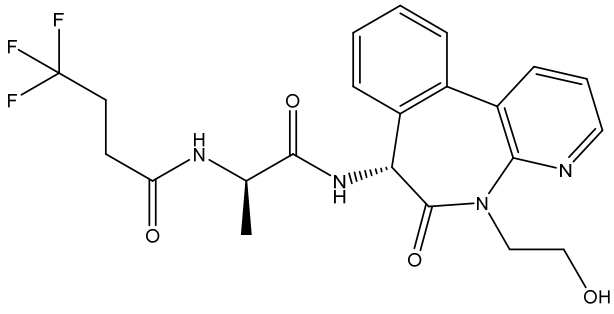
or a pharmaceutically acceptable salt or hydrate thereof.

[0273] In some embodiments, the gamma secretase inhibitor is a compound of the structure:



or a pharmaceutically acceptable salt or hydrate thereof.

[0274] In some embodiments, the gamma secretase inhibitor is a compound of the structure:



or a pharmaceutically acceptable salt or hydrate thereof.

[0275] In some embodiments, the compound is Compound 1. In some embodiments, Compound 1 is named: 4,4,4-trifluoro-N-[(1 S)-2-[[(7S)-5-(2-hydroxyethyl)-6-oxo-7H-pyrido[2,3-d][3]benzazepin-7-yl]amino]-1 -methyl-2-oxo-ethyl]butanamide; and may also be named: N-[(1S)-2-[[(7S)-6,7-dihydro-5-(2-hydroxyethyl)-6-oxo-5H-pyrido[3,2-a][3]benzazepin-7-yl]amino]-1-methyl-2-oxoethyl]-4,4,4-trifluorobutanamide; and other names may be used to unambiguously identify Compound 1. In some embodiments, Compound 1 is known as LY3039478. See also published PCT App. No. WO2013/016081 for the synthesis and preparion of Compound 1.

1. Compositions and formulations

[0276] In some embodiments of the combination therapy methods, compositions, combinations, kits and uses provided herein, the combination therapy can be administered in one or more compositions, *e.g.*, a pharmaceutical composition containing a gamma secretase inhibitor, or a pharmaceutically acceptable salt of hydrate thereof.

[0277] In some embodiments, the composition, *e.g.*, a pharmaceutical composition containing the gamma secretase inhibitor, or a pharmaceutically acceptable salt of hydrate thereof, can include carriers such as a diluent, adjuvant, excipient, or vehicle with which gamma secretase inhibitor, or a pharmaceutically acceptable salt of hydrate thereof, and/or the cells are administered. Examples of suitable pharmaceutical carriers are described in "Remington's Pharmaceutical Sciences" by E. W. Martin. Such compositions will contain a therapeutically effective amount of the gamma secretase inhibitor, or a pharmaceutically acceptable salt of hydrate thereof, generally in purified form, together with a suitable amount of carrier so as to provide the form for proper administration to the patient. Such pharmaceutical carriers can be sterile liquids, such as water and oils, including those of petroleum, animal, vegetable or synthetic origin, such as peanut oil, soybean oil, mineral oil, and sesame oil. Saline solutions and aqueous dextrose and glycerol solutions also can be employed as liquid carriers, particularly for injectable solutions. The pharmaceutical compositions can contain any one or more of a diluents(s), adjuvant(s), antiadherent(s), binder(s), coating(s), filler(s), flavor(s), color(s), lubricant(s), glidant(s), preservative(s), detergent(s), sorbent(s), emulsifying agent(s), pharmaceutical excipient(s), pH buffering agent(s), or sweetener(s) and a combination thereof. In some embodiments, the pharmaceutical composition can be liquid, solid, a lyophilized

powder, in gel form, and/or combination thereof. In some aspects, the choice of carrier is determined in part by the particular inhibitor and/or by the method of administration.

[0278] Pharmaceutically acceptable carriers are generally nontoxic to recipients at the dosages and concentrations employed, and include, but are not limited to: buffers such as phosphate, citrate, and other organic acids; antioxidants including ascorbic acid and methionine; preservatives (such as octadecyldimethylbenzyl ammonium chloride; hexamethonium chloride; benzalkonium chloride; benzethonium chloride; phenol, butyl or benzyl alcohol; alkyl parabens such as methyl or propyl paraben; catechol; resorcinol; cyclohexanol; 3-pentanol; and m-cresol); low molecular weight (less than about 10 residues) polypeptides; proteins, such as serum albumin, gelatin, or immunoglobulins; hydrophilic polymers such as polyvinylpyrrolidone; amino acids such as glycine, glutamine, asparagine, histidine, arginine, or lysine; monosaccharides, disaccharides, and other carbohydrates including glucose, mannose, or dextrins; chelating agents such as EDTA; sugars such as sucrose, mannitol, trehalose or sorbitol; salt-forming counter-ions such as sodium; metal complexes (*e.g.* Zn-protein complexes); and/or non-ionic surfactants such as polyethylene glycol (PEG), stabilizers and/or preservatives. The compositions containing gamma secretase inhibitor or a pharmaceutically acceptable salt of hydrate thereof can also be lyophilized.

[0279] In some embodiments, the pharmaceutical compositions can be formulated for administration by any route known to those of skill in the art including intramuscular, intravenous, intradermal, intralesional, intraperitoneal injection, subcutaneous, intratumoral, epidural, nasal, oral, vaginal, rectal, topical, local, otic, inhalational, buccal (*e.g.*, sublingual), and transdermal administration or any route. In some embodiments, other modes of administration also are contemplated. In some embodiments, the administration is by bolus infusion, by injection, *e.g.*, intravenous or subcutaneous injections, intraocular injection, periocular injection, subretinal injection, intravitreal injection, trans-septal injection, subscleral injection, intrachoroidal injection, intracameral injection, subconjectval injection, subconjuntival injection, sub-Tenon's injection, retrobulbar injection, peribulbar injection, or posterior juxtascleral delivery. In some embodiments, administration is by parenteral, intrapulmonary, and intranasal, and, if desired for local treatment, intralesional administration. Parenteral infusions include intramuscular, intravenous, intraarterial, intraperitoneal, or subcutaneous administration. In some embodiments, a given dose is administered by a single bolus administration. In some

embodiments, it is administered by multiple bolus administrations, for example, over a period of no more than 3 days, or by continuous infusion administration.

[0280] In some embodiments, the administration can be local, topical or systemic depending upon the locus of treatment. In some embodiments local administration to an area in need of treatment can be achieved by, for example, but not limited to, local infusion during surgery, topical application, *e.g.*, in conjunction with a wound dressing after surgery, by injection, by means of a catheter, by means of a suppository, or by means of an implant. In some embodiments, compositions also can be administered with other biologically active agents, either sequentially, intermittently or in the same composition. In some embodiments, administration also can include controlled release systems including controlled release formulations and device controlled release, such as by means of a pump. In some embodiments, the administration is oral.

[0281] In some embodiments, pharmaceutically and therapeutically active compounds and derivatives thereof are typically formulated and administered in unit dosage forms or multiple dosage forms. Each unit dose contains a predetermined quantity of therapeutically active compound sufficient to produce the desired therapeutic effect, in association with the required pharmaceutical carrier, vehicle or diluent. In some embodiments, unit dosage forms, include, but are not limited to, tablets, capsules, pills, powders, granules, sterile parenteral solutions or suspensions, and oral solutions or suspensions, and oil water emulsions containing suitable quantities of the compounds or pharmaceutically acceptable derivatives thereof. Unit dose forms can be contained ampoules and syringes or individually packaged tablets or capsules. Unit dose forms can be administered in fractions or multiples thereof. In some embodiments, a multiple dose form is a plurality of identical unit dosage forms packaged in a single container to be administered in segregated unit dose form. Examples of multiple dose forms include vials, bottles of tablets or capsules or bottles of pints or gallons.

2. Dosage Schedule of Gamma Secretase Inhibitor

[0282] In some embodiments, the provided combination therapy method involves administering to the subject a therapeutically effective amount of a gamma secretase inhibitor, or a pharmaceutically acceptable salt of hydrate thereof, and the cell therapy, such as a T cell therapy (*e.g.* CAR-expressing T cells).

[0283] In some embodiments, the administration of the gamma secretase inhibitor, or a pharmaceutically acceptable salt of hydrate thereof, is initiated prior to, subsequently to, during, during the course of, simultaneously, near simultaneously, sequentially and/or intermittently with the administration of the cell therapy, such as a T cell therapy (*e.g.* CAR-expressing T cells). In some embodiments, the method involves initiating the administration of gamma secretase inhibitor, or a pharmaceutically acceptable salt of hydrate thereof, prior to administration of the T cell therapy. In other embodiments, the method involves initiating the administration of the gamma secretase inhibitor, or a pharmaceutically acceptable salt of hydrate thereof, after administration of a cell therapy (e.g., a CAR-expressing T cell therapy). In some embodiments, the dosage schedule comprises initiating the administration of gamma secretase inhibitor, or a pharmaceutically acceptable salt of hydrate thereof, concurrently or simultaneously with the administration of a cell therapy (e.g., a CAR-expressing T cell therapy).

[0284] In some embodiments, the gamma secretase inhibitor, or a pharmaceutically acceptable salt of hydrate thereof is administered in a cycle. In some embodiments, the cycle comprises an administration period in which the gamma secretase inhibitor is administered followed by a rest period during which the gamma secretase inhibitor is not administered. In some embodiments, the total number of days of the cycle, e.g. from the beginning of initiating administration of the gamma secretase inhibitor, is greater than or greater than about or is about 1 day, 3, days, 7 days, 14 days, 21 days, 28 days, 30 days, 40 days, 50 days, 60 days or more.

[0285] In some embodiments, the initiation of the administration of the gamma secretase inhibitor is carried out in at least one cycle and initiation of administration of the cell therapy (e.g., CAR-expressing T cells) are carried out on the same day, optionally concurrently. In some embodiments, the initiation of the administration of the gamma secretase inhibitor in at least one cycle is prior to initiation of administration of the cell therapy. In some embodiments, the initiation of the administration of the gamma secretase inhibitor in at least one cycle is concurrent with or on the same day as initiation of administration of the cell therapy. In some embodiments, the gamma secretase inhibitor is administered from or from about 0 to 30 days, such as 0 to 15 days, 0 to 6 days, 0 to 96 hours, 0 to 24 hours, 0 to 12 hours, 0 to 6 hours, or 0 to 2 hours, 2 hours to 15 days, 2 hours to 6 days, 2 hours to 96 hours, 2 hours to 24 hours, 2 hours to 12 hours, 2 hours to 6 hours, 6 hours to 30 days, 6 hours to 15 days, 6 hours to 6 days, 6 hours to 96 hours, 6 hours to 24 hours, 6 hours to 12 hours, 12 hours to 30 days, 12 hours to 15 days,

12 hours to 6 days, 12 hours to 96 hours, 12 hours to 24 hours, 24 hours to 30 days, 24 hours to 15 days, 24 hours to 6 days, 24 hours to 96 hours, 96 hours to 30 days, 96 hours to 15 days, 96 hours to 6 days, 6 days to 30 days, 6 days to 15 days, or 15 days to 30 days prior to initiation of the cell therapy (e.g., CAR-expressing T cell therapy). In some aspects, the gamma secretase inhibitor is administered no more than about 96 hours, 72 hours, 48 hours, 24 hours, 12 hours, 6 hours, 2 hours or 1 hour prior to initiation of the T cell therapy.

[0286] In some of any such embodiments in which the gamma secretase inhibitor, or a pharmaceutically acceptable salt of hydrate thereof, is given prior to the cell therapy (*e.g.* T cell therapy, such as CAR-T cell therapy), the administration of the gamma secretase inhibitor, continues at regular intervals until the initiation of the cell therapy and/or for a time after the initiation of the cell therapy.

[0287] In some embodiments, the gamma secretase inhibitor is administered, or is further administered, after administration of the cell therapy (*e.g.* T cell therapy, such as CAR-T cell therapy). In some embodiments, the gamma secretase inhibitor is administered within or within about 1 hours, 2 hours, 6 hours, 12 hours, 24 hours, 48 hours, 96 hours, 4 days, 5 days, 6 days or 7 days, 14 days, 15 days, 21 days, 24 days, 28 days, 30 days, 36 days, 42 days, 60 days, 72 days or 90 days after initiation of administration of the cell therapy (*e.g.* T cell therapy).

[0288] In some embodiments, the provided methods involve continued administration, such as at regular intervals, of the gamma secretase inhibitor after initiation of administration of the cell therapy. For example, the gamma secretase inhibitor is administered at regular intervals of daily, every other day, every third day, three times a week, or once a week for a period of time.

[0289] In some embodiments, the gamma secretase inhibitor is administered, such as at regular intervals as described, for up to or up to about 1 day, up to or up to about 2 days, up to or up to about 3 days, up to or up to about 4 days, up to or up to about 5 days, up to or up to about 6 days, up to or up to about 7 days, up to or up to about 12 days, up to or up to about 14 days, up to or up to about 21 days, up to or up to about 24 days, up to or up to about 28 days, up to or up to about 30 days, up to or up to about 35 days, up to or up to about 42 days, up to or up to about 60 days or up to or up to about 90 days, up to or up to about 120 days, up to or up to about 180 days, up to or up to about 240 days, up to or up about 360 days, or up to or up to about 720 days or more after the initiation of administration of the cell therapy (*e.g.* T cell therapy, such as CAR-T cell therapy).

[0290] In some of any such above embodiments, the gamma secretase inhibitor is administered prior to and after initiation of administration of the cell therapy (*e.g.* T cell therapy, such as CAR-T cell therapy).

[0291] In some embodiments, the initiation of the administration of the gamma secretase inhibitor is carried out at or after, optionally immediately after or within 1 to 3 days after: (i) peak or maximum level of the cells of the T cell therapy are detectable in the blood of the subject; (ii) the number of cells of the T cell therapy detectable in the blood, after having been detectable in the blood, is not detectable or is reduced, optionally reduced compared to a preceding time point after administration of the T cell therapy; (iii) the number of cells of the T cell therapy detectable in the blood is decreased by or more than 1.2-fold, 1.5-fold, 2.0-fold, 3.0-fold, 4.0-fold, 5.0-fold, 10-fold or more the peak or maximum number cells of the T cell therapy detectable in the blood of the subject after initiation of administration of the T cell therapy; (iv) at a time after a peak or maximum level of the cells of the T cell therapy are detectable in the blood of the subject, the number of cells of or derived from the T cells detectable in the blood from the subject is less than 10%, less than 5%, less than 1% or less than 0.1% of total peripheral blood mononuclear cells (PBMCs) in the blood of the subject; (v) the subject exhibits disease progression and/or has relapsed following remission after treatment with the T cell therapy; and/or (iv) the subject exhibits increased tumor burden as compared to tumor burden at a time prior to or after administration of the T cells and prior to initiation of administration of the gamma secretase inhibitor.

[0292] In some aspects, initiation of administration of the gamma secretase inhibitior is at a time after the subject has relapsed or is suspected to or likely to relapse following administration of the cell therapy (e.g. T cell therapy, such as CAR T-cell therapy). In some cases, the cell therapy is an anti-BCMA CAR-T cell therapy and the gamma secretase inhibitor is administered at a time after the subject has relapsed or is suspected to or likely to relapse following administration of the anti-BCMA CAR-T cell therapy.

[0293] In some embodiments, the initiation of the administration of the gamma secretase inhibitor in at least one cycle is after initiation of administration of the cell therapy. In some embodiments, the initiation of the administration of the gamma secretase inhibitor is at least or about at least 1 day, at least or about at least 2 days, at least or about at least 3 days, at least or about at least 4 days, at least or about at least 5 days, at least or about at least 6 days, at least or

about at least 7 days, at least or about at least 8 days, at least or about at least 9 days, at least or about at least 10 days, at least or at least about 12 days, at least or about at least 14 days, at least or at least about 15 days, at least or about at least 21 days, at least or at least about 24 days, at least or about at least 28 days, at least or about at least 30 days, at least or about at least 35 days or at least or about at least 42 days, at least or about at least 60 days, or at least or about at least 90 days after initiation of the administration of the cell therapy. In some embodiments, the initiation of the administration of the gamma secretase inhibitor is carried out at least 2 days after, at least 1 week after, at least 2 weeks after, at least 3 weeks after, or at least 4 weeks after, the initiation of the administration of the cell therapy. In some embodiments, the initiation of the administration of the gamma secretase inhibitor is carried out 2 to 28 days or 7 to 21 days after initiation of administration of the cell therapy. In some embodiments, the initiation of the administration of the gamma secretase inhibitor is carried out at a time that is greater than or greater than about 14 days, 15 days, 16 days, 17 days, 18 days, 19, days, 20 days, 21 days, 24 days, or 28 days after initiation of the administration of the cell therapy.

[0294] In some embodiments, the gamma secretase inhibitor is administered several times a day, twice a day, daily, every other day, three times a week, twice a week, or once a week after initiation of the cell therapy. In some embodiments, the gamma secretase inhibitor is administered daily. In some embodiments the gamma secretase inhibitor is administered twice a day. In some embodiments, the gamma secretase inhibitor is administered three times a day. In other embodiments, the gamma secretase inhibitor is administered every other day.

[0295] In some embodiments, the gamma secretase inhibitor is administered daily. In some embodiments, the gamma secretase inhibitor is administered during the administration period for a plurality of consecutive days, such as for up to about 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, 30, or more than 30 consecutive days. In some embodiments, the gamma secretase inhibitor, is administered for greater than or greater than about 7 consecutive days, greater than or greater than about 14 consecutive days, greater than or greater than about 21 consecutive days, greater than or greater than about 21 consecutive days, or greater than or greater than about 28 consecutive days. In some embodiments, the gamma secretase inhibitor, is administered during the administration period for up to 21 consecutive days. In some embodiments, the gamma secretase inhibitor, is administered during the

administration period for up to 21 consecutive days, wherein the cycle comprises greater than 30 days beginning upon initiation of the administration of the gamma secretase inhibitor.

[0296] In some embodiments, the gamma secretase inhibitor is administered during the administration period for no more than about 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, 30, or no more than 30 consecutive days. In certain embodiments, the gamma secretase inhibitor is administered once daily for 14 days over a 21 day treatment cycle. In certain embodiments, the gamma secretase inhibitor is administered once daily for 21 days over a 28 day treatment cycle. In some embodiments, the gamma secretase inhibitor is administered during the administration period for no more than 14 consecutive days.

[0297] In some embodiments, the gamma secretase inhibitor is administered in a cycle, wherein the cycle comprises the administration of the gamma secretase inhibitor for a plurality of consecutive days followed by a rest period during which the gamma secretase inhibitor is not administered. In some embodiments, the rest period is greater than about 1 day, greater than about 3 consecutive days, greater than about 5 consecutive days, greater than about 7 consecutive days, greater than about 8 consecutive days, greater than about 9 consecutive days, greater than about 10 consecutive days, greater than about 11 consecutive days, greater than about 12 consecutive days, greater than about 13 consecutive days, greater than about 14 consecutive days, greater than about 15 consecutive days, greater than about 16 consecutive days, greater than about 17 consecutive days, greater than about 18 consecutive days, greater than about 19 consecutive days, greater than about 20 consecutive days, or greater than about 21 or more consecutive days. In some embodiments, the rest period is greater than 7 consecutive days, greater than 14 consecutive days, greater than 21 days, or greater than 28 days. In some embodiments, the rest period is greater than about 14 consecutive days. In some embodiments, the cycle of administration of the gamma secretase inhibitor does not contain a rest period.

[0298] In some embodiments, the gamma secretase inhibitor is administered in a cycle, wherein the cycle is repeated at least one time. In some embodiments, the gamma secretase inhibitor is administered for at least 2 cycles, at least 3 cycles, at least 4 cycles, at least 5 cycles, at least 6 cycles, at least 7 cycles, at least 8 cycles, at least 9 cycles, at least 10 cycles, at least 11 cycles, or at least 12 cycles. In some embodiments, the gamma secretase inhibitor is

administered for 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, or 24 cycles.

[0299] In some embodiments, the gamma secretase inhibitor is administered at least six times daily, five times daily, four times daily, three times daily, twice daily, once daily, every other day, every three days, three times a week, twice weekly, once weekly or only one time prior to or subsequently to initiation of administration of the cell therapy. In some embodiments, the gamma secretase inhibitor is administered three times a week. In some embodiments, the administration of the inhibitor is carried out in a treatment cycle that is at least or at least about or 14 days, at least or at least about or 21 days or at least or at least about or 28 days. In some embodiments, the gamma secretase inhibitor is administered in multiple doses in regular intervals prior to, during, during the course of, and/or after the period of administration of the cell therapy. In some embodiments, the gamma secretase inhibitor is administered in one or more doses in regular intervals prior to the administration of the cell therapy. In some embodiments, the gamma secretase inhibitor is administered in one or more doses in regular intervals after the administration of the cell therapy. In some embodiments, one or more of the doses of the gamma secretase inhibitor, can occur simultaneously with the administration of a dose of the cell therapy.

[0300] In some embodiments, the dose, frequency, duration, timing and/or order of administration of the gamma secretase inhibitor, is determined, based on particular thresholds or criteria of results of the screening step and/or assessment of treatment outcomes described herein, *e.g.*, those described in Section III herein.

[0301] In some embodiments, the method involves administering the cell therapy to a subject that has been previously administered a therapeutically effective amount of the gamma secretase inhibitor. In some embodiments, the gamma secretase inhibitor is administered to a subject before administering a dose of cells expressing a recombinant receptor to the subject. In some embodiments, the treatment with the gamma secretase inhibitor occurs at the same time as the administration of the dose of cells. In some embodiments, the gamma secretase inhibitor is administered after the administration of the dose of cells.

[0302] In some embodiments, administration of the gamma secretase inhibitor is carried out for an administration period of 2 to 28 days, such as 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, or more than 21 days.In some embodiments, the gamma secretase inhibitor is

administered daily for 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, or more than 21 days. In some embodiments, the gamma secretase inhibitor is administered twice a day for 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, or more than 21 days. In some embodiments, the gamma secretase inhibitor is administered three times a day for 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, or more than 21 days. In some embodiments, the gamma secretase inhibitor is administered every other day for 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, or more than 21 days. In some embodiments, the gamma secretase inhibitor is administered three times a week for 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, or more than 21 days.

[0303] In some embodiments, the gamma secretase inhibitor is administered on days 2, 4, 7, 9, 11, 14, 16, and 18 after initiation of administration of the cell therapy (*e.g.* T cell therapy).

[0304] In some embodiments of the methods provided herein, the gamma secretase inhibitor, and the cell therapy are administered simultaneously or near simultaneously.

[0305] In some embodiments, gamma secretase inhibitor is administered at a total daily dosage amount of at most or at most about 50 mg/day, 100 mg/day, 150 mg/day, 175 mg/day, 200 mg/day, 250 mg/day, 300 mg/day, 350 mg/day, 400 mg/day, 450 mg/day, 500 mg/day, 600 mg/day, 700 mg/day,, 800 mg/day or less. In some embodiments, the gamma secretase inhibitor is administered, or each administration of the inhibitor is independently administered, in an amount of 0.5 mg to 500 mg, 0.5 mg to 250 mg, 0.5 mg to 100 mg, 0.5 mg to 50 mg, 0.5 mg to 25 mg, 0.5 mg to 10 mg, 0.5 mg to 5.0 mg, 0.5 mg to 2.5 mg, 0.5 mg to 1.0 mg, 1.0 mg to 500 mg, 1.0 mg to 250 mg, 1.0 mg to 100 mg, 1.0 mg to 50 mg, 1.0 mg to 25 mg, 1.0 mg to 10 mg, 1.0 mg to 5.0 mg, 1.0 mg to 2.5 mg, 2.5 mg to 500 mg, 2.5 mg to 250 mg, 2.5 mg to 100 mg, 2.5 mg to 50 mg, 2.5 mg to 25 mg, 2.5 mg to 10 mg, 2.5 mg to 5.0 mg, 5.0 mg to 500 mg, 5.0 mg to 250 mg, 5.0 mg to 100 mg, 5.0 mg to 50 mg, 5.0 mg to 25 mg, 5.0 mg to 10 mg, 10 mg to 500 mg, 10 mg to 250 mg, 10 mg to 100 mg, 10 mg to 50 mg, 10 mg to 25 mg, 25 mg to 500 mg, 25 mg to 250 mg, 25 mg to 100 mg, 25 mg to 50 mg, 50 mg to 500 mg, 50 mg to 250 mg, 50 mg to 100 mg, 100 mg to 500 mg, 100 mg to 250 mg or 250 mg to 500 mg. In some embodiments, the inhibitor is administered, or each administration of the inhibitor is independently administered, in an amount that is at least or at least about or is or is about 0.5 mg, 1.0 mg, 2.5 mg, 5.0 mg, 10.0 mg, 25 mg, 50 mg, 100 mg, 250 mg or 500 mg. In some embodiments, the amount is a once daily amount of the gamma secretase inhibitor. Such dosage amounts can be administered at regular intervals as described, such as every other day, every

third day, three times a week or once a week for an administration period. In some embodiments, the administration period is 2 to 28 days, such as 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, or more than 21 days.

[0306] In some embodiments, the gamma secretase inhibitor is administered at a dosage of from about 1 mg to about 20 mg, e.g., from about 1 mg to about 10 mg, from about 2.5 mg to about 7.5 mg, from about 5 mg to about 15 mg, such as about 5 mg, 10 mg, 15 mg or 20 mg. In some embodiments, the gamma secretase inhibitor is administered at a dose of from at or about 10 µg/kg to at or about 5 mg/kg, such as at or about 50 µg/kg to at or about 2 mg/kg , at or about 50 µg/kg to at or about 1 mg/kg, at or about 50 µg/kg to at or about 500 µg/kg, at or about 50 µg/kg to at or about 250 µg/kg, at or about 100 µg/kg to at or about 200 µg/kg, at or about 50 to at or bout 100 µg/kg, at or about 100 µg/kg to about 2 mg/kg, at or about 100 µg/kg to at or about 1 mg/kg, at or about 100 µg/kg to at or about 500 µg/kg, at or about 100 µg/kg to at or about 250 µg/kg, at or about 100 µg/kg to at or about 200 µg/kg, at or about 100 µg/kg, at or about 200 µg/kg to about 2 mg/kg, at or about 200 µg/kg to at or about 1 mg/kg, at or about 200 µg/kg to at or about at or about 500 µg/kg, at or about 200 µg/kg to at or about 250 µg/kg or at or about 250 µg/kg to at or about 2 mg/kg, at or about 250 µg/kg to at or about 1 mg/kg, at or about 250 µg/kg to at or about at or about 500 µg/kg, In some embodiments, the gamma secretase inhibitor is administered at a dose of from at or about 400 µg/kg to at or about 600 µg/kg, such as at or about 500 µg/kg. In some embodiments, the amount is a once daily amount of the gamma secretase inhibitor. Such dosage amounts can be administered at regular intervals as described, such as every other day, every third day, three times a week or once a week for an administration period. In some embodiments, the administration period is 2 to 28 days, such as 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, or more than 21 days.

[0307] In some embodiments, the gamma secretase inhibitor is administered at a total daily dosage amount of at least or at least about 0.1 mg per day, 0.5 mg per day, 1.0 mg per day, 2.5 mg per day, 5 mg per day, 10 mg per day, 25 mg per day, 50 mg per day or 100 mg per day. In some embodiments, the dose of the inhibitor is about 25 mg per day. In particular embodiments, the dose of the inhibitor is or is about10 mg per day. Such dosage amounts can be administered at regular intervals as described, such as every other day, every third day, three times a week or once a week for an administration period. In some embodiments, the administration period is 2 to 28 days, such as 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, or more than 21 days.

[0308] In some embodiments, the gamma secretase inhibitor is administered in an amount greater than or greater than about 1 mg, 2.5 mg, 5 mg, 7.5 mg, 10 mg, 15 mg and less than 25 mg. In some embodiments, the gamma secretase inhibitor is administered in an amount greater than or greater than about 1 mg per day, 2.5 mg per day, 5 mg per day, 7.5 mg per day, 10 mg per day, 15 mg per day and less than 25 mg per day. Such dosage amounts can be administered at regular intervals as described, such as every other day, every third day, three times a week or once a week for an administration period. In some embodiments, the administration period is 2 to 28 days, such as 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, or more than 21 days.

[0309] In some embodiments, the gamma secretase inhibitor is administered on days 2, 4, 7, 9, 11, 14, 16, and 18 after initiation of administration of the cell therapy (*e.g.* T cell therapy).

[0310] In some embodiments, dosages, such as daily dosages, are administered in one or more divided doses, such as 2, 3, or 4 doses, or in a single formulation. The gamma secretase inhibitor can be administered alone, in the presence of a pharmaceutically acceptable carrier, or in the presence of other therapeutic agents.

[0311] One skilled in the art will recognize that higher or lower dosages of the gamma secretase inhibitor could be used, for example depending on the particular agent and the route of administration. In some embodiments, the gamma secretase inhibitor may be administered alone or in the form of a pharmaceutical composition wherein the compound is in admixture or mixture with one or more pharmaceutically acceptable carriers, excipients, or diluents. In some embodiments, the gamma secretase inhibitor may be administered either systemically or locally to the organ or tissue to be treated. Exemplary routes of administration include, but are not limited to, topical, injection (such as subcutaneous, intramuscular, intradermal, intraperitoneal, intratumoral, and intravenous), oral, sublingual, rectal, transdermal, intranasal, vaginal and inhalation routes. In some embodiments, the route of administration is oral, parenteral, rectal, nasal, topical, or ocular routes, or by inhalation. In some embodiments, the gamma secretase inhibitor is administered orally. In some embodiments, the gamma secretase inhibitor is administered orally in solid dosage forms, such as capsules, tablets and powders, or in liquid dosage forms, such as elixirs, syrups and suspensions.

[0312] Once improvement of the patient's disease has occurred, the dose may be adjusted for preventative or maintenance treatment. For example, the dosage or the frequency of administration, or both, may be reduced as a function of the symptoms, to a level at which the

desired therapeutic or prophylactic effect is maintained. If symptoms have been alleviated to an appropriate level, treatment may cease. Patients may, however, require intermittent treatment on a long-term basis upon any recurrence of symptoms. Patients may also require chronic treatment on a long-term basis.

3. Additional Therapy

[0313] In some aspects, the provided methods can further include administering one or more lymphodepleting therapies, such as prior to or simultaneous with initiation of administration of the cell therapy (e.g., T cell therapy). In some embodiments, the lymphodepleting therapy comprises administration of a phosphamide, such as cyclophosphamide. In some embodiments, the lymphodepleting therapy can include administration of fludarabine.

[0314] In some aspects, preconditioning subjects with immunodepleting (e.g., lymphodepleting) therapies can improve the effects of adoptive cell therapy (ACT). Preconditioning with lymphodepleting agents, including combinations of cyclosporine and fludarabine, have been effective in improving the efficacy of transferred tumor infiltrating lymphocyte (TIL) cells in cell therapy, including to improve response and/or persistence of the transferred cells. *See, e.g.,* Dudley et al., *Science*, 298, 850-54 (2002); Rosenberg et al., *Clin Cancer Res*, 17(13):4550–4557 (2011). Likewise, in the context of CAR+ T cells, several studies have incorporated lymphodepleting agents, most commonly cyclophosphamide, fludarabine, bendamustine, or combinations thereof, sometimes accompanied by low-dose irradiation. *See* Han et al. *Journal of Hematology & Oncology*, 6:47 (2013); Kochenderfer et al., *Blood,* 119: 2709-2720 (2012); Kalos et al., *Sci Transl Med*, 3(95):95ra73 (2011); Clinical Trial Study Record Nos.: NCT02315612; NCT01822652.

[0315] Such preconditioning can be carried out with the goal of reducing the risk of one or more of various outcomes that could dampen efficacy of the therapy. These include the phenomenon known as "cytokine sink," by which T cells, B cells, NK cells compete with TILs for homeostatic and activating cytokines, such as IL-2, IL-7, and/or IL-15; suppression of TILs by regulatory T cells, NK cells, or other cells of the immune system; impact of negative regulators in the tumor microenvironment. Muranski et al., *Nat Clin Pract Oncol.* December; 3(12): 668–681 (2006).

[0316] Thus in some embodiments, the provided method further involves administering a lymphodepleting therapy to the subject. In some embodiments, the method involves

administering the lymphodepleting therapy to the subject prior to the administration of the dose of cells. In some embodiments, the lymphodepleting therapy contains a chemotherapeutic agent such as fludarabine and/or cyclophosphamide. In some embodiments, the administration of the cells and/or the lymphodepleting therapy is carried out via outpatient delivery.

[0317] In some embodiments, the methods include administering a preconditioning agent, such as a lymphodepleting or chemotherapeutic agent, such as cyclophosphamide, fludarabine, or combinations thereof, to a subject prior to the administration of the dose of cells. For example, the subject may be administered a preconditioning agent at least 2 days prior, such as at least 3, 4, 5, 6, or 7 days prior, to the first or subsequent dose. In some embodiments, the subject is administered a preconditioning agent no more than 7 days prior, such as no more than 6, 5, 4, 3, or 2 days prior, to the administration of the dose of cells.

[0318] In some embodiments, the subject is preconditioned with cyclophosphamide at a dose between or between about 20 mg/kg and 100 mg/kg of body surface area of the subject, such as between or between about 40 mg/kg and 80 mg/kg. In some aspects, the subject is preconditioned with or with about 60 mg/kg of cyclophosphamide. In some embodiments, the fludarabine can be administered in a single dose or can be administered in a plurality of doses, such as given daily, every other day or every three days. In some embodiments, the cyclophosphamide is administered once daily for one or two days. In some embodiments, where the lymphodepleting agent comprises cyclophosphamide, the subject is administered cyclophosphamide at a dose between or between about 100 mg/m^2 and 500 mg/m^2 body surface area of the subject, such as between or between about 200 mg/m^2 and 400 mg/m^2, or 250 mg/m^2 and 350 mg/m^2, inclusive. In some instances, the subject is administered about 300 mg/m^2 of cyclophosphamide. In some embodiments, the cyclophosphamide can be administered in a single dose or can be administered in a plurality of doses, such as given daily, every other day or every three days. In some embodiments, cyclophosphamide is administered daily, such as for 1-5 days, for example, for 2 to 4 days. In some instances, the subject is administered about 300 mg/m^2 body surface area of the subject, of cyclophosphamide, daily for 3 days, prior to initiation of the cell therapy.In some embodiments, where the lymphodepleting agent comprises fludarabine, the subject is administered fludarabine at a dose between or between about 1 mg/m^2 and 100 mg/m^2 of body surface area of the subject, such as between or between about 10 mg/m^2 and 75 mg/m^2, 15 mg/m^2 and 50 mg/m^2, 20 mg/m^2 and 30 mg/m^2, 20 mg/m^2 and 40 mg/m^2, 24

mg/m^2 and 35 mg/m^2, or 24 mg/m^2 and 26 mg/m^2, each inclusive. In some instances, the subject is administered about 30 mg/m^2 of fludarabine. In some instances, the subject is administered 25 mg/m^2 of fludarabine. In some embodiments, the fludarabine can be administered in a single dose or can be administered in a plurality of doses, such as given daily, every other day or every three days. In some embodiments, fludarabine is administered daily, such as for 1-5 days, for example, for 3 to 5 days or for 3-4 days. In some instances, the subject is administered about 30 mg/m^2 body surface area of the subject, of fludarabine, daily for 3 days, prior to initiation of the cell therapy.

[0319] In some embodiments, the lymphodepleting agent comprises a combination of agents, such as a combination of cyclophosphamide and fludarabine. Thus, the combination of agents may include cyclophosphamide at any dose or administration schedule, such as those described above, and fludarabine at any dose or administration schedule, such as those described above. For example, in some aspects, the subject is administered 60 mg/kg (~2 g/m^2) of cyclophosphamide and 3 to 5 doses of 25 mg/m^2 fludarabine prior to the dose of cells. In some aspects, the subject is administered fludarabine at or about 30 mg/m^2 body surface area of the subject, daily, and cyclophosphamide at or about 300 mg/m^2 body surface area of the subject, daily, for 3 days.

[0320] In one exemplary dosage regime, prior to receiving the first dose, subjects receive a gamma secretase inhibitor 1 day before the administration of cells and a lymphodepleting preconditioning chemotherapy of cyclophosphamide and fludarabine (CY/FLU), which is administered at least two days before the first dose of CAR-expressing cells and generally no more than 7 days before administration of cells. In another exemplary dosage regime, subjects receive the gamma secretase inhibitor concurrently with the administration of cells, such as on the same day. In yet another exemplary dosage regime, subjects receive the gamma secretase inhibitor several days after the administration of cells, such as 7, 8, 9, 10, 11, 12, 13, 14, or more than 14 days after. In some cases, for example, cyclophosphadmide is given from 24 to 27 days after the administration of the gamma secretase inhibitor. After preconditioning treatment, subjects are administered the dose of CAR-expressing T cells as described above.

[0321] In some embodiments, the administration of the preconditioning agent prior to infusion of the dose of cells improves an outcome of the treatment. For example, in some aspects, preconditioning improves the efficacy of treatment with the dose or increases the

persistence of the recombinant receptor-expressing cells (e.g., CAR-expressing cells, such as CAR-expressing T cells) in the subject. In some embodiments, preconditioning treatment increases disease-free survival, such as the percent of subjects that are alive and exhibit no minimal residual or molecularly detectable disease after a given period of time following the dose of cells. In some embodiments, the time to median disease-free survival is increased.

[0322] Once the cells are administered to the subject (e.g., human), the biological activity of the engineered cell populations in some aspects is measured by any of a number of known methods. Parameters to assess include specific binding of an engineered or natural T cell or other immune cell to antigen, in vivo, e.g., by imaging, or ex vivo, e.g., by ELISA or flow cytometry. In certain embodiments, the ability of the engineered cells to destroy target cells can be measured using any suitable method known in the art, such as cytotoxicity assays described in, for example, Kochenderfer et al., *J. Immunotherapy,* 32(7): 689-702 (2009), and Herman et al. *J. Immunological Methods*, 285(1): 25-40 (2004). In certain embodiments, the biological activity of the cells also can be measured by assaying expression and/or secretion of certain cytokines, such as CD 107a, IFNγ, IL-2, and TNF. In some aspects the biological activity is measured by assessing clinical outcome, such as reduction in tumor burden or load. In some aspects, toxic outcomes, persistence and/or expansion of the cells, and/or presence or absence of a host immune response, are assessed.

[0323] In some embodiments, the administration of the preconditioning agent prior to infusion of the dose of cells improves an outcome of the treatment such as by improving the efficacy of treatment with the dose or increases the persistence of the recombinant receptor-expressing cells (e.g., CAR-expressing cells, such as CAR-expressing T cells) in the subject. Therefore, in some embodiments, the dose of preconditioning agent given in the method which is a combination therapy with the gamma secretase inhibitor and cell therapy is higher than the dose given in the method without the gamma secretase inhibitor.

II. CELL THERAPY AND ENGINEERING CELLS

[0324] In some embodiments, the cell therapy (e.g., T cell therapy) for use in accord with the provided combination therapy methods includes administering engineered cells expressing recombinant receptors designed to recognize and/or specifically bind to molecules associated with the disease or condition and result in a response, such as an immune response against such molecules upon binding to such molecules. The receptors may include chimeric receptors, *e.g.*,

chimeric antigen receptors (CARs), and other transgenic antigen receptors including transgenic T cell receptors (TCRs).

[0325] In some embodiments, the cells contain or are engineered to contain an engineered receptor, *e.g.*, an engineered antigen receptor, such as a chimeric antigen receptor (CAR), or a T cell receptor (TCR). Also provided are populations of such cells, compositions containing such cells and/or enriched for such cells, such as in which cells of a certain type such as T cells or $CD8^+$ or $CD4^+$ cells are enriched or selected. Among the compositions are pharmaceutical compositions and formulations for administration, such as for adoptive cell therapy. Also provided are therapeutic methods for administering the cells and compositions to subjects, *e.g.*, patients.

[0326] Thus, in some embodiments, the cells include one or more nucleic acids introduced via genetic engineering, and thereby express recombinant or genetically engineered products of such nucleic acids. In some embodiments, gene transfer is accomplished by first stimulating the cells, such as by combining it with a stimulus that induces a response such as proliferation, survival, and/or activation, *e.g.*, as measured by expression of a cytokine or activation marker, followed by transduction of the activated cells, and expansion in culture to numbers sufficient for clinical applications.

A. RECOMBINANT RECEPTORS

[0327] The cells generally express recombinant receptors, such as antigen receptors including functional non-TCR antigen receptors, *e.g.*, chimeric antigen receptors (CARs), and other antigen-binding receptors such as transgenic T cell receptors (TCRs). Also among the receptors are other chimeric receptors.

[0328] In some embodiments of the provided methods and uses, recombinant receptors, including chimeric receptors, such as a chimeric antigen receptors, contain one or more domains that combine a ligand-binding domain (e.g. antibody or antibody fragment) that provides specificity for a desired antigen (e.g., tumor antigen) with intracellular signaling domains. In some embodiments, the intracellular signaling domain is an activating intracellular domain portion, such as a T cell activating domain, providing a primary activation signal. In some embodiments, the intracellular signaling domain contains or additionally contains a costimulatory signaling domain to facilitate effector functions. In some embodiments, chimeric receptors when genetically engineered into immune cells can modulate T cell activity, and, in

some cases, can modulate T cell differentiation or homeostasis, thereby resulting in genetically engineered cells with improved longevity, survival and/or persistence *in vivo*, such as for use in adoptive cell therapy methods.

1. Chimeric Antigen Receptors (CARs)

[0329] In some embodiments, engineered cells, such as T cells, are provided that express a CAR with specificity for a particular antigen (or marker or ligand or receptor), such as an antigen expressed on the surface of a particular cell type.

[0330] In particular embodiments, the recombinant receptor, such as a chimeric receptor, contains an intracellular signaling region, which includes a cytoplasmic signaling domain (also interchangeably called an intracellular signaling domain), such as a cytoplasmic (intracellular) region capable of inducing a primary activation signal in a T cell, for example, a cytoplasmic signaling domain of a T cell receptor (TCR) component (e.g. a cytoplasmic signaling domain of a zeta chain of a CD3-zeta (CD3ζ) chain or a functional variant or signaling portion thereof) and/or that comprises an immunoreceptor tyrosine-based activation motif (ITAM).

[0331] In some embodiments, the chimeric receptor further contains an extracellular ligand-binding domain that specifically binds to a ligand (e.g. antigen) antigen. In some embodiments, the chimeric receptor is a CAR that contains an extracellular antigen-recognition domain that specifically binds to an antigen. In some embodiments, the ligand, such as an antigen, is a protein expressed on the surface of cells. The chimeric receptors, such as CARs, generally include an extracellular antigen binding domain, such as a portion of an antibody molecule, generally a variable heavy (V_H) chain region and/or variable light (V_L) chain region of the antibody, *e.g.,* an scFv antibody fragment. In some embodiments, the CAR is a TCR-like CAR and the antigen is a processed peptide antigen, such as a peptide antigen of an intracellular protein, which, like a TCR, is recognized on the cell surface in the context of a major histocompatibility complex (MHC) molecule.

[0332] Exemplary antigen receptors, including CARs, and methods for engineering and introducing such receptors into cells, include those described, for example, in international patent application publication numbers WO200014257, WO2013126726, WO2012/129514, WO2014031687, WO2013/166321, WO2013/071154, WO2013/123061, WO2016/0046724, WO2016/014789, WO2016/090320, WO2016/094304, WO2017/025038, WO2017/173256, U.S. patent application publication numbers US2002131960, US2013287748, US20130149337,

U.S. Patent Nos. 6,451,995, 7,446,190, 8,252,592, 8,339,645, 8,398,282, 7,446,179, 6,410,319, 7,070,995, 7,265,209, 7,354,762, 7,446,191, 8,324,353, 8,479,118, and 9,765,342, and European patent application number EP2537416, and/or those described by Sadelain *et al.*, *Cancer Discov*., 3(4): 388–398 (2013); Davila *et al.*, *PLoS ONE* 8(4): e61338 (2013); Turtle *et al.*, *Curr. Opin. Immunol.*, 24(5): 633-39 (2012); Wu *et al.*, *Cancer,* 18(2): 160-75 (2012). In some aspects, the antigen receptors include a CAR as described in U.S. Patent No. 7,446,190, and those described in International Patent Application Publication No. WO/2014055668 A1. Examples of the CARs include CARs as disclosed in any of the aforementioned publications, such as WO2014031687, US 8,339,645, US 7,446,179, US 2013/0149337, U.S. Patent No. 7,446,190, US Patent No. 8,389,282, Kochenderfer *et al.*, *Nature Reviews Clinical Oncology*, 10, 267-276 (2013); Wang *et al.*, *J. Immunother.* 35(9): 689-701 (2012); and Brentjens *et al.*, *Sci Transl Med.* 5(177) (2013). See also WO2014031687, US 8,339,645, US 7,446,179, US 2013/0149337, U.S. Patent No. 7,446,190, US Patent No. 8,389,282, WO 2016/090320, WO2016090327, WO2010104949A2 and WO2017173256.

[0333] In some embodiments, the CAR is constructed with a specificity for a particular antigen (or marker or ligand or receptor), such as an antigen expressed in a particular cell type to be targeted by adoptive therapy, *e.g.*, a cancer marker, and/or an antigen intended to induce a dampening response, such as an antigen expressed on a normal or non-diseased cell type. Thus, the CAR typically includes in its extracellular portion one or more antigen binding molecules, such as one or more antigen-binding fragment, domain, or portion, or one or more antibody variable domains, and/or antibody molecules. In some embodiments, the CAR includes an antigen-binding portion or portions of an antibody molecule, such as a single-chain antibody fragment (scFv) derived from the variable heavy (VH) and variable light (VL) chains of a monoclonal antibody (mAb).

[0334] In some embodiments, the antibody or antigen-binding portion thereof is expressed on cells as part of a recombinant receptor, such as an antigen receptor. Among the antigen receptors are functional non-TCR antigen receptors, such as chimeric antigen receptors (CARs). Generally, a CAR containing an antibody or antigen-binding fragment that exhibits TCR-like specificity directed against peptide-MHC complexes also may be referred to as a TCR-like CAR. In some embodiments, the extracellular antigen binding domain specific for an MHC-peptide complex of a TCR-like CAR is linked to one or more intracellular signaling

components, in some aspects via linkers and/or transmembrane domain(s). In some embodiments, such molecules can typically mimic or approximate a signal through a natural antigen receptor, such as a TCR, and, optionally, a signal through such a receptor in combination with a costimulatory receptor.

[0335] In some embodiments, the recombinant receptor, such as a chimeric receptor (e.g. CAR), includes a ligand-binding domain that binds, such as specifically binds, to an antigen (or a ligand or receptor). Among the antigens targeted by the chimeric receptors are those expressed in the context of a disease, condition, or cell type to be targeted via the adoptive cell therapy. Among the diseases and conditions are proliferative, neoplastic, and malignant diseases and disorders, including cancers and tumors, including hematologic cancers, cancers of the immune system, such as lymphomas, leukemias, and/or myelomas, such as B, T, and myeloid leukemias, lymphomas, and multiple myelomas.

[0336] In some embodiments, the antigen targeted by the receptor is a polypeptide. In some embodiments, it is a carbohydrate or other molecule. In some embodiments, the antigen is selectively expressed or overexpressed on cells of the disease or condition, *e.g.*, the tumor or pathogenic cells, as compared to normal or non-targeted cells or tissues. In other embodiments, the antigen is expressed on normal cells and/or is expressed on the engineered cells.

[0337] In some embodiments, the CAR contains an antibody or an antigen-binding fragment (*e.g.* scFv) that specifically recognizes an antigen, such as an intact antigen, expressed on the surface of a cell.

[0338] In some embodiments, the antigen (or a ligand or receptor) is a tumor antigen or cancer marker. In certain embodiments, the antigen is or includes αvβ6 integrin (avb6 integrin), B cell maturation antigen (BCMA), B7-H3, B7-H6, carbonic anhydrase 9 (CA9, also known as CAIX or G250), a cancer-testis antigen, cancer/testis antigen 1B (CTAG, also known as NY-ESO-1 and LAGE-2), carcinoembryonic antigen (CEA), a cyclin, cyclin A2, C-C Motif Chemokine Ligand 1 (CCL-1), CD19, CD20, CD22, CD23, CD24, CD30, CD33, CD38, CD44, CD44v6, CD44v7/8, CD123, CD133, CD138, CD171, chondroitin sulfate proteoglycan 4 (CSPG4), epidermal growth factor protein (EGFR), type III epidermal growth factor receptor mutation (EGFR vIII), epithelial glycoprotein 2 (EPG-2), epithelial glycoprotein 40 (EPG-40), ephrinB2, ephrine receptor A2 (EPHa2), estrogen receptor, Fc receptor like 5 (FCRL5; also known as Fc receptor homolog 5 or FCRH5), fetal acetylcholine receptor (fetal AchR), a folate

binding protein (FBP), folate receptor alpha, ganglioside GD2, O-acetylated GD2 (OGD2), ganglioside GD3, glycoprotein 100 (gp100), glypican-3 (GPC3), G Protein Coupled Receptor 5D (GPCR5D), Her2/neu (receptor tyrosine kinase erb-B2), Her3 (erb-B3), Her4 (erb-B4), erbB dimers, Human high molecular weight-melanoma-associated antigen (HMW-MAA), hepatitis B surface antigen, Human leukocyte antigen A1 (HLA-A1), Human leukocyte antigen A2 (HLA-A2), IL-22 receptor alpha(IL-22Rα), IL-13 receptor alpha 2 (IL-13Rα2), kinase insert domain receptor (kdr), kappa light chain, L1 cell adhesion molecule (L1-CAM), CE7 epitope of L1-CAM, Leucine Rich Repeat Containing 8 Family Member A (LRRC8A), Lewis Y, Melanoma-associated antigen (MAGE)-A1, MAGE-A3, MAGE-A6, MAGE-A10, mesothelin (MSLN), c-Met, murine cytomegalovirus (CMV), mucin 1 (MUC1), MUC16, natural killer group 2 member D (NKG2D) ligands, melan A (MART-1), neural cell adhesion molecule (NCAM), oncofetal antigen, Preferentially expressed antigen of melanoma (PRAME), progesterone receptor, a prostate specific antigen, prostate stem cell antigen (PSCA), prostate specific membrane antigen (PSMA), Receptor Tyrosine Kinase Like Orphan Receptor 1 (ROR1), survivin, Trophoblast glycoprotein (TPBG also known as 5T4), tumor-associated glycoprotein 72 (TAG72), Tyrosinase related protein 1 (TRP1, also known as TYRP1 or gp75), Tyrosinase related protein 2 (TRP2, also known as dopachrome tautomerase, dopachrome delta-isomerase or DCT), vascular endothelial growth factor receptor (VEGFR), vascular endothelial growth factor receptor 2 (VEGFR2), Wilms Tumor 1 (WT-1), a pathogen-specific or pathogen-expressed antigen, or an antigen associated with a universal tag, and/or biotinylated molecules, and/or molecules expressed by HIV, HCV, HBV or other pathogens. Antigens targeted by the receptors in some embodiments include antigens associated with a B cell malignancy, such as any of a number of known B cell marker. In some embodiments, the antigen is or includes BCMA, CD20, CD19, CD22, ROR1, CD45, CD21, CD5, CD33, Igkappa, Iglambda, CD79a, CD79b or CD30.

[0339] In some embodiments, the antigen (or a ligand) is or includes orphan tyrosine kinase receptor ROR1, B cell maturation antigen (BCMA), carbonic anhydrase 9 (CAIX), Her2/neu (receptor tyrosine kinase erbB2), L1-CAM, CD19, CD20, CD22, mesothelin, CEA, and hepatitis B surface antigen, anti-folate receptor, CD23, CD24, CD30, CD33, CD38, CD44, EGFR, epithelial glycoprotein 2 (EPG-2), epithelial glycoprotein 40 (EPG-40), EPHa2, erb-B2, erb-B3, erb-B4, erbB dimers, EGFR vIII, folate binding protein (FBP), FCRL5, FCRH5, fetal

acetylcholine receptor, GD2, GD3, HMW-MAA, IL-22R-alpha, IL-13R-alpha2, kinase insert domain receptor (kdr), kappa light chain, Lewis Y, L1-cell adhesion molecule, (L1-CAM), Melanoma-associated antigen (MAGE)-A1, MAGE-A3, MAGE-A6, Preferentially expressed antigen of melanoma (PRAME), survivin, TAG72, B7-H6, IL-13 receptor alpha 2 (IL-13Ra2), CA9, GD3, HMW-MAA, CD171, G250/CAIX, HLA-AI MAGE Al, HLA-A2 NY-ESO-1, PSCA, folate receptor-a, CD44v6, CD44v7/8, avb6 integrin, 8H9, NCAM, VEGF receptors, 5T4, Foetal AchR, NKG2D ligands, CD44v6, dual antigen, a cancer-testes antigen, mesothelin, murine CMV, mucin 1 (MUC1), MUC16, PSCA, NKG2D, NY-ESO-1, MART-1, gp100, oncofetal antigen, ROR1, TAG72, VEGF-R2, carcinoembryonic antigen (CEA), Her2/neu, estrogen receptor, progesterone receptor, ephrinB2, CD123, c-Met, GD-2, O-acetylated GD2 (OGD2), CE7, Wilms Tumor 1 (WT-1), a cyclin, cyclin A2, CCL-1, CD138, a pathogen-specific antigen and an antigen associated with a universal tag, and/or biotinylated molecules, and/or molecules expressed by HIV, HCV, HBV or other pathogens. Antigens targeted by the receptors in some embodiments include antigens associated with a B cell malignancy, such as any of a number of known B cell marker. In some embodiments, the antigen targeted by the receptor is CD20, CD19, CD22, ROR1, CD45, CD21, CD5, CD33, Igkappa, Iglambda, CD79a, CD79b or CD30.

[0340] Antigens targeted by the receptors in some embodiments include orphan tyrosine kinase receptor ROR1, Her2, L1-CAM, CD19, CD20, CD22, mesothelin, CEA, and hepatitis B surface antigen, anti-folate receptor, CD23, CD24, CD30, CD33, CD38, CD44, EGFR, EGP-2, EGP-4, EPHa2, ErbB2, 3, or 4, FBP, fetal acethycholine e receptor, GD2, GD3, HMW-MAA, IL-22R-alpha, IL-13R-alpha2, kdr, kappa light chain, Lewis Y, L1-cell adhesion molecule, MAGE-A1, mesothelin, MUC1, MUC16, PSCA, NKG2D Ligands, NY-ESO-1, MART-1, gp100, oncofetal antigen, ROR1, TAG72, VEGF-R2, carcinoembryonic antigen (CEA), prostate specific antigen, PSMA, Her2/neu, estrogen receptor, progesterone receptor, ephrinB2, CD123, c-Met, GD-2, and MAGE A3, CE7, Wilms Tumor 1 (WT-1), a cyclin, such as cyclin A1 (CCNA1), and/or biotinylated molecules, and/or molecules expressed by HIV, HCV, HBV or other pathogens.

[0341] In some embodiments, the CAR binds a pathogen-specific or pathogen-expressed antigen. In some embodiments, the CAR is specific for viral antigens (such as HIV, HCV, HBV, etc.), bacterial antigens, and/or parasitic antigens.

[0342] In some embodiments, the CAR contains a TCR-like antibody, such as an antibody or an antigen-binding fragment (*e.g.* scFv) that specifically recognizes an intracellular antigen, such as a tumor-associated antigen, presented on the cell surface as a MHC-peptide complex. In some embodiments, an antibody or antigen-binding portion thereof that recognizes an MHC-peptide complex can be expressed on cells as part of a recombinant receptor, such as an antigen receptor. Among the antigen receptors are functional non-TCR antigen receptors, such as chimeric antigen receptors (CARs). Generally, a CAR containing an antibody or antigen-binding fragment that exhibits TCR-like specificity directed against peptide-MHC complexes also may be referred to as a TCR-like CAR.

[0343] Reference to "Major histocompatibility complex" (MHC) refers to a protein, generally a glycoprotein, that contains a polymorphic peptide binding site or binding groove that can, in some cases, complex with peptide antigens of polypeptides, including peptide antigens processed by the cell machinery. In some cases, MHC molecules can be displayed or expressed on the cell surface, including as a complex with peptide, *i.e.* MHC-peptide complex, for presentation of an antigen in a conformation recognizable by an antigen receptor on T cells, such as a TCRs or TCR-like antibody. Generally, MHC class I molecules are heterodimers having a membrane spanning α chain, in some cases with three α domains, and a non-covalently associated β2 microglobulin. Generally, MHC class II molecules are composed of two transmembrane glycoproteins, α and β, both of which typically span the membrane. An MHC molecule can include an effective portion of an MHC that contains an antigen binding site or sites for binding a peptide and the sequences necessary for recognition by the appropriate antigen receptor. In some embodiments, MHC class I molecules deliver peptides originating in the cytosol to the cell surface, where a MHC-peptide complex is recognized by T cells, such as generally $CD8^+$ T cells, but in some cases CD4+ T cells. In some embodiments, MHC class II molecules deliver peptides originating in the vesicular system to the cell surface, where they are typically recognized by $CD4^+$ T cells. Generally, MHC molecules are encoded by a group of linked loci, which are collectively termed H-2 in the mouse and human leukocyte antigen (HLA) in humans. Hence, typically human MHC can also be referred to as human leukocyte antigen (HLA).

[0344] The term "MHC-peptide complex" or "peptide-MHC complex" or variations thereof, refers to a complex or association of a peptide antigen and an MHC molecule, such as,

generally, by non-covalent interactions of the peptide in the binding groove or cleft of the MHC molecule. In some embodiments, the MHC-peptide complex is present or displayed on the surface of cells. In some embodiments, the MHC-peptide complex can be specifically recognized by an antigen receptor, such as a TCR, TCR-like CAR or antigen-binding portions thereof.

[0345] In some embodiments, a peptide, such as a peptide antigen or epitope, of a polypeptide can associate with an MHC molecule, such as for recognition by an antigen receptor. Generally, the peptide is derived from or based on a fragment of a longer biological molecule, such as a polypeptide or protein. In some embodiments, the peptide typically is about 8 to about 24 amino acids in length. In some embodiments, a peptide has a length of from or from about 9 to 22 amino acids for recognition in the MHC Class II complex. In some embodiments, a peptide has a length of from or from about 8 to 13 amino acids for recognition in the MHC Class I complex. In some embodiments, upon recognition of the peptide in the context of an MHC molecule, such as MHC-peptide complex, the antigen receptor, such as TCR or TCR-like CAR, produces or triggers an activation signal to the T cell that induces a T cell response, such as T cell proliferation, cytokine production, a cytotoxic T cell response or other response.

[0346] In some embodiments, a TCR-like antibody or antigen-binding portion, are known or can be produced by methods known in the art (see e.g. US Published Application Nos. US 2002/0150914; US 2003/0223994; US 2004/0191260; US 2006/0034850; US 2007/00992530; US20090226474; US20090304679; and International PCT Publication No. WO 03/068201).

[0347] In some embodiments, an antibody or antigen-binding portion thereof that specifically binds to a MHC-peptide complex, can be produced by immunizing a host with an effective amount of an immunogen containing a specific MHC-peptide complex. In some cases, the peptide of the MHC-peptide complex is an epitope of antigen capable of binding to the MHC, such as a tumor antigen, for example a universal tumor antigen, myeloma antigen or other antigen. In some embodiments, an effective amount of the immunogen is then administered to a host for eliciting an immune response, wherein the immunogen retains a three-dimensional form thereof for a period of time sufficient to elicit an immune response against the three-dimensional presentation of the peptide in the binding groove of the MHC molecule. Serum collected from the host is then assayed to determine if desired antibodies that recognize a three-dimensional

presentation of the peptide in the binding groove of the MHC molecule is being produced. In some embodiments, the produced antibodies can be assessed to confirm that the antibody can differentiate the MHC-peptide complex from the MHC molecule alone, the peptide of interest alone, and a complex of MHC and irrelevant peptide. The desired antibodies can then be isolated.

[0348] In some embodiments, an antibody or antigen-binding portion thereof that specifically binds to an MHC-peptide complex can be produced by employing antibody library display methods, such as phage antibody libraries. In some embodiments, phage display libraries of mutant Fab, scFv or other antibody forms can be generated, for example, in which members of the library are mutated at one or more residues of a CDR or CDRs. See *e.g.* US published application No. US20020150914, US2014/0294841; and Cohen CJ. *et al*. (2003) *J Mol. Recogn.* 16:324-332.

[0349] The term "antibody" herein is used in the broadest sense and includes polyclonal and monoclonal antibodies, including intact antibodies and functional (antigen-binding) antibody fragments, including fragment antigen binding (Fab) fragments, $F(ab')_2$ fragments, Fab' fragments, Fv fragments, recombinant IgG (rIgG) fragments, variable heavy chain (V_H) regions capable of specifically binding the antigen, single chain antibody fragments, including single chain variable fragments (scFv), and single domain antibodies (*e.g.*, sdAb, sdFv, nanobody) fragments. The term encompasses genetically engineered and/or otherwise modified forms of immunoglobulins, such as intrabodies, peptibodies, chimeric antibodies, fully human antibodies, humanized antibodies, and heteroconjugate antibodies, multispecific, *e.g.*, bispecific, antibodies, diabodies, triabodies, and tetrabodies, tandem di-scFv, tandem tri-scFv. Unless otherwise stated, the term "antibody" should be understood to encompass functional antibody fragments thereof. The term also encompasses intact or full-length antibodies, including antibodies of any class or sub-class, including IgG and sub-classes thereof, IgM, IgE, IgA, and IgD.

[0350] In certain embodiments, multispecific binding molecules, e.g., multispecific chimeric receptors, such as multispecific CARs, can contain any of the multispecific antibodies, including, *e.g.* bispecific antibodies, multispecific single-chain antibodies, *e.g.*, diabodies, triabodies, and tetrabodies, tandem di-scFvs, and tandem tri-scFvs.

[0351] In some embodiments, the antigen-binding proteins, antibodies and antigen binding fragments thereof specifically recognize an antigen of a full-length antibody. In some

embodiments, the heavy and light chains of an antibody can be full-length or can be an antigen-binding portion (a Fab, F(ab')2, Fv or a single chain Fv fragment (scFv)). In other embodiments, the antibody heavy chain constant region is chosen from, *e.g.*, IgG1, IgG2, IgG3, IgG4, IgM, IgA1, IgA2, IgD, and IgE, particularly chosen from, *e.g.*, IgG1, IgG2, IgG3, and IgG4, more particularly, IgG1 (*e.g.*, human IgG1). In another embodiment, the antibody light chain constant region is chosen from, *e.g.*, kappa or lambda, particularly kappa.

[0352] Among the provided antibodies are antibody fragments. An "antibody fragment" or "antigen-binding fragment" refers to a molecule other than an intact antibody that comprises a portion of an intact antibody that binds the antigen to which the intact antibody binds. Examples of antibody fragments include but are not limited to Fv, Fab, Fab', Fab'-SH, $F(ab')_2$; diabodies; linear antibodies; variable heavy chain (V_H) regions, single-chain antibody molecules such as scFvs and single-domain V_H single antibodies; and multispecific antibodies formed from antibody fragments. In some embodiments, the antigen-binding domain in the provided CARs is or comprises an antibody fragment comprising a variable heavy chain (V_H) and a variable light chain (V_L) region. In particular embodiments, the antibodies are single-chain antibody fragments comprising a variable heavy chain region and/or a variable light chain region, such as scFvs.

[0353] The term "variable region" or "variable domain" refers to the domain of an antibody heavy or light chain that is involved in binding the antibody to antigen. The variable domains of the heavy chain and light chain (V_H and V_L, respectively) of a native antibody generally have similar structures, with each domain comprising four conserved framework regions (FRs) and three CDRs. (See, *e.g.*, Kindt et al. Kuby Immunology, 6th ed., W.H. Freeman and Co., page 91 (2007). A single V_H or V_L domain may be sufficient to confer antigen-binding specificity. Furthermore, antibodies that bind a particular antigen may be isolated using a V_H or V_L domain from an antibody that binds the antigen to screen a library of complementary V_L or V_H domains, respectively. See, *e.g.*, Portolano et al., J. Immunol. 150:880-887 (1993); Clarkson et al., Nature 352:624-628 (1991).

[0354] The terms "complementarity determining region," and "CDR," synonymous with "hypervariable region" or "HVR," are known in the art to refer to non-contiguous sequences of amino acids within antibody variable regions, which confer antigen specificity and/or binding affinity. In general, there are three CDRs in each heavy chain variable region (CDR-H1, CDR-H2, CDR-H3) and three CDRs in each light chain variable region (CDR-L1, CDR-L2, CDR-

L3). "Framework regions" and "FR" are known in the art to refer to the non-CDR portions of the variable regions of the heavy and light chains. In general, there are four FRs in each full-length heavy chain variable region (FR-H1, FR-H2, FR-H3, and FR-H4), and four FRs in each full-length light chain variable region (FR-L1, FR-L2, FR-L3, and FR-L4).

[0355] The precise amino acid sequence boundaries of a given CDR or FR can be readily 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); MacCallum *et al.*, J. Mol. Biol. 262:732-745 (1996), "Antibody-antigen interactions: Contact analysis and binding site topography," J. Mol. Biol. 262, 732-745." ("Contact" numbering scheme); Lefranc MP *et al.*, "IMGT unique numbering for immunoglobulin and T cell receptor variable domains and Ig superfamily V-like domains," Dev Comp Immunol, 2003 Jan;27(1):55-77 ("IMGT" numbering scheme); Honegger A and Plückthun A, "Yet another numbering scheme for immunoglobulin variable domains: an automatic modeling and analysis tool," J Mol Biol, 2001 Jun 8;309(3):657-70, ("Aho" numbering scheme); and Martin *et al.*, "Modeling antibody hypervariable loops: a combined algorithm," PNAS, 1989, 86(23):9268-9272, ("AbM" numbering scheme).

[0356] The boundaries of a given CDR or FR may vary depending on the scheme used for identification. For example, the Kabat scheme is based on structural alignments, while the Chothia scheme is based on structural information. Numbering for both the Kabat and Chothia schemes is based upon the most common antibody region sequence lengths, with insertions accommodated by insertion letters, for example, "30a," and deletions appearing in some antibodies. The two schemes place certain insertions and deletions ("indels") at different positions, resulting in differential numbering. The Contact scheme is based on analysis of complex crystal structures and is similar in many respects to the Chothia numbering scheme. The AbM scheme is a compromise between Kabat and Chothia definitions based on that used by Oxford Molecular's AbM antibody modeling software.

[0357] **Table 1**, below, lists exemplary position boundaries of CDR-L1, CDR-L2, CDR-L3 and CDR-H1, CDR-H2, CDR-H3 as identified by Kabat, Chothia, AbM, and Contact schemes, respectively. For CDR-H1, residue numbering is listed using both the Kabat and Chothia numbering schemes. FRs are located between CDRs, for example, with FR-L1 located before

CDR-L1, FR-L2 located between CDR-L1 and CDR-L2, FR-L3 located between CDR-L2 and CDR-L3 and so forth. It is noted that because the shown Kabat numbering scheme places insertions at H35A and H35B, the end of the Chothia CDR-H1 loop when numbered using the shown Kabat numbering convention varies between H32 and H34, depending on the length of the loop.

Table 1. Boundaries of CDRs according to various numbering schemes.

CDR	**Kabat**	**Chothia**	**AbM**	**Contact**
CDR-L1	L24--L34	L24--L34	L24--L34	L30--L36
CDR-L2	L50--L56	L50--L56	L50--L56	L46--L55
CDR-L3	L89--L97	L89--L97	L89--L97	L89--L96
CDR-H1 (Kabat Numbering[1])	H31--H35B	H26--H32..34	H26--H35B	H30--H35B
CDR-H1 (Chothia Numbering[2])	H31--H35	H26--H32	H26--H35	H30--H35
CDR-H2	H50--H65	H52--H56	H50--H58	H47--H58
CDR-H3	H95--H102	H95--H102	H95--H102	H93--H101

1 - Kabat *et al.* (1991), "Sequences of Proteins of Immunological Interest," 5th Ed. Public Health Service, National Institutes of Health, Bethesda, MD
2 - Al-Lazikani *et al.*, (1997) JMB 273,927-948

[0358] Thus, unless otherwise specified, a "CDR" or "complementary determining region," or individual specified CDRs (*e.g.*, CDR-H1, CDR-H2, CDR-H3), of a given antibody or region thereof, such as a variable region thereof, should be understood to encompass a (or the specific) complementary determining region as defined by any of the aforementioned schemes, or other known schemes. For example, where it is stated that a particular CDR (*e.g.*, a CDR-H3) contains the amino acid sequence of a corresponding CDR in a given V_H or V_L region amino acid sequence, it is understood that such a CDR has a sequence of the corresponding CDR (*e.g.*, CDR-H3) within the variable region, as defined by any of the aforementioned schemes, or other known schemes. In some embodiments, specific CDR sequences are specified. Exemplary CDR sequences of provided antibodies are described using various numbering schemes, although it is understood that a provided antibody can include CDRs as described according to any of the other aforementioned numbering schemes or other numbering schemes known to a skilled artisan.

[0359] Likewise, unless otherwise specified, a FR or individual specified FR(s) (*e.g.*, FR-H1, FR-H2, FR-H3, FR-H4), of a given antibody or region thereof, such as a variable region thereof, should be understood to encompass a (or the specific) framework region as defined by

any of the known schemes. In some instances, the scheme for identification of a particular CDR, FR, or FRs or CDRs is specified, such as the CDR as defined by the Kabat, Chothia, AbM or Contact method, or other known schemes. In other cases, the particular amino acid sequence of a CDR or FR is given.

[0360] Single-domain antibodies (sdAbs) are antibody fragments comprising all or a portion of the heavy chain variable domain or all or a portion of the light chain variable domain of an antibody. In certain embodiments, a single-domain antibody is a human single-domain antibody. In some embodiments, the CAR comprises an antibody heavy chain domain that specifically binds the antigen, such as a cancer marker or cell surface antigen of a cell or disease to be targeted, such as a tumor cell or a cancer cell, such as any of the target antigens described herein or known in the art.

[0361] Antibody fragments can be made by various techniques, including but not limited to proteolytic digestion of an intact antibody as well as production by recombinant host cells. In some embodiments, the antibodies are recombinantly-produced fragments, such as fragments comprising arrangements that do not occur naturally, such as those with two or more antibody regions or chains joined by synthetic linkers, *e.g.*, peptide linkers, and/or that are may not be produced by enzyme digestion of a naturally-occurring intact antibody. In some embodiments, the antibody fragments are scFvs.

[0362] A "humanized" antibody is an antibody in which all or substantially all CDR amino acid residues are derived from non-human CDRs and all or substantially all FR amino acid residues are derived from human FRs. A humanized antibody optionally may include at least a portion of an antibody constant region derived from a human antibody. A "humanized form" of a non-human antibody, refers to a variant of the non-human antibody that has undergone humanization, typically to reduce immunogenicity to humans, while retaining the specificity and affinity of the parental non-human antibody. In some embodiments, some FR residues in a humanized antibody are substituted with corresponding residues from a non-human antibody (*e.g.*, the antibody from which the CDR residues are derived), *e.g.*, to restore or improve antibody specificity or affinity.

[0363] Thus, in some embodiments, the chimeric antigen receptor, including TCR-like CARs, includes an extracellular portion containing an antibody or antibody fragment. In some embodiments, the antibody or fragment includes an scFv. In some embodiments, the antibody

or antigen-binding fragment thereof is a single-chain antibody fragment, such as a single chain variable fragment (scFv) or a diabody or a single domain antibody (sdAb). In some embodiments, the antibody or antigen-binding fragment is a single domain antibody comprising only the V_H region. In some embodiments, the antibody or antigen binding fragment is an scFv comprising a heavy chain variable (V_H) region and a light chain variable (V_L) region.

[0364] In some embodiments, the antibody is an antigen-binding fragment, such as a scFv, that includes one or more linkers joining two antibody domains or regions, such as a heavy chain variable (V_H) region and a light chain variable (V_L) region. The linker typically is a peptide linker, *e.g.*, a flexible and/or soluble peptide linker. Among the linkers are those rich in glycine and serine and/or in some cases threonine. In some embodiments, the linkers further include charged residues such as lysine and/or glutamate, which can improve solubility. In some embodiments, the linkers further include one or more proline. In some aspects, the linkers rich in glycine and serine (and/or threonine) include at least 80%, 85%, 90%, 91%, 92%, 93%, 94%, 95%, 96%, 97%, 98%, or 99% such amino acid(s). In some embodiments, they include at least at or about 50%, 55%, 60%, 70%, or 75%, glycine, serine, and/or threonine. In some embodiments, the linker is comprised substantially entirely of glycine, serine, and/or threonine. The linkers generally are between about 5 and about 50 amino acids in length, typically between at or about 10 and at or about 30, *e.g.*, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, or 30, and in some examples between 10 and 25 amino acids in length. Exemplary linkers include linkers having various numbers of repeats of the sequence GGGGS (4GS; SEQ ID NO:26) or GGGS (3GS; SEQ ID NO:27), such as between 2, 3, 4, and 5 repeats of such a sequence. Exemplary linkers include those having or consisting of an sequence set forth in SEQ ID NO:28 (GGGGSGGGGSGGGGS), SEQ ID NO:29 (GSTSGSGKPGSGEGSTKG), SEQ ID NO: 30 (SRGGGGSGGGGSGGGGSLEMA), or SEQ ID NO:38 (ASGGGGSGGRASGGGGS).

[0365] In some embodiments, the CAR is an anti-BCMA CAR that is specific for BCMA, e.g. human BCMA. Chimeric antigen receptors containing anti-BCMA antibodies, including mouse anti-human BCMA antibodies and human anti-human BCMA antibodies, and cells expressing such chimeric receptors have been previously described. *See* Carpenter *et al.*, *Clin Cancer Res*., 2013, 19(8):2048-2060, US 9,765,342, WO 2016/090320, WO2016090327, WO2010104949A2, WO2016/0046724, WO2016/014789, WO2016/090320, WO2016/094304,

WO2017/025038, and WO2017173256. In some embodiments, the anti-BCMA CAR contains an antigen-binding domain, such as an scFv, containing a variable heavy (VH) and/or a variable light (VL) region derived from an antibody described in WO 2016/090320 or WO2016090327. In some embodiments, the anti-BCMA CAR contains an antigen-binding domain, such as an scFv, containing a variable heavy (V_H) and/or a variable light (V_L) region derived from an antibody described in WO 2016/090320 or WO2016090327. In some embodiments, the antigen-binding domain is an antibody fragment containing a variable heavy chain (V_H) and a variable light chain (V_L) region. In some aspects, the V_H region is or includes an amino acid sequence having at least 90%, 91%, 92%, 93%, 94%, 95%, 96%, 97%, 98% or 99% sequence identity to the V_H region amino acid sequence set forth in any of SEQ ID NOs: 18, 20, 22, 24, 32, 34, 36, 41, 43, 45, 47, 49, 51, 53, 55, 57, 59, 61, 63, 65, 67, 69, 71, 73, 75, 145, 147, 149 and 151; and/or the V_L region is or includes an amino acid sequence having at least 90%, 91%, 92%, 93%, 94%, 95%, 96%, 97%, 98%, or 99% sequence identity to the V_L region amino acid sequence set forth in any of SEQ ID NOs: 19, 21, 23, 25, 33, 35, 37, 42, 44, 46, 48, 50, 52, 54, 56, 58, 60, 62, 64, 66, 68, 70, 72, 74, 76, 146, 148, 150 and 152.

[0366] In some embodiments, the antigen-binding domain, such as an scFv, contains a V_H set forth in SEQ ID NO: 18 and a V_L set forth in SEQ ID NO:19. In some embodiments, the antigen-binding domain, such as an scFv, contains a V_H set forth in SEQ ID NO: 20 and a V_L set forth in SEQ ID NO:21. In some embodiments, the antigen-binding domain, such as an scFv, contains a V_H set forth in SEQ ID NO: 22 and a V_L set forth in SEQ ID NO:23. In some embodiments, the antigen-binding domain, such as an scFv, contains a V_H set forth in SEQ ID NO: 24 and a V_L set forth in SEQ ID NO:25. In some embodiment the antigen-binding domain, such as an scFv, contains a V_H set forth in SEQ ID NO: 32 and a V_L set forth in SEQ ID NO:33. In some embodiments, the antigen-binding domain, such as an scFv, contains a V_H set forth in SEQ ID NO:34 and a V_L set forth in SEQ ID NO:35. In some embodiments, the antigen-binding domain, such as an scFv, contains a V_H set forth in SEQ ID NO: 36 and a V_L set forth in SEQ ID NO:37. In some embodiments, the antigen-binding domain, such as an scFv, contains a V_H set forth in SEQ ID NO: 41 and a V_L set forth in SEQ ID NO: 42. In some embodiments, the antigen-binding domain, such as an scFv, contains a V_H set forth in SEQ ID NO: 43 and a V_L set forth in SEQ ID NO: 44. In some embodiments, the antigen-binding domain, such as an scFv, contains a V_H set forth in SEQ ID NO: 45 and a V_L set forth in SEQ ID NO: 46. In some

embodiments, the antigen-binding domain, such as an scFv, contains a V_H set forth in SEQ ID NO: 47 and a V_L set forth in SEQ ID NO: 48. In some embodiments, the antigen-binding domain, such as an scFv, contains a V_H set forth in SEQ ID NO: 49 and a V_L set forth in SEQ ID NO: 50. In some embodiments, the antigen-binding domain, such as an scFv, contains a V_H set forth in SEQ ID NO: 51 and a V_L set forth in SEQ ID NO: 52. In some embodiments, the antigen-binding domain, such as an scFv, contains a V_H set forth in SEQ ID NO: 53 and a V_L set forth in SEQ ID NO: 54. In some embodiments, the antigen-binding domain, such as an scFv, contains a V_H set forth in SEQ ID NO: 55 and a V_L set forth in SEQ ID NO: 56. In some embodiments, the antigen-binding domain, such as an scFv, contains a V_H set forth in SEQ ID NO: 57 and a V_L set forth in SEQ ID NO: 58. In some embodiments, the antigen-binding domain, such as an scFv, contains a V_H set forth in SEQ ID NO: 59 and a V_L set forth in SEQ ID NO: 60. In some embodiments, the antigen-binding domain, such as an scFv, contains a V_H set forth in SEQ ID NO: 61 and a V_L set forth in SEQ ID NO: 62. In some embodiments, the antigen-binding domain, such as an scFv, contains a V_H set forth in SEQ ID NO: 63 and a V_L set forth in SEQ ID NO: 64. In some embodiments, the antigen-binding domain, such as an scFv, contains a V_H set forth in SEQ ID NO: 65 and a V_L set forth in SEQ ID NO: 66. In some embodiments, the antigen-binding domain, such as an scFv, contains a V_H set forth in SEQ ID NO: 67 and a V_L set forth in SEQ ID NO: 68. In some embodiments, the antigen-binding domain, such as an scFv, contains a V_H set forth in SEQ ID NO: 69 and a V_L set forth in SEQ ID NO: 70. In some embodiments, the antigen-binding domain, such as an scFv, contains a V_H set forth in SEQ ID NO: 71 and a V_L set forth in SEQ ID NO: 72. In some embodiments, the antigen-binding domain, such as an scFv, contains a V_H set forth in SEQ ID NO: 73 and a V_L set forth in SEQ ID NO: 74. In some embodiments, the antigen-binding domain, such as an scFv, contains a V_H set forth in SEQ ID NO: 75 and a V_L set forth in SEQ ID NO: 76. In some embodiments, the antigen-binding domain, such as an scFv, contains a V_H set forth in SEQ ID NO: 145 and a V_L set forth in SEQ ID NO: 146. In some embodiments, the antigen-binding domain, such as an scFv, contains a V_H set forth in SEQ ID NO: 147 and a V_L set forth in SEQ ID NO: 148. In some embodiments, the antigen-binding domain, such as an scFv, contains a V_H set forth in SEQ ID NO: 149 and a V_L set forth in SEQ ID NO: 150. In some embodiments, the antigen-binding domain, such as an scFv, contains a V_H set forth in SEQ ID NO: 151 and a V_L set forth in SEQ ID NO: 152. In some embodiments, the V_H or V_L has a sequence of amino acids

that exhibits at least 85%, 86%, 87%, 88%, 89%, 90%, 91%, 92%, 93%, 94%, 95%, 96%, 97%, 98%, 99% or more sequence identity to any of the foregoing V_H or V_L sequences, and retains binding to BCMA. In some embodiments, the V_H region is amino-terminal to the V_L region. In some embodiments, the V_H region is carboxy-terminal to the V_L region. In some embodiments, the variable heavy and variable light chains are connected by a linker. In some embodiments, the linker is set forth in SEQ ID NO: 28, 29, 30, or 38.

[0367] Among a provided anti-BCMA CAR is a CAR in which the antibody or antigen-binding fragment contains a V_H region comprising the sequence set forth in SEQ ID NO: 24 or an amino acid sequence having at least at or about 90%, at or about 91%, at or about 92%, at or about 93%, at or about 94%, at or about 95%, at or about 96%, at or about 97%, at or about 98%, or at or about 99% identity to SEQ ID NO:24; and contains a V_L region comprising the sequence set forth in SEQ ID NO:25 or an amino acid sequence having at least at or about 90%, at or about 91%, at or about 92%, at or about 93%, at or about 94%, at or about 95%, at or about 96%, at or about 97%, at or about 98%, or at or about 99% identity to SEQ ID NO:25. In some embodiments, the antibody or antigen-binding fragment of the provided CAR contains a V_H region that has a CDRH1, a CDRH2 and a CDRH3 comprising the amino acid sequence of SEQ ID NOS: 173, 174 and 175, respectively and a V_L region that has a CDRL1, a CDRL2 and a CDRL3 comprising the amino acid sequence of SEQ ID NOS: 183, 184 and 185, respectively. In some embodiments, the antibody or antigen-binding fragment of the provided CAR contains a V_H region that has a CDRH1, a CDRH2 and a CDRH3 comprising the amino acid sequence of SEQ ID NOS: 176, 177 and 175, respectively and a V_L region that has a CDRL1, a CDRL2 and a CDRL3 comprising the amino acid sequence of SEQ ID NOS: 183, 184 and 185, respectively. In some embodiments, the antibody or antigen-binding fragment of the provided CAR contains a V_H region that has a CDRH1, a CDRH2 and a CDRH3 comprising the amino acid sequence of SEQ ID NOS: 178, 179 and 175, respectively and a V_L region that has a CDRL1, a CDRL2 and a CDRL3 comprising the amino acid sequence of SEQ ID NOS: 183, 184 and 185, respectively. In some embodiments, the antibody or antigen-binding fragment of the provided CAR contains a V_H region that has a CDRH1, a CDRH2 and a CDRH3 comprising the amino acid sequence of SEQ ID NOS: 180, 181 and 182, respectively and a V_L region that has a CDRL1, a CDRL2 and a CDRL3 comprising the amino acid sequence of SEQ ID NOS: 186, 187 and 185, respectively. In some embodiments, the V_H region comprises the sequence set forth in SEQ ID NO:24 and the

V_L region comprises the sequence set forth in SEQ ID NO:25. In some embodiments, the antibody or antigen-binding fragment is a single-chain antibody fragment, such as an scFv. In some embodiments, the scFv comprises the sequence of amino acids set forth in SEQ ID NO:188 or a sequence of amino acids at least at or about 90%, at or about 91%, at or about 92%, at or about 93%, at or about 94%, at or about 95%, at or about 96%, at or about 97%, at or about 98%, or at or about 99% identity to SEQ ID NO:188. In some embodiments, the anti-BCMA CAR has the sequence of amino acids set forth in SEQ NO: 124 or a sequence of amino acids at least at or about 90%, at or about 91%, at or about 92%, at or about 93%, at or about 94%, at or about 95%, at or about 96%, at or about 97%, at or about 98%, or at or about 99% identity to SEQ ID NO:124. In some embodiments, the anti-BCMA CAR has the sequence of amino acids set forth in SEQ NO: 125 or a sequence of amino acids at least at or about 90%, at or about 91%, at or about 92%, at or about 93%, at or about 94%, at or about 95%, at or about 96%, at or about 97%, at or about 98%, or at or about 99% identity to SEQ ID NO:125.

[0368] In some embodiments, the antigen-binding domain comprises an sdAb. In some embodiments, the antigen-binding domain contains the sequence set forth by SEQ ID NO:77. In some embodiments, the antigen-binding domain comprises a sequence at least or about 50, 60, 70, 80, 85, 90, 95, 96, 97, 98, 99, or 100% identical to the sequence set forth by SEQ ID NO:77.

[0369] In some embodiments, among such antibodies or antigen-binding domains in the provided CARs are antibodies capable of binding BCMA protein, such as human BCMA protein, with at least a certain affinity, as measured by any of a number of known methods. In some embodiments, the affinity is represented by an equilibrium dissociation constant (K_D); in some embodiments, the affinity is represented by EC50.

[0370] In some embodiments, among such antibodies or antigen-binding domains in the provided CARs, are antibodies or antigen-binding domains or CARs in which binding to soluble or shed BCMA protein is less than or at or about 10% of the binding of the antibody to membrane-bound BCMA protein.

[0371] A variety of assays are known for assessing binding affinity and/or determining whether a binding molecule (e.g., an antibody or fragment thereof) specifically binds to a particular ligand (e.g., an antigen, such as a BCMA protein). It is within the level of a skilled artisan to determine the binding affinity of a binding molecule, e.g., an antibody, for an antigen, e.g., BCMA. For example, in some embodiments, a BIAcore® instrument can be used to

determine the binding kinetics and constants of a complex between two proteins (e.g., an antibody or fragment thereof, and an antigen, such as a BCMA cell surface protein, soluble BCMA protein), using surface plasmon resonance (SPR) analysis (see, e.g., Scatchard et al., Ann. N.Y. Acad. Sci. 51:660, 1949; Wilson, Science 295:2103, 2002; Wolff et al., Cancer Res. 53:2560, 1993; and U.S. Patent Nos. 5,283,173, 5,468,614, or the equivalent).

[0372] SPR measures changes in the concentration of molecules at a sensor surface as molecules bind to or dissociate from the surface. The change in the SPR signal is directly proportional to the change in mass concentration close to the surface, thereby allowing measurement of binding kinetics between two molecules. The dissociation constant for the complex can be determined by monitoring changes in the refractive index with respect to time as buffer is passed over the chip. Other suitable assays for measuring the binding of one protein to another include, for example, immunoassays such as enzyme linked immunosorbent assays (ELISA) and radioimmunoassays (RIA), or determination of binding by monitoring the change in the spectroscopic or optical properties of the proteins through fluorescence, UV absorption, circular dichroism, or nuclear magnetic resonance (NMR). Other exemplary assays include, but are not limited to, Western blot, ELISA, analytical ultracentrifugation, spectroscopy, flow cytometry, sequencing and other methods for detection of expressed polynucleotides or binding of proteins.

[0373] In some embodiments, the binding molecule, e.g., antibody or fragment thereof or antigen-binding domain of a CAR, binds, such as specifically binds, to an antigen, e.g., a cell surface BCMA protein or soluble BCMA protein or an epitope therein, with an affinity or K_A (i.e., an equilibrium association constant of a particular binding interaction with units of 1/M; equal to the ratio of the on-rate [k_{on} or k_a] to the off-rate [k_{off} or k_d] for this association reaction, assuming bimolecular interaction) equal to or greater than 105 M^{-1}. In some embodiments, the antibody or fragment thereof or antigen-binding domain of a CAR exhibits a binding affinity for the peptide epitope with a K_D (i.e., an equilibrium dissociation constant of a particular binding interaction with units of M; equal to the ratio of the off-rate [k_{off} or k_d] to the on-rate [k_{on} or k_a] for this association reaction, assuming bimolecular interaction) of equal to or less than 10^{-5} M. For example, the equilibrium dissociation constant K_D ranges from 10^{-5} M to 10^{-13} M, such as 10^{-7} M to 10^{-11} M, 10^{-8} M to 10^{-10} M, or 10^{-9} M to 10^{-10} M. The on-rate (association rate constant; k_{on} or k_a; units of 1/Ms) and the off-rate (dissociation rate constant; k_{off} or k_d; units of

1/s) can be determined using any of the assay methods known in the art, for example, surface plasmon resonance (SPR).

[0374] In some embodiments, the binding affinity (EC50) and/or the dissociation constant of the antibody (e.g. antigen-binding fragment) or antigen-binding domain of a CAR to a BCMA protein, such as human BCMA protein, is from or from about 0.01 nM to about 500 nM, from or from about 0.01 nM to about 400 nM, from or from about 0.01 nM to about 100 nM, from or from about 0.01 nM to about 50 nM, from or from about 0.01 nM to about 10 nM, from or from about 0.01 nM to about 1 nM, from or from about 0.01 nM to about 0.1 nM, is from or from about 0.1 nM to about 500 nM, from or from about 0.1 nM to about 400 nM, from or from about 0.1 nM to about 100 nM, from or from about 0.1 nM to about 50 nM, from or from about 0.1 nM to about 10 nM, from or from about 0.1 nM to about 1 nM, from or from about 0.5 nM to about 200 nM, from or from about 1 nM to about 500 nM, from or from about 1 nM to about 100 nM, from or from about 1 nM to about 50 nM, from or from about 1 nM to about 10 nM, from or from about 2 nM to about 50 nM, from or from about 10 nM to about 500 nM, from or from about 10 nM to about 100 nM, from or from about 10 nM to about 50 nM, from or from about 50 nM to about 500 nM, from or from about 50 nM to about 100 nM or from or from about 100 nM to about 500 nM. In certain embodiments, the binding affinity (EC50) and/or the equilibrium dissociation constant, K_D, of the antibody to a BCMA protein, such as human BCMA protein, is at or less than or about 400 nM, 300 nM, 200 nM, 100 nM, 50 nM, 40 nM, 30 nM, 25 nM, 20 nM, 19 nM, 18 nM, 17 nM, 16 nM, 15 nM, 14 nM, 13 nM, 12 nM, 11 nM, 10 nM, 9 nM, 8 nM, 7 nM, 6 nM, 5 nM, 4 nM, 3 nM, 2 nM, or 1 nM or less. In some embodiments, the antibodies bind to a BCMA protein, such as human BCMA protein, with a sub-nanomolar binding affinity, for example, with a binding affinity less than about 1 nM, such as less than about 0.9 nM, about 0.8 nM, about 0.7 nM, about 0.6 nM, about 0.5 nM, about 0.4 nM, about 0.3 nM, about 0.2 nM or about 0.1 nM or less.

[0375] In some embodiments, the binding affinity may be classified as high affinity or as low affinity. In some cases, the binding molecule (e.g. antibody or fragment thereof) or antigen-binding domain of a CAR that exhibits low to moderate affinity binding exhibits a K_A of up to 10^7 M^{-1}, up to 10^6 M^{-1}, up to 10^5 M^{-1}. In some cases, a binding molecule (e.g. antibody or fragment thereof) that exhibits high affinity binding to a particular epitope interacts with such epitope with a K_A of at least 10^7 M^{-1}, at least 10^8 M^{-1}, at least 10^9 M^{-1}, at least 10^{10} M^{-1}, at least

10^{11} M^{-1}, at least 10^{12} M^{-1}, or at least 10^{13} M^{-1}. In some embodiments, the binding affinity (EC50) and/or the equilibrium dissociation constant, K_D, of the binding molecule, e.g., anti-BCMA antibody or fragment thereof or antigen-binding domain of a CAR, to a BCMA protein, is from or from about 0.01 nM to about 1 μM, 0.1 nM to 1 μM, 1 nM to 1 μM, 1 nM to 500 nM, 1 nM to 100 nM, 1 nM to 50 nM, 1 nM to 10 nM, 10 nM to 500 nM, 10 nM to 100 nM, 10 nM to 50 nM, 50 nM to 500 nM, 50 nM to 100 nM or 100 nM to 500 nM. In certain embodiments, the binding affinity (EC50) and/or the dissociation constant of the equilibrium dissociation constant, K_D, of the binding molecule, e.g., anti-BCMA antibody or fragment thereof or antigen-binding domain of a CAR, to a BCMA protein, is at or about or less than at or about 1 μM, 500 nM, 100 nM, 50 nM, 40 nM, 30 nM, 25 nM, 20 nM, 19 nM, 18 nM, 17 nM, 16 nM, 15 nM, 14 nM, 13 nM, 12 nM, 11 nM, 10 nM, 9 nM, 8 nM, 7 nM, 6 nM, 5 nM, 4 nM, 3 nM, 2 nM, or 1 nM or less. The degree of affinity of a particular antibody can be compared with the affinity of a known antibody, such as a reference antibody (e.g., anti-BCMA reference antibody).

[0376] In some embodiments, the binding affinity of the anti-BCMA antibody or antigen-binding domain of a CAR, for different form or topological type of antigens, e.g., soluble or shed BCMA protein compared to the binding affinity to a membrane-bound BCMA, to determine the preferential binding or relative affinity for a particular form or topological type. For example, in some aspects, an anti-BCMA antibodies or antigen-binding domain of a CAR can exhibit preferential binding to membrane-bound BCMA as compared to soluble or shed BCMA and/or exhibit greater binding affinity for, membrane-bound BCMA compared to soluble or shed BCMA. In some embodiments, the equilibrium dissociation constant, K_D, for different form or topological type of BCMA proteins, can be compared to determine preferential binding or relative binding affinity. In some embodiments, the preferential binding or relative affinity to a membrane-bound BCMA compared to soluble or shed BCMA can be high. For example, in some cases, the ratio of K_D for soluble or shed BCMA and the K_D for membrane-bound BCMA is more than 10, 15, 20, 25, 30, 40, 50, 60, 70, 80, 90, 100, 200, 500, 1000, 2000 or more and the antibody or antigen-binding domain preferentially binds or has higher binding affinity for membrane-bound BCMA. In some cases, the ratio of K_A for membrane-bound BCMA and the K_A for soluble or shed BCMA is more than 10, 15, 20, 25, 30, 40, 50, 60, 70, 80, 90, 100, 200, 500, 1000, 2000 or more and the antibody or antigen-binding domain preferentially binds or has higher binding affinity for membrane-bound BCMA. In some cases,

the antibody or antigen-binding domain of CAR binds soluble or shed BCMA and membrane-bound BCMA to a similar degree, e.g., the ratio of K_D for soluble BCMA and KD for membrane-bound BCMA is or is about 1. In some cases, the antibody or antigen-binding domain of CAR binds soluble or shed BCMA and membrane-bound BCMA to a similar degree, e.g., the ratio of K_A for soluble BCMA and K_A for membrane-bound BCMA is or is about 1. The degree of preferential binding or relative affinity for membrane-bound BCMA or soluble or shed BCMA can be compared with that of a known antibody, such as a reference antibody (e.g., reference anti-BCMA CAR). In some embodiments, the reference antibody (e.g., reference anti-BCMA CAR) binds to membrane-bound and soluble or shed BCMA protein.

[0377] In some aspects, the chimeric antigen receptor includes an extracellular portion containing the antibody or fragment and an intracellular signaling region. In some embodiments, the intracellular signaling region comprises an intracellular signaling domain. In some embodiments, the intracellular signaling domain is or comprises a primary signaling domain, a signaling domain that is capable of inducing a primary activation signal in a T cell, a signaling domain of a T cell receptor (TCR) component, and/or a signaling domain comprising an immunoreceptor tyrosine-based activation motif (ITAM).

[0378] In some embodiments, the antibody portion of the recombinant receptor, *e.g.*, CAR, further includes a spacer, which may be or include at least a portion of an immunoglobulin constant region or variant or modified version thereof, such as a hinge region, *e.g.*, an IgG4 hinge region, an IgG1 hinge region, a C_H1/C_L, and/or Fc region. In some embodiments, the recombinant receptor further comprises a spacer and/or a hinge region. In some embodiments, the constant region or portion is of a human IgG, such as IgG4 or IgG1. In some aspects, the portion of the constant region serves as a spacer region between the antigen-recognition component, *e.g.*, scFv, and transmembrane domain.

[0379] The spacer can be of a length that provides for increased responsiveness of the cell following antigen binding, as compared to in the absence of the spacer. Exemplary spacers, *e.g.*, hinge regions, include those described in international patent application publication number WO2014031687. In some examples, the spacer is or is about 12 amino acids in length or is no more than 12 amino acids in length. Exemplary spacers include those having at least about 10 to 229 amino acids, about 10 to 200 amino acids, about 10 to 175 amino acids, about 10 to 150 amino acids, about 10 to 125 amino acids, about 10 to 100 amino acids, about 10 to 75 amino

acids, about 10 to 50 amino acids, about 10 to 40 amino acids, about 10 to 30 amino acids, about 10 to 20 amino acids, or about 10 to 15 amino acids, and including any integer between the endpoints of any of the listed ranges. In some embodiments, a spacer region has about 12 amino acids or less, about 119 amino acids or less, or about 229 amino acids or less. In some embodiments, the spacer is a spacer having at least a particular length, such as having a length that is at least 100 amino acids, such as at least 110, 125, 130, 135, 140, 145, 150, 160, 170, 180, 190, 200, 210, 220, 230, 240, or 250 amino acids in length. Exemplary spacers include IgG4 hinge alone, IgG4 hinge linked to C_H2 and C_H3 domains, or IgG4 hinge linked to the C_H3 domain. Exemplary spacers include IgG4 hinge alone, IgG4 hinge linked to C_H2 and C_H3 domains, or IgG4 hinge linked to the C_H3 domain. Exemplary spacers include IgG4 hinge alone, IgG4 hinge linked to C_H2 and C_H3 domains, or IgG4 hinge linked to the C_H3 domain. Exemplary spacers include, but are not limited to, those described in Hudecek *et al.*, *Clin. Cancer Res*., 19:3153 (2013), Hudecek et al. (2015) Cancer Immunol Res. 3(2): 125–135, international patent application publication number WO2014031687, U.S. Patent No. 8,822,647 or published app. No. US2014/0271635. In some embodiments, the spacer includes a sequence of an immunoglobulin hinge region, a C_H2 and C_H3 region. In some embodiments, one of more of the hinge, C_H2 and C_H3 is derived all or in part from IgG4 or IgG2. In some cases, the hinge, C_H2 and C_H3 is derived from IgG4. In some aspects, one or more of the hinge, C_H2 and C_H3 is chimeric and contains sequence derived from IgG4 and IgG2. In some examples, the spacer contains an IgG4/2 chimeric hinge, an IgG2/4 C_H2, and an IgG4 C_H3 region.

[0380] In some embodiments, the spacer can be derived all or in part from IgG4 and/or IgG2 and can contain mutations, such as one or more single amino acid mutations in one or more domains. In some examples, the amino acid modification is a substitution of a proline (P) for a serine (S) in the hinge region of an IgG4. In some embodiments, the amino acid modification is a substitution of a glutamine (Q) for an asparagine (N) to reduce glycosylation heterogeneity, such as an N177Q mutation at position 177, in the C_H2 region, of the full-length IgG4 Fc sequence or an N176Q at position 176, in the C_H2 region, of the full-length IgG4 Fc sequence.

[0381] In some embodiments, the spacer has the sequence ESKYGPPCPPCP (set forth in SEQ ID NO: 1), and is encoded by the sequence set forth in SEQ ID NO: 2. In some embodiments, the spacer has the sequence set forth in SEQ ID NO: 3. In some embodiments, the spacer has the sequence set forth in SEQ ID NO: 4. In some embodiments, the encoded spacer

is or contains the sequence set forth in SEQ ID NO: 31. In some embodiments, the constant region or portion is of IgD. In some embodiments, the spacer has the sequence set forth in SEQ ID NO: 5. In some embodiments, the spacer has the sequence set forth in SEQ ID NO: 89.

[0382] Other exemplary spacer regions include hinge regions derived from CD8a, CD28, CTLA4, PD-1, or FcγRIIIa. In some embodiments, the spacer contains a truncated extracellular domain or hinge region of a CD8a, CD28, CTLA4, PD-1, or FcγRIIIa. In some embodiments, the spacer is a truncated CD28 hinge region. In some embodiments, a short oligo- or polypeptide linker, for example, a linker of between 2 and 10 amino acids in length, such as one containing alanines or alanine and arginine, *e.g.*, alanine triplet (AAA) or RAAA (SEQ ID NO: 144), is present and forms a linkage between the scFv and the spacer region of the CAR. In some embodiments, the spacer has the sequence set forth in SEQ ID NO: 78. In some embodiments, the spacer has the sequence set forth in SEQ ID NO: 80. In some embodiments, the spacer has the sequence set forth in any of SEQ ID NOs: 81-83, In some embodiments, the spacer has the sequence set forth in SEQ ID NO: 84. In some embodiments, the spacer has the sequence set forth in SEQ ID NO: 86. In some embodiments, the spacer has the sequence set forth in SEQ ID NO: 88.

[0383] In some embodiments, the spacer has a sequence of amino acids that exhibits at least 85%, 86%, 87%, 88%, 89%, 90%, 91%, 92%, 93%, 94%, 95%, 96%, 97%, 98%, 99% or more sequence identity to any of SEQ ID NOS: 1, 3, 4, 5, 31, 78, 80, 81, 82, 83, 84, 86, 88, or 89.

[0384] In some embodiments, the spacer has the sequence set forth in SEQ ID NOS: 157-165. In some embodiments, the spacer has a sequence of amino acids that exhibits at least 85%, 86%, 87%, 88%, 89%, 90%, 91%, 92%, 93%, 94%, 95%, 96%, 97%, 98%, 99% or more sequence identity to any of SEQ ID NOS: 157-165.

[0385] This antigen recognition domain generally is linked to one or more intracellular signaling components, such as signaling components that mimic stimulation and/or activation through an antigen receptor complex, such as a TCR complex, in the case of a CAR, and/or signal via another cell surface receptor. Thus, in some embodiments, the antigen-binding component (*e.g.*, antibody) is linked to one or more transmembrane and intracellular signaling domains. In some embodiments, the chimeric antigen receptor includes a transmembrane domain linking the extracellular domain and the intracellular signaling domain. In some embodiments, the transmembrane domain is fused to the extracellular domain, such as linked or

fused between the extracellular domain (e.g. scFv) and intracellular signaling domain.. In one embodiment, a transmembrane domain that naturally is associated with one of the domains in the receptor, *e.g.*, CAR, is used. In some instances, the transmembrane domain is 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 to minimize interactions with other members of the receptor complex.

[0386] The transmembrane domain in some embodiments is derived either from a natural or from a synthetic source. Where the source is natural, the domain in some aspects is derived from any membrane-bound or transmembrane protein. Transmembrane regions include those derived from (*i.e.* comprise at least the transmembrane region(s) of) the alpha, beta or zeta chain of the T-cell receptor, CD28, CD3 epsilon, CD45, CD4, CD5, CD8, CD8a, CD9, CD16, CD22, CD33, CD37, CD64, CD80, CD86, CD134, CD137 (4-1BB), CD154, CTLA-4 or PD-1. Alternatively the transmembrane domain in some embodiments is synthetic. In some aspects, the synthetic transmembrane domain comprises predominantly hydrophobic residues such as leucine and valine. In some aspects, a triplet of phenylalanine, tryptophan and valine will be found at each end of a synthetic transmembrane domain. In some embodiments, the linkage is by linkers, spacers, and/or transmembrane domain(s). In some aspects, the transmembrane domain contains a transmembrane portion of CD28. Exemplary sequences of transmembrane domains are or comprise the sequences set forth in SEQ ID NOs: 8, 79, 85, 87, 142, or 143.

[0387] Among the intracellular signaling domains are those that mimic or approximate a signal through a natural antigen receptor, a signal through such a receptor in combination with a costimulatory receptor, and/or a signal through a costimulatory receptor alone. In some embodiments, a short oligo- or polypeptide linker, for example, a linker of between 2 and 10 amino acids in length, such as one containing glycines and serines, *e.g.*, glycine-serine doublet, is present and forms a linkage between the transmembrane domain and the cytoplasmic signaling domain of the CAR.

[0388] The receptor, *e.g.*, the CAR, generally includes at least one intracellular signaling component or components. In some aspects, the CAR includes a primary cytoplasmic signaling sequence that regulates primary activation of the TCR complex. Primary cytoplasmic signaling sequences 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 cytoplasmic signaling sequences include those derived from TCR CD3 chain that mediates T-cell stimulation and/or activation and cytotoxicity, *e.g.*, CD3 zeta chain, CD3 gamma, CD3 delta, CD3 epsilon, FcR gamma, FcR beta, CDS, CD22, CD79a, CD79b and CD66d. In some cxamples of ITAM containing primary cytoplasmic signaling sequences include those derived from CD3 zeta chain, FcR gamma, CD3 gamma, CD3 delta and CD3 epsilon. In some embodiments, cytoplasmic signaling molecule(s) in the CAR contain(s) a cytoplasmic signaling domain, portion thereof, or sequence derived from CD3 zeta.

[0389] In some embodiments, the receptor includes an intracellular component of a TCR complex, such as a TCR CD3 chain that mediates T-cell stimulation and/or activation and cytotoxicity, *e.g.*, CD3 zeta chain. Thus, in some aspects, the antigen-binding portion is linked to one or more cell signaling modules. In some embodiments, cell signaling modules include CD3 transmembrane domain, CD3 intracellular signaling domains, and/or other CD transmembrane domains. In some embodiments, the receptor, *e.g.*, CAR, further includes a portion of one or more additional molecules such as Fc receptor γ, CD8, CD4, CD25 or CD16. For example, in some aspects, the CAR or other chimeric receptor includes a chimeric molecule between CD3-zeta (CD3-ζ) or Fc receptor γ and CD8, CD4, CD25 or CD16.

[0390] In some embodiments, upon ligation of the CAR or other chimeric receptor, the cytoplasmic domain or intracellular signaling domain of the receptor stimulates and/or activates at least one of the normal effector functions or responses of the immune cell, *e.g.*, T cell engineered to express the CAR. For example, in some contexts, the CAR induces a function of a T cell such as cytolytic activity or T-helper activity, such as secretion of cytokines or other factors. In some embodiments, a truncated portion of an intracellular signaling domain of an antigen receptor component or costimulatory molecule is used in place of an intact immunostimulatory chain, for example, if it transduces the effector function signal. In some embodiments, the intracellular signaling domain or domains include the cytoplasmic sequences of the T cell receptor (TCR), and in some aspects also those of co-receptors that in the natural context act in concert with such receptors to initiate signal transduction following antigen receptor engagement, and/or any derivative or variant of such molecules, and/or any synthetic sequence that has the same functional capability.

[0391] In the context of a natural TCR, full activation generally requires not only signaling through the TCR, but also a costimulatory signal. Thus, in some embodiments, to promote full

activation, a component for generating secondary or co-stimulatory signal is also included in the CAR. In other embodiments, the CAR does not include a component for generating a costimulatory signal. In some aspects, an additional CAR is expressed in the same cell and provides the component for generating the secondary or costimulatory signal.

[0392] T cell stimulation and/or activation is in some aspects described as being mediated by two classes of cytoplasmic signaling sequences: those that initiate antigen-dependent primary stimulation and/or activation through the TCR (primary cytoplasmic signaling regions, domains or sequences), and those that act in an antigen-independent manner to provide a secondary or co-stimulatory signal (secondary cytoplasmic signaling regions, domains or sequences). In some aspects, the CAR includes one or both of such signaling components.

[0393] In some embodiments, the CAR includes a signaling region and/or transmembrane portion of a costimulatory receptor, such as CD28, 4-1BB, OX40 (CD134), CD27, DAP10, DAP12, ICOS and/or other costimulatory receptors. In some aspects, the same CAR includes both the primary cytoplasmic signaling region and costimulatory signaling components. In some embodiments, the chimeric antigen receptor contains an intracellular domain derived from a T cell costimulatory molecule or a functional variant thereof, such as between the transmembrane domain and intracellular signaling domain. In some aspects, the T cell costimulatory molecule is CD28 or 41BB.

[0394] In some embodiments, one or more different recombinant receptors can contain one or more different intracellular signaling region(s) or domain(s). In some embodiments, the primary cytoplasmic signaling region is included within one CAR, whereas the costimulatory component is provided by another receptor, e.g., another CAR recognizing another antigen. In some embodiments, the CARs include activating or stimulatory CARs, and costimulatory CARs, both expressed on the same cell (*see* WO2014/055668).

[0395] In some aspects, the cells include one or more stimulatory or activating CAR and/or a costimulatory CAR. In some embodiments, the cells further include inhibitory CARs (iCARs, see Fedorov *et al.*, *Sci. Transl. Medicine*, 5(215) (2013), such as a CAR recognizing an antigen other than the one associated with and/or specific for the disease or condition whereby an activating signal delivered through the disease-targeting CAR is diminished or inhibited by binding of the inhibitory CAR to its ligand, *e.g.*, to reduce off-target effects.

[0396] In some embodiments, the two receptors induce, respectively, an activating and an inhibitory signal to the cell, such that ligation of one of the receptor to its antigen activates the cell or induces a response, but ligation of the second inhibitory receptor to its antigen induces a signal that suppresses or dampens that response. Examples are combinations of activating CARs and inhibitory CARs (iCARs). Such a strategy may be used, for example, to reduce the likelihood of off-target effects in the context in which the activating CAR binds an antigen expressed in a disease or condition but which is also expressed on normal cells, and the inhibitory receptor binds to a separate antigen which is expressed on the normal cells but not cells of the disease or condition.

[0397] In some aspects, the chimeric receptor is or includes an inhibitory CAR (e.g. iCAR) and includes intracellular components that dampen or suppress an immune response, such as an ITAM- and/or co stimulatory-promoted response in the cell. Exemplary of such intracellular signaling components are those found on immune checkpoint molecules, including PD-1, CTLA4, LAG3, BTLA, OX2R, TIM-3, TIGIT, LAIR-1, PGE2 receptors, EP2/4 Adenosine receptors including A2AR. In some aspects, the engineered cell includes an inhibitory CAR including a signaling domain of or derived from such an inhibitory molecule, such that it serves to dampen the response of the cell, for example, that induced by an activating and/or costimulatory CAR

[0398] In certain embodiments, the intracellular signaling domain comprises a CD28 transmembrane and signaling domain linked to a CD3 (*e.g.*, CD3-zeta) intracellular domain. In some embodiments, the intracellular signaling domain comprises a chimeric CD28 and CD137 (4-1BB, TNFRSF9) co-stimulatory domains, linked to a CD3 zeta intracellular domain.

[0399] In some embodiments, the CAR encompasses one or more, e.g., two or more, costimulatory domains and primary cytoplasmic signaling region, in the cytoplasmic portion. Exemplary CARs include intracellular components, such as intracellular signaling region(s) or domain(s), of CD3-zeta, CD28, CD137 (4-1BB), OX40 (CD134), CD27, DAP10, DAP12, NKG2D and/or ICOS. In some embodiments, the chimeric antigen receptor contains an intracellular signaling region or domain of a T cell costimulatory molecule, e.g., from CD28, CD137 (4-1BB), OX40 (CD134), CD27, DAP10, DAP12, NKG2D and/or ICOS, in some cases, between the transmembrane domain and intracellular signaling region or domain. In some

aspects, the T cell costimulatory molecule is one or more of CD28, CD137 (4-1BB), OX40 (CD134), CD27, DAP10, DAP12, NKG2D and/or ICOS.

[0400] In some cases, CARs are referred to as first, second, and/or third generation CARs. In some aspects, a first generation CAR is one that solely provides a CD3-chain induced signal upon antigen binding; in some aspects, a second-generation CARs is one that provides such a signal and costimulatory signal, such as one including an intracellular signaling domain from a costimulatory receptor such as CD28 or CD137; in some aspects, a third generation CAR is one that includes multiple costimulatory domains of different costimulatory receptors.

[0401] In some embodiments, the chimeric antigen receptor includes an extracellular portion containing an antibody or antibody fragment. In some aspects, the chimeric antigen receptor includes an extracellular portion containing the antibody or fragment and an intracellular signaling domain. In some embodiments, the antibody or fragment includes an scFv and the intracellular domain contains an ITAM. In some aspects, the intracellular signaling domain includes a signaling domain of a zeta chain of a CD3-zeta (CD3ζ) chain. In some embodiments, the chimeric antigen receptor includes a transmembrane domain linking the extracellular domain and the intracellular signaling domain. In some aspects, the transmembrane domain contains a transmembrane portion of CD28. In some embodiments, the chimeric antigen receptor contains an intracellular domain of a T cell costimulatory molecule. The extracellular domain and transmembrane domain can be linked directly or indirectly. In some embodiments, the extracellular domain and transmembrane are linked by a spacer, such as any described herein. In some embodiments, the receptor contains extracellular portion of the molecule from which the transmembrane domain is derived, such as a CD28 extracellular portion. In some embodiments, the chimeric antigen receptor contains an intracellular domain derived from a T cell costimulatory molecule or a functional variant thereof, such as between the transmembrane domain and intracellular signaling domain. In some aspects, the T cell costimulatory molecule is CD28 or 41BB.

[0402] In some embodiments, the CAR contains an antibody, *e.g.*, an antibody fragment, a transmembrane domain that is or contains a transmembrane portion of CD28 or a functional variant thereof, and an intracellular signaling domain containing a signaling portion of CD28 or functional variant thereof and a signaling portion of CD3 zeta or functional variant thereof. In some embodiments, the CAR contains an antibody, *e.g.*, antibody fragment, a transmembrane

domain that is or contains a transmembrane portion of CD28 or a functional variant thereof, and an intracellular signaling domain containing a signaling portion of a 4-1BB or functional variant thereof and a signaling portion of CD3 zeta or functional variant thereof. In some such embodiments, the receptor further includes a spacer containing a portion of an Ig molecule, such as a human Ig molecule, such as an Ig hinge, *e.g.* an IgG4 hinge, such as a hinge-only spacer.

[0403] In some embodiments, the transmembrane domain of the recombinant receptor, *e.g.*, the CAR, is or includes a transmembrane domain of human CD28 (*e.g.* Accession No. P10747.1), or CD8a (Accession No. P01732.1), or variant thereof, such as a transmembrane domain that comprises the sequence of amino acids set forth in SEQ ID NO: 8, 79, 142, or 143 or a sequence of amino acids that exhibits at least 85%, 86%, 87%, 88%, 89%, 90%, 91%, 92%, 93%, 94%, 95%, 96%, 97%, 98%, 99% or more sequence identity to SEQ ID NO: 8, 79, 142, or 143; in some embodiments, the transmembrane-domain containing portion of the recombinant receptor comprises the sequence of amino acids set forth in SEQ ID NO: 9 or a sequence of amino acids having at least at or about 85%, 86%, 87%, 88%, 89%, 90%, 91%, 92%, 93%, 94%, 95%, 96%, 97%, 98%, 99% or more sequence identity thereto.

[0404] In some embodiments, the intracellular signaling component(s) of the recombinant receptor, *e.g.* the CAR, contains an intracellular costimulatory signaling domain of human CD28 or a functional variant or portion thereof, such as a domain with an LL to GG substitution at positions 186-187 of a native CD28 protein. For example, the intracellular signaling domain can comprise the sequence of amino acids set forth in SEQ ID NO: 10 or 11 or a sequence of amino acids that exhibits at least 85%, 86%, 87%, 88%, 89%, 90%, 91%, 92%, 93%, 94%, 95%, 96%, 97%, 98%, 99% or more sequence identity to SEQ ID NO: 10 or 11. In some embodiments, the intracellular domain comprises an intracellular costimulatory signaling domain of 4-1BB (*e.g.* Accession No. Q07011.1) or functional variant or portion thereof, such as the sequence of amino acids set forth in SEQ ID NO: 12 or a sequence of amino acids that exhibits at least 85%, 86%, 87%, 88%, 89%, 90%, 91%, 92%, 93%, 94%, 95%, 96%, 97%, 98%, 99% or more sequence identity to SEQ ID NO: 12.

[0405] In some embodiments, the intracellular signaling domain of the recombinant receptor, *e.g.* the CAR, comprises a human CD3 zeta stimulatory signaling domain or functional variant thereof, such as a 112 AA cytoplasmic domain of isoform 3 of human CD3ζ (Accession No. P20963.2) or a CD3 zeta signaling domain as described in U.S. Patent No. 7,446,190 or

U.S. Patent No. 8,911,993. For example, in some embodiments, the intracellular signaling domain comprises the sequence of amino acids as set forth in SEQ ID NO: 13, 14 or 15 or a sequence of amino acids that exhibits at least 85%, 86%, 87%, 88%, 89%, 90%, 91%, 92%, 93%, 94%, 95%, 96%, 97%, 98%, 99% or more sequence identity to SEQ ID NO: 13, 14 or 15.

[0406] In some aspects, the spacer contains only a hinge region of an IgG, such as only a hinge of IgG4 or IgG1, such as the hinge only spacer set forth in SEQ ID NO: 1 or SEQ ID NO: 89. In other embodiments, the spacer is or contains an Ig hinge, *e.g.*, an IgG4-derived hinge, optionally linked to a C_H2 and/or C_H3 domains. In some embodiments, the spacer is an Ig hinge, *e.g.*, an IgG4 hinge, linked to C_H2 and C_H3 domains, such as set forth in SEQ ID NO: 4. In some embodiments, the spacer is an Ig hinge, *e.g.*, an IgG4 hinge, linked to a C_H3 domain only, such as set forth in SEQ ID NO: 3. In some embodiments, the spacer is or comprises a glycine-serine rich sequence or other flexible linker such as known flexible linkers. In some embodiments, the spacer is a CD8a hinge, such as set forth in any of SEQ ID NOs: 81-83, an FcγRIIIa hinge, such as set forth in SEQ ID NO: 88, a CTLA4 hinge, such as set forth in SEQ ID NO: 84, or a PD-1 hinge, such as set forth in SEQ ID NO: 86.

[0407] For example, in some embodiments, the CAR includes an antibody such as an antibody fragment, including scFvs, a spacer, such as a spacer containing a portion of an immunoglobulin molecule, such as a hinge region and/or one or more constant regions of a heavy chain molecule, such as an Ig-hinge containing spacer, a transmembrane domain containing all or a portion of a CD28-derived transmembrane domain, a CD28-derived intracellular signaling domain, and a CD3 zeta signaling domain. In some embodiments, the CAR includes an antibody or fragment, such as scFv, a spacer such as any of the Ig-hinge containing spacers, a CD28-derived transmembrane domain, a 4-1BB-derived intracellular signaling domain, and a CD3 zeta-derived signaling domain.

[0408] In some embodiments, the antigen receptor further includes a marker and/or cells expressing the CAR or other antigen receptor further includes a surrogate marker, such as a cell surface marker, which may be used to confirm transduction or engineering of the cell to express the receptor. In some embodiments, the marker is a molecule, e.g., cell surface protein, not naturally found on T cells or not naturally found on the surface of T cells, or a portion thereof. In some embodiments, the molecule is a non-self molecule, e.g., non-self protein, i.e., one that is not recognized as "self" by the immune system of the host into which the cells will be

adoptively transferred. In some embodiments, the marker serves no therapeutic function and/or produces no effect other than to be used as a marker for genetic engineering, e.g., for selecting cells successfully engineered. In other embodiments, the marker may be a therapeutic molecule or molecule otherwise exerting some desired effect, such as a ligand for a cell to be encountered *in vivo*, such as a costimulatory or immune checkpoint molecule to enhance and/or dampen responses of the cells upon adoptive transfer and encounter with ligand. In some aspects, the marker includes all or part (e.g., truncated form) of CD34, a NGFR, or epidermal growth factor receptor, such as truncated version of such a cell surface receptor (e.g., tEGFR). In some embodiments, the nucleic acid encoding the marker is operably linked to a polynucleotide encoding for a linker sequence, such as a cleavable linker sequence, e.g., T2A. For example, a marker, and optionally a linker sequence, can be any as disclosed in published patent application No. WO2014031687. For example, the marker can be a truncated EGFR (tEGFR) that is, optionally, linked to a linker sequence, such as a T2A cleavable linker sequence.

[0409] An exemplary polypeptide for a truncated EGFR (e.g. tEGFR) comprises the sequence of amino acids set forth in SEQ ID NO: 7 or 166 or a sequence of amino acids that exhibits at least 85%, 86%, 87%, 88%, 89%, 90%, 91%, 92%, 93%, 94%, 95%, 96%, 97%, 98%, 99% or more sequence identity to SEQ ID NO: 7 or 166. An exemplary T2A linker sequence comprises the sequence of amino acids set forth in SEQ ID NO: 6 or 167 or a sequence of amino acids that exhibits at least 85%, 86%, 87%, 88%, 89%, 90%, 91%, 92%, 93%, 94%, 95%, 96%, 97%, 98%, 99% or more sequence identity to SEQ ID NO: 6 or 167

[0410] In some embodiments, nucleic acid molecules encoding such CAR constructs further includes a sequence encoding a T2A ribosomal skip element and/or a tEGFR sequence, *e.g.*, downstream of the sequence encoding the CAR. In some embodiments, the sequence encodes a T2A ribosomal skip element set forth in SEQ ID NO: 6 or 167, or a sequence of amino acids that exhibits at least 85%, 86%, 87%, 88%, 89%, 90%, 91%, 92%, 93%, 94%, 95%, 96%, 97%, 98%, 99% or more sequence identity to SEQ ID NO: 6 or 167. In some embodiments, T cells expressing an antigen receptor (*e.g.* CAR) can also be generated to express a truncated EGFR (EGFRt) as a non-immunogenic selection epitope (*e.g.* by introduction of a construct encoding the CAR and EGFRt separated by a T2A ribosome switch to express two proteins from the same construct), which then can be used as a marker to detect such cells (see *e.g.* U.S. Patent No. 8,802,374). In some embodiments, the sequence encodes an tEGFR sequence set forth in SEQ

ID NO: 7 or 166, or a sequence of amino acids that exhibits at least 85%, 86%, 87%, 88%, 89%, 90%, 91%, 92%, 93%, 94%, 95%, 96%, 97%, 98%, 99% or more sequence identity to SEQ ID NO: 7.

[0411] In some embodiments, the encoded CAR can sequence can further include a signal sequence or signal peptide that directs or delivers the CAR to the surface of the cell in which the CAR is expressed. In some embodiments, the signal peptide is derived from a transmembrane protein. In some examples the signal peptide is derived from CD8a, CD33, or an IgG. Exemplary signal peptides include the sequences set forth in SEQ ID NOs: 39, 40 and 153.

[0412] In some embodiments, the CAR includes an anti-BCMA antibody or fragment, such as any of the anti-human BCMA antibodies, including sdAbs and scFvs, described herein, a spacer such as any of the Ig-hinge containing spacers or other spacers described herein, a CD28 transmembrane domain, a CD28 intracellular signaling domain, and a CD3 zeta signaling domain. In some embodiments, the CAR includes an anti-BCMA antibody or fragment, such as any of the anti-human BCMA antibodies, including sdAbs and scFvs described herein, a spacer such as any of the Ig-hinge containing spacers or other spacers described herein, a CD28 transmembrane domain, a 4-1BB intracellular signaling domain, and a CD3 zeta signaling domain. In some embodiments, such CAR constructs further includes a T2A ribosomal skip element and/or a tEGFR sequence, *e.g.*, downstream of the CAR.

[0413] The recombinant receptors, such as CARs, expressed by the cells administered to the subject generally recognize or specifically bind to a molecule that is expressed in, associated with, and/or specific for the disease or condition or cells thereof being treated. Upon specific binding to the molecule, *e.g.*, antigen, the receptor generally delivers an immunostimulatory signal, such as an ITAM-transduced signal, into the cell, thereby promoting an immune response targeted to the disease or condition. For example, in some embodiments, the cells express a CAR that specifically binds to an antigen expressed by a cell or tissue of the disease or condition or associated with the disease or condition. In some embodiments, the CAR specifically binds to BCMA, such as human BCMA, and includes an anti-human BCMA antibody or fragment as described. Non-limiting exemplary CAR sequences, including anti-BCMA CAR sequences, are set forth in SEQ ID NOs: 90-141. In some embodiments, an anti-BCMA CAR includes the amino acid sequence set forth in any of SEQ ID NOS: 90-141 or an amino acid sequence that exhibits at least at or about 90%, at or about 91%, at or about 92%, at or about 93%, at or about

94%, at or about 95%, at or about 96%, at or about 97%, at or about 96%, at or about 97%, at or about 98%, at or about 99% sequence identity to any one of SEQ ID NOS: 90-141, and wherein the CAR specifically binds BCMA, e.g. human BCMA.

2. Chimeric Auto-Antibody Receptor (CAAR)

[0414] In some embodiments, the recombinant receptor is a chimeric autoantibody receptor (CAAR). In some embodiments, the CAAR binds, e.g., specifically binds, or recognizes, an autoantibody. In some embodiments, a cell expressing the CAAR, such as a T cell engineered to express a CAAR, can be used to bind to and kill autoantibody-expressing cells, but not normal antibody expressing cells. In some embodiments, CAAR-expressing cells can be used to treat an autoimmune disease associated with expression of self-antigens, such as autoimmune diseases. In some embodiments, CAAR-expressing cells can target B cells that ultimately produce the autoantibodies and display the autoantibodies on their cell surfaces, mark these B cells as disease-specific targets for therapeutic intervention. In some embodiments, CAAR-expressing cells can be used to efficiently targeting and killing the pathogenic B cells in autoimmune diseases by targeting the disease-causing B cells using an antigen-specific chimeric autoantibody receptor. In some embodiments, the recombinant receptor is a CAAR, such as any described in U.S. Patent Application Pub. No. US 2017/0051035.

[0415] In some embodiments, the CAAR comprises an autoantibody binding domain, a transmembrane domain, and one or more intracellular signaling region or domain (also interchangeably called a cytoplasmic signaling domain or region). In some embodiments, the intracellular signaling region comprises an intracellular signaling domain. In some embodiments, the intracellular signaling domain is or comprises a primary signaling region, a signaling domain that is capable of stimulating and/or inducing a primary activation signal in a T cell, a signaling domain of a T cell receptor (TCR) component (e.g. an intracellular signaling domain or region of a zeta chain of a CD3-zeta (CD3ζ) chain or a functional variant or signaling portion thereof), and/or a signaling domain comprising an immunoreceptor tyrosine-based activation motif (ITAM).

[0416] In some embodiments, the autoantibody binding domain comprises an autoantigen or a fragment thereof. The choice of autoantigen can depend upon the type of autoantibody being targeted. For example, the autoantigen may be chosen because it recognizes an autoantibody on a target cell, such as a B cell, associated with a particular disease state, e.g. an autoimmune

disease, such as an autoantibody-mediated autoimmune disease. In some embodiments, the autoimmune disease includes pemphigus vulgaris (PV). Exemplary autoantigens include desmoglein 1 (Dsg1) and Dsg3.

3. TCRs

[0417] In some embodiments, engineered cells, such as T cells, are provided that express a T cell receptor (TCR) or antigen-binding portion thereof that recognizes an peptide epitope or T cell epitope of a target polypeptide, such as an antigen of a tumor, viral or autoimmune protein. In some aspects, the TCR is or includes a recombinant TCR.

[0418] In some embodiments, a "T cell receptor" or "TCR" is a molecule that contains a variable α and β chains (also known as TCRα and TCRβ, respectively) or a variable γ and δ chains (also known as TCRα and TCRβ, respectively), or antigen-binding portions thereof, and which is capable of specifically binding to a peptide bound to an MHC molecule. In some embodiments, the TCR is in the αβ form. Typically, TCRs that exist in αβ and γδ forms are generally structurally similar, but T cells expressing them may have distinct anatomical locations or functions. A TCR can be found on the surface of a cell or in soluble form. Generally, a TCR is found on the surface of T cells (or T lymphocytes) where it is generally responsible for recognizing antigens bound to major histocompatibility complex (MHC) molecules.

[0419] Unless otherwise stated, the term "TCR" should be understood to encompass full TCRs as well as antigen-binding portions or antigen-binding fragments thereof. In some embodiments, the TCR is an intact or full-length TCR, including TCRs in the αβ form or γδ form. In some embodiments, the TCR is an antigen-binding portion that is less than a full-length TCR but that binds to a specific peptide bound in an MHC molecule, such as binds to an MHC-peptide complex. In some cases, an antigen-binding portion or fragment of a TCR can contain only a portion of the structural domains of a full-length or intact TCR, but yet is able to bind the peptide epitope, such as MHC-peptide complex, to which the full TCR binds. In some cases, an antigen-binding portion contains the variable domains of a TCR, such as variable α chain and variable β chain of a TCR, sufficient to form a binding site for binding to a specific MHC-peptide complex. Generally, the variable chains of a TCR contain complementarity determining regions involved in recognition of the peptide, MHC and/or MHC-peptide complex.

[0420] In some embodiments, the variable domains of the TCR contain hypervariable loops, or complementarity determining regions (CDRs), which generally are the primary contributors to antigen recognition and binding capabilities and specificity. In some embodiments, a CDR of a TCR or combination thereof forms all or substantially all of the antigen-binding site of a given TCR molecule. The various CDRs within a variable region of a TCR chain generally are separated by framework regions (FRs), which generally display less variability among TCR molecules as compared to the CDRs (see, *e.g.*, Jores *et al.*, Proc. Nat'l Acad. Sci. U.S.A. 87:9138, 1990; Chothia *et al.*, EMBO J. 7:3745, 1988; see also Lefranc *et al.*, Dev. Comp. Immunol. 27:55, 2003). In some embodiments, CDR3 is the main CDR responsible for antigen binding or specificity, or is the most important among the three CDRs on a given TCR variable region for antigen recognition, and/or for interaction with the processed peptide portion of the peptide-MHC complex. In some contexts, the CDR1 of the alpha chain can interact with the N-terminal part of certain antigenic peptides. In some contexts, CDR1 of the beta chain can interact with the C-terminal part of the peptide. In some contexts, CDR2 contributes most strongly to or is the primary CDR responsible for the interaction with or recognition of the MHC portion of the MHC-peptide complex. In some embodiments, the variable region of the β-chain can contain a further hypervariable region (CDR4 or HVR4), which generally is involved in superantigen binding and not antigen recognition (Kotb (1995) Clinical Microbiology Reviews, 8:411-426).

[0421] In some embodiments, a TCR also can contain a constant domain, a transmembrane domain and/or a short cytoplasmic tail (see, *e.g.*, Janeway *et al.*, *Immunobiology: The Immune System in Health and Disease,* 3rd Ed., *Current Biology Publications*, p. 4:33, 1997). In some aspects, each chain of the TCR can possess one N-terminal immunoglobulin variable domain, one immunoglobulin constant domain, a transmembrane region, and a short cytoplasmic tail at the C-terminal end. In some embodiments, a TCR is associated with invariant proteins of the CD3 complex involved in mediating signal transduction.

[0422] In some embodiments, a TCR chain contains one or more constant domain. For example, the extracellular portion of a given TCR chain (*e.g.*, α-chain or β-chain) can contain two immunoglobulin-like domains, such as a variable domain (*e.g.*, Vα or Vβ; typically amino acids 1 to 116 based on Kabat numbering Kabat *et al.*, "Sequences of Proteins of Immunological Interest, US Dept. Health and Human Services, Public Health Service National Institutes of Health, 1991, 5th ed.) and a constant domain (*e.g.*, α-chain constant domain or Cα, typically

positions 117 to 259 of the chain based on Kabat numbering or β chain constant domain or Cβ, typically positions 117 to 295 of the chain based on Kabat) adjacent to the cell membrane. For example, in some cases, the extracellular portion of the TCR formed by the two chains contains two membrane-proximal constant domains, and two membrane-distal variable domains, which variable domains each contain CDRs. The constant domain of the TCR may contain short connecting sequences in which a cysteine residue forms a disulfide bond, thereby linking the two chains of the TCR. In some embodiments, a TCR may have an additional cysteine residue in each of the α and β chains, such that the TCR contains two disulfide bonds in the constant domains.

[0423] In some embodiments, the TCR chains contain a transmembrane domain. In some embodiments, the transmembrane domain is positively charged. In some cases, the TCR chain contains a cytoplasmic tail. In some cases, the structure allows the TCR to associate with other molecules like CD3 and subunits thereof. For example, a TCR containing constant domains with a transmembrane region may anchor the protein in the cell membrane and associate with invariant subunits of the CD3 signaling apparatus or complex. The intracellular tails of CD3 signaling subunits (*e.g.* CD3γ, CD3δ, CD3ε and CD3ζ chains) contain one or more immunoreceptor tyrosine-based activation motif or ITAM that are involved in the signaling capacity of the TCR complex.

[0424] In some embodiments, the TCR may be a heterodimer of two chains α and β (or optionally γ and δ) or it may be a single chain TCR construct. In some embodiments, the TCR is a heterodimer containing two separate chains (α and β chains or γ and δ chains) that are linked, such as by a disulfide bond or disulfide bonds.

[0425] In some embodiments, the TCR can be generated from a known TCR sequence(s), such as sequences of Vα,β chains, for which a substantially full-length coding sequence is readily available. Methods for obtaining full-length TCR sequences, including V chain sequences, from cell sources are well known. In some embodiments, nucleic acids encoding the TCR can be obtained from a variety of sources, such as by polymerase chain reaction (PCR) amplification of TCR-encoding nucleic acids within or isolated from a given cell or cells, or synthesis of publicly available TCR DNA sequences.

[0426] In some embodiments, the recombinant receptors include recombinant TCRs and/or TCRs cloned from naturally occurring T cells. In some embodiments, a high-affinity T cell

clone for a target antigen (e.g., a cancer antigen) is identified, isolated from a patient, and introduced into the cells. In some embodiments, the TCR clone for a target antigen has been generated in transgenic mice engineered with human immune system genes (e.g., the human leukocyte antigen system, or HLA). See, e.g., tumor antigens (see, e.g., Parkhurst et al. (2009) Clin Cancer Res. 15:169–180 and Cohen et al. (2005) J Immunol. 175:5799–5808. In some embodiments, phage display is used to isolate TCRs against a target antigen (see, e.g., Varela-Rohena et al. (2008) Nat Med. 14:1390–1395 and Li (2005) Nat Biotechnol. 23:349–354.

[0427] In some embodiments, the TCR is obtained from a biological source, such as from cells such as from a T cell (*e.g.* cytotoxic T cell), T-cell hybridomas or other publicly available source. In some embodiments, the T-cells can be obtained from *in vivo* isolated cells. In some embodiments, the TCR is a thymically selected TCR. In some embodiments, the TCR is a neoepitope-restricted TCR. In some embodiments, the T- cells can be a cultured T-cell hybridoma or clone. In some embodiments, the TCR or antigen-binding portion thereof can be synthetically generated from knowledge of the sequence of the TCR.

[0428] In some embodiments, the TCR is generated from a TCR identified or selected from screening a library of candidate TCRs against a target polypeptide antigen, or target T cell epitope thereof. TCR libraries can be generated by amplification of the repertoire of Vα and Vβ from T cells isolated from a subject, including cells present in PBMCs, spleen or other lymphoid organ. In some cases, T cells can be amplified from tumor-infiltrating lymphocytes (TILs). In some embodiments, TCR libraries can be generated from $CD4^+$ or $CD8^+$ cells. In some embodiments, the TCRs can be amplified from a T cell source of a normal of healthy subject, *i.e.* normal TCR libraries. In some embodiments, the TCRs can be amplified from a T cell source of a diseased subject, *i.e.* diseased TCR libraries. In some embodiments, degenerate primers are used to amplify the gene repertoire of Vα and Vβ, such as by RT-PCR in samples, such as T cells, obtained from humans. In some embodiments, scTv libraries can be assembled from naïve Vα and Vβ libraries in which the amplified products are cloned or assembled to be separated by a linker. Depending on the source of the subject and cells, the libraries can be HLA allele-specific. Alternatively, in some embodiments, TCR libraries can be generated by mutagenesis or diversification of a parent or scaffold TCR molecule. In some aspects, the TCRs are subjected to directed evolution, such as by mutagenesis, *e.g.*, of the α or β chain. In some aspects, particular residues within CDRs of the TCR are altered. In some embodiments, selected

TCRs can be modified by affinity maturation. In some embodiments, antigen-specific T cells may be selected, such as by screening to assess CTL activity against the peptide. In some aspects, TCRs, *e.g.* present on the antigen-specific T cells, may be selected, such as by binding activity, *e.g.*, particular affinity or avidity for the antigen.

[0429] In some embodiments, the TCR or antigen-binding portion thereof is one that has been modified or engineered. In some embodiments, directed evolution methods are used to generate TCRs with altered properties, such as with higher affinity for a specific MHC-peptide complex. In some embodiments, directed evolution is achieved by display methods including, but not limited to, yeast display (Holler *et al.*, (2003) *Nat Immunol*, 4, 55-62; Holler *et al.*, (2000) *Proc Natl Acad Sci U S A,* 97, 5387-92), phage display (Li *et al.*, (2005) *Nat Biotechnol,* 23, 349-54), or T cell display (Chervin *et al.*, (2008) *J Immunol Methods,* 339, 175-84). In some embodiments, display approaches involve engineering, or modifying, a known, parent or reference TCR. For example, in some cases, a wild-type TCR can be used as a template for producing mutagenized TCRs in which in one or more residues of the CDRs are mutated, and mutants with an desired altered property, such as higher affinity for a desired target antigen, are selected.

[0430] In some embodiments, peptides of a target polypeptide for use in producing or generating a TCR of interest are known or can be readily identified. In some embodiments, peptides suitable for use in generating TCRs or antigen-binding portions can be determined based on the presence of an HLA-restricted motif in a target polypeptide of interest, such as a target polypeptide described herein. In some embodiments, peptides are identified using available computer prediction models. In some embodiments, for predicting MHC class I binding sites, such models include, but are not limited to, ProPred1 (Singh and Raghava (2001) *Bioinformatics* 17(12):1236-1237, and SYFPEITHI (see Schuler *et al.*, (2007) *Immunoinformatics Methods in Molecular Biology*, 409(1): 75-93 2007). In some embodiments, the MHC-restricted epitope is HLA-A0201, which is expressed in approximately 39-46% of all Caucasians and therefore, represents a suitable choice of MHC antigen for use preparing a TCR or other MHC-peptide binding molecule.

[0431] HLA-A0201-binding motifs and the cleavage sites for proteasomes and immune-proteasomes using computer prediction models. For predicting MHC class I binding sites, such models include, but are not limited to, ProPred1 (described in more detail in Singh and Raghava,

ProPred: prediction of HLA-DR binding sites. *BIOINFORMATICS* 17(12):1236-1237 2001), and SYFPEITHI (see Schuler *et al.*, SYFPEITHI, Database for Searching and T-Cell Epitope Prediction. in *Immunoinformatics Methods in Molecular Biology*, vol 409(1): 75-93 2007)

[0432] In some embodiments, the TCR or antigen binding portion thereof may be a recombinantly produced natural protein or mutated form thereof in which one or more property, such as binding characteristic, has been altered. In some embodiments, a TCR may be derived from one of various animal species, such as human, mouse, rat, or other mammal. A TCR may be cell-bound or in soluble form. In some embodiments, for purposes of the provided methods, the TCR is in cell-bound form expressed on the surface of a cell.

[0433] In some embodiments, the TCR is a full-length TCR. In some embodiments, the TCR is an antigen-binding portion. In some embodiments, the TCR is a dimeric TCR (dTCR). In some embodiments, the TCR is a single-chain TCR (sc-TCR). In some embodiments, a dTCR or scTCR have the structures as described in WO 03/020763, WO 04/033685, WO2011/044186.

[0434] In some embodiments, the TCR contains a sequence corresponding to the transmembrane sequence. In some embodiments, the TCR does contain a sequence corresponding to cytoplasmic sequences. In some embodiments, the TCR is capable of forming a TCR complex with CD3. In some embodiments, any of the TCRs, including a dTCR or scTCR, can be linked to signaling domains that yield an active TCR on the surface of a T cell. In some embodiments, the TCR is expressed on the surface of cells.

[0435] In some embodiments a dTCR contains a first polypeptide wherein a sequence corresponding to a TCR α chain variable region sequence is fused to the N terminus of a sequence corresponding to a TCR α chain constant region extracellular sequence, and a second polypeptide wherein a sequence corresponding to a TCR β chain variable region sequence is fused to the N terminus a sequence corresponding to a TCR β chain constant region extracellular sequence, the first and second polypeptides being linked by a disulfide bond. In some embodiments, the bond can correspond to the native inter-chain disulfide bond present in native dimeric αβ TCRs. In some embodiments, the interchain disulfide bonds are not present in a native TCR. For example, in some embodiments, one or more cysteines can be incorporated into the constant region extracellular sequences of dTCR polypeptide pair. In some cases, both a native and a non-native disulfide bond may be desirable. In some embodiments, the TCR contains a transmembrane sequence to anchor to the membrane.

[0436] In some embodiments, a dTCR contains a TCR α chain containing a variable α domain, a constant α domain and a first dimerization motif attached to the C-terminus of the constant α domain, and a TCR β chain comprising a variable β domain, a constant β domain and a first dimerization motif attached to the C-terminus of the constant β domain, wherein the first and second dimerization motifs easily interact to form a covalent bond between an amino acid in the first dimerization motif and an amino acid in the second dimerization motif linking the TCR α chain and TCR β chain together.

[0437] In some embodiments, the TCR is a scTCR. Typically, a scTCR can be generated using methods known, See *e.g.*, Soo Hoo, W. F. *et al.*, *PNAS (USA)* 89, 4759 (1992); Wülfing, C. and Plückthun, A., *J. Mol. Biol.* 242, 655 (1994); Kurucz, I. *et al.*, *PNAS (USA)* 90 3830 (1993); International published PCT Nos. WO 96/13593, WO 96/18105, WO99/60120, WO99/18129, WO 03/020763, WO2011/044186; and Schlueter, C. J. *et al.*, *J. Mol. Biol.* 256, 859 (1996). In some embodiments, a scTCR contains an introduced non-native disulfide interchain bond to facilitate the association of the TCR chains (see *e.g.* International published PCT No. WO 03/020763). In some embodiments, a scTCR is a non-disulfide linked truncated TCR in which heterologous leucine zippers fused to the C-termini thereof facilitate chain association (see *e.g.* International published PCT No. WO99/60120). In some embodiments, a scTCR contains a TCRα variable domain covalently linked to a TCRβ variable domain via a peptide linker (see *e.g.*, International published PCT No. WO99/18129).

[0438] In some embodiments, a scTCR contains a first segment constituted by an amino acid sequence corresponding to a TCR α chain variable region, a second segment constituted by an amino acid sequence corresponding to a TCR β chain variable region sequence fused to the N terminus of an amino acid sequence corresponding to a TCR β chain constant domain extracellular sequence, and a linker sequence linking the C terminus of the first segment to the N terminus of the second segment.

[0439] In some embodiments, a scTCR contains a first segment constituted by an α chain variable region sequence fused to the N terminus of an α chain extracellular constant domain sequence, and a second segment constituted by a β chain variable region sequence fused to the N terminus of a sequence β chain extracellular constant and transmembrane sequence, and, optionally, a linker sequence linking the C terminus of the first segment to the N terminus of the second segment.

[0440] In some embodiments, a scTCR contains a first segment constituted by a TCR β chain variable region sequence fused to the N terminus of a β chain extracellular constant domain sequence, and a second segment constituted by an α chain variable region sequence fused to the N terminus of a sequence α chain extracellular constant and transmembrane sequence, and, optionally, a linker sequence linking the C terminus of the first segment to the N terminus of the second segment.

[0441] In some embodiments, the linker of a scTCRs that links the first and second TCR segments can be any linker capable of forming a single polypeptide strand, while retaining TCR binding specificity. In some embodiments, the linker sequence may, for example, have the formula -P-AA-P- wherein P is proline and AA represents an amino acid sequence wherein the amino acids are glycine and serine. In some embodiments, the first and second segments are paired so that the variable region sequences thereof are orientated for such binding. Hence, in some cases, the linker has a sufficient length to span the distance between the C terminus of the first segment and the N terminus of the second segment, or vice versa, but is not too long to block or reduces bonding of the scTCR to the target ligand. In some embodiments, the linker can contain from 10 to 45 amino acids or from about 10 to 45 amino acids, such as 10 to 30 amino acids or 26 to 41 amino acids residues, for example 29, 30, 31 or 32 amino acids. In some embodiments, the linker has the formula -PGGG-(SGGGG)5-P- wherein P is proline, G is glycine and S is serine (SEQ ID NO: 16). In some embodiments, the linker has the sequence GSADDAKKDAAKKDGKS (SEQ ID NO: 17)

[0442] In some embodiments, the scTCR contains a covalent disulfide bond linking a residue of the immunoglobulin region of the constant domain of the α chain to a residue of the immunoglobulin region of the constant domain of the β chain. In some embodiments, the interchain disulfide bond in a native TCR is not present. For example, in some embodiments, one or more cysteines can be incorporated into the constant region extracellular sequences of the first and second segments of the scTCR polypeptide. In some cases, both a native and a non-native disulfide bond may be desirable.

[0443] In some embodiments of a dTCR or scTCR containing introduced interchain disulfide bonds, the native disulfide bonds are not present. In some embodiments, the one or more of the native cysteines forming a native interchain disulfide bonds are substituted to another residue, such as to a serine or alanine. In some embodiments, an introduced disulfide

bond can be formed by mutating non-cysteine residues on the first and second segments to cysteine. Exemplary non-native disulfide bonds of a TCR are described in published International PCT No. WO2006/000830.

[0444] In some embodiments, the TCR or antigen-binding fragment thereof exhibits an affinity with an equilibrium binding constant for a target antigen of between or between about 10^{-5} and 10^{-12} M and all individual values and ranges therein. In some embodiments, the target antigen is an MHC-peptide complex or ligand.

[0445] In some embodiments, nucleic acid or nucleic acids encoding a TCR, such as α and β chains, can be amplified by PCR, cloning or other suitable means and cloned into a suitable expression vector or vectors. The expression vector can be any suitable recombinant expression vector, and can be used to transform or transfect any suitable host. Suitable vectors include those designed for propagation and expansion or for expression or both, such as plasmids and viruses.

[0446] In some embodiments, the vector can a vector of the pUC series (Fermentas Life Sciences), the pBluescript series (Stratagene, LaJolla, Calif.), the pET series (Novagen, Madison, Wis.), the pGEX series (Pharmacia Biotech, Uppsala, Sweden), or the pEX series (Clontech, Palo Alto, Calif.). In some cases, bacteriophage vectors, such as λG10, λGT11, λZapII (Stratagene), λEMBL4, and λNM1149, also can be used. In some embodiments, plant expression vectors can be used and include pBI01, pBI101.2, pBI101.3, pBI121 and pBIN19 (Clontech). In some embodiments, animal expression vectors include pEUK-Cl, pMAM and pMAMneo (Clontech). In some embodiments, a viral vector is used, such as a retroviral vector.

[0447] In some embodiments, the recombinant expression vectors can be prepared using standard recombinant DNA techniques. In some embodiments, vectors can contain regulatory sequences, such as transcription and translation initiation and termination codons, which are specific to the type of host (*e.g.*, bacterium, fungus, plant, or animal) into which the vector is to be introduced, as appropriate and taking into consideration whether the vector is DNA- or RNA-based. In some embodiments, the vector can contain a nonnative promoter operably linked to the nucleotide sequence encoding the TCR or antigen-binding portion (or other MHC-peptide binding molecule). In some embodiments, the promoter can be a non-viral promoter or a viral promoter, such as a cytomegalovirus (CMV) promoter, an SV40 promoter, an RSV promoter, and a promoter found in the long-terminal repeat of the murine stem cell virus. Other known promoters also are contemplated.

[0448] In some embodiments, to generate a vector encoding a TCR, the α and β chains are PCR amplified from total cDNA isolated from a T cell clone expressing the TCR of interest and cloned into an expression vector. In some embodiments, the α and β chains are cloned into the same vector. In some embodiments, the α and β chains are cloned into different vectors. In some embodiments, the generated α and β chains are incorporated into a retroviral, *e.g.* lentiviral, vector.

4. Multi-targeting

[0449] In some embodiments, the cells and methods include multi-targeting strategies, such as expression of two or more genetically engineered receptors on the cell, each recognizing the same or a different antigen and typically each including a different intracellular signaling component. Such multi-targeting strategies are described, for example, in PCT Pub. No. WO 2014055668 A1 (describing combinations of activating and costimulatory CARs, *e.g.*, targeting two different antigens present individually on off-target, *e.g.*, normal cells, but present together only on cells of the disease or condition to be treated) and Fedorov *et al.*, *Sci. Transl. Medicine,* 5(215) (2013) (describing cells expressing an activating and an inhibitory CAR, such as those in which the activating CAR binds to one antigen expressed on both normal or non-diseased cells and cells of the disease or condition to be treated, and the inhibitory CAR binds to another antigen expressed only on the normal cells or cells which it is not desired to treat).

[0450] For example, in some embodiments, the cells include a receptor expressing a first genetically engineered antigen receptor (*e.g.*, CAR or TCR) which is capable of inducing an activating signal to the cell, generally upon specific binding to the antigen recognized by the first receptor, *e.g.*, a first antigen. In some embodiments, the cell further includes a second genetically engineered antigen receptor (*e.g.*, CAR or TCR), *e.g.*, a chimeric costimulatory receptor, which is capable of inducing a costimulatory signal to the immune cell, generally upon specific binding to a second antigen recognized by the second receptor. In some embodiments, the first antigen and second antigen are the same. In some embodiments, the first antigen and second antigen are different.

[0451] In some embodiments, the first and/or second genetically engineered antigen receptor (*e.g.* CAR or TCR) is capable of inducing an activating signal to the cell. In some embodiments, the receptor includes an intracellular signaling component containing ITAM or ITAM-like motifs. In some embodiments, the activation induced by the first receptor involves a signal

transduction or change in protein expression in the cell resulting in initiation of an immune response, such as ITAM phosphorylation and/or initiation of ITAM-mediated signal transduction cascade, formation of an immunological synapse and/or clustering of molecules near the bound receptor (*e.g.* CD4 or CD8, etc.), activation of one or more transcription factors, such as NF-κB and/or AP-1, and/or induction of gene expression of factors such as cytokines, proliferation, and/or survival.

[0452] In some embodiments, the first and/or second receptor includes intracellular signaling domains of costimulatory receptors such as CD28, CD137 (4-1BB), OX40, and/or ICOS. In some embodiments, the first and second receptors include an intracellular signaling domain of a costimulatory receptor that are different. In one embodiment, the first receptor contains a CD28 costimulatory signaling region and the second receptor contain a 4-1BB co-stimulatory signaling region or vice versa.

[0453] In some embodiments, the first and/or second receptor includes both an intracellular signaling domain containing ITAM or ITAM-like motifs and an intracellular signaling domain of a costimulatory receptor.

[0454] In some embodiments, the first receptor contains an intracellular signaling domain containing ITAM or ITAM-like motifs and the second receptor contains an intracellular signaling domain of a costimulatory receptor. The costimulatory signal in combination with the activating signal induced in the same cell is one that results in an immune response, such as a robust and sustained immune response, such as increased gene expression, secretion of cytokines and other factors, and T cell mediated effector functions such as cell killing.

[0455] In some embodiments, neither ligation of the first receptor alone nor ligation of the second receptor alone induces a robust immune response. In some aspects, if only one receptor is ligated, the cell becomes tolerized or unresponsive to antigen, or inhibited, and/or is not induced to proliferate or secrete factors or carry out effector functions. In some such embodiments, however, when the plurality of receptors are ligated, such as upon encounter of a cell expressing the first and second antigens, a desired response is achieved, such as full immune activation or stimulation, *e.g.*, as indicated by secretion of one or more cytokine, proliferation, persistence, and/or carrying out an immune effector function such as cytotoxic killing of a target cell.

[0456] In some embodiments, the cells expressing the recombinant receptor further include inhibitory CARs (iCARs, see Fedorov *et al.*, *Sci. Transl. Medicine*, 5(215) (2013), such as a CAR recognizing an antigen other than the one associated with and/or specific for the disease or condition whereby an activating signal delivered through the disease-targeting CAR is diminished or inhibited by binding of the inhibitory CAR to its ligand, *e.g.*, to reduce off-target effects.

[0457] In some embodiments, the two receptors induce, respectively, an activating and an inhibitory signal to the cell, such that binding by one of the receptor to its antigen activates the cell or induces a response, but binding by the second inhibitory receptor to its antigen induces a signal that suppresses or dampens that response. Examples are combinations of activating CARs and inhibitory CARs or iCARs. Such a strategy may be used, for example, in which the activating CAR binds an antigen expressed in a disease or condition but which is also expressed on normal cells, and the inhibitory receptor binds to a separate antigen which is expressed on the normal cells but not cells of the disease or condition.

[0458] In some aspects, the chimeric receptor is or includes an inhibitory CAR (e.g. iCAR) and includes intracellular components that dampen or suppress an immune response, such as an ITAM- and/or co stimulatory-promoted response in the cell. Examples of such intracellular signaling components are those found on immune checkpoint molecules, including PD-1, CTLA4, LAG3, BTLA, OX2R, TIM-3, TIGIT, LAIR-1, PGE2 receptors, EP2/4 Adenosine receptors including A2AR. In some aspects, the engineered cell includes an inhibitory CAR including a signaling domain of or derived from such an inhibitory molecule, such that it serves to dampen the response of the cell, for example, that induced by an activating and/or costimulatory CAR.

[0459] In some embodiments, the multi-targeting strategy is employed in a case where an antigen associated with a particular disease or condition is expressed on a non-diseased cell and/or is expressed on the engineered cell itself, either transiently (*e.g.*, upon stimulation in association with genetic engineering) or permanently. In such cases, by requiring ligation of two separate and individually specific antigen receptors, specificity, selectivity, and/or efficacy may be improved.

[0460] In some embodiments, the plurality of antigens, *e.g.*, the first and second antigens, are expressed on the cell, tissue, or disease or condition being targeted, such as on the cancer

cell. In some aspects, the cell, tissue, disease or condition is multiple myeloma or a multiple myeloma cell. In some embodiments, one or more of the plurality of antigens generally also is expressed on a cell which it is not desired to target with the cell therapy, such as a normal or non-diseased cell or tissue, and/or the engineered cells themselves. In such embodiments, by requiring ligation of multiple receptors to achieve a response of the cell, specificity and/or efficacy is achieved.

B. CELLS AND PREPARATION OF CELLS FOR GENETIC ENGINEERING

[0461] Among the cells expressing the receptors and administered by the provided methods are engineered cells. The genetic engineering generally involves introduction of a nucleic acid encoding the recombinant or engineered component into a composition containing the cells, such as by retroviral transduction, transfection, or transformation.

[0462] In some embodiments, the nucleic acids are heterologous, *i.e.*, normally not present in a cell or sample obtained from the cell, such as one obtained from another organism or cell, which for example, is not ordinarily found in the cell being engineered and/or an organism from which such cell is derived. In some embodiments, the nucleic acids are not naturally occurring, such as a nucleic acid not found in nature, including one comprising chimeric combinations of nucleic acids encoding various domains from multiple different cell types.

[0463] The cells generally are eukaryotic cells, such as mammalian cells, and typically are human cells. In some embodiments, the cells are derived from the blood, bone marrow, lymph, or lymphoid organs, are cells of the immune system, such as cells of the innate or adaptive immunity, *e.g.*, myeloid or lymphoid cells, including lymphocytes, typically T cells and/or NK cells. Other exemplary cells include stem cells, such as multipotent and pluripotent stem cells, including induced pluripotent stem cells (iPSCs). The cells typically are primary cells, such as those isolated directly from a subject and/or isolated from a subject and frozen. In some embodiments, the cells include one or more subsets of T cells or other cell types, such as whole T cell populations, $CD4^+$ cells, $CD8^+$ cells, and subpopulations thereof, such as those defined by function, activation state, maturity, potential for differentiation, expansion, recirculation, localization, and/or persistence capacities, antigen-specificity, type of antigen receptor, presence in a particular organ or compartment, marker or cytokine secretion profile, and/or degree of differentiation. With reference to the subject to be treated, the cells may be allogeneic and/or autologous. Among the methods include off-the-shelf methods. In some aspects, such as for off-

the-shelf technologies, the cells are pluripotent and/or multipotent, such as stem cells, such as induced pluripotent stem cells (iPSCs). In some embodiments, the methods include isolating cells from the subject, preparing, processing, culturing, and/or engineering them, and re-introducing them into the same subject, before or after cryopreservation.

[0464] Among the sub-types and subpopulations of T cells and/or of $CD4^+$ and/or of $CD8^+$ T cells are naïve T (T_N) cells, effector T cells (T_{EFF}), memory T cells and sub-types thereof, such as stem cell memory T (T_{SCM}), central memory T (T_{CM}), effector memory T (T_{EM}), or terminally differentiated effector memory T cells, tumor-infiltrating lymphocytes (TIL), immature T cells, mature T cells, helper T cells, cytotoxic T cells, mucosa-associated invariant T (MAIT) cells, naturally occurring and adaptive regulatory T (Treg) cells, helper T cells, such as TH1 cells, TH2 cells, TH3 cells, TH17 cells, TH9 cells, TH22 cells, follicular helper T cells, alpha/beta T cells, and delta/gamma T cells.

[0465] In some embodiments, the cells are natural killer (NK) cells. In some embodiments, the cells are monocytes or granulocytes, *e.g.*, myeloid cells, macrophages, neutrophils, dendritic cells, mast cells, eosinophils, and/or basophils.

[0466] In some embodiments, the cells include one or more nucleic acids introduced via genetic engineering, and thereby express recombinant or genetically engineered products of such nucleic acids. In some embodiments, the nucleic acids are heterologous, *i.e.*, normally not present in a cell or sample obtained from the cell, such as one obtained from another organism or cell, which for example, is not ordinarily found in the cell being engineered and/or an organism from which such cell is derived. In some embodiments, the nucleic acids are not naturally occurring, such as a nucleic acid not found in nature, including one comprising chimeric combinations of nucleic acids encoding various domains from multiple different cell types.

[0467] In some embodiments, preparation of the engineered cells includes one or more culture and/or preparation steps. The cells for introduction of the nucleic acid encoding the transgenic receptor such as the CAR, may be isolated from a sample, such as a biological sample, *e.g.*, one obtained from or derived from a subject. In some embodiments, the subject from which the cell is isolated is one having the disease or condition or in need of a cell therapy or to which cell therapy will be administered. The subject in some embodiments is a human in

need of a particular therapeutic intervention, such as the adoptive cell therapy for which cells are being isolated, processed, and/or engineered.

[0468] Accordingly, the cells in some embodiments are primary cells, *e.g.*, primary human cells. The samples include tissue, fluid, and other samples taken directly from the subject, as well as samples resulting from one or more processing steps, such as separation, centrifugation, genetic engineering (*e.g.* transduction with viral vector), washing, and/or incubation. The biological sample can be a sample obtained directly from a biological source or a sample that is processed. Biological samples include, but are not limited to, body fluids, such as blood, plasma, serum, cerebrospinal fluid, synovial fluid, urine and sweat, tissue and organ samples, including processed samples derived therefrom.

[0469] In some aspects, the sample from which the cells are derived or isolated is blood or a blood-derived sample, or is or is derived from an apheresis or leukapheresis product. Exemplary samples include whole blood, peripheral blood mononuclear cells (PBMCs), leukocytes, bone marrow, thymus, tissue biopsy, tumor, leukemia, lymphoma, lymph node, gut associated lymphoid tissue, mucosa associated lymphoid tissue, spleen, other lymphoid tissues, liver, lung, stomach, intestine, colon, kidney, pancreas, breast, bone, prostate, cervix, testes, ovaries, tonsil, or other organ, and/or cells derived therefrom. Samples include, in the context of cell therapy, *e.g.*, adoptive cell therapy, samples from autologous and allogeneic sources.

[0470] In some embodiments, the cells are derived from cell lines, *e.g.*, T cell lines. The cells in some embodiments are obtained from a xenogeneic source, for example, from mouse, rat, non-human primate, and pig.

[0471] In some embodiments, isolation of the cells includes one or more preparation and/or non-affinity based cell separation steps. In some examples, cells are washed, centrifuged, and/or incubated in the presence of one or more reagents, for example, to remove unwanted components, enrich for desired components, lyse or remove cells sensitive to particular reagents. In some examples, cells are separated based on one or more property, such as density, adherent properties, size, sensitivity and/or resistance to particular components.

[0472] In some examples, cells from the circulating blood of a subject are obtained, *e.g.*, by apheresis or leukapheresis. The samples, in some aspects, contain lymphocytes, including T cells, monocytes, granulocytes, B cells, other nucleated white blood cells, red blood cells, and/or platelets, and in some aspects contain cells other than red blood cells and platelets.

[0473] In some embodiments, the blood cells collected from the subject are washed, *e.g.*, to remove the plasma fraction and to place the cells in an appropriate buffer or media for subsequent processing steps. In some embodiments, the cells are washed with phosphate buffered saline (PBS). In some embodiments, the wash solution lacks calcium and/or magnesium and/or many or all divalent cations. In some aspects, a washing step is accomplished a semi-automated "flow-through" centrifuge (for example, the Cobe 2991 cell processor, Baxter) according to the manufacturer's instructions. In some aspects, a washing step is accomplished by tangential flow filtration (TFF) according to the manufacturer's instructions. In some embodiments, the cells are resuspended in a variety of biocompatible buffers after washing, such as, for example, Ca^{++}/Mg^{++} free PBS. In certain embodiments, components of a blood cell sample are removed and the cells directly resuspended in culture media.

[0474] In some embodiments, the methods include density-based cell separation methods, such as the preparation of white blood cells from peripheral blood by lysing the red blood cells and centrifugation through a Percoll or Ficoll gradient.

[0475] In some embodiments, the isolation methods include the separation of different cell types based on the expression or presence in the cell of one or more specific molecules, such as surface markers, *e.g.*, surface proteins, intracellular markers, or nucleic acid. In some embodiments, any known method for separation based on such markers may be used. In some embodiments, the separation is affinity- or immunoaffinity-based separation. For example, the isolation in some aspects includes separation of cells and cell populations based on the cells' expression or expression level of one or more markers, typically cell surface markers, for example, by incubation with an antibody or binding partner that specifically binds to such markers, followed generally by washing steps and separation of cells having bound the antibody or binding partner, from those cells having not bound to the antibody or binding partner.

[0476] Such separation steps can be based on positive selection, in which the cells having bound the reagents are retained for further use, and/or negative selection, in which the cells having not bound to the antibody or binding partner are retained. In some examples, both fractions are retained for further use. In some aspects, negative selection can be particularly useful where no antibody is available that specifically identifies a cell type in a heterogeneous population, such that separation is best carried out based on markers expressed by cells other than the desired population.

[0477] The separation need not result in 100% enrichment or removal of a particular cell population or cells expressing a particular marker. For example, positive selection of or enrichment for cells of a particular type, such as those expressing a marker, refers to increasing the number or percentage of such cells, but need not result in a complete absence of cells not expressing the marker. Likewise, negative selection, removal, or depletion of cells of a particular type, such as those expressing a marker, refers to decreasing the number or percentage of such cells, but need not result in a complete removal of all such cells.

[0478] In some examples, multiple rounds of separation steps are carried out, where the positively or negatively selected fraction from one step is subjected to another separation step, such as a subsequent positive or negative selection. In some examples, a single separation step can deplete cells expressing multiple markers simultaneously, such as by incubating cells with a plurality of antibodies or binding partners, each specific for a marker targeted for negative selection. Likewise, multiple cell types can simultaneously be positively selected by incubating cells with a plurality of antibodies or binding partners expressed on the various cell types.

[0479] For example, in some aspects, specific subpopulations of T cells, such as cells positive or expressing high levels of one or more surface markers, *e.g.*, $CD28^+$, $CD62L^+$, $CCR7^+$, $CD27^+$, $CD127^+$, $CD4^+$, $CD8^+$, $CD45RA^+$, and/or $CD45RO^+$ T cells, are isolated by positive or negative selection techniques.

[0480] For example, $CD3^+$, $CD28^+$ T cells can be positively selected using CD3/CD28 conjugated magnetic beads (*e.g.*, DYNABEADS® M-450 CD3/CD28 T Cell Expander).

[0481] In some embodiments, isolation is carried out by enrichment for a particular cell population by positive selection, or depletion of a particular cell population, by negative selection. In some embodiments, positive or negative selection is accomplished by incubating cells with one or more antibodies or other binding agent that specifically bind to one or more surface markers expressed or expressed ($marker^+$) at a relatively higher level (markerhigh) on the positively or negatively selected cells, respectively.

[0482] In some embodiments, T cells are separated from a PBMC sample by negative selection of markers expressed on non-T cells, such as B cells, monocytes, or other white blood cells, such as CD14. In some aspects, a $CD4^+$ or $CD8^+$ selection step is used to separate $CD4^+$ helper and $CD8^+$ cytotoxic T cells. Such $CD4^+$ and $CD8^+$ populations can be further sorted into

sub-populations by positive or negative selection for markers expressed or expressed to a relatively higher degree on one or more naive, memory, and/or effector T cell subpopulations.

[0483] In some embodiments, $CD8^+$ cells are further enriched for or depleted of naive, central memory, effector memory, and/or central memory stem cells, such as by positive or negative selection based on surface antigens associated with the respective subpopulation. In some embodiments, enrichment for central memory T (T_{CM}) cells is carried out to increase efficacy, such as to improve long-term survival, expansion, and/or engraftment following administration, which in some aspects is particularly robust in such sub-populations. See Terakura *et al.*, *Blood*.1:72–82 (2012); Wang *et al.*, *J Immunother.* 35(9):689-701 (2012). In some embodiments, combining T_{CM}-enriched $CD8^+$ T cells and $CD4^+$ T cells further enhances efficacy.

[0484] In embodiments, memory T cells are present in both $CD62L^+$ and CD62L- subsets of $CD8^+$ peripheral blood lymphocytes. PBMC can be enriched for or depleted of CD62L-$CD8^+$ and/or $CD62L^+CD8^+$ fractions, such as using anti-CD8 and anti-CD62L antibodies.

[0485] In some embodiments, the enrichment for central memory T (TCM) cells is based on positive or high surface expression of CD45RO, CD62L, CCR7, CD28, CD3, and/or CD 127; in some aspects, it is based on negative selection for cells expressing or highly expressing CD45RA and/or granzyme B. In some aspects, isolation of a $CD8^+$ population enriched for TCM cells is carried out by depletion of cells expressing CD4, CD14, CD45RA, and positive selection or enrichment for cells expressing CD62L. In one aspect, enrichment for central memory T (T_{CM}) cells is carried out starting with a negative fraction of cells selected based on CD4 expression, which is subjected to a negative selection based on expression of CD14 and CD45RA, and a positive selection based on CD62L. Such selections in some aspects are carried out simultaneously and in other aspects are carried out sequentially, in either order. In some aspects, the same CD4 expression-based selection step used in preparing the $CD8^+$ cell population or subpopulation, also is used to generate the $CD4^+$ cell population or sub-population, such that both the positive and negative fractions from the CD4-based separation are retained and used in subsequent steps of the methods, optionally following one or more further positive or negative selection steps.

[0486] In a particular example, a sample of PBMCs or other white blood cell sample is subjected to selection of $CD4^+$ cells, where both the negative and positive fractions are retained.

The negative fraction then is subjected to negative selection based on expression of CD14 and CD45RA or CD19, and positive selection based on a marker characteristic of central memory T cells, such as CD62L or CCR7, where the positive and negative selections are carried out in either order.

[0487] $CD4^+$ T helper cells are sorted into naïve, central memory, and effector cells by identifying cell populations that have cell surface antigens. $CD4^+$ lymphocytes can be obtained by standard methods. In some embodiments, naive $CD4^+$ T lymphocytes are CD45RO-, $CD45RA^+$, $CD62L^+$, $CD4^+$ T cells. In some embodiments, central memory $CD4^+$ cells are $CD62L^+$ and $CD45RO^+$. In some embodiments, effector $CD4^+$ cells are CD62L- and CD45RO-.

[0488] In one example, to enrich for $CD4^+$ cells by negative selection, a monoclonal antibody cocktail typically includes antibodies to CD14, CD20, CD11b, CD16, HLA-DR, and CD8. In some embodiments, the antibody or binding partner is bound to a solid support or matrix, such as a magnetic bead or paramagnetic bead, to allow for separation of cells for positive and/or negative selection. For example, in some embodiments, the cells and cell populations are separated or isolated using immunomagnetic (or affinitymagnetic) separation techniques (reviewed in Methods in Molecular Medicine, vol. 58: Metastasis Research Protocols, Vol. 2: Cell Behavior *In vitro* and *In vivo*, p 17-25 Edited by: S. A. Brooks and U. Schumacher © Humana Press Inc., Totowa, NJ).

[0489] In some aspects, the sample or composition of cells to be separated is incubated with small, magnetizable or magnetically responsive material, such as magnetically responsive particles or microparticles, such as paramagnetic beads (*e.g.*, such as Dynalbeads or MACS beads). The magnetically responsive material, *e.g.*, particle, generally is directly or indirectly attached to a binding partner, *e.g.*, an antibody, that specifically binds to a molecule, *e.g.*, surface marker, present on the cell, cells, or population of cells that it is desired to separate, *e.g.*, that it is desired to negatively or positively select.

[0490] In some embodiments, the magnetic particle or bead comprises a magnetically responsive material bound to a specific binding member, such as an antibody or other binding partner. There are many well-known magnetically responsive materials used in magnetic separation methods. Suitable magnetic particles include those described in Molday, U.S. Pat. No. 4,452,773, and in European Patent Specification EP 452342 B, which are hereby

incorporated by reference. Colloidal sized particles, such as those described in Owen U.S. Pat. No. 4,795,698, and Liberti *et al.*, U.S. Pat. No. 5,200,084 are other examples.

[0491] The incubation generally is carried out under conditions whereby the antibodies or binding partners, or molecules, such as secondary antibodies or other reagents, which specifically bind to such antibodies or binding partners, which are attached to the magnetic particle or bead, specifically bind to cell surface molecules if present on cells within the sample.

[0492] In some aspects, the sample is placed in a magnetic field, and those cells having magnetically responsive or magnetizable particles attached thereto will be attracted to the magnet and separated from the unlabeled cells. For positive selection, cells that are attracted to the magnet are retained; for negative selection, cells that are not attracted (unlabeled cells) are retained. In some aspects, a combination of positive and negative selection is performed during the same selection step, where the positive and negative fractions are retained and further processed or subject to further separation steps.

[0493] In certain embodiments, the magnetically responsive particles are coated in primary antibodies or other binding partners, secondary antibodies, lectins, enzymes, or streptavidin. In certain embodiments, the magnetic particles are attached to cells via a coating of primary antibodies specific for one or more markers. In certain embodiments, the cells, rather than the beads, are labeled with a primary antibody or binding partner, and then cell-type specific secondary antibody- or other binding partner (*e.g.*, streptavidin)-coated magnetic particles, are added. In certain embodiments, streptavidin-coated magnetic particles are used in conjunction with biotinylated primary or secondary antibodies.

[0494] In some embodiments, the magnetically responsive particles are left attached to the cells that are to be subsequently incubated, cultured and/or engineered; in some aspects, the particles are left attached to the cells for administration to a patient. In some embodiments, the magnetizable or magnetically responsive particles are removed from the cells. Methods for removing magnetizable particles from cells are known and include, *e.g.*, the use of competing non-labeled antibodies, and magnetizable particles or antibodies conjugated to cleavable linkers. In some embodiments, the magnetizable particles are biodegradable.

[0495] In some embodiments, the affinity-based selection is via magnetic-activated cell sorting (MACS) (Miltenyi Biotec, Auburn, CA). Magnetic Activated Cell Sorting (MACS) systems are capable of high-purity selection of cells having magnetized particles attached

thereto. In certain embodiments, MACS operates in a mode wherein the non-target and target species are sequentially eluted after the application of the external magnetic field. That is, the cells attached to magnetized particles are held in place while the unattached species are eluted. Then, after this first elution step is completed, the species that were trapped in the magnetic field and were prevented from being eluted are freed in some manner such that they can be eluted and recovered. In certain embodiments, the non-target cells are labelled and depleted from the heterogeneous population of cells.

[0496] In certain embodiments, the isolation or separation is carried out using a system, device, or apparatus that carries out one or more of the isolation, cell preparation, separation, processing, incubation, culture, and/or formulation steps of the methods. In some aspects, the system is used to carry out each of these steps in a closed or sterile environment, for example, to minimize error, user handling and/or contamination. In one example, the system is a system as described in PCT Pub. Number WO2009/072003, or US 20110003380 A1.

[0497] In some embodiments, the system or apparatus carries out one or more, *e.g.*, all, of the isolation, processing, engineering, and formulation steps in an integrated or self-contained system, and/or in an automated or programmable fashion. In some aspects, the system or apparatus includes a computer and/or computer program in communication with the system or apparatus, which allows a user to program, control, assess the outcome of, and/or adjust various aspects of the processing, isolation, engineering, and formulation steps.

[0498] In some aspects, the separation and/or other steps is carried out using CliniMACS system (Miltenyi Biotec), for example, for automated separation of cells on a clinical-scale level in a closed and sterile system. Components can include an integrated microcomputer, magnetic separation unit, peristaltic pump, and various pinch valves. The integrated computer in some aspects controls all components of the instrument and directs the system to perform repeated procedures in a standardized sequence. The magnetic separation unit in some aspects includes a movable permanent magnet and a holder for the selection column. The peristaltic pump controls the flow rate throughout the tubing set and, together with the pinch valves, ensures the controlled flow of buffer through the system and continual suspension of cells.

[0499] The CliniMACS system in some aspects uses antibody-coupled magnetizable particles that are supplied in a sterile, non-pyrogenic solution. In some embodiments, after labelling of cells with magnetic particles the cells are washed to remove excess particles. A cell

preparation bag is then connected to the tubing set, which in turn is connected to a bag containing buffer and a cell collection bag. The tubing set consists of pre-assembled sterile tubing, including a pre-column and a separation column, and are for single use only. After initiation of the separation program, the system automatically applies the cell sample onto the separation column. Labelled cells are retained within the column, while unlabeled cells are removed by a series of washing steps. In some embodiments, the cell populations for use with the methods described herein are unlabeled and are not retained in the column. In some embodiments, the cell populations for use with the methods described herein are labeled and are retained in the column. In some embodiments, the cell populations for use with the methods described herein are eluted from the column after removal of the magnetic field, and are collected within the cell collection bag.

[0500] In certain embodiments, separation and/or other steps are carried out using the CliniMACS Prodigy system (Miltenyi Biotec). The CliniMACS Prodigy system in some aspects is equipped with a cell processing unity that permits automated washing and fractionation of cells by centrifugation. The CliniMACS Prodigy system can also include an onboard camera and image recognition software that determines the optimal cell fractionation endpoint by discerning the macroscopic layers of the source cell product. For example, peripheral blood is automatically separated into erythrocytes, white blood cells and plasma layers. The CliniMACS Prodigy system can also include an integrated cell cultivation chamber which accomplishes cell culture protocols such as, *e.g.*, cell differentiation and expansion, antigen loading, and long-term cell culture. Input ports can allow for the sterile removal and replenishment of media and cells can be monitored using an integrated microscope. See, *e.g.*, Klebanoff *et al.*, *J Immunother.* 35(9): 651–660 (2012), Terakura *et al.*, *Blood.*1:72–82 (2012), and Wang *et al.*, *J Immunother.* 35(9):689-701 (2012).

[0501] In some embodiments, a cell population described herein is collected and enriched (or depleted) via flow cytometry, in which cells stained for multiple cell surface markers are carried in a fluidic stream. In some embodiments, a cell population described herein is collected and enriched (or depleted) via preparative scale (FACS)-sorting. In certain embodiments, a cell population described herein is collected and enriched (or depleted) by use of microelectromechanical systems (MEMS) chips in combination with a FACS-based detection system (see, *e.g.*, WO 2010/033140, Cho *et al.*, *Lab Chip* 10, 1567-1573 (2010); and Godin *et*

al., *J Biophoton.* 1(5):355–376 (2008). In both cases, cells can be labeled with multiple markers, allowing for the isolation of well-defined T cell subsets at high purity.

[0502] In some embodiments, the antibodies or binding partners are labeled with one or more detectable marker, to facilitate separation for positive and/or negative selection. For example, separation may be based on binding to fluorescently labeled antibodies. In some examples, separation of cells based on binding of antibodies or other binding partners specific for one or more cell surface markers are carried in a fluidic stream, such as by fluorescence-activated cell sorting (FACS), including preparative scale (FACS) and/or microelectromechanical systems (MEMS) chips, *e.g.*, in combination with a flow-cytometric detection system. Such methods allow for positive and negative selection based on multiple markers simultaneously.

[0503] In some embodiments, the preparation methods include steps for freezing, *e.g.*, cryopreserving, the cells, either before or after isolation, incubation, and/or engineering. In some embodiments, the freeze and subsequent thaw step removes granulocytes and, to some extent, monocytes in the cell population. In some embodiments, the cells are suspended in a freezing solution, *e.g.*, following a washing step to remove plasma and platelets. Any of a variety of known freezing solutions and parameters in some aspects may be used. One example involves using PBS containing 20% DMSO and 8% human serum albumin (HSA), or other suitable cell freezing media. This is then diluted 1:1 with media so that the final concentration of DMSO and HSA are 10% and 4%, respectively. The cells are generally then frozen to −80° C. at a rate of 1° per minute and stored in the vapor phase of a liquid nitrogen storage tank.

[0504] In some embodiments, the cells are incubated and/or cultured prior to or in connection with genetic engineering. The incubation steps can include culture, cultivation, stimulation, activation, and/or propagation. The incubation and/or engineering may be carried out in a culture vessel, such as a unit, chamber, well, column, tube, tubing set, valve, vial, culture dish, bag, or other container for culture or cultivating cells. In some embodiments, the compositions or cells are incubated in the presence of stimulating conditions or a stimulatory agent. Such conditions include those designed to induce proliferation, expansion, activation, and/or survival of cells in the population, to mimic antigen exposure, and/or to prime the cells for genetic engineering, such as for the introduction of a recombinant antigen receptor.

[0505] The conditions can include one or more of particular media, temperature, oxygen content, carbon dioxide content, time, agents, *e.g.*, nutrients, amino acids, antibiotics, ions, and/or stimulatory factors, such as cytokines, chemokines, antigens, binding partners, fusion proteins, recombinant soluble receptors, and any other agents designed to activate the cells.

[0506] In some embodiments, the stimulating conditions or agents include one or more agent, *e.g.*, ligand, which is capable of activating an intracellular signaling domain of a TCR complex. In some aspects, the agent turns on or initiates TCR/CD3 intracellular signaling cascade in a T cell. Such agents can include antibodies, such as those specific for a TCR component and/or costimulatory receptor, *e.g.*, anti-CD3, anti-CD28, for example, bound to solid support such as a bead, and/or one or more cytokines. Optionally, the expansion method may further comprise the step of adding anti-CD3 and/or anti CD28 antibody to the culture medium (*e.g.*, at a concentration of at least about 0.5 ng/ml). In some embodiments, the stimulating agents include IL-2 and/or IL-15, for example, an IL-2 concentration of at least about 10 units/mL.

[0507] In some aspects, incubation is carried out in accordance with techniques such as those described in US Patent No. 6,040,1 77 to Riddell *et al.*, Klebanoff *et al.*, *J Immunother.* 35(9): 651–660 (2012), Terakura *et al.*, *Blood.*1:72–82 (2012), and/or Wang *et al.*, *J Immunother.* 35(9):689-701 (2012).

[0508] In some embodiments, the T cells are expanded by adding to a culture-initiating composition feeder cells, such as non-dividing peripheral blood mononuclear cells (PBMC), (*e.g.*, such that the resulting population of cells contains at least about 5, 10, 20, or 40 or more PBMC feeder cells for each T lymphocyte in the initial population to be expanded); and incubating the culture (*e.g.* for a time sufficient to expand the numbers of T cells). In some aspects, the non-dividing feeder cells can comprise gamma-irradiated PBMC feeder cells. In some embodiments, the PBMC are irradiated with gamma rays in the range of about 3000 to 3600 rads to prevent cell division. In some aspects, the feeder cells are added to culture medium prior to the addition of the populations of T cells.

[0509] In some embodiments, the stimulating conditions include temperature suitable for the growth of human T lymphocytes, for example, at least about 25 degrees Celsius, generally at least about 30 degrees, and generally at or about 37 degrees Celsius. Optionally, the incubation may further comprise adding non-dividing EBV-transformed lymphoblastoid cells (LCL) as

feeder cells. LCL can be irradiated with gamma rays in the range of about 6000 to 10,000 rads. The LCL feeder cells in some aspects is provided in any suitable amount, such as a ratio of LCL feeder cells to initial T lymphocytes of at least about 10:1.

[0510] In embodiments, antigen-specific T cells, such as antigen-specific $CD4^+$ and/or $CD8^+$ T cells, are obtained by stimulating naive or antigen specific T lymphocytes with antigen. For example, antigen-specific T cell lines or clones can be generated to cytomegalovirus antigens by isolating T cells from infected subjects and stimulating the cells *in vitro* with the same antigen.

C. NUCLEIC ACIDS, VECTORS AND METHODS FOR GENETIC ENGINEERING

[0511] In some embodiments, the cells, e.g., T cells, are genetically engineered to express a recombinant receptor. In some embodiments, the engineering is carried out by introducing nucleic acid molecules that encode the recombinant receptor. Also provided are nucleic acid molecules encoding a recombinant receptor, and vectors or constructs containing such nucleic acids and/or nucleic acid molecules.

[0512] In some cases, the nucleic acid sequence encoding the recombinant receptor, e.g., chimeric antigen receptor (CAR), contains a signal sequence that encodes a signal peptide. In some aspects, the signal sequence may encode a signal peptide derived from a native polypeptide. In other aspects, the signal sequence may encode a heterologous or non-native signal peptide. In some embodiments, the signal peptide is derived from a transmembrane protein. In some examples the signal peptide is derived from CD8a, CD33, or an IgG. Non-limiting exemplary examples of signal peptides include, for example, the CD33 signal peptide set forth in SEQ ID NO:153, CD8a signal peptide set forth in SEQ ID NO:39, or the signal peptide set forth in SEQ ID NO:40 or modified variant thereof.

[0513] In some embodiments, the nucleic acid molecule encoding the recombinant receptor contains at least one promoter that is operatively linked to control expression of the recombinant receptor. In some examples, the nucleic acid molecule contains two, three, or more promoters operatively linked to control expression of the recombinant receptor. In some embodiments, nucleic acid molecule can contain regulatory sequences, such as transcription and translation initiation and termination codons, which are specific to the type of host (e.g., bacterium, fungus, plant, or animal) into which the nucleic acid molecule is to be introduced, as appropriate and taking into consideration whether the nucleic acid molecule is DNA- or RNA-based. In some

embodiments, the nucleic acid molecule can contain regulatory/control elements, such as a promoter, an enhancer, an intron, a polyadenylation signal, a Kozak consensus sequence, and splice acceptor or donor. In some embodiments, the nucleic acid molecule can contain a nonnative promoter operably linked to the nucleotide sequence encoding the recombinant receptor and/or one or more additional polypeptide(s). In some embodiments, the promoter is selected from among an RNA pol I, pol II or pol III promoter. In some embodiments, the promoter is recognized by RNA polymerase II (e.g., a CMV, SV40 early region or adenovirus major late promoter). In another embodiment, the promoter is recognized by RNA polymerase III (e.g., a U6 or H1 promoter). In some embodiments, the promoter can be a non-viral promoter or a viral promoter, such as a cytomegalovirus (CMV) promoter, an SV40 promoter, an RSV promoter, and a promoter found in the long-terminal repeat of the murine stem cell virus. Other known promoters also are contemplated.

[0514] In some embodiments, the promoter is or comprises a constitutive promoter. Exemplary constitutive promoters include, e.g., simian virus 40 early promoter (SV40), cytomegalovirus immediate-early promoter (CMV), human Ubiquitin C promoter (UBC), human elongation factor 1α promoter (EF1α), mouse phosphoglycerate kinase 1 promoter (PGK), and chicken β-Actin promoter coupled with CMV early enhancer (CAGG). In some embodiments, the constitutive promoter is a synthetic or modified promoter. In some embodiments, the promoter is or comprises an MND promoter, a synthetic promoter that contains the U3 region of a modified MoMuLV LTR with myeloproliferative sarcoma virus enhancer (see Challita et al. (1995) J. Virol. 69(2):748-755). In some embodiments, the promoter is a tissue-specific promoter. In another embodiment, the promoter is a viral promoter. In another embodiment, the promoter is a non-viral promoter.

[0515] In another embodiment, the promoter is a regulated promoter (e.g., inducible promoter). In some embodiments, the promoter is an inducible promoter or a repressible promoter. In some embodiments, the promoter comprises a Lac operator sequence, a tetracycline operator sequence, a galactose operator sequence or a doxycycline operator sequence, or is an analog thereof or is capable of being bound by or recognized by a Lac repressor or a tetracycline repressor, or an analog thereof. In some embodiments, the nucleic acid molecule does not include a regulatory element, e.g. promoter.

[0516] In some embodiments, the nucleic acid molecule encoding the recombinant receptor, e.g., CAR or other antigen receptor, further includes nucleic acid sequences encoding a marker and/or cells expressing the CAR or other antigen receptor further includes a marker, e.g., a surrogate marker, such as a cell surface marker, which may be used to confirm transduction or engineering of the cell to express the receptor, such as a truncated version of a cell surface receptor, such as truncated EGFR (tEGFR). In some embodiments, the one or more marker(s) is a transduction marker, surrogate marker and/or a selection marker.

[0517] In some embodiments, the marker is a transduction marker or a surrogate marker. A transduction marker or a surrogate marker can be used to detect cells that have been introduced with the nucleic acid molecule, e.g., a nucleic acid molecule encoding a recombinant receptor. In some embodiments, the transduction marker can indicate or confirm modification of a cell. In some embodiments, the surrogate marker is a protein that is made to be co-expressed on the cell surface with the recombinant receptor, e.g. CAR. In particular embodiments, such a surrogate marker is a surface protein that has been modified to have little or no activity. In certain embodiments, the surrogate marker is encoded on the same nucleic acid molecule that encodes the recombinant receptor. In some embodiments, the nucleic acid sequence encoding the recombinant receptor is operably linked to a nucleic acid sequence encoding a marker, optionally separated by an internal ribosome entry site (IRES), or a nucleic acid encoding a self-cleaving peptide or a peptide that causes ribosome skipping, such as a 2A sequence, such as a T2A (*e.g.*, SEQ ID NO: 6 or 167), a P2A (*e.g.*, SEQ ID NO: 168 or 169), an E2A (*e.g.*, SEQ ID NO: 170) or an F2A (*e.g.*, SEQ ID NO: 172). Extrinsic marker genes may in some cases be utilized in connection with engineered cell to permit detection or selection of cells and, in some cases, also to promote cell suicide.

[0518] Exemplary surrogate markers can include truncated forms of cell surface polypeptides, such as truncated forms that are non-functional and to not transduce or are not capable of transducing a signal or a signal ordinarily transduced by the full-length form of the cell surface polypeptide, and/or do not or are not capable of internalizing. Exemplary truncated cell surface polypeptides including truncated forms of growth factors or other receptors such as a truncated human epidermal growth factor receptor 2 (tHER2), a truncated epidermal growth factor receptor (tEGFR, exemplary tEGFR sequence set forth in SEQ ID NO:7 or 166) or a prostate-specific membrane antigen (PSMA) or modified form thereof. tEGFR may contain an

epitope recognized by the antibody cetuximab (Erbitux®) or other therapeutic anti-EGFR antibody or binding molecule, which can be used to identify or select cells that have been engineered with the tEGFR construct and an encoded exogenous protein, and/or to eliminate or separate cells expressing the encoded exogenous protein. See U.S. Patent No. 8,802,374 and Liu et al., Nature Biotech. 2016 April; 34(4): 430–434). In some aspects, the marker, *e.g.* surrogate marker, includes all or part (*e.g.*, truncated form) of CD34, a NGFR, a CD19 or a truncated CD19, e.g., a truncated non-human CD19, or epidermal growth factor receptor (*e.g.*, tEGFR). In some embodiments, the marker is or comprises a fluorescent protein, such as green fluorescent protein (GFP), enhanced green fluorescent protein (EGFP), such as super-fold GFP (sfGFP), red fluorescent protein (RFP), such as tdTomato, mCherry, mStrawberry, AsRed2, DsRed or DsRed2, cyan fluorescent protein (CFP), blue green fluorescent protein (BFP), enhanced blue fluorescent protein (EBFP), and yellow fluorescent protein (YFP), and variants thereof, including species variants, monomeric variants, and codon-optimized and/or enhanced variants of the fluorescent proteins. In some embodiments, the marker is or comprises an enzyme, such as a luciferase, the lacZ gene from *E. coli*, alkaline phosphatase, secreted embryonic alkaline phosphatase (SEAP), chloramphenicol acetyl transferase (CAT). Exemplary light-emitting reporter genes include luciferase (luc), β-galactosidase, chloramphenicol acetyltransferase (CAT), β-glucuronidase (GUS) or variants thereof.

[0519] In some embodiments, the marker is a selection marker. In some embodiments, the selection marker is or comprises a polypeptide that confers resistance to exogenous agents or drugs. In some embodiments, the selection marker is an antibiotic resistance gene. In some embodiments, the selection marker is an antibiotic resistance gene confers antibiotic resistance to a mammalian cell. In some embodiments, the selection marker is or comprises a Puromycin resistance gene, a Hygromycin resistance gene, a Blasticidin resistance gene, a Neomycin resistance gene, a Geneticin resistance gene or a Zeocin resistance gene or a modified form thereof.

[0520] In some aspects, the marker, *e.g.* surrogate marker, includes all or part (*e.g.*, truncated form) of CD34, a NGFR, or epidermal growth factor receptor (*e.g.*, tEGFR). In some embodiments, the nucleic acid encoding the marker is operably linked to a polynucleotide encoding for a linker sequence, such as a cleavable linker sequence, *e.g.*, T2A. For example, a marker, and optionally a linker sequence, can be any as disclosed in PCT Pub. No.

WO2014031687. For example, the marker can be a truncated EGFR (tEGFR) that is, optionally, linked to a linker sequence, such as a T2A cleavable linker sequence. An exemplary polypeptide for a truncated EGFR (*e.g.* tEGFR) comprises the sequence of amino acids set forth in SEQ ID NO: 7 or 166, or a sequence of amino acids that exhibits at least 85%, 86%, 87%, 88%, 89%, 90%, 91%, 92%, 93%, 94%, 95%, 96%, 97%, 98%, 99% or more sequence identity to SEQ ID NO: 7 or 166. An exemplary T2A linker sequence comprises the sequence of amino acids set forth in SEQ ID NO: 6 or 167 or a sequence of amino acids that exhibits at least 85%, 86%, 87%, 88%, 89%, 90%, 91%, 92%, 93%, 94%, 95%, 96%, 97%, 98%, 99% or more sequence identity to SEQ ID NO: 6 or 167.

[0521] In some embodiments, the marker is a molecule, *e.g.*, cell surface protein, not naturally found on T cells or not naturally found on the surface of T cells, or a portion thereof. In some embodiments, the molecule is a non-self molecule, *e.g.*, non-self protein, *i.e.*, one that is not recognized as "self" by the immune system of the host into which the cells will be adoptively transferred.

[0522] In some embodiments, the marker serves no therapeutic function and/or produces no effect other than to be used as a marker for genetic engineering, *e.g.*, for selecting cells successfully engineered. In other embodiments, the marker may be a therapeutic molecule or molecule otherwise exerting some desired effect, such as a ligand for a cell to be encountered *in vivo*, such as a costimulatory or immune checkpoint molecule to enhance and/or dampen responses of the cells upon adoptive transfer and encounter with ligand.

[0523] In some embodiments, a single promoter may direct expression of an RNA that contains, in a single open reading frame (ORF), two or three genes separated from one another by sequences encoding a self-cleavage peptide (*e.g.*, 2A sequences) or a protease recognition site (*e.g.*, furin). The ORF thus encodes a single polypeptide, which, either during (in the case of 2A) or after translation, is processed into the individual proteins. In some cases, the peptide, such as T2A, can cause the ribosome to skip (ribosome skipping) synthesis of a peptide bond at the C-terminus of a 2A element, leading to separation between the end of the 2A sequence and the next peptide downstream (*see*, for example, de Felipe. *Genetic Vaccines and Ther.* 2:13 (2004) and deFelipe *et al. Traffic* 5:616-626 (2004)). Many 2A elements are known in the art. Examples of 2A sequences that can be used in the methods and nucleic acids disclosed herein, without limitation, 2A sequences from the foot-and-mouth disease virus (F2A), equine rhinitis A

virus (E2A), Thosea asigna virus (T2A, *e.g.*, SEQ ID NO: 6), and porcine teschovirus-1 (P2A) as described in U.S. Patent Publication No. 20070116690.

[0524] Introduction of the nucleic acid molecules encoding the recombinant receptor in the cell may be carried out using any of a number of known vectors. Such vectors include viral and non-viral systems, including lentiviral and gammaretroviral systems, as well as transposon-based systems such as PiggyBac or Sleeping Beauty-based gene transfer systems. Exemplary methods include those for transfer of nucleic acids encoding the receptors, including via viral, *e.g.*, retroviral or lentiviral, transduction, transposons, and electroporation.

[0525] In some embodiments, gene transfer is accomplished by first stimulating the cell, such as by combining it with a stimulus that induces a response such as proliferation, survival, and/or activation, e.g., as measured by expression of a cytokine or activation marker, followed by transduction of the activated cells, and expansion in culture to numbers sufficient for clinical applications.

[0526] In some embodiments, prior to or during gene transfer, the cells are incubated or cultured in the presence of an gamma secretase inhibitor, including any as described herein. In some embodiments, the gamma secretase inhibitor is added during the cell manufacturing process, for example, during the process of engineering CAR-T cells. In some aspects, the presence of the gamma secretase inhibitor can improve the quality of the population of cells produced. In some aspects, the gamma secretase inhibitor may increase the proliferation or expansion of cells or may alter one or more signaling pathways thereby resulting in cells with a less-differentiated or less activated surface phenotype, despite exhibiting substantial expansion and/or effector function.

[0527] In some contexts, overexpression of a stimulatory factor (for example, a lymphokine or a cytokine) may be toxic to a subject. Thus, in some contexts, the engineered cells include gene segments that cause the cells to be susceptible to negative selection in vivo, such as upon administration in adoptive immunotherapy. For example in some aspects, the cells are engineered so that they can be eliminated as a result of a change in the *in vivo* condition of the patient to which they are administered. The negative selectable phenotype may result from the insertion of a gene that confers sensitivity to an administered agent, for example, a compound. Negative selectable genes include the Herpes simplex virus type I thymidine kinase (HSV-I TK) gene (Wigler et al., Cell II :223, I977) which confers ganciclovir sensitivity; the cellular

hypoxanthine phosphribosyltransferase (HPRT) gene, the cellular adenine phosphoribosyltransferase (APRT) gene, bacterial cytosine deaminase, (Mullen et al., Proc. Natl. Acad. Sci. USA. 89:33 (1992)).

[0528] In some embodiments, recombinant nucleic acids are transferred into cells using recombinant infectious virus particles, such as, e.g., vectors derived from simian virus 40 (SV40), adenoviruses, adeno-associated virus (AAV). In some embodiments, recombinant nucleic acids are transferred into T cells using recombinant lentiviral vectors or retroviral vectors, such as gamma-retroviral vectors (see, e.g., Koste et al. (2014) Gene Therapy 2014 Apr 3. doi: 10.1038/gt.2014.25; Carlens et al. (2000) Exp Hematol 28(10): 1137-46; Alonso-Camino et al. (2013) Mol Ther Nucl Acids 2, e93; Park et al., Trends Biotechnol. 2011 November 29(11): 550–557.

[0529] In some embodiments, the retroviral vector has a long terminal repeat sequence (LTR), *e.g*., a retroviral vector derived from the Moloney murine leukemia virus (MoMLV), myeloproliferative sarcoma virus (MPSV), murine embryonic stem cell virus (MESV), murine stem cell virus (MSCV), spleen focus forming virus (SFFV), or adeno-associated virus (AAV). Most retroviral vectors are derived from murine retroviruses. In some embodiments, the retroviruses include those derived from any avian or mammalian cell source. The retroviruses typically are amphotropic, meaning that they are capable of infecting host cells of several species, including humans. In one embodiment, the gene to be expressed replaces the retroviral gag, pol and/or env sequences. A number of illustrative retroviral systems have been described (e.g., U.S. Pat. Nos. 5,219,740; 6,207,453; 5,219,740; Miller and Rosman (1989) BioTechniques 7:980-990; Miller, A. D. (1990) Human Gene Therapy 1:5-14; Scarpa et al. (1991) Virology 180:849-852; Burns et al. (1993) Proc. Natl. Acad. Sci. USA 90:8033-8037; and Boris-Lawrie and Temin (1993) Cur. Opin. Genet. Develop. 3:102-109.

[0530] Methods of lentiviral transduction are known. Exemplary methods are described in, e.g., Wang et al. (2012) *J. Immunother.* 35(9): 689-701; Cooper et al. (2003) *Blood.* 101:1637–1644; Verhoeyen et al. (2009) *Methods Mol Biol.* 506: 97-114; and Cavalieri et al. (2003) Blood. 102(2): 497-505.

[0531] In some embodiments, recombinant nucleic acids are transferred into T cells via electroporation (*see*, e.g., Chicaybam et al, (2013) *PLoS ONE* 8(3): e60298 and Van Tedeloo et al. (2000) *Gene Therapy* 7(16): 1431-1437). In some embodiments, recombinant nucleic acids

are transferred into T cells via transposition (see, e.g., Manuri et al. (2010) Hum Gene Ther 21(4): 427-437; Sharma et al. (2013) *Molec Ther Nucl Acids* 2, e74; and Huang et al. (2009) *Methods Mol Biol* 506: 115-126). Other methods of introducing and expressing genetic material in immune cells include calcium phosphate transfection (e.g., as described in Current Protocols in Molecular Biology, John Wiley & Sons, New York. N.Y.), protoplast fusion, cationic liposome-mediated transfection; tungsten particle-facilitated microparticle bombardment (Johnston, Nature, 346: 776-777 (1990)); and strontium phosphate DNA co-precipitation (Brash et al., Mol. Cell Biol., 7: 2031-2034 (1987)).

[0532] Other approaches and vectors for transfer of the nucleic acids encoding the recombinant products are those described, e.g., in international patent application, Publication No.: WO2014055668, and U.S. Patent No. 7,446,190.

[0533] In some embodiments, the cells, e.g., T cells, may be transfected either during or after expansion e.g. with a T cell receptor (TCR) or a chimeric antigen receptor (CAR). This transfection for the introduction of the gene of the desired receptor can be carried out with any suitable retroviral vector, for example. The genetically modified cell population can then be liberated from the initial stimulus (the CD3/CD28 stimulus, for example) and subsequently be stimulated with a second type of stimulus e.g. via a de novo introduced receptor). This second type of stimulus may include an antigenic stimulus in form of a peptide/MHC molecule, the cognate (cross-linking) ligand of the genetically introduced receptor (e.g. natural ligand of a CAR) or any ligand (such as an antibody) that directly binds within the framework of the new receptor (e.g. by recognizing constant regions within the receptor). See, for example, Cheadle et al, "Chimeric antigen receptors for T-cell based therapy" Methods Mol Biol. 2012; 907:645-66 or Barrett et al., Chimeric Antigen Receptor Therapy for Cancer Annual Review of Medicine Vol. 65: 333-347 (2014).

[0534] In some cases, a vector may be used that does not require that the cells, *e.g.*, T cells, are activated. In some such instances, the cells may be selected and/or transduced prior to activation. Thus, the cells may be engineered prior to, or subsequent to culturing of the cells, and in some cases at the same time as or during at least a portion of the culturing.

[0535] In some aspects, the cells further are engineered to promote expression of cytokines or other factors. Among additional nucleic acids, e.g., genes for introduction are those to improve the efficacy of therapy, such as by promoting viability and/or function of transferred

cells; genes to provide a genetic marker for selection and/or evaluation of the cells, such as to assess *in vivo* survival or localization; genes to improve safety, for example, by making the cell susceptible to negative selection in vivo as described by Lupton S. D. et al., *Mol. and Cell Biol.,* 11:6 (1991); and Riddell et al., *Human Gene Therapy* 3:319-338 (1992); see also the publications of PCT/US91/08442 and PCT/US94/05601 by Lupton et al. describing the use of bifunctional selectable fusion genes derived from fusing a dominant positive selectable marker with a negative selectable marker. *See,* e.g.*,* Riddell et al., US Patent No. 6,040,177, at columns 14-17.

III. EXEMPLARY TREATMENT OUTCOMES AND METHODS FOR ASSESSING SAME

[0536] In some embodiments of the methods, compositions, combinations, kits and uses provided herein, the provided combination therapy results in one or more treatment outcomes, such as a feature associated with any one or more of the parameters associated with the therapy or treatment, as described herein. In some embodiments, the method includes assessment of the exposure, persistence and proliferation of the T cells, *e.g.*, T cells administered for the T cell based therapy. In some embodiments, the exposure, or prolonged expansion and/or persistence of the cells, and/or changes in cell phenotypes or functional activity of the cells, *e.g.*, cells administered for immunotherapy, *e.g.* T cell therapy, in the methods provided herein, can be measured by assessing the characteristics of the T cells *in vitro* or *ex vivo*. In some embodiments, such assays can be used to determine or confirm the function of the T cells, *e.g.* T cell therapy, before or after administering the combination therapy provided herein.

[0537] In some embodiments, the combination therapy can further include one or more screening steps to identify subjects for treatment with the combination therapy and/or continuing the combination therapy, and/or a step for assessment of treatment outcomes and/or monitoring treatment outcomes. In some embodiments, subjects suitable for treatment have low surface expression of BCMA. In some embodiments, low surface expression of BCMA is determined by comparing BCMA expression in a subject against average surface BCMA expression in a healthy population of subjects. In some embodiments, low BCMA expression is a BCMA expression level that is less than the population average. In some embodiments, low BCMA expression is a BCMA expression level that is, is about, is at least 1.1, 1.2, 1.3, 1.4, 1.5, 1.6, 1.7, 1.8, 1.9, 2, 3, 4, 5, 10, 15, 20, 30, 40, 50, 100, 150, 200, 250-fold or more less than, or 10%,

20%, 30%, 40%, 50%, 60%, 70%, 80%, 90%, or more below, the average BCMA expression level in a healthy population of subjects. In some embodiments, low surface expression of BCMA is determined by comparing BCMA expression in a subject against median surface BCMA expression in a healthy population of subjects. In some embodiments, low BCMA expression is a BCMA expression level that is less than the population median. In some embodiments, low BCMA expression is a BCMA expression level that is, is about, is at least 1.1, 1.2, 1.3, 1.4, 1.5, 1.6, 1.7, 1.8, 1.9, 2, 3, 4, 5, 10, 15, 20, 30, 40, 50, 100, 150, 200, 250-fold or more less than, or 10%, 20%, 30%, 40%, 50%, 60%, 70%, 80%, 90%, or more below, the median BCMA expression level in a healthy population of subjects. In some embodiments, low surface expression of BCMA is determined by comparing BCMA expression in a subject against average surface BCMA expression in a patient population (e.g., multiple myeloma pateient population). In some embodiments, low BCMA expression is a BCMA expression level that is less than the patient population average. In some embodiments, low BCMA expression is a BCMA expression level that is, is about, is at least 1.1, 1.2, 1.3, 1.4, 1.5, 1.6, 1.7, 1.8, 1.9, 2, 3, 4, 5, 10, 15, 20, 30, 40, 50, 100, 150, 200, 250-fold or more less than, or 10%, 20%, 30%, 40%, 50%, 60%, 70%, 80%, 90%, or more below,the average BCMA expression level in a patient population (e.g., multiple myeloma pateient population). In some embodiments, low surface expression of BCMA is determined by comparing BCMA expression in a subject against median surface BCMA expression in a patient population (e.g., multiple myeloma pateient population). In some embodiments, low BCMA expression is a BCMA expression level that is less than the patient population median. In some embodiments, low BCMA expression is a BCMA expression level that is, is about, is at least 1.1, 1.2, 1.3, 1.4, 1.5, 1.6, 1.7, 1.8, 1.9, 2, 3, 4, 5, 10, 15, 20, 30, 40, 50, 100, 150, 200, 250-fold or more less than, or 10%, 20%, 30%, 40%, 50%, 60%, 70%, 80%, 90%, or more below, the median BCMA expression level in a patient population (e.g., multiple myeloma patient population). In some embodiments, determination of low BCMA expression in a subject can be determined by comparing BCMA expression in the subject at two or more different time point (e.g., before diagnosis and after diagnosis, at diagnosis and at a time point following diagnosis, before treatment and after treatment, etc.). In some embodiments according to this method, low BCMA expression would be considered about or at least 10%, 20%, 30%, 40%, 50%, 60%, 70%, 80%, 90%, or more decrease in BCMA expression compared to the earlier time point.

[0538] In some embodiments, low BCMA expression is considered BCMA expression below a threshold level. In some embodiments, the threshold level is, is about, is at least, 10%, 20%, 30%, 40%, 50%, 60%, 70%, 80%, 90% or more below the average or median BCMA expression level ascertained from a population of healthy subjects. In some embodiments, the threshold level is, is about, is at least, 10%, 20%, 30%, 40%, 50%, 60%, 70%, 80%, 90% or more below the average or median BCMA expression level ascertained from a population of patients (e.g., multiple myeloma patients). In some embodiments, the threshold level is, is about, is at least, 10%, 20%, 30%, 40%, 50%, 60%, 70%, 80%, 90% or more below a BCMA expression level obtained from the subject.

[0539] In some embodiments, the step for assessment of treatment outcomes can include steps to evaluate and/or to monitor treatment and/or to identify subjects for administration of further or remaining steps of the therapy and/or for repeat therapy. In some embodiments, the screening step and/or assessment of treatment outcomes can be used to determine the dose, frequency, duration, timing and/or order of the combination therapy provided herein.

[0540] In some embodiments, any of the screening steps and/or assessment of treatment of outcomes described herein can be used prior to, during, during the course of, or subsequent to administration of one or more steps of the provided combination therapy, *e.g.*, administration of the T cell therapy (*e.g.* CAR-expressing T cells), and/or an gamma secretase inhibitor. In some embodiments, assessment is made prior to, during, during the course of, or after performing any of the methods provided herein. In some embodiments, the assessment is made prior to performing the methods provided herein. In some embodiments, assessment is made after performing one or more steps of the methods provided herein. In some embodiments, the assessment is performed prior to administration of administration of one or more steps of the provided combination therapy, for example, to screen and identify patients suitable and/or susceptible to receive the combination therapy. In some embodiments, the assessment is performed during, during the course of, or subsequent to administration of one or more steps of the provided combination therapy, for example, to assess the intermediate or final treatment outcome, *e.g.*, to determine the efficacy of the treatment and/or to determine whether to continue or repeat the treatments and/or to determine whether to administer the remaining steps of the combination therapy.

[0541] In some embodiments, treatment of outcomes includes improved immune function, *e.g.*, immune function of the T cells administered for cell based therapy and/or of the endogenous T cells in the body. In some embodiments, exemplary treatment outcomes include, but are not limited to, enhanced T cell proliferation, enhanced T cell functional activity, changes in immune cell phenotypic marker expression, such as such features being associated with the engineered T cells, e.g. CAR-T cells, administered to the subject. In some embodiments, exemplary treatment outcomes include decreased disease burden, *e.g.*, tumor burden, improved clinical outcomes and/or enhanced efficacy of therapy.

[0542] In some embodiments, the screening step and/or assessment of treatment of outcomes includes assessing the survival and/or function of the T cells administered for cell based therapy. In some embodiments, the screening step and/or assessment of treatment of outcomes includes assessing the levels of cytokines or growth factors. In some embodiments, the screening step and/or assessment of treatment of outcomes includes assessing disease burden and/or improvements, *e.g.*, assessing tumor burden and/or clinical outcomes. In some embodiments, either of the screening step and/or assessment of treatment of outcomes can include any of the assessment methods and/or assays described herein and/or known in the art, and can be performed one or more times, *e.g.*, prior to, during, during the course of, or subsequently to administration of one or more steps of the combination therapy. Exemplary sets of parameters associated with a treatment outcome, which can be assessed in some embodiments of the methods provided herein, include peripheral blood immune cell population profile and/or tumor burden.

[0543] In some embodiments, the methods affect efficacy of the cell therapy in the subject. In some embodiments, the persistence, expansion, and/or presence of recombinant receptor-expressing, *e.g.*, CAR-expressing, cells in the subject following administration of the dose of cells in the method with the gamma secretase inhibitor is greater as compared to that achieved via a method without the administration of the gamma secretase inhibitor. In some embodiments of the immunotherapy methods provided herein, such as a T cell therapy (*e.g.* CAR-expressing T cells), assessment of the parameter includes assessing the expansion and/or persistence in the subject of the administered T cells for the immunotherapy, *e.g.*, T cell therapy, as compared to a method in which the immunotherapy is administered to the subject in the absence of the gamma secretase inhibitor. In some embodiments, the methods result in the

administered T cells exhibiting increased or prolonged expansion and/or persistence in the subject as compared to a method in which the T cell therapy is administered to the subject in the absence of the gamma secretase inhibitor.

[0544] In some embodiments, the administration of the gamma secretase inhibitor decreases disease burden, *e.g.*, tumor burden, in the subject as compared to a method in which the dose of cells expressing the recombinant receptor is administered to the subject in the absence of the gamma secretase inhibitor. In some embodiments, the administration of the gamma secretase inhibitor decreases blast marrow in the subject as compared to a method in which the dose of cells expressing the recombinant receptor is administered to the subject in the absence of the gamma secretase inhibitor. In some embodiments, the administration of the gamma secretase inhibitor results in improved clinical outcomes, *e.g.*, objective response rate (ORR), progression-free survival (PFS) and overall survival (OS), compared to a method in which the dose of cells expressing the recombinant receptor is administered to the subject in the absence of the gamma secretase inhibitor.

[0545] In some embodiments, the subject can be screened prior to the administration of one or more steps of the combination therapy. For example, the subject can be screened for characteristics of the disease and/or disease burden, *e.g.*, tumor burden, prior to administration of the combination therapy, to determine suitability, responsiveness and/or susceptibility to administering the combination therapy. In some embodiments, the screening step and/or assessment of treatment outcomes can be used to determine the dose, frequency, duration, timing and/or order of the combination therapy provided herein.

[0546] In some embodiments, the subject can be screened after administration of one of the steps of the combination therapy, to determine and identify subjects to receive the remaining steps of the combination therapy and/or to monitor efficacy of the therapy. In some embodiments, the number, level or amount of administered T cells and/or proliferation and/or activity of the administered T cells is assessed prior to administration and/or after administration of the gamma secretase inhibitor.

[0547] In some embodiments, the gamma secretase inhibitor is administered until the concentration or number of engineered cells in the blood of the subject is (i) at least at or about 10 engineered cells per microliter, (ii) at least 20%, 30%, 40% or 50% of the total number of peripheral blood mononuclear cells (PBMCs), (iii) at least or at least about 1 x 10^5 engineered

cells; or (iv) at least 5,000 copies of recombinant receptor-encoding DNA per micrograms DNA; and/or at day 90 following the initiation of the administration in (a), CAR-expressing cells are detectable in the blood or serum of the subject; and/or at day 90 following the initiation of the administration in (a), the blood of the subject contains at least 20% CAR-expressing cells, at least 10 CAR-expressing cells per microliter or at least 1 x 10^4 CAR-expressing cells.

[0548] In some embodiments, the gamma secretase inhibitor is administered until there is a clinical benefit to the treatment, such as at least or greater than a 50% decrease in the total tumor volume, a complete response (CR) in which detectable tumor has disappeared, progression free survival or disease free survival for greater than 6 months or greater than 1 year or more.

[0549] In some embodiments, a change and/or an alteration, *e.g.*, an increase, an elevation, a decrease or a reduction, in levels, values or measurements of a parameter or outcome compared to the levels, values or measurements of the same parameter or outcome in a different time point of assessment, a different condition, a reference point and/or a different subject is determined or assessed. For example, in some embodiments, a fold change, *e.g.*, an increase or decrease, in particular parameters, e.g., number of engineered T cells in a sample, compared to the same parameter in a different condition, e.g., before or after administration of the gamma secretase inhibitor can be determined. In some embodiments, the levels, values or measurements of two or more parameters are determined, and relative levels are compared. In some embodiments, the determined levels, values or measurements of parameters are compared to the levels, values or measurements from a control sample or an untreated sample. In some embodiments, the determined levels, values or measurements of parameters are compared to the levels from a sample from the same subject but at a different time point. The values obtained in the quantification of individual parameter can be combined for the purpose of disease assessment, *e.g.*, by forming an arithmetical or logical operation on the levels, values or measurements of parameters by using multi-parametric analysis. In some embodiments, a ratio of two or more specific parameters can be calculated.

A. T cell exposure, persistence and proliferation

[0550] In some embodiments, the parameter associated with therapy or a treatment outcome, which include parameters that can be assessed for the screening steps and/or assessment of treatment of outcomes and/or monitoring treatment outcomes, is or includes assessment of the exposure, persistence and proliferation of the T cells, *e.g.*, T cells administered for the T cell

based therapy. In some embodiments, the increased exposure, or prolonged expansion and/or persistence of the cells, and/or changes in cell phenotypes or functional activity of the cells, *e.g.*, cells administered for immunotherapy, *e.g.* T cell therapy, in the methods provided herein, can be measured by assessing the characteristics of the T cells *in vitro* or *ex vivo*. In some embodiments, such assays can be used to determine or confirm the function of the T cells used for the immunotherapy, *e.g.* T cell therapy, before or after administering one or more steps of the combination therapy provided herein.

[0551] In some embodiments, the administration of the gamma secretase inhibitor are designed to promote exposure of the subject to the cells, *e.g.*, T cells administered for T cell based therapy, such as by promoting their expansion and/or persistence over time. In some embodiments, the T cell therapy exhibits increased or prolonged expansion and/or persistence in the subject as compared to a method in which the T cell therapy is administered to the subject in the absence of the gamma secretase inhibitor.

[0552] In some embodiments, the provided methods increase exposure of the subject to the administered cells (*e.g.*, increased number of cells or duration over time) and/or improve efficacy and therapeutic outcomes of the immunotherapy, *e.g.* T cell therapy. In some aspects, the methods are advantageous in that a greater and/or longer degree of exposure to the cells expressing the recombinant receptors, *e.g.*, CAR-expressing cells, improves treatment outcomes as compared with other methods. Such outcomes may include patient survival and remission, even in individuals with severe tumor burden.

[0553] In some embodiments, the administration of the gamma secretase inhibitor can increase the maximum, total, and/or duration of exposure to the cells, *e.g.* T cells administered for the T cell based therapy, in the subject as compared to administration of the T cells alone in the absence of the gamma secretase inhibitor. In some aspects, administration of the gamma secretase inhibitor, in the context of high disease burden (and thus higher amounts of antigen) and/or a more aggressive or resistant cancer enhances efficacy as compared with administration of the T cells alone in the absence of the gamma secretase inhibitor in the same context, which may result in immunosuppression, anergy and/or exhaustion which may prevent expansion and/or persistence of the cells.

[0554] In some embodiments, the presence and/or amount of cells expressing the recombinant receptor (*e.g.*, CAR-expressing cells administered for T cell based therapy) in the

subject following the administration of the T cells and before, during and/or after the administration of the gamma secretase inhibitor is detected. In some aspects, quantitative PCR (qPCR) is used to assess the quantity of cells expressing the recombinant receptor (*e.g.*, CAR-expressing cells administered for T cell based therapy) in the blood or serum or organ or tissue sample (*e.g.*, disease site, *e.g.*, tumor sample) of the subject. In some aspects, persistence is quantified as copies of DNA or plasmid encoding the receptor, *e.g.*, CAR, per microgram of DNA, or as the number of receptor-expressing, *e.g.*, CAR-expressing, cells per microliter of the sample, *e.g.*, of blood or serum, or per total number of peripheral blood mononuclear cells (PBMCs) or white blood cells or T cells per microliter of the sample.

[0555] In some embodiments, the cells are detected in the subject at or at least at 4, 14, 15, 27, or 28 days following the administration of the T cells, *e.g.*, CAR-expressing T cells. In some aspects, the cells are detected at or at least at 2, 4, or 6 weeks following, or 3, 6, 12, 18, 24, 30 or 36 months, or 1, 2, 3, 4, 5, or more years, following the administration of the T cells, *e.g.*, CAR-expressing T cells and/or the gamma secretase inhibitor.

[0556] In some embodiments, the persistence of receptor-expressing cells (*e.g.* CAR-expressing cells) in the subject by the methods, following the administration of the T cells, *e.g.*, CAR-expressing T cells and/or the gamma secretase inhibitor, is greater as compared to that which would be achieved by alternative methods such as those involving the administration of the immunotherapy alone, *e.g.*, administration the T cells, *e.g.*, CAR-expressing T cells, in the absence of the gamma secretase inhibitor.

[0557] The exposure, *e.g.*, number of cells, *e.g.* T cells administered for T cell therapy, indicative of expansion and/or persistence, may be stated in terms of maximum numbers of the cells to which the subject is exposed, duration of detectable cells or cells above a certain number or percentage, area under the curve for number of cells over time, and/or combinations thereof and indicators thereof. Such outcomes may be assessed using known methods, such as qPCR to detect copy number of nucleic acid encoding the recombinant receptor compared to total amount of nucleic acid or DNA in the particular sample, *e.g.*, blood, serum, plasma or tissue, such as a tumor sample, and/or flow cytometric assays detecting cells expressing the receptor generally using antibodies specific for the receptors. Cell-based assays may also be used to detect the number or percentage of functional cells, such as cells capable of binding to and/or neutralizing

and/or inducing responses, *e.g.*, cytotoxic responses, against cells of the disease or condition or expressing the antigen recognized by the receptor.

[0558] In some aspects, increased exposure of the subject to the cells includes increased expansion of the cells. In some embodiments, the receptor expressing cells, *e.g.* CAR-expressing cells, expand in the subject following administration of the T cells, *e.g.*, CAR-expressing T cells, and/or following administration of gamma secretase inhibitor. In some aspects, the methods result in greater expansion of the cells compared with other methods, such as those involving the administration of the T cells, *e.g.*, CAR-expressing T cells, in the absence of administering the gamma secretase inhibitor.

[0559] In some aspects, the method results in high *in vivo* proliferation of the administered cells, for example, as measured by flow cytometry. In some aspects, high peak proportions of the cells are detected. For example, in some embodiments, at a peak or maximum level following the administration of the T cells, *e.g.*, CAR-expressing T cells and/or the gamma secretase inhibitor, in the blood or disease-site of the subject or white blood cell fraction thereof, *e.g.*, PBMC fraction or T cell fraction, at least about 10%, at least about 20%, at least about 30%, at least about 40%, at least about 50%, at least about 60%, at least about 70%, at least about 80%, or at least about 90% of the cells express the recombinant receptor, *e.g.*, the CAR.

[0560] In some embodiments, the method results in a maximum concentration, in the blood or serum or other bodily fluid or organ or tissue of the subject, of at least 100, 500, 1000, 1500, 2000, 5000, 10,000 or 15,000 copies of or nucleic acid encoding the receptor, *e.g.*, the CAR, per microgram of DNA, or at least 0.1, 0.2, 0.3, 0.4, 0.5, 0.6, 0.7, 0.8, or 0.9 receptor-expressing, *e.g.*, CAR,-expressing cells per total number of peripheral blood mononuclear cells (PBMCs), total number of mononuclear cells, total number of T cells, or total number of microliters. In some embodiments, the cells expressing the receptor are detected as at least 10, 20, 30, 40, 50, or 60% of total PBMCs in the blood of the subject, and/or at such a level for at least 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 24, 36, 48, or 52 weeks following the T cells, *e.g.*, CAR-expressing T cells and/or the gamma secretase inhibitor, or for 1, 2, 3, 4, or 5, or more years following such administration.

[0561] In some aspects, the method results in at least a 2-fold, at least a 4-fold, at least a 10-fold, or at least a 20-fold increase in copies of nucleic acid encoding the recombinant receptor,

e.g., CAR, per microgram of DNA, *e.g.*, in the serum, plasma, blood or tissue, *e.g.*, tumor sample, of the subject.

[0562] In some embodiments, cells expressing the receptor are detectable in the serum, plasma, blood or tissue, *e.g.*, tumor sample, of the subject, *e.g.*, by a specified method, such as qPCR or flow cytometry-based detection methods, at least 20, 21, 22, 23, 24, 25, 26, 27, 28, 29, 30, 31, 32, 33, 34, 35, 36, 37, 38, 39, 40, 41, 42, 43, 44, 45, 46, 47, 48, 49, 50, 51, 52, 53, 54, 55, 56, 57, 58, 59, or 60 or more days following administration of the T cells, *e.g.*, CAR-expressing T cells, or after administration of the gamma secretase inhibitor, for at least at or about 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 18, 19, 20, 21, 22, 23, or 24 or more weeks following the administration of the T cells, *e.g.*, CAR-expressing T cells, and/or the gamma secretase inhibitor.

[0563] In some aspects, at least about 1 x 10^2, at least about 1 x 10^3, at least about 1 x 10^4, at least about 1 x 10^5, or at least about 1 x 10^6 or at least about 5 x 10^6 or at least about 1 x 10^7 or at least about 5 x 10^7 or at least about 1 x 10^8 recombinant receptor-expressing, *e.g.*, CAR-expressing cells, and/or at least 10, 25, 50, 100, 200, 300, 400, or 500, or 1000 receptor-expressing cells per microliter, *e.g.*, at least 10 per microliter, are detectable or are present in the subject or fluid, plasma, serum, tissue, or compartment thereof, such as in the blood, *e.g.*, peripheral blood, or disease site, *e.g.*, tumor, thereof. In some embodiments, such a number or concentration of cells is detectable in the subject for at least about 20 days, at least about 40 days, or at least about 60 days, or at least about 3, 4, 5, 6, 7, 8, 9, 10, 11, or 12 months, or at least 2 or 3 years, following administration of the T cells, *e.g.*, CAR-expressing T cells, and/or following the administration of the gamma secretase inhibitor. Such cell numbers may be as detected by flow cytometry-based or quantitative PCR-based methods and extrapolation to total cell numbers using known methods. *See, e.g.,* Brentjens *et al.*, *Sci Transl Med.* 2013 5(177), Park et al, Molecular Therapy 15(4):825–833 (2007), Savoldo *et al.*, *JCI* 121(5):1822-1826 (2011), Davila *et al.*, (2013) *PLoS ONE* 8(4):e61338, Davila *et al.*, *Oncoimmunology* 1(9):1577-1583 (2012), Lamers, *Blood* 2011 117:72-82, Jensen *et al.*, *Biol Blood Marrow Transplant* 2010 September; 16(9): 1245–1256, Brentjens *et al.*, *Blood* 2011 118(18):4817-4828.

[0564] In some aspects, the copy number of nucleic acid encoding the recombinant receptor, *e.g.*, vector copy number, per 100 cells, for example in the peripheral blood or bone marrow or other compartment, as measured by immunohistochemistry, PCR, and/or flow cytometry, is at

least 0.01, at least 0.1, at least 1, or at least 10, at about 1 week, about 2 weeks, about 3 weeks, about 4 weeks, about 5 weeks, or at least about 6 weeks, or at least about 2, 3, 4, 5, 6, 7, 8. 9, 10, 11, or 12 months or at least 2 or 3 years following administration of the cells, *e.g.*, CAR-expressing T cells, and/or the gamma secretase inhibitor. In some embodiments, the copy number of the vector expressing the receptor, *e.g.* CAR, per microgram of genomic DNA is at least 100, at least 1000, at least 5000, or at least 10,000, or at least 15,000 or at least 20,000 at a time about 1 week, about 2 weeks, about 3 weeks, or at least about 4 weeks following administration of the T cells, *e.g.*, CAR-expressing T cells, or gamma secretase inhibitor, or at least 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, or 12 months or at least 2 or 3 years following such administration.

[0565] In some aspects, the receptor, *e.g.* CAR, expressed by the cells, is detectable by quantitative PCR (qPCR) or by flow cytometry in the subject, plasma, serum, blood, tissue and/or disease site thereof, *e.g.*, tumor site, at a time that is at least about 3 months, at least about 6 months, at least about 12 months, at least about 1 year, at least about 2 years, at least about 3 years, or more than 3 years, following the administration of the cells, *e.g.*, following the initiation of the administration of the T cells, *e.g.*, CAR-expressing T cells, and/or the gamma secretase inhibitor.

[0566] In some embodiments, the area under the curve (AUC) for concentration of receptor-(*e.g.*, CAR-) expressing cells in a fluid, plasma, serum, blood, tissue, organ and/or disease site, *e.g.* tumor site, of the subject over time following the administration of the T cells, *e.g.*, CAR-expressing T cells and/or gamma secretase inhibitor, is greater as compared to that achieved via an alternative dosing regimen where the subject is administered the T cells, *e.g.*, CAR-expressing T cells, in the absence of administering the gamma secretase inhibitor.

[0567] In some aspects, the method results in high *in vivo* proliferation of the administered cells, for example, as measured by flow cytometry. In some aspects, high peak proportions of the cells are detected. For example, in some embodiments, at a peak or maximum level following the T cells, *e.g.*, CAR-expressing T cells and/or gamma secretase inhibitor, in the blood, plasma, serum, tissue or disease site of the subject or white blood cell fraction thereof, *e.g.*, PBMC fraction or T cell fraction, at least about 10%, at least about 20%, at least about 30%, at least about 40%, at least about 50%, at least about 60%, at least about 70%, at least about 80%, or at least about 90% of the cells express the recombinant receptor, *e.g.*, the CAR.

[0568] In some aspects, the increased or prolonged expansion and/or persistence of the dose of cells in the subject administered with the gamma secretase inhibitor is associated with a benefit in tumor related outcomes in the subject. In some embodiments, the tumor related outcome includes a decrease in tumor burden or a decrease in blast marrow in the subject. In some embodiments, the tumor burden is decreased by or by at least at or about 10, 20, 30, 40, 50, 60, 70, 80, 90, or 100 percent after administration of the method. In some embodiments, disease burden, tumor size, tumor volume, tumor mass, and/or tumor load or bulk is reduced following the dose of cells by at least at or about 50%, 60%, 70%, 80%, 90% or more compared a subject that has been treated with a method that does not involve the administration of an gamma secretase inhibitor.

B. T Cell Functional Activity

[0569] In some embodiments, parameters associated with therapy or a treatment outcome, which include parameters that can be assessed for the screening steps and/or assessment of treatment of outcomes and/or monitoring treatment outcomes, includes one or more of activity, phenotype, proliferation or function of T cells. In some embodiments, any of the known assays in the art for assessing the activity, phenotypes, proliferation and/or function of the T cells, *e.g.*, T cells administered for T cell therapy, can be used. Prior to and/or subsequent to administration of the cells and/or gamma secretase inhibitor, the biological activity of the engineered cell populations in some embodiments is measured, *e.g.*, by any of a number of known methods. Parameters to assess include specific binding of an engineered or natural T cell or other immune cell to antigen, *in vivo*, *e.g.*, by imaging, or *ex vivo*, *e.g.*, by ELISA or flow cytometry. In certain embodiments, the ability of the engineered cells to destroy target cells can be measured using any suitable method known in the art, such as cytotoxicity assays described in, for example, Kochenderfer *et al.*, *J. Immunotherapy,* 32(7): 689-702 (2009), and Herman *et al.*, *J. Immunological Methods*, 285(1): 25-40 (2004). In certain embodiments, the biological activity of the cells is measured by assaying expression and/or secretion of one or more cytokines, such as CD107a, IFNγ, IL-2, GM-CSF and TNFα, and/or by assessing cytolytic activity.

[0570] In some embodiments, assays for the activity, phenotypes, proliferation and/or function of the T cells, *e.g.*, T cells administered for T cell therapy include, but are not limited to, ELISPOT, ELISA, cellular proliferation, cytotoxic lymphocyte (CTL) assay, binding to the T cell epitope, antigen or ligand, or intracellular cytokine staining, proliferation assays,

lymphokine secretion assays, direct cytotoxicity assays, and limiting dilution assays. In some embodiments, proliferative responses of the T cells can be measured, *e.g.* by incorporation of ^{3}H-thymidine, BrdU (5-Bromo-2'-Deoxyuridine) or 2'-deoxy-5-ethynyluridine (EdU) into their DNA or dye dilution assays, using dyes such as carboxyfluorescein diacetate succingamma secretase inhibitoryl ester (CFSE), CellTrace Violet, or membrane dye PKH26.

[0571] In some embodiments, assessing the activity, phenotypes, proliferation and/or function of the T cells, *e.g.*, T cells administered for T cell therapy, include measuring cytokine production from T cells, and/or measuring cytokine production in a biological sample from the subject, *e.g.*, plasma, serum, blood, and/or tissue samples, *e.g.*, tumor samples. In some cases, such measured cytokines can include, without limitation, interleukin-2 (IL-2), interferon-gamma (IFNγ), interleukin-4 (IL-4), TNF-alpha (TNFα), interleukin-6 (IL-6), interleukin-10 (IL-10), interleukin-12 (IL-12), granulocyte-macrophage colony-stimulating factor (GM-CSF), CD107a, and/or TGF-beta (TGFβ). Assays to measure cytokines are well known in the art, and include but are not limited to, ELISA, intracellular cytokine staining, cytometric bead array, RT-PCR, ELISPOT, flow cytometry and bio-assays in which cells responsive to the relevant cytokine are tested for responsiveness (*e.g.* proliferation) in the presence of a test sample.

[0572] In some embodiments, assessing the activity, phenotypes, proliferation and/or function of the T cells, *e.g.*, T cells administered for T cell therapy, include assessing cell phenotypes, *e.g.*, expression of particular cell surface markers. In some embodiments, the T cells, *e.g.*, T cells administered for T cell therapy, are assessed for expression of T cell activation markers, T cell exhaustion markers, and/or T cell differentiation markers. In some embodiments, the cell phenotype is assessed before administration. In some embodiments, the cell phenotype is assessed after administration. T cell activation markers, T cell exhaustion markers, and/or T cell differentiation markers for assessment include any markers known in the art for particular subsets of T cells, *e.g.*, CD25, CD38, human leukocyte antigen-DR (HLA-DR), CD69, CD44, CD137, KLRG1, CD62L^{low}, CCR7^{low}, CD71, CD2, CD54, CD58, CD244, CD160, programmed cell death protein 1 (PD-1), lymphocyte activation gene 3 protein (LAG-3), T-cell immunoglobulin domain and mucin domain protein 3 (TIM-3), cytotoxic T lymphocyte antigen-4 (CTLA-4), band T lymphocyte attenuator (BTLA) and/or T-cell immunoglobulin and immunoreceptor tyrosine-based inhibitory motif domain (TIGIT) (see, *e.g.*, Liu *et al.*, Cell

Death and Disease (2015) 6, e1792). In some embodiments, the assessed cell surface marker is CD25, PD-1 and/or TIM-3. In some embodiments, the assessed cell surface marker is CD25.

[0573] In some aspects, detecting the expression levels includes performing an *in vitro* assay. In some embodiments, the *in vitro* assay is an immunoassay, an aptamer-based assay, a histological or cytological assay, or an mRNA expression level assay. In some embodiments, the parameter or parameters for one or more of each of the one or more factors, effectors, enzymes and/or surface markers are detected by an enzyme linked immunosorbent assay (ELISA), immunoblotting, immunoprecipitation, radioimmunoassay (RIA), immunostaining, flow cytometry assay, surface plasmon resonance (SPR), chemiluminescence assay, lateral flow immunoassay, inhibition assay or avidity assay. In some embodiments, detection of cytokines and/or surface markers is determined using a binding reagent that specifically binds to at least one biomarker. In some cases, the binding reagent is an antibody or antigen-binding fragment thereof, an aptamer or a nucleic acid probe.

[0574] In some embodiments, the administration of the gamma secretase inhibitor increases the level of circulating CAR T cells.

C. Disease burden

[0575] In some embodiments, parameters associated with therapy or a treatment outcome, which include parameters that can be assessed for the screening steps and/or assessment of treatment of outcomes and/or monitoring treatment outcomes, includes tumor or disease burden. The administration of the immunotherapy, such as a T cell therapy (*e.g.* CAR-expressing T cells) and/or the gamma secretase inhibitor, can reduce or prevent the expansion or burden of the disease or condition in the subject. For example, where the disease or condition is a tumor, the methods generally reduce tumor size, bulk, metastasis, percentage of blasts in the bone marrow or molecularly detectable cancer and/or improve prognosis or survival or other symptom associated with tumor burden.

[0576] In some embodiments, the provided methods result in a decreased tumor burden in treated subjects compared to alternative methods in which the immunotherapy, such as a T cell therapy (*e.g.* CAR-expressing T cells) is given without administration of the gamma secretase inhibitor. It is not necessary that the tumor burden actually be reduced in all subjects receiving the combination therapy, but that tumor burden is reduced on average in subjects treated, such as based on clinical data, in which a majority of subjects treated with such a combination

therapy exhibit a reduced tumor burden, such as at least 50%, 60%, 70%, 80%, 90%, 95% or more of subjects treated with the combination therapy, exhibit a reduced tumor burden.

[0577] Disease burden can encompass a total number of cells of the disease in the subject or in an organ, tissue, or bodily fluid of the subject, such as the organ or tissue of the tumor or another location, *e.g.*, which would indicate metastasis. For example, tumor cells may be detected and/or quantified in the blood, lymph or bone marrow in the context of certain hematological malignancies. Disease burden can include, in some embodiments, the mass of a tumor, the number or extent of metastases and/or the percentage of blast cells present in the bone marrow.

[0578] In some embodiments, the subject has a myeloma, a lymphoma or a leukemia. In some embodiments, the subject has a non-Hodgkin lymphoma (NHL), an acute lymphoblastic leukemia (ALL), a chronic lymphocytic leukemia (CLL), a diffuse large B-cell lymphoma (DLBCL) or a myeloma, *e.g.*, a multiple myeloma (MM). In some embodiments, the subject has a MM or a DBCBL.

[0579] In some embodiments, the subject has a solid tumor.

[0580] In some embodiments, the subject has a MM. In some cases, the MM is a relapsed and/or refractory MM. In the case of MM, exemplary parameters to assess the extent of disease burden include such parameters as number of clonal plasma cells (*e.g.*, >10% on bone marrow biopsy or in any quantity in a biopsy from other tissues; plasmacytoma), presence of monoclonal protein (paraprotein) in either serum or urine, evidence of end-organ damage felt related to the plasma cell disorder (*e.g.*, hypercalcemia (corrected calcium >2.75 mmol/l); renal insufficiency attributable to myeloma; anemia (hemoglobin <10 g/dl); and/or bone lesions (lytic lesions or osteoporosis with compression fractures)).

[0581] In the case of DLBCL, exemplary parameters to assess the extent of disease burden include such parameters as cellular morphology (*e.g.*, centroblastic, immunoblastic, and anaplastic cells), gene expression, miRNA expression and protein expression (*e.g.*, expression of BCL2, BCL6, MUM1, LMO2, MYC, and p21).

[0582] In the case of leukemia, the extent of disease burden can be determined by assessment of residual leukemia in blood or bone marrow. In some embodiments, a subject exhibits morphologic disease if there are greater than or equal to 5% blasts in the bone marrow,

for example, as detected by light microscopy. In some embodiments, a subject exhibits complete or clinical remission if there are less than 5% blasts in the bone marrow.

[0583] In some embodiments, for leukemia, a subject may exhibit complete remission, but a small proportion of morphologically undetectable (by light microscopy techniques) residual leukemic cells are present. A subject is said to exhibit minimum residual disease (MRD) if the subject exhibits less than 5% blasts in the bone marrow and exhibits molecularly detectable cancer. In some embodiments, molecularly detectable cancer can be assessed using any of a variety of molecular techniques that permit sensitive detection of a small number of cells. In some aspects, such techniques include PCR assays, which can determine unique Ig/T-cell receptor gene rearrangements or fusion transcripts produced by chromosome translocations. In some embodiments, flow cytometry can be used to identify cancer cells based on leukemia-specific immunophenotypes. In some embodiments, molecular detection of cancer can detect as few as 1 leukemia cell in 100,000 normal cells. In some embodiments, a subject exhibits MRD that is molecularly detectable if at least or greater than 1 leukemia cell in 100,000 cells is detected, such as by PCR or flow cytometry. In some embodiments, the disease burden of a subject is molecularly undetectable or MRD, such that, in some cases, no leukemia cells are able to be detected in the subject using PCR or flow cytometry techniques.

[0584] In some embodiments, the methods and/or administration of an immunotherapy, such as a T cell therapy (*e.g.* CAR-expressing T cells) and/or gamma secretase inhibitor decrease(s) disease burden as compared with disease burden at a time immediately prior to the administration of the immunotherapy, *e.g.*, T cell therapy and/or gamma secretase inhibitor.

[0585] In some aspects, administration of the immunotherapy, *e.g.* T cell therapy and/or gamma secretase inhibitor, may prevent an increase in disease burden, and this may be evidenced by no change in disease burden.

[0586] In some embodiments, the method reduces the burden of the disease or condition, *e.g.*, number of tumor cells, size of tumor, duration of patient survival or event-free survival, to a greater degree and/or for a greater period of time as compared to the reduction that would be observed with a comparable method using an alternative therapy, such as one in which the subject receives immunotherapy, *e.g.* T cell therapy alone, in the absence of administration of the gamma secretase inhibitor In some embodiments, disease burden is reduced to a greater extent or for a greater duration following the combination therapy of administration of the

immunotherapy, *e.g.*, T cell therapy, and the gamma secretase inhibitor, compared to the reduction that would be effected by administering each of the agent alone, *e.g.*, administering the gamma secretase inhibitor to a subject having not received the immunotherapy, *e.g.* T cell therapy; or administering the immunotherapy, *e.g.* T cell therapy, to a subject having not received the gamma secretase inhibitor.

[0587] In some embodiments, the burden of a disease or condition in the subject is detected, assessed, or measured. Disease burden may be detected in some aspects by detecting the total number of disease or disease-associated cells, *e.g.*, tumor cells, in the subject, or in an organ, tissue, or bodily fluid of the subject, such as blood or serum. In some embodiments, disease burden, *e.g.* tumor burden, is assessed by measuring the mass of a solid tumor and/or the number or extent of metastases. In some aspects, survival of the subject, survival within a certain time period, extent of survival, presence or duration of event-free or symptom-free survival, or relapse-free survival, is assessed. In some embodiments, any symptom of the disease or condition is assessed. In some embodiments, the measure of disease or condition burden is specified. In some embodiments, exemplary parameters for determination include particular clinical outcomes indicative of amelioration or improvement in the disease or condition, *e.g.*, tumor. Such parameters include: duration of disease control, including complete response (CR), partial response (PR) or stable disease (SD) (see, *e.g.*, Response Evaluation Criteria In Solid Tumors (RECIST) guidelines), objective response rate (ORR), progression-free survival (PFS) and overall survival (OS). Specific thresholds for the parameters can be set to determine the efficacy of the method of combination therapy provided herein.

[0588] In some aspects, disease burden is measured or detected prior to administration of the immunotherapy, *e.g.* T cell therapy, following the administration of the immunotherapy, *e.g.* T cell therapy but prior to administration of the gamma secretase inhibitor, following administration of the gamma secretase inhibitor but prior to the administration of the immunotherapy, *e.g.*, T cell therapy, and/or following the administration of both the immunotherapy, *e.g.* T cell therapy and the gamma secretase inhibitor. In the context of multiple administration of one or more steps of the combination therapy, disease burden in some embodiments may be measured prior to or following administration of any of the steps, doses and/or cycles of administration, or at a time between administration of any of the steps, doses and/or cycles of administration.

[0589] In some embodiments, the burden is decreased by or by at least at or about 10, 20, 30, 40, 50, 60, 70, 80, 90, or 100 percent by the provided methods compared to immediately prior to the administration of the gamma secretase inhibitor and the immunotherapy, *e.g.* T cell therapy. In some embodiments, disease burden, tumor size, tumor volume, tumor mass, and/or tumor load or bulk is reduced following administration of the immunotherapy, *e.g.* T cell therapy and the gamma secretase inhibitor, by at least at or about 10, 20, 30, 40, 50, 60, 70, 80, 90 % or more compared to that immediately prior to the administration of the immunotherapy, *e.g.* T cell therapy and/or the gamma secretase inhibitor.

[0590] In some embodiments, reduction of disease burden by the method comprises an induction in morphologic complete remission, for example, as assessed at 1 month, 2 months, 3 months, or more than 3 months, after administration of, *e.g.*, initiation of, the combination therapy.

[0591] In some aspects, an assay for minimal residual disease, for example, as measured by multiparametric flow cytometry, is negative, or the level of minimal residual disease is less than about 0.3%, less than about 0.2%, less than about 0.1%, or less than about 0.05%.

[0592] In some embodiments, the event-free survival rate or overall survival rate of the subject is improved by the methods, as compared with other methods. For example, in some embodiments, event-free survival rate or probability for subjects treated by the methods at 6 months following the method of combination therapy provided herein, is greater than about 40%, greater than about 50%, greater than about 60%, greater than about 70%, greater than about 80%, greater than about 90%, or greater than about 95%. In some aspects, overall survival rate is greater than about 40%, greater than about 50%, greater than about 60%, greater than about 70%, greater than about 80%, greater than about 90%, or greater than about 95%. In some embodiments, the subject treated with the methods exhibits event-free survival, relapse-free survival, or survival to at least 6 months, or at least 1, 2, 3, 4, 5, 6, 7, 8, 9, or 10 years. In some embodiments, the time to progression is improved, such as a time to progression of greater than at or about 6 months, or at least 1, 2, 3, 4, 5, 6, 7, 8, 9, or 10 years.

[0593] In some embodiments, following treatment by the method, the probability of relapse is reduced as compared to other methods. For example, in some embodiments, the probability of relapse at 6 months following the method of combination therapy, is less than about 80%, less

than about 70%, less than about 60%, less than about 50%, less than about 40%, less than about 30%, less than about 20%, or less than about 10%.

IV. ARTICLES OF MANUFACTURE AND KITS

[0594] Also provided are articles of manufacture containing a gamma secretase inhibitor, and components for the immunotherapy, *e.g.*, antibody or antigen binding fragment thereof or T cell therapy, *e.g.* engineered cells, and/or compositions thereof. The articles of manufacture may include a container and a label or package insert on or associated with the container. Suitable containers include, for example, bottles, vials, syringes, IV solution bags, etc. The containers may be formed from a variety of materials such as glass or plastic. The container in some embodiments holds a composition which is by itself or combined with another composition effective for treating, preventing and/or diagnosing the condition. In some embodiments, the container has a sterile access port. Exemplary containers include an intravenous solution bags, vials, including those with stoppers pierceable by a needle for injection, or bottles or vials for orally administered agents. The label or package insert may indicate that the composition is used for treating a disease or condition.

[0595] The article of manufacture may include (a) a first container with a composition contained therein, wherein the composition includes the antibody or engineered cells used for the immunotherapy, *e.g.* T cell therapy; and (b) a second container with a composition contained therein, wherein the composition includes the second agent, such as a gamma secretase inhibitor. The article of manufacture may further include a package insert indicating that the compositions can be used to treat a particular condition. Alternatively, or additionally, the article of manufacture may further include another or the same container comprising a pharmaceutically-acceptable buffer. It may further include other materials such as other buffers, diluents, filters, needles, and/or syringes.

V. DEFINITIONS

[0596] Unless defined otherwise, all terms of art, notations and other technical and scientific terms or terminology used herein are intended to have the same meaning as is commonly understood by one of ordinary skill in the art to which the claimed subject matter pertains. In some cases, terms with commonly understood meanings are defined herein for clarity and/or for

ready reference, and the inclusion of such definitions herein should not necessarily be construed to represent a substantial difference over what is generally understood in the art.

[0597] As used herein, a “subject” is a mammal, such as a human or other animal, and typically is human. In some embodiments, the subject, e.g., patient, to whom the immunomodulatory polypeptides, engineered cells, or compositions, such as containing a compound, e.g. gamma secretase inhibitor, are administered, is a mammal, typically a primate, such as a human. In some embodiments, the primate is a monkey or an ape. The subject can be male or female and can be any suitable age, including infant, juvenile, adolescent, adult, and geriatric subjects. In some embodiments, the subject is a non-primate mammal, such as a rodent. In particular embodiments, the subject is a human.

[0598] As used herein, “treatment” (and grammatical variations thereof such as “treat” or “treating”) refers to complete or partial amelioration or reduction of a disease or condition or disorder, or a symptom, adverse effect or outcome, or phenotype associated therewith. Desirable effects of treatment include, but are not limited to, preventing occurrence or recurrence of disease, alleviation of symptoms, diminishment of any direct or indirect pathological consequences of the disease, preventing metastasis, decreasing the rate of disease progression, amelioration or palliation of the disease state, and remission or improved prognosis. The terms do not imply complete curing of a disease or complete elimination of any symptom or effect(s) on all symptoms or outcomes.

[0599] As used herein, “delaying development of a disease" means to defer, hinder, slow, retard, stabilize, suppress and/or postpone development of the disease (such as cancer). This delay can be of varying lengths of time, depending on the history of the disease and/or individual being treated. As is evident to one skilled in the art, a sufficient or significant delay can, in effect, encompass prevention, in that the individual does not develop the disease. For example, a late stage cancer, such as development of metastasis, may be delayed.

[0600] “Preventing,” as used herein, includes providing prophylaxis with respect to the occurrence or recurrence of a disease in a subject that may be predisposed to the disease but has not yet been diagnosed with the disease. In some embodiments, the provided cells and compositions are used to delay development of a disease or to slow the progression of a disease.

[0601] As used herein, to “suppress” a function or activity is to reduce the function or activity when compared to otherwise same conditions except for a condition or parameter of

interest, or alternatively, as compared to another condition. For example, cells that suppress tumor growth reduce the rate of growth of the tumor compared to the rate of growth of the tumor in the absence of the cells.

[0602] An "effective amount" of an agent, e.g., a pharmaceutical formulation, cells, or composition, in the context of administration, refers to an amount effective, at dosages/amounts and for periods of time necessary, to achieve a desired result, such as a therapeutic or prophylactic result.

[0603] A "therapeutically effective amount" of an agent, e.g., a pharmaceutical formulation or engineered cells, refers to an amount effective, at dosages and for periods of time necessary, to achieve a desired therapeutic result, such as for treatment of a disease, condition, or disorder, and/or pharmacokinetic or pharmacodynamic effect of the treatment. The therapeutically effective amount may vary according to factors such as the disease state, age, sex, and weight of the subject, and the immunomodulatory polypeptides or engineered cells administered. In some embodiments, the provided methods involve administering the immunomodulatory polypeptides, engineered cells, or compositions at effective amounts, e.g., therapeutically effective amounts.

[0604] A "prophylactically effective amount" refers to an amount effective, at dosages and for periods of time necessary, to achieve the desired prophylactic result. Typically but not necessarily, since a prophylactic dose is used in subjects prior to or at an earlier stage of disease, the prophylactically effective amount will be less than the therapeutically effective amount.

[0605] The term "pharmaceutical formulation" refers to a preparation which is in such form as to permit the biological activity of an active ingredient contained therein to be effective, and which contains no additional components which are unacceptably toxic to a subject to which the formulation would be administered.

[0606] A "pharmaceutically acceptable carrier" refers to an ingredient in a pharmaceutical formulation, other than an active ingredient, which is nontoxic to a subject. A pharmaceutically acceptable carrier includes, but is not limited to, a buffer, excipient, stabilizer, or preservative.

[0607] As used herein, recitation that nucleotides or amino acid positions "correspond to" nucleotides or amino acid positions in a disclosed sequence, such as set forth in the Sequence listing, refers to nucleotides or amino acid positions identified upon alignment with the disclosed sequence to maximize identity using a standard alignment algorithm, such as the GAP

algorithm. By aligning the sequences, one skilled in the art can identify corresponding residues, for example, using conserved and identical amino acid residues as guides. In general, to identify corresponding positions, the sequences of amino acids are aligned so that the highest order match is obtained (see, e.g. : Computational Molecular Biology, Lesk, A.M., ed., Oxford University Press, New York, 1988; Biocomputing: Informatics and Genome Projects, Smith, D.W., ed., Academic Press, New York, 1993; Computer Analysis of Sequence Data, Part I, Griffin, A.M., and Griffin, H.G., eds., Humana Press, New.Jersey, 1994; Sequence Analysis in Molecular Biology, von Heinje, G., Academic Press, 1987; and Sequence Analysis Primer, Gribskov, M. and Devereux, J., eds., M Stockton Press, New York, 1991; Carrillo et al. (1988) SIAM J Applied Math 48: 1073).

[0608] The term "vector," as used herein, refers to a nucleic acid molecule capable of propagating another nucleic acid to which it is linked. The term includes the vector as a self-replicating nucleic acid structure as well as the vector incorporated into the genome of a host cell into which it has been introduced. Certain vectors are capable of directing the expression of nucleic acids to which they are operatively linked. Such vectors are referred to herein as "expression vectors." Among the vectors are viral vectors, such as retroviral, e.g., gammaretroviral and lentiviral vectors.

[0609] The terms "host cell," "host cell line," and "host cell culture" are used interchangeably and refer to cells into which exogenous nucleic acid has been introduced, including the progeny of such cells. Host cells include "transformants" and "transformed cells," which include the primary transformed cell and progeny derived therefrom without regard to the number of passages. Progeny may not be completely identical in nucleic acid content to a parent cell, but may contain mutations. Mutant progeny that have the same function or biological activity as screened or selected for in the originally transformed cell are included herein.

[0610] As used herein, a statement that a cell or population of cells is "positive" for a particular marker refers to the detectable presence on or in the cell of a particular marker, typically a surface marker. When referring to a surface marker, the term refers to the presence of surface expression as detected by flow cytometry, for example, by staining with an antibody that specifically binds to the marker and detecting said antibody, wherein the staining is detectable by flow cytometry at a level substantially above the staining detected carrying out the same procedure with an isotype-matched control under otherwise identical conditions and/or at a

level substantially similar to that for cell known to be positive for the marker, and/or at a level substantially higher than that for a cell known to be negative for the marker.

[0611] As used herein, a statement that a cell or population of cells is "negative" for a particular marker refers to the absence of substantial detectable presence on or in the cell of a particular marker, typically a surface marker. When referring to a surface marker, the term refers to the absence of surface expression as detected by flow cytometry, for example, by staining with an antibody that specifically binds to the marker and detecting said antibody, wherein the staining is not detected by flow cytometry at a level substantially above the staining detected carrying out the same procedure with an isotype-matched control under otherwise identical conditions, and/or at a level substantially lower than that for cell known to be positive for the marker, and/or at a level substantially similar as compared to that for a cell known to be negative for the marker.

[0612] As used herein, "percent (%) amino acid sequence identity" and "percent identity" when used with respect to an amino acid sequence (reference polypeptide sequence) is defined as the percentage of amino acid residues in a candidate sequence (e.g., the subject antibody or fragment) that are identical with the amino acid residues in the reference polypeptide sequence, after aligning the sequences and introducing gaps, if necessary, to achieve the maximum percent sequence identity, and not considering any conservative substitutions as part of the sequence identity. Alignment for purposes of determining percent amino acid sequence identity can be achieved in various ways that are within the skill in the art, for instance, using publicly available computer software such as BLAST, BLAST-2, ALIGN or Megalign (DNASTAR) software. Those skilled in the art can determine appropriate parameters for aligning sequences, including any algorithms needed to achieve maximal alignment over the full length of the sequences being compared.

[0613] As used herein, the singular forms "a," "an," and "the" include plural referents unless the context clearly dictates otherwise. For example, "a" or "an" means "at least one" or "one or more." It is understood that aspects and variations described herein include "consisting" and/or "consisting essentially of" aspects and variations.

[0614] Throughout this disclosure, various aspects of the claimed subject matter are 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 claimed subject matter. Accordingly, the description of a range should be considered to have specifically disclosed all the possible sub-ranges as well as individual numerical values within that range. For example, where a range of values is provided, it is understood that each intervening value, between the upper and lower limit of that range and any other stated or intervening value in that stated range is encompassed within the claimed subject matter. The upper and lower limits of these smaller ranges may independently be included in the smaller ranges, and are also encompassed within the claimed subject matter, subject to any specifically excluded limit in the stated range. Where the stated range includes one or both of the limits, ranges excluding either or both of those included limits are also included in the claimed subject matter. This applies regardless of the breadth of the range.

[0615] The term "about" as used herein refers to the usual error range for the respective value readily known to the skilled person in this technical field. Reference to "about" a value or parameter herein includes (and describes) embodiments that are directed to that value or parameter per se. For example, description referring to "about X" includes description of "X".

[0616] As used herein, a composition refers to any mixture of two or more products, substances, or compounds, including cells. It may be a solution, a suspension, liquid, powder, a paste, aqueous, non-aqueous or any combination thereof.

VI. EXEMPLARY EMBODIMENTS

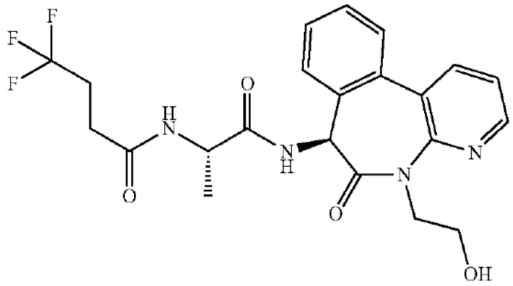
[0617] Among the provided embodiments are:

1. A method of treatment, the method comprising:

(a) administering a cell therapy to a subject having a disease or disorder, said cell therapy comprising a dose of genetically engineered cells expressing a recombinant receptor; and

(b) administering to the subject LY3039478 or a compound of the structure:

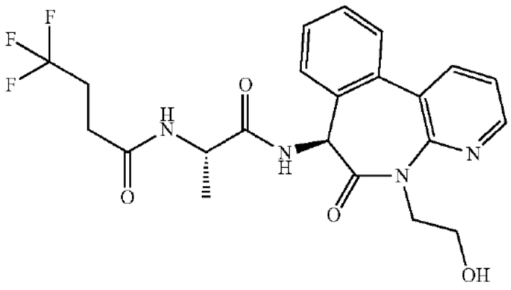
Compound 1



or a stereoisomer or a pharmaceutically acceptable salt or hydrate of any of the foregoing.

2. A method of treatment, the method comprising administering a cell therapy to a subject having a disease or disorder, said cell therapy comprising a dose of genetically engineered cells expressing a recombinant receptor, wherein, at the time of initiation of the administration of the cell therapy, the subject has been previously administered, and/or is undergoing treatment with, LY3039478 or a compound of the structure:

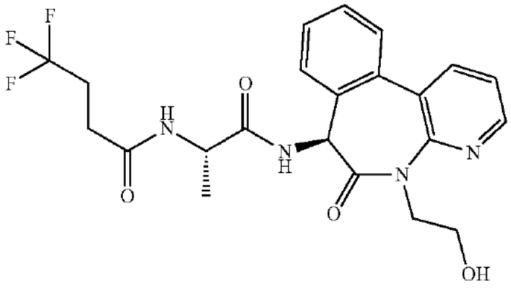
Compound 1



or a stereoisomer or a pharmaceutically acceptable salt or hydrate of any of the foregoing.

3. A method of treatment, the method comprising administering to a subject having a disease or disorder LY3039478 or a compound of the structure:

Compound 1



or a stereoisomer or a pharmaceutically acceptable salt or hydrate of any of the foregoing,

wherein, at the time of initiation of the administration, the subject has been previously administered, and/or is undergoing treatment with, a cell therapy, said cell therapy comprising a dose of genetically engineered cells expressing a recombinant receptor.

4. The method of any of embodiments 1-3, wherein the recombinant receptor specifically binds to a target antigen associated with the disease or disorder.

5. The method of embodiment 4, wherein the target antigen is selected from among, B cell maturation antigen (BCMA), carbonic anhydrase 9 (CAIX), Her2/neu (receptor tyrosine kinase erbB2), L1-CAM, CD19, CD20, CD22, mesothelin, CEA, and hepatitis B surface antigen, anti-folate receptor, CD23, CD24, CD30, CD33, CD38, CD44, EGFR, epithelial glycoprotein 2 (EPG-2), epithelial glycoprotein 40 (EPG-40), EPHa2, erb-B2, erb-B3, erb-B4, erbB dimers, EGFR vIII, folate binding protein (FBP), FCRL5, FCRH5, fetal acetylcholine receptor, GD2, GD3, HMW-MAA, IL-22R-alpha, IL-13R-alpha2, kinase insert domain receptor (kdr), kappa light chain, Lewis Y, L1-cell adhesion molecule, (L1-CAM), Melanoma-associated antigen (MAGE)-A1, MAGE-A3, MAGE-A6, Preferentially expressed antigen of melanoma (PRAME), survivin, TAG72, B7-H6, IL-13 receptor alpha 2 (IL-13Ra2), CA9, GD3, HMW-MAA, CD171, G250/CAIX, HLA-AI MAGE Al, HLA-A2 NY-ESO-1, PSCA, folate receptor-a, CD44v6, CD44v7/8, avb6 integrin, 8H9, NCAM, VEGF receptors, 5T4, Foetal AchR, NKG2D ligands, CD44v6, dual antigen, a cancer-testes antigen, mesothelin, murine CMV, mucin 1 (MUC1), MUC16, PSCA, NKG2D, NY-ESO-1, MART-1, gp100, oncofetal antigen, ROR1, TAG72, VEGF-R2, carcinoembryonic antigen (CEA), Her2/neu, estrogen receptor, progesterone receptor, ephrinB2, CD123, c-Met, GD-2, O-acetylated GD2 (OGD2), CE7, Wilms Tumor 1 (WT-1), a cyclin, cyclin A2, CCL-1, CD138, a pathogen-specific antigen and an antigen associated with a universal tag.

6. The method of embodiment 4 or embodiment 5, wherein the target antigen is Muc1.

7. The method of embodiment 4 or embodiment 5, wherein the target antigen is BCMA.

8. The method of embodiment 7, wherein the BCMA is surface BCMA.

9. The method of embodiment 8, wherein binding of the recombinant receptor to surface BCMA or a measure indicative of function or activity of recombinant receptor-expressing cells, following exposure to cells expressing surface BCMA, is not reduced or blocked or is not substantially reduced or blocked in the presence of a soluble or shed form of the BCMA.

10. The method of embodiment 9, wherein the concentration or amount of the soluble or shed form of the BCMA corresponds to a concentration or amount present in serum or blood or plasma of the subject or of a multiple myeloma patient, or on average in a patient population for the disease or disorder, or at a concentration or amount of the soluble or shed BCMA at which the binding or measure is reduced or blocked, or is substantially reduced or blocked, for cells expressing a reference anti-BCMA recombinant receptor, optionally a reference anti-BCMA CAR, in the same assay.

11. A method of treatment, the method comprising:

(a) administering a cell therapy to a subject having a disease or disorder, said cell therapy comprising a dose of genetically engineered cells expressing a chimeric antigen receptor (CAR) comprises an antigen-binding domain that specifically binds to surface B cell maturation antigen (BCMA),; and

(b) administering to the subject an inhibitor of gamma secretase,

wherein binding of the CAR to surface BCMA or a measure indicative of function or activity of the CAR-expressing cells, following exposure to cells expressing surface BCMA, is not reduced or blocked or is not substantially reduced or blocked in the presence of a soluble or shed form of the BCMA,

optionally at a concentration or amount of the soluble or shed form of the BCMA corresponding to a concentration or amount present in serum or blood or plasma of the subject or of a multiple myeloma patient, or on average in a patient population for the disease or disorder, or at a concentration or amount of the soluble or shed BCMA at which the binding or measure is reduced or blocked, or is substantially reduced or blocked, for cells expressing a reference anti-BCMA CAR in the same assay.

12. A method of treatment, the method comprising administering a cell therapy to a subject having a disease or disorder, said cell therapy comprising a dose of genetically engineered cells expressing a chimeric antigen receptor that specifically binds to surface B cell maturation antigen (BCMA), wherein:

binding of the CAR to surface BCMA or a measure indicative of function or activity of the CAR-expressing cells, following exposure to cells expressing surface BCMA, is not reduced or blocked or is not substantially reduced or blocked in the presence of a soluble or shed form of the BCMA, optionally at a concentration or amount of the soluble or shed form of the BCMA

corresponding to a concentration or amount present in serum or blood or plasma of the subject or of a multiple myeloma patient, or on average in a patient population for the disease or disorder, or at a concentration or amount of the soluble or shed BCMA at which the binding or measure is reduced or blocked, or is substantially reduced or blocked, for cells expressing a reference anti-BCMA CAR in the same assay; and

at the time of initiation of the administration of the cell therapy, the subject has been previously administered, and/or is undergoing treatment with, an inhibitor of gamma secretase.

13. A method of treatment, the method comprising administering an inhibitor of gamma secretase to a subject having a disease or disorder, wherein, at the time of initiation of the administration of the inhibitor, the subject has been previously administered, and/or is undergoing treatment with, a cell therapy, said cell therapy comprising a dose of genetically engineered cells expressing a chimeric antigen receptor that specifically binds surface B cell maturation antigen (BCMA),

wherein binding of the CAR to surface BCMA or a measure indicative of function or activity of the CAR-expressing cells, following exposure to cells expressing surface BCMA, is not reduced or blocked or is not substantially reduced or blocked in the presence of a soluble or shed form of the BCMA,

optionally at a concentration or amount of the soluble or shed form of the BCMA corresponding to a concentration or amount present in serum or blood or plasma of the subject or of a multiple myeloma patient, or on average in a patient population for the disease or disorder, or at a concentration or amount of the soluble or shed BCMA at which the binding or measure is reduced or blocked, or is substantially reduced or blocked, for cells expressing a reference anti-BCMA CAR in the same assay.

14. The method of any of embodiments 1-13, wherein the method comprises selecting a subject having a cancer in which cells of the cancer in the subject (i) express CD138, surface CD38 or a surface plasma cell marker or are derived from plasma cells and (ii) comprise low expression of surface B cell maturation antigen (BCMA) and/or a level of expression of surface BCMA below a threshold level;

15. The method of embodiment 14, wherein the cell therapy and/or the gamma secretase inhibitor is administered to the selected subject.

16. The method of embodiment 14 or embodiment 15, wherein the threshold level of expression of surface BCMA is lower than the average or median expression or level of surface BCMA on plasma cells in a plurality of control subjects, optionally wherein the control subjects are a group of healthy or normal subjects.

17. The method of any of embodiments 14-16, wherein;

the low expression of surface BCMA is present when less than or less than about 60%, less than or less than about 50%, less than or less than about 40%, less than or less than about 30%, less than or less than about 20% or less than or less than about 10% of the plasma cells, or cells with a plasma marker or phenotype or the cancer cells, in the subject express surface BCMA; or

the threshold level of surface BCMA is less than or less than about 60%, less than or less than about 50%, less than or less than about 40%, less than or less than about 30%, less than or less than about 20% or less than or less than about 10% of the plasma cells, or cells with a plasma marker or phenotype or the cancer cells, in the subject that express surface BCMA.

18. The method of any of embodiments 14-17, wherein expression of surface BCMA is determined by flow cytometry and/or an immunoassay.

19. The method of any of embodiments 14-18, wherein binding of the recombinant receptor to surface BCMA or a measure indicative of function or activity of the recombinant receptor-expressing cells, following exposure to cells expressing surface BCMA, is not reduced or blocked or is not substantially reduced or blocked in the presence of a soluble or shed form of the BCMA.

20. The method of embodiment 19, wherein the concentration or amount of the soluble or shed form of the BCMA corresponding to a concentration or amount present in serum or blood or plasma of the subject or of a multiple myeloma patient, or on average in a patient population for the disease or disorder, or at a concentration or amount of the soluble or shed BCMA at which the binding or measure is reduced or blocked, or is substantially reduced or blocked, for cells expressing a reference anti-BCMA recombinant receptor, optionally a reference anti-BCMA CAR, in the same assay.

21. The method of any one of embodiments 1-20, wherein the gamma secretase inhibitor inhibits intramembrane cleavage of BCMA.

22. The method of embodiment 21, wherein the gamma secretase inhibitor inhibits intramembrane cleavage of BCMA with a half-maximal inhibitory concentration (IC_{50}) of 0.01 nM to 10 nM, 0.01 nM to 5 nM, 0.01 nM to 1 nM, 0.01 nM to 0.5 nM, 0.01 nM to 0.35 nM, 0.01 nM to 0.25 nM, 0.01 nM to 0.1 nM, 0.01 nM to 0.05 nM, 0.05 nM to 10 nM, 0.05 nM to 5 nM, 0.05 nM to 1 nM, 0.05 nM to 0.5 nM, 0.05 nM to 0.35 nM, 0.05 nM to 0.25 nM, 0.05 nM to 0.1 nM, 0.1 nM to 10 nM, 0.1 nM to 5 nM, 0.1 nM to 1 nM, 0.1 nM, to 0.5 nM, 0.1 nM to 0.35 nM, 0.1 nM to 0.25 nM, 0.25 nM to 10 nM, 0.25 nM to 5 nM, 0.25 nM to 1 nM, 0.25 nM to 0.5 nM, 0.25 nM to 0.35 nM, 0.35 nM to 10 nM, 0.35 nM to 5 nM, 0.35 nM to 1 nM, 0.35 nM to 0.5 nM, 0.5 nM to 10 nM, 0.5 nM to 5 nM, 0.5 nM to 1 nM, 1 nM to 10 nM, 1 nM to 5 nM or 5 nM to 10 nM.

23. The method of embodiment 21 or embodiment 22, wherein the gamma secretase inhibitor inhibits intramembrane cleavage of BCMA with a half-maximal inhibitory concentration (IC_{50}) of less than or less than about 10 nM, 5 nM, 1 nM., 0.5 nM, 0.35 nM, 0.25 nM, 0.1 nM, 0.05 nM or 0.01 nM or less.

24. A method of treatment, the method comprising:

(a) administering a cell therapy to a subject having a disease or disorder, said cell therapy comprising a dose of genetically engineered cells expressing a recombinant receptor, wherein the recombinant receptor does not specifically bind to surface B cell maturation antigen (BCMA); and

(b) administering an inhibitor of gamma secretase.

25. A method of treatment, the method comprising administering a cell therapy to a subject having a disease or disorder, said cell therapy comprising a dose of genetically engineered cells expressing a recombinant receptor, wherein the recombinant receptor does not specifically bind to surface B cell maturation antigen (BCMA), wherein, at the time of initiation of the administration of the cell therapy, the subject has been previously administered, and/or is undergoing treatment with, an inhibitor of gamma secretase.

26. A method of treatment, the method comprising administering an inhibitor of gamma secretase to a subject having a disease or disorder, wherein, at the time of initiation of the administration of the inhibitor, the subject has been previously administered, and/or is undergoing treatment with, a cell therapy, said cell therapy comprising a dose of genetically

engineered cells expressing a recombinant receptor, wherein the recombinant receptor does not specifically bind to surface B cell maturation antigen (BCMA).

27. The method of any of embodiments 24-26, wherein the recombinant receptor specifically binds to a target antigen associated with the disease or disorder.

28. The method of embodiment 27, wherein the target antigen is selected from among, carbonic anhydrase 9 (CAIX), Her2/neu (receptor tyrosine kinase erbB2), L1-CAM, CD19, CD20, CD22, mesothelin, CEA, and hepatitis B surface antigen, anti-folate receptor, CD23, CD24, CD30, CD33, CD38, CD44, EGFR, epithelial glycoprotein 2 (EPG-2), epithelial glycoprotein 40 (EPG-40), EPHa2, erb-B2, erb-B3, erb-B4, erbB dimers, EGFR vIII, folate binding protein (FBP), FCRL5, FCRH5, fetal acetylcholine receptor, GD2, GD3, HMW-MAA, IL-22R-alpha, IL-13R-alpha2, kinase insert domain receptor (kdr), kappa light chain, Lewis Y, L1-cell adhesion molecule, (L1-CAM), Melanoma-associated antigen (MAGE)-A1, MAGE-A3, MAGE-A6, Preferentially expressed antigen of melanoma (PRAME), survivin, TAG72, B7-H6, IL-13 receptor alpha 2 (IL-13Ra2), CA9, GD3, HMW-MAA, CD171, G250/CAIX, HLA-AI MAGE Al, HLA-A2 NY-ESO-1, PSCA, folate receptor-a, CD44v6, CD44v7/8, avb6 integrin, 8H9, NCAM, VEGF receptors, 5T4, Foetal AchR, NKG2D ligands, CD44v6, dual antigen, a cancer-testes antigen, mesothelin, murine CMV, mucin 1 (MUC1), MUC16, PSCA, NKG2D, NY-ESO-1, MART-1, gp100, oncofetal antigen, ROR1, TAG72, VEGF-R2, carcinoembryonic antigen (CEA), Her2/neu, estrogen receptor, progesterone receptor, ephrinB2, CD123, c-Met, GD-2, O-acetylated GD2 (OGD2), CE7, Wilms Tumor 1 (WT-1), a cyclin, cyclin A2, CCL-1, CD138, a pathogen-specific antigen and an antigen associated with a universal tag.

29. A method of modulating activity of a cell therapy, the method comprising

(a) administering a cell therapy to a subject having a disease or disorder, said cell therapy comprising a dose of genetically engineered cells expressing a recombinant receptor; and

(b) subsequent to the administration in (a) administering to the subject an inhibitor of gamma secretase.

30. A method of modulating activity of a cell therapy, the method comprising administering an inhibitor of gamma secretase to a subject having a disease or disorder, wherein, at the time of initiation of the administration of the inhibitor, the subject has been previously administered, and/or is undergoing treatment with, a cell therapy, said cell therapy comprising a dose of genetically engineered cells expressing recombinant receptor.

31. The method of embodiment 29 or embodiment 30, wherein the recombinant receptor specifically binds to a target antigen associated with the disease or disorder.

32. The method of embodiment 31, wherein the target antigen is selected from among, B cell maturation antigen (BCMA), carbonic anhydrase 9 (CAIX), Her2/neu (receptor tyrosine kinase erbB2), L1-CAM, CD19, CD20, CD22, mesothelin, CEA, and hepatitis B surface antigen, anti-folate receptor, CD23, CD24, CD30, CD33, CD38, CD44, EGFR, epithelial glycoprotein 2 (EPG-2), epithelial glycoprotein 40 (EPG-40), EPHa2, erb-B2, erb-B3, erb-B4, erbB dimers, EGFR vIII, folate binding protein (FBP), FCRL5, FCRH5, fetal acetylcholine receptor, GD2, GD3, HMW-MAA, IL-22R-alpha, IL-13R-alpha2, kinase insert domain receptor (kdr), kappa light chain, Lewis Y, L1-cell adhesion molecule, (L1-CAM), Melanoma-associated antigen (MAGE)-A1, MAGE-A3, MAGE-A6, Preferentially expressed antigen of melanoma (PRAME), survivin, TAG72, B7-H6, IL-13 receptor alpha 2 (IL-13Ra2), CA9, GD3, HMW-MAA, CD171, G250/CAIX, HLA-AI MAGE Al, HLA-A2 NY-ESO-1, PSCA, folate receptor-a, CD44v6, CD44v7/8, avb6 integrin, 8H9, NCAM, VEGF receptors, 5T4, Foetal AchR, NKG2D ligands, CD44v6, dual antigen, a cancer-testes antigen, mesothelin, murine CMV, mucin 1 (MUC1), MUC16, PSCA, NKG2D, NY-ESO-1, MART-1, gp100, oncofetal antigen, ROR1, TAG72, VEGF-R2, carcinoembryonic antigen (CEA), Her2/neu, estrogen receptor, progesterone receptor, ephrinB2, CD123, c-Met, GD-2, O-acetylated GD2 (OGD2), CE7, Wilms Tumor 1 (WT-1), a cyclin, cyclin A2, CCL-1, CD138, a pathogen-specific antigen and an antigen associated with a universal tag.

33. The method of embodiment 27, 28, 31 or 32, wherein the target antigen is Muc1.

34. The method of embodiment 31 or embodiment 32, wherein the target antigen is BCMA.

35. The method of embodiment 34, wherein the BCMA is surface BCMA.

36. The method of any of embodiments 14-35, wherein the gamma secretase inhibitor inhibits or reduces or is capable of inhibiting or reducing cleavage of one or more targets selected from BCMA, Notch 1, Notch 2, Notch 3, Notch 4, Muc1, Ephrin B2, Betaglycan (TGFBR3), CD43, CD44, CSF1R, CXCR1, CXCL16, Delta1, E-cadherein, N-cadherin, HLA-A2, IFNaR2, IL1R1, IL1R2, IL6R or ameloid precursor protein (APP).

37. The method of embodiment 36, wherein the one or more target of the gamma secretase inhibitor comprises Muc1.

38. The method of embodiment 36, wherein the one or more target of the gamma secretase inhibitor comprises BCMA.

39. The method of any of embodiments 36-38, wherein the gamma secretase inhibitor inhibits the target with a half-maximal inhibitory concentration (IC_{50}) of 0.01 nM to 1 μM, 0.01 nM to 100 nM, 0.01 nM to 10 nM, 0.01 nM to 5 nM, 0.01 nM to 1 nM, 0.01 nM to 0.5 nM, 0.01 nM to 0.25 nM, 0.01 nM to 0.1 nM, 0.01 nM to 0.05 nM, 0.05 nM to 1 μM, 0.05 nM to 100 nM, 0.05 nM to 10 nM, 0.05 nM to 5 nM, 0.05 nM to 1 nM, 0.05 nM to 0.5 nM, 0.05 nM to 0.25 nM, 0.05 nM to 0.1 nM, 0.1 nM to 1 μM, 0.1 nM to 100 nM, 0.1 nM to 10 nM, 0.1 nM to 5 nM, 0.1 nM to 1 nM, 0.1 nM to 0.5 nM, 0.1 nM to 0.25 nM, 0.25 nM to 1 μM, 0.25 nM to 100 nM, 0.25 nM to 10 nM, 0.25 nM to 5 nM, 0.25 nM to 1 nM, 0.25 nM to 0.5 nM, 0.5 nM to 1 μM, 0.5 nM to 100 nM, 0.5 nM to 10 nM, 0.5 nM to 5 nM, 0.5 nM to 1 nM, 1 nM to 1 μM, 1 nM to 100 nM, 1 nM to 10 nM, 1 nM to 5 nM, 5 nM to 1 μM, 5 nM to 100 nM, 5 nM to 10 nM, 10 nM to 1 μM, 10 nM to 100 nM or 100 nM to 1 μM.

40. The method of any of embodiments 36-39, wherein the gamma secretase inhibitor inhibits the target with a half-maximal inhibitory concentration (IC_{50}) of less than or less than about 1 μM, 100 nM, 10 nM, 5 nM, 1 nM, 0.5 nM, 0.35 nM, 0.25 nM, 0.1 nM, 0.05 nM or 0.01 nM or less.

41. The method of any of embodiments 11-40, wherein the gamma secretase inhibitor is a peptide inhibitor or non-peptide inhibitor.

42. The method of embodiment 41, wherein the gamma secretase inhibitor is a peptide inhibitor and the peptide inhibitor is selected from among peptide aldehydes derivatives, difluoroketones derivatives, hydroxyethylene dipeptide isotere derivatives, alpha-helical peptide derivatives and dipeptide analogs.

43. The method of embodiment 41, wherein the gamma secretase inhibitor is a non-peptide inhibitor and the non-peptide inhibitor is a benzodiazepines derivative or a sulfonamides derivative.

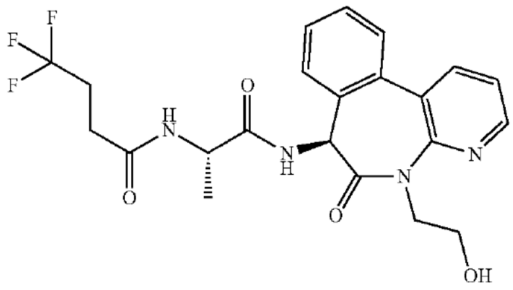
44. The method of any of embodiments 41-43, wherein the gamma secretase inhibitor is a transition state inhibitor or non-transition state inhibitor.

45. The method of any of embodiments 41-44, wherein the gamma secretase inhibitor is a nonsteroidal anti-inflammatory drug.

46. The method of any of embodiments 41-45, wherein the gamma secretase inhibitor is selected from LY3039478, secretase inhibitor I (GSI I) Z-Leu-Leu- Norleucine; γ-secretase inhibitor II (GSI II); γ- secretase inhibitor III (GSI III), N- Benzyloxycarbonyl-Leu-leucinal, N-(2-Naphthoyl)-Val- phenylalaninal; γ-secretase inhibitor III (GSI IV); γ-secretase inhibitor III (GSI V), N- Benzyloxycarbonyl-Leu- phenylalaninal; γ- secretase inhibitor III (GSI VI), l-(S)-endo-N- (1,3,3)- Trimethylbicyclo[2.2.1]hept-2-yl)-4- fluorophenyl Sulfonamide; γ-secretase inhibitor III (GSI VII), Menthyloxycarbonyl-LL-CHO; γ- secretase inhibitor III (GSI IX), (DAPT), N- [N-(3,5- Difluorophenacetyl-L- alanyl)]-S- phenylglycine t- Butyl Ester; γ-secretase inhibitor X (GSI X), { 1 S-Benzyl-4R-[1-(1S- carbamoyl- 2- phenethylcarbamoyl)-lS-3- methylbutylcarb-amoyl]-2R- hydroxy-5- phenylpentyl}carbamic Acid tert-butyl Ester; γ-secretase inhibitor XI (GSI XI), 7-Amino-4-chloro-3- methoxyisocoumarin; γ-secretase inhibitor XII (GSI XII), Z-Ile-Leu-CHO; γ-secretase inhibitor XIII (GSI XIII), Z-Tyr-Ile-Leu- CHO; γ-secretase inhibitor XIV (GSI XIV), Z-Cys(t-Bu)-Ile- Leu-CHO; γ-secretase inhibitor XVI (GSI XVI), N-[N-3,5- Difluorophenacetyl]-L- alanyl-S- phenylglycine Methyl Ester; γ- secretase inhibitor XVII (GSI XVII); γ-secretase inhibitor XIX (GSI XIX), benzo[e][l,4]diazepin-3-yl)-butyramide; γ-secretase inhibitor XX (GSI XX), (S,S)- 2-[2-(3,5- Difluorophenyl)acetylamino]-N-(5-methyl-6-oxo-6,7- dihydro-5H- dibenzo[b,d]azepin-7- yl)propionamide; γ-secretase inhibitor XXI (GSI XXI), (S,S)-2-[2-(3,5- Difluorophenyl)- acetylamino]-N-(l-methyl-2- oxo-5-phenyl-2-,3-dihydro-lH- benzo[e][l,4]diazepin-3- yl)- propionamide; Gamma40 secretase inhibitor I, N-trans-3,5- Dimethoxycinnamoyl-Ile- leucinal; Gamma40 secretase inhibitor II, N-tert-Butyloxycarbonyl- Gly-Val-Valinal Isovaleryl-V V-Sta-A-Sta- OCH3; MK-0752; MRK-003 (Merck); semagacestat/LY450139; RO4929097; PF-03084,014; BMS-708163; MPC-7869 (γ- secretase modifier), YO-01027 (Dibenzazepine), Compound E ([(2S)-2-{[(3,5-Difluorophenyl)acetyl]amino}-N-[(3S)-l-methyl-2-oxo-5-phenyl-2,3-dihydro-lH-l,4-benzodiazepin-3-yl]propanamide], LY411575, L-685,458, BMS-289948 (4-chloro-N-(2,5-difluorophenyl)-N-((lR)- {4-fluoro-2-[3-(lH-imidazol-l-yl)propyl]phenyl}ethyl)benzenesulfonamide hydrochloride) and BMS-299897 (4-[2-((lR)-l-{[(4-chlorophenyl)sulfonyl]-2,5-difluoroanilino}ethyl)-5- fluorophenyljbutanoic acid).

47. The method of any of embodiments 11-46, wherein the gamma secretase inhibitor is LY3039478 or a compound of the structure:

Compound 1



or a stereoisomer or a pharmaceutically acceptable salt or hydrate of any of the foregoing.

48. The method of any of embodiments 1-47, wherein the subject comprises plasma cells, or cancer cells or myeloma cells or cells expressing plasma cell markers, expressing surface BCMA.

49. The method of any of embodiments 21-48, wherein:

the subject has a cancer in which cells of the cancer in the subject (i) express CD138, surface CD38 or a surface plasma cell marker or are derived from plasma cells and (ii) comprise low expression of surface B cell maturation antigen (BCMA) and/or a level of expression of surface BCMA below a threshold level; and/or

the method further comprises selecting a subject that has a cancer in which cells of the cancer in the subject (i) express CD138, surface CD38 or a surface plasma cell marker or are derived from plasma cells and (ii) comprise low expression of surface B cell maturation antigen (BCMA) and/or a level of expression of surface BCMA below a threshold level.

50. The method of any of embodiments 4-8, 27-29 and 31-49, wherein the target of the gamma secretase inhibitor is the target antigen and the gamma secretase inhibitor inhibits cleavage of the target antigen.

51. The method of any of embodiments 7-23 and 34-50, wherein administration of the inhibitor:

decreases BCMA cleavage or shedding from cells, optionally plasma cells, by greater than or greater than about 5%, 10%, 20%, 30%, 40%, 50%, 60%, 70%, 80%, 90% or more

compared to the level of BCMA cleavage or shedding from cells in the subject prior to administration of the inhibitor;

decreases the level or amount of BCMA detected in the serum of a subject by greater than or greater than about 5%, 10%, 20%, 30%, 40%, 50%, 60%, 70%, 80%, 90% or more compared to the level or amount of BCMA in the serum of the subject prior to administration of the inhibitor; and/or

increases expression of surface BCMA on cells, optionally plasma cells by greater than or greater than about 5%, 10%, 20%, 30%, 40%, 50%, 60%, 70%, 80%, 90% or more compared to the level of surface BCMA on the cells in the subject prior to administration of the inhibitor.

52. The method of any of embodiments1-51, wherein administration of the inhibitor:

decreases cleavage or shedding of the target or the target antigen, optionally BCMA or Muc1, from cells, optionally plasma cells, by greater than or greater than about 5%, 10%, 20%, 30%, 40%, 50%, 60%, 70%, 80%, 90% or more compared to the level of cleavage or shedding of the target or target antigen from cells in the subject prior to administration of the inhibitor;

decreases the level or amount of the target or target antigen, optionally BCMA or Muc1, detected in the serum of a subject by greater than or greater than about 5%, 10%, 20%, 30%, 40%, 50%, 60%, 70%, 80%, 90% or more compared to the level or amount of the target or target antigen in the serum of the subject prior to administration of the inhibitor; and/or

increases expression of surface target or target antigen, optionally BCMA or Muc1, on cells, optionally plasma cells, by greater than or greater than about 5%, 10%, 20%, 30%, 40%, 50%, 60%, 70%, 80%, 90% or more compared to the level of surface target or target antigen on the cells in the subject prior to administration of the inhibitor.

53. The method of any of embodiments 1-52, wherein the disease or disorder is a cancer.

54. The method of embodiment 53, wherein the cancer is a B cell malignancy.

55. The method of embodiment 53 or embodiment 54, wherein the cancer is multiple myeloma,, plasmacytoma, a cancer of plasma cell origin and/or a cancer of B cell origin.

56. The method of any one of embodiments 1-55, wherein the inhibitor is administered orally, subcutaneously or intravenously.

57. The method of any of embodiments 1-56, wherein the inhibitor is administered orally.

58. The method of any one of embodiments 1-57, wherein the inhibitor is administered at least or is administered six times daily, five times daily, four times daily, three times daily, twice daily, once daily, every other day, three times a week, at least once a week, or only one time.

59. The method of any of embodiments 1-58, wherein the inhibitor is administered three times a week.

60. The method of embodiment 58 or embodiment 59, wherein the administration of the inhibitor is carried out in a treatment cycle that is at least or at least about or 14 days, at least or at least about or 21 days or at least or at least about or 28 days.

61. The method of any of embodiments 1-60, wherein the inhibitor is administered, or each administration of the inhibitor is independently administered, in an amount of 0.5 mg to 500 mg, 0.5 mg to 250 mg, 0.5 mg to 100 mg, 0.5 mg to 50 mg, 0.5 mg to 25 mg, 0.5 mg to 10 mg, 0.5 mg to 5.0 mg, 0.5 mg to 2.5 mg, 0.5 mg to 1.0 mg, 1.0 mg to 500 mg, 1.0 mg to 250 mg, 1.0 mg to 100 mg, 1.0 mg to 50 mg, 1.0 mg to 25 mg, 1.0 mg to 10 mg, 1.0 mg to 5.0 mg, 1.0 mg to 2.5 mg, 2.5 mg to 500 mg, 2.5 mg to 250 mg, 2.5 mg to 100 mg, 2.5 mg to 50 mg, 2.5 mg to 25 mg, 2.5 mg to 10 mg, 2.5 mg to 5.0 mg, 5.0 mg to 500 mg, 5.0 mg to 250 mg, 5.0 mg to 100 mg, 5.0 mg to 50 mg, 5.0 mg to 25 mg, 5.0 mg to 10 mg, 10 mg to 500 mg, 10 mg to 250 mg, 10 mg to 100 mg, 10 mg to 50 mg, 10 mg to 25 mg, 25 mg to 500 mg, 25 mg to 250 mg, 25 mg to 100 mg, 25 mg to 50 mg, 50 mg to 500 mg, 50 mg to 250 mg, 50 mg to 100 mg, 100 mg to 500 mg, 100 mg to 250 mg or 250 mg to 500 mg.

62. The method of any of embodiments 1-61, wherein the inhibitor is administered, or each administration of the inhibitor is independently administered, in an amount that is at least or at least about or is or is about 0.5 mg, 1.0 mg, 2.5 mg, 5.0 mg, 10.0 mg, 25 mg, 50 mg, 100 mg, 250 mg or 500 mg.

63. The method of any one of embodiments of 14-62, wherein the recombinant receptor is a transgenic T cell receptor (TCR) or a functional non-T cell receptor.

64. The method of any one of embodiments of 14-63, wherein the recombinant receptor is a chimeric receptor, which optionally is a chimeric antigen receptor (CAR).

65. The method of any of embodiments 1-13 and 64, wherein the chimeric antigen receptor (CAR) comprises an extracellular antigen-recognition domain that specifically binds to the antigen and an intracellular signaling domain comprising an ITAM.

66. The method of embodiment 65, wherein the intracellular signaling domain comprises and intracellular domain of a CD3-zeta (CD3ζ) chain.

67. The method of embodiment 65 or embodiment 66, wherein the chimeric antigen receptor (CAR) further comprises a costimulatory signaling region.

68. The method of embodiment 67, wherein the costimulatory signaling region comprises a signaling domain derived from CD28 or 4-1BB, optionally human CD28 or human 4-1BB.

69. The method of embodiment 67 or embodiment 68, wherein the costimulatory signaling region is a domain derived from 4-1BB, optionally human 4-1BB.

70. The method of any one of embodiments 1-69, wherein the subject is a human.

71. The method of any of embodiments 1-70, wherein the BCMA is human BCMA or the target antigen is a human antigen.

72. The method of any of embodiments 1-72, wherein the genetically engineered cells comprise T cells or NK cells.

73. The method of embodiment 72, wherein the cell therapy is a T cell therapy and the dose of genetically engineered cells comprises T cells.

74. The method of embodiment 72 or embodiment 73, wherein the T cells are CD4+ and/or CD8+.

75. The method of any of the embodiments 72-74, wherein the T cells are primary T cells obtained from a subject.

76. The method of any of embodiments 1-75, wherein the cell therapy comprises cells that are autologous to the subject.

77. The method of any of embodiments 1-76, wherein the cell therapy comprises cells that are allogeneic to the subject.

78. The method of any of embodiments 1-77, wherein the cell therapy comprises the administration of from or from about 1×10^5 to 5×10^8 total recombinant receptor-expressing cells, total T cells, or total peripheral blood mononuclear cells (PBMCs), the cell therapy comprises the administration of from or from about 1×10^5 to 1×10^8 total recombinant

receptor-expressing cells, total T cells, or total peripheral blood mononuclear cells (PBMCs), from or from about 5 x 10^5 to 1 x 10^7 total recombinant receptor-expressing cells, total T cells, or total peripheral blood mononuclear cells (PBMCs) or from or from about 1 x 10^6 to 1 x 10^7 total recombinant receptor-expressing cells, total T cells, or total peripheral blood mononuclear cells (PBMCs), each inclusive.

79. The method of any of embodiments 1-78, wherein the cell therapy comprises the administration of no more than 5 x 10^8 total recombinant receptor-expressing cells, total T cells, or total peripheral blood mononuclear cells (PBMCs), no more than 2.5 x 10^8 total recombinant receptor-expressing cells, total T cells, or total peripheral blood mononuclear cells (PBMCs), no more than 1 x 10^8 total recombinant receptor-expressing cells, total T cells, or total peripheral blood mononuclear cells (PBMCs), no more than 1 x 10^7 total recombinant receptor-expressing cells, total T cells, or total peripheral blood mononuclear cells (PBMCs), no more than 0.5 x 10^7 total recombinant receptor-expressing cells, total T cells, or total peripheral blood mononuclear cells (PBMCs), no more than 1 x 10^6 total recombinant receptor-expressing cells, total T cells, or total peripheral blood mononuclear cells (PBMCs), no more than 0.5 x 10^6 total recombinant receptor-expressing cells, total T cells, or total peripheral blood mononuclear cells (PBMCs).

80. The method of any one of embodiments 1, 4, 5, 14, and 38, wherein the initiation of administration of the inhibitor is prior to, concurrently with or subsequently to initiation of administration of the cell therapy.

81. The method of embodiment 80, wherein the inhibitor is administered prior to initiation of administration of the cell therapy.

82. The method of 1, 2, 4, 5, 14, 15, 38, 39 and 81, wherein initiation of administration of the inhibitor is within, or within about, 1 hours, 2 hour, 6 hours, 12 hours, 24 hours, 48 hours, 72 hours, 96 hours or 1 week prior to the initiation of the administration of the cell therapy.

83. The method of embodiment 80, wherein the inhibitor is administered subsequently to initiation of administration of the cell therapy.

84. The method of 1, 3, 4, 5, 14, 16, 19, 20, 38, 40 and 83, wherein initiation of administration of the inhibitor is within, or within about, 1 hours, 2 hour, 6 hours, 12 hours, 24 hours, 48 hours, 72 hours, 96 hours,1 week, 14 days, 21 days, 28 days or more after the initiation of the administration of the cell therapy.

85. The method of any of embodiments 1, 3, 4, 5, 14, 16, 19, 20, 38, 40, 83 and 84, wherein the inhibitor is administered at a time in which:

the number of cells of the cell therapy detectable in the blood from the subject is decreased compared to in the subject at a preceding time point after initiation of the administration of the cells;

the number of cells of the cell therapy detectable in the blood is less than or less than about 1.5-fold, 2-fold, 3-fold, 4-fold, 5-fold, 10-fold, 50-fold or 100-fold or less the peak or maximum number of the cells of the cell therapy detectable in the blood of the subject after initiation of administration of the administration of the cells; and/or

at a time after a peak or maximum level of the cells of the cell therapy are detectable in the blood of the subject, the number of cells of or derived from the cells detectable in the blood from the subject is less than less than 10%, less than 5%, less than 1% or less than 0.1% of total peripheral blood mononuclear cells (PBMCs) in the blood of the subject.

86. The method of any of embodiments 1, 3, 4, 5, 14, 16, 19, 20, 38, 40, 83-854, wherein the inhibitor is administered up to 7 days, 14 days, 21 days, 28 days or more after initiation of the administration of the cells.

87. The method of any of embodiments 1-86, wherein the method further comprises administering a lymphodepleting chemotherapy prior to administration of the cell therapy and/or wherein the subject has received a lymphodepleting chemotherapy prior to administration of the cells.

88. The method of embodiment 87, wherein the lymphodepleting chemotherapy comprises administering fludarabine and/or cyclophosphamide to the subject.

89. The method of any of embodiments 1-88, wherein the method further comprises administering a steroid, optionally wherein the steroid is administered prior to, concurrently with and/or subsequently to initiation of administration of the inhibitor, optionally wherein the steroid is administered during the cycle of treatment with the inhibitor.

90. The method of embodiment 89, wherein the steroid is or comprises dexamethasone.

91. The method of any one of embodiments 1-90, wherein the cell therapy exhibits increased or prolonged expansion and/or persistence in the subject as compared to a method in

which the cell therapy is administered to the subject in the absence of the gamma secretase inhibitor.

92. The method of any one of embodiments 1-91, wherein the method thereby prevents, reduces or ameliorates one or more symptoms or outcomes of the disease or disorder.

93. A method of treatment, the method comprising:

(a) administering, to a subject having a disease or disorder, a therapeutic agent or therapy that targets or is specific for B cell maturation antigen (BCMA); and

(b) administering a gamma secretase inhibitor to the subject.

94. The method of embodiment 93, wherein the therapeutic agent or therapy is or comprises an antibody or antigen-binding fragment thereof, optionally a bispecific antibody.

95. The method of embodiment 93 or embodiment 94, wherein the therapeutic agent or therapy is a bispecific antibody that further targets or specifically binds to a T cell antigen, optionally CD2 or CD3.

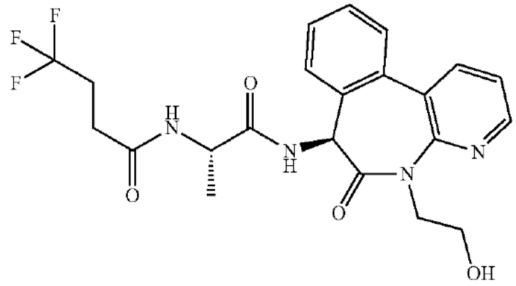
96. The method of embodiment 93 or embodiment 94, wherein the therapeutic agent or therapy is a bispecific antibody that further targets a second antigen, optionally wherein the second antigen is selected from CD19, CD20, CD22, CD33, CD38, CS1, ROR1, GPC3, CD123, IL-13R, CD138, c-Met, EGFRvIII, GD-2, NY-ESO-1, MAGE A3, and glycolipid F77.

97. The method of any of embodiments 93-96, wherein the gamma secretase inhibitor is selected from LY3039478, secretase inhibitor I (GSI I) Z-Leu-Leu- Norleucine; γ-secretase inhibitor II (GSI II); γ- secretase inhibitor III (GSI III), N- Benzyloxycarbonyl-Leu-leucinal, N-(2-Naphthoyl)-Val- phenylalaninal; γ-secretase inhibitor III (GSI IV); γ-secretase inhibitor III (GSI V), N- Benzyloxycarbonyl-Leu- phenylalaninal; γ- secretase inhibitor III (GSI VI), l-(S)-endo-N- (1,3,3)- Trimethylbicyclo[2.2.1]hept-2-yl)-4- fluorophenyl Sulfonamide; γ-secretase inhibitor III (GSI VII), Menthyloxycarbonyl-LL-CHO; γ- secretase inhibitor III (GSI IX), (DAPT), N- [N-(3,5- Difluorophenacetyl-L- alanyl)]-S- phenylglycine t- Butyl Ester; γ-secretase inhibitor X (GSI X), { 1 S-Benzyl-4R-[1-(1S- carbamoyl- 2- phenethylcarbamoyl)-lS-3- methylbutylcarb-amoyl]-2R- hydroxy-5- phenylpentyl}carbamic Acid tert-butyl Ester; γ-secretase inhibitor XI (GSI XI), 7-Amino-4-chloro-3- methoxyisocoumarin; γ-secretase inhibitor XII (GSI XII), Z-Ile-Leu-CHO; γ-secretase inhibitor XIII (GSI XIII), Z-Tyr-Ile-Leu- CHO; γ-secretase inhibitor XIV (GSI XIV), Z-Cys(t-Bu)-Ile- Leu-CHO; γ-secretase inhibitor XVI (GSI XVI), N-[N-3,5- Difluorophenacetyl]-L- alanyl-S- phenylglycine Methyl Ester; γ- secretase

inhibitor XVII (GSI XVII); γ-secretase inhibitor XIX (GSI XIX), benzo[e][l,4]diazepin-3-yl)-butyramide; γ-secretase inhibitor XX (GSI XX), (S,S)- 2-[2-(3,5- Difluorophenyl)acetylamino]-N-(5-methyl-6-oxo-6,7- dihydro-5H- dibenzo[b,d]azepin-7- yl)propionamide; γ-secretase inhibitor XXI (GSI XXI), (S,S)-2-[2-(3,5- Difluorophenyl)- acetylamino]-N-(l-methyl-2- oxo-5-phenyl-2-,3-dihydro-lH- benzo[e][l,4]diazepin-3- yl)- propionamide; Gamma40 secretase inhibitor I, N-trans-3,5- Dimethoxycinnamoyl-Ile- leucinal; Gamma40 secretase inhibitor II, N-tert-Butyloxycarbonyl- Gly-Val-Valinal Isovaleryl-V V-Sta-A-Sta- OCH3; MK-0752; MRK-003 (Merck); semagacestat/LY450139; RO4929097; PF-03084,014; BMS-708163; MPC-7869 (γ- secretase modifier), YO-01027 (Dibenzazepine), Compound E ([(2S)-2-{[(3,5-Difluorophenyl)acetyl]amino}-N-[(3S)-l-methyl-2-oxo-5-phenyl-2,3-dihydro-lH-l,4-benzodiazepin-3-yl]propanamide], LY411575, L-685,458, BMS-289948 (4-chloro-N-(2,5-difluorophenyl)-N-((lR)- {4-fluoro-2-[3-(lH-imidazol-l-yl)propyl]phenyl}ethyl)benzenesulfonamide hydrochloride) and BMS-299897 (4-[2-((lR)-l-{[(4-chlorophenyl)sulfonyl]-2,5-difluoroanilino}ethyl)-5- fluorophenyljbutanoic acid).

98. The method of any of embodiments 93-97, wherein the gamma secretase inhibitor is LY3039478 or a compound of the structure:

Compound 1



or a stereoisomer or a pharmaceutically acceptable salt or hydrate of any of the foregoing.

99. A combination, comprising:

(a) genetically engineered cells expressing a chimeric antigen receptor (CAR) comprising an antigen binding domain that specifically binds to surface B cell maturation antigen (BCMA) ; and

(b) an inhibitor of gamma secretase,

wherein binding of the CAR to surface BCMA or a measure indicative of function or activity of the CAR-expressing cells, following exposure to cells expressing surface BCMA, is not reduced or blocked or is not substantially reduced or blocked in the presence of a soluble or shed form of the BCMA,

optionally at a concentration or amount of the soluble or shed form of the BCMA corresponding to a concentration or amount present in serum or blood or plasma of the subject or of a multiple myeloma patient, or on average in a patient population for the disease or disorder, or at a concentration or amount of the soluble or shed BCMA at which the binding or measure is reduced or blocked, or is substantially reduced or blocked, for cells expressing a reference anti-BCMA CAR in the same assay.

100. The combination of embodiment 99, wherein the gamma secretase inhibitor inhibits or reduces or is capable of inhibiting or reducing intramembrane cleavage of BCMA.

101. The combination of embodiment 100, wherein the gamma secretase inhibitor inhibits intramembrane cleavage of BCMA with a half-maximal inhibitory concentration (IC_{50}) of 0.01 nM to 10 nM, 0.01 nM to 5 nM, 0.01 nM to 1 nM, 0.01 nM to 0.5 nM, 0.01 nM to 0.35 nM, 0.01 nM to 0.25 nM, 0.01 nM to 0.1 nM, 0.01 nM to 0.05 nM, 0.05 nM to 10 nM, 0.05 nM to 5 nM, 0.05 nM to 1 nM, 0.05 nM to 0.5 nM, 0.05 nM to 0.35 nM, 0.05 nM to 0.25 nM, 0.05 nM to 0.1 nM, 0.1 nM to 10 nM, 0.1 nM to 5 nM, 0.1 nM to 1 nM, 0.1 nM, to 0.5 nM, 0.1 nM to 0.35 nM, 0.1 nM to 0.25 nM, 0.25 nM to 10 nM, 0.25 nM to 5 nM, 0.25 nM to 1 nM, 0.25 nM to 0.5 nM, 0.25 nM to 0.35 nM, 0.35 nM to 10 nM, 0.35 nM to 5 nM, 0.35 nM to 1 nM, 0.35 nM to 0.5 nM, 0.5 nM to 10 nM, 0.5 nM to 5 nM, 0.5 nM to 1 nM, 1 nM to 10 nM, 1 nM to 5 nM or 5 nM to 10 nM.

102. The combination of embodiment 100 or embodiment 101, wherein the gamma secretase inhibitor inhibits intramembrane cleavage of BCMA with a half-maximal inhibitory concentration (IC_{50}) of less than or less than about 10 nM, 5 nM, 1 nM, 0.5 nM, 0.35 nM, 0.25 nM, 0.1 nM, 0.05 nM or 0.01 nM or less.

103. A combination, comprising:

(a) genetically engineered cells expressing a recombinant receptor, wherein the recombinant receptor does not specifically binds to surface B cell maturation antigen (BCMA); and

(b) an inhibitor of gamma secretase.

104. The method of embodiment 103, wherein the recombinant receptor specifically binds to a target antigen associated with the disease or disorder.

105. The combination of embodiment 104, wherein the target antigen is selected from among, carbonic anhydrase 9 (CAIX), Her2/neu (receptor tyrosine kinase erbB2), L1-CAM, CD19, CD20, CD22, mesothelin, CEA, and hepatitis B surface antigen, anti-folate receptor, CD23, CD24, CD30, CD33, CD38, CD44, EGFR, epithelial glycoprotein 2 (EPG-2), epithelial glycoprotein 40 (EPG-40), EPHa2, erb-B2, erb-B3, erb-B4, erbB dimers, EGFR vIII, folate binding protein (FBP), FCRL5, FCRH5, fetal acetylcholine receptor, GD2, GD3, HMW-MAA, IL-22R-alpha, IL-13R-alpha2, kinase insert domain receptor (kdr), kappa light chain, Lewis Y, L1-cell adhesion molecule, (L1-CAM), Melanoma-associated antigen (MAGE)-A1, MAGE-A3, MAGE-A6, Preferentially expressed antigen of melanoma (PRAME), survivin, TAG72, B7-H6, IL-13 receptor alpha 2 (IL-13Ra2), CA9, GD3, HMW-MAA, CD171, G250/CAIX, HLA-AI MAGE Al, HLA-A2 NY-ESO-1, PSCA, folate receptor-a, CD44v6, CD44v7/8, avb6 integrin, 8H9, NCAM, VEGF receptors, 5T4, Foetal AchR, NKG2D ligands, CD44v6, dual antigen, a cancer-testes antigen, mesothelin, murine CMV, mucin 1 (MUC1), MUC16, PSCA, NKG2D, NY-ESO-1, MART-1, gp100, oncofetal antigen, ROR1, TAG72, VEGF-R2, carcinoembryonic antigen (CEA), Her2/neu, estrogen receptor, progesterone receptor, ephrinB2, CD123, c-Met, GD-2, O-acetylated GD2 (OGD2), CE7, Wilms Tumor 1 (WT-1), a cyclin, cyclin A2, CCL-1, CD138, a pathogen-specific antigen and an antigen associated with a universal tag.

106. The combination of embodiment 104 or embodiment 105, wherein the target antigen is Muc1.

107. The combination of any of embodiments 99-106, wherein the gamma secretase inhibitor inhibits or reduces or is capable of inhibiting or reducing cleavage of one or more targets selected from BCMA, Notch 1, Notch 2, Notch 3, Notch 4, Muc1, Ephrin B2, Betaglycan (TGFBR3), CD43, CD44, CSF1R, CXCR1, CXCL16, Delta1, E-cadherein, N-cadherin, HLA-A2, IFNaR2, IL1R1, IL1R2, IL6R or ameloid precursor protein (APP).

108. The combination of embodiment 107, wherein the one or more target of the gamma secretase inhibitor comprises Muc1.

109. The combination of embodiment 107, wherein the one or more target of the gamma secretase inhibitor comprises BCMA.

110. The combination of any of embodiments 103-109, wherein the gamma secretase inhibitor inhibits the target with a half-maximal inhibitory concentration (IC_{50}) of 0.01 nM to 1 μM, 0.01 nM to 100 nM, 0.01 nM to 10 nM, 0.01 nM to 5 nM, 0.01 nM to 1 nM, 0.01 nM to 0.5 nM, 0.01 nM to 0.25 nM, 0.01 nM to 0.1 nM, 0.01 nM to 0.05 nM, 0.05 nM to 1 μM, 0.05 nM to 100 nM, 0.05 nM to 10 nM, 0.05 nM to 5 nM, 0.05 nM to 1 nM, 0.05 nM to 0.5 nM, 0.05 nM to 0.25 nM, 0.05 nM to 0.1 nM, 0.1 nM to 1 μM, 0.1 nM to 100 nM, 0.1 nM to 10 nM, 0.1 nM to 5 nM, 0.1 nM to 1 nM, 0.1 nM to 0.5 nM, 0.1 nM to 0.25 nM, 0.25 nM to 1 μM, 0.25 nM to 100 nM, 0.25 nM to 10 nM, 0.25 nM to 5 nM, 0.25 nM to 1 nM, 0.25 nM to 0.5 nM, 0.5 nM to 1 μM, 0.5 nM to 100 nM, 0.5 nM to 10 nM, 0.5 nM to 5 nM, 0.5 nM to 1 nM, 1 nM to 1 μM, 1 nM to 100 nM, 1 nM to 10 nM, 1 nM to 5 nM, 5 nM to 1 μM, 5 nM to 100 nM, 5 nM to 10 nM, 10 nM to 1 μM, 10 nM to 100 nM or 100 nM to 1 μM.

111. The combination of any of embodiments 103-110, wherein the gamma secretase inhibitor inhibits the target with a half-maximal inhibitory concentration (IC_{50}) of less than or less than about 1 μM, 100 nM, 10 nM, 5 nM, 1 nM, 0.5 nM, 0.35 nM, 0.25 nM, 0.1 nM, 0.05 nM or 0.01 nM or less.

112. The combination of any of embodiments 99-111, wherein the gamma secretase inhibitor is a peptide inhibitor or non-peptide inhibitor.

113. The combination of embodiment 112, wherein the gamma secretase inhibitor is a peptide inhibitor and the peptide inhibitor is selected from among peptide aldehydes derivatives, difluoroketones derivatives, hydroxyethylene dipeptide isotere derivatives, alpha-helical peptide derivatives and dipeptide analogs.

114. The combination of embodiment 113, wherein the gamma secretase inhibitor is a non-peptide inhibitor and the non-peptide inhibitor is a benzodiazepines derivative or a sulfonamides derivative.

115. The combination of any of embodiments 99-114, wherein the gamma secretase inhibitor is a transition state inhibitor or non-transition state inhibitor.

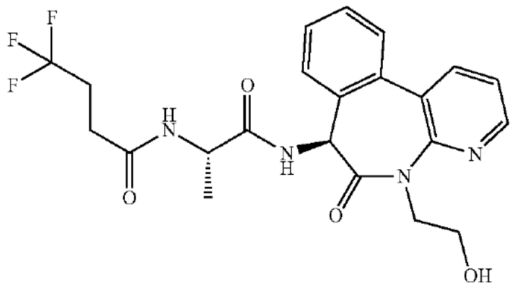
116. The combination of any of embodiments 99-115, wherein the gamma secretase inhibitor is a nonsteroidal anti-inflammatory drug.

117. The combination of any of embodiments 99-116, wherein the gamma secretase inhibitor is selected from LY3039478, secretase inhibitor I (GSI I) Z-Leu-Leu- Norleucine; γ-secretase inhibitor II (GSI II); γ- secretase inhibitor III (GSI III), N- Benzyloxycarbonyl-Leu-

leucinal, N-(2-Naphthoyl)-Val- phenylalaninal; γ-secretase inhibitor III (GSI IV); γ-secretase inhibitor III (GSI V), N- Benzyloxycarbonyl-Leu- phenylalaninal; γ- secretase inhibitor III (GSI VI), l-(S)-endo-N- (1,3,3)- Trimethylbicyclo[2.2.1]hept-2-yl)-4- fluorophenyl Sulfonamide; γ-secretase inhibitor III (GSI VII), Menthyloxycarbonyl-LL-CHO; γ- secretase inhibitor III (GSI IX), (DAPT), N- [N-(3,5- Difluorophenacetyl-L- alanyl)]-S- phenylglycine t- Butyl Ester; γ-secretase inhibitor X (GSI X), { 1 S-Benzyl-4R-[1-(1S- carbamoyl- 2- phenethylcarbamoyl)-lS-3- methylbutylcarb-amoyl]-2R- hydroxy-5- phenylpentyl}carbamic Acid tert-butyl Ester; γ-secretase inhibitor XI (GSI XI), 7-Amino-4-chloro-3- methoxyisocoumarin; γ-secretase inhibitor XII (GSI XII), Z-Ile-Leu-CHO; γ-secretase inhibitor XIII (GSI XIII), Z-Tyr-Ile-Leu- CHO; γ-secretase inhibitor XIV (GSI XIV), Z-Cys(t-Bu)-Ile- Leu-CHO; γ-secretase inhibitor XVI (GSI XVI), N-[N-3,5- Difluorophenacetyl]-L- alanyl-S- phenylglycine Methyl Ester; γ- secretase inhibitor XVII (GSI XVII); γ-secretase inhibitor XIX (GSI XIX), benzo[e][l,4]diazepin-3-yl)-butyramide; γ-secretase inhibitor XX (GSI XX), (S,S)- 2-[2-(3,5- Difluorophenyl)acetylamino]-N-(5-methyl-6-oxo-6,7- dihydro-5H- dibenzo[b,d]azepin-7- yl)propionamide; γ-secretase inhibitor XXI (GSI XXI), (S,S)-2-[2-(3,5- Difluorophenyl)- acetylamino]-N-(l-methyl-2- oxo-5-phenyl-2-,3-dihydro-lH- benzo[e][l,4]diazepin-3- yl)- propionamide; Gamma40 secretase inhibitor I, N-trans-3,5- Dimethoxycinnamoyl-Ile- leucinal; Gamma40 secretase inhibitor II, N-tert-Butyloxycarbonyl- Gly-Val-Valinal Isovaleryl-V V-Sta-A-Sta- OCH3; MK-0752; MRK-003 (Merck); semagacestat/LY450139; RO4929097; PF-03084,014; BMS-708163; MPC-7869 (γ- secretase modifier), YO-01027 (Dibenzazepine), Compound E ([(2S)-2-{[(3,5-Difluorophenyl)acetyl]amino}-N-[(3S)-l-methyl-2-oxo-5-phenyl-2,3-dihydro-lH-l,4-benzodiazepin-3-yl]propanamide], LY411575, L-685,458, BMS-289948 (4-chloro-N-(2,5-difluorophenyl)-N-((lR)- {4-fluoro-2-[3-(lH-imidazol-l-yl)propyl]phenyl}ethyl)benzenesulfonamide hydrochloride) and BMS-299897 (4-[2-((lR)-l-{[(4-chlorophenyl)sulfonyl]-2,5-difluoroanilino}ethyl)-5- fluorophenyljbutanoic acid).

118. The combination of any of embodiments 99-117, wherein the gamma secretase inhibitor is LY3039478 or a compound of the structure:

Compound 1



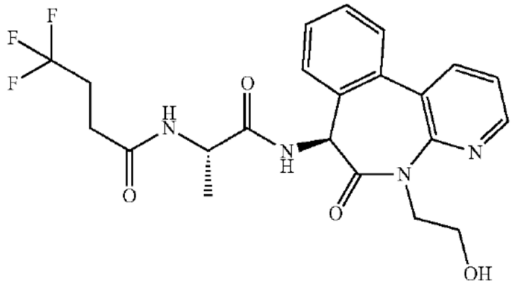
or a stereoisomer or a pharmaceutically acceptable salt or hydrate of any of the foregoing.

119. A combination, comprising:

(a) genetically engineered cells expressing a recombinant receptor; and

(b) LY3039478 or a compound of the structure:

Compound 1



or a stereoisomer or a pharmaceutically acceptable salt or hydrate of any of the foregoing.

120. The combination of embodiment 119, wherein the recombinant receptor specifically binds to a target antigen associated with the disease or disorder.

121. The combination of embodiment 120, wherein the target antigen is selected from among, B cell maturation antigen (BCMA), carbonic anhydrase 9 (CAIX) , Her2/neu (receptor tyrosine kinase erbB2), L1-CAM, CD19, CD20, CD22, mesothelin, CEA, and hepatitis B surface antigen, anti-folate receptor, CD23, CD24, CD30, CD33, CD38, CD44, EGFR, epithelial glycoprotein 2 (EPG-2), epithelial glycoprotein 40 (EPG-40), EPHa2, erb-B2, erb-B3, erb-B4, erbB dimers, EGFR vIII, folate binding protein (FBP), FCRL5, FCRH5, fetal acetylcholine receptor, GD2, GD3, HMW-MAA, IL-22R-alpha, IL-13R-alpha2, kinase insert domain receptor (kdr), kappa light chain, Lewis Y, L1-cell adhesion molecule, (L1-CAM),

Melanoma-associated antigen (MAGE)-A1, MAGE-A3, MAGE-A6, Preferentially expressed antigen of melanoma (PRAME), survivin, TAG72, B7-H6, IL-13 receptor alpha 2 (IL-13Ra2), CA9, GD3, HMW-MAA, CD171, G250/CAIX, HLA-AI MAGE Al, HLA-A2 NY-ESO-1, PSCA, folate receptor-a, CD44v6, CD44v7/8, avb6 integrin, 8H9, NCAM, VEGF receptors, 5T4, Foetal AchR, NKG2D ligands, CD44v6, dual antigen, a cancer-testes antigen, mesothelin, murine CMV, mucin 1 (MUC1), MUC16, PSCA, NKG2D, NY-ESO-1, MART-1, gp100, oncofetal antigen, ROR1, TAG72, VEGF-R2, carcinoembryonic antigen (CEA), Her2/neu, estrogen receptor, progesterone receptor, ephrinB2, CD123, c-Met, GD-2, O-acetylated GD2 (OGD2), CE7, Wilms Tumor 1 (WT-1), a cyclin, cyclin A2, CCL-1, CD138, a pathogen-specific antigen and an antigen associated with a universal tag.

122. The combination of embodiment 120 or embodiment 121, wherein the target antigen is Muc1.

123. The combination of embodiment 120 or embodiment 121, wherein the target antigen is BCMA.

124. The combination of embodiment 123, wherein the BCMA is surface BCMA.

125. The combination of embodiment 124, wherein binding of the recombinant receptor, optionally the CAR, to surface BCMA or a measure indicative of function or activity of the recombinant receptor-expressing cells, optionally CAR-expressing cells, following exposure to cells expressing surface BCMA, is not reduced or blocked or is not substantially reduced or blocked in the presence of a soluble or shed form of the BCMA, optionally at a concentration or amount of the soluble or shed form of the BCMA corresponding to a concentration or amount present in serum or blood or plasma of the subject or of a multiple myeloma patient, or on average in a patient population for the disease or disorder, or at a concentration or amount of the soluble or shed BCMA at which the binding or measure is reduced or blocked, or is substantially reduced or blocked, for cells expressing a reference anti-BCMA recombinant receptor, optionally, anti-BCMA CAR, in the same assay.

126. The combination of any of embodiments 103-125, wherein the recombinant receptor is a transgenic T cell receptor (TCR) or a functional non-T cell receptor.

127. The combination of any of embodiments 103-126, wherein the recombinant receptor is a chimeric receptor, which optionally is a chimeric antigen receptor (CAR).

128. The combination of any of embodiments 99-102 and 127, wherein the chimeric antigen receptor (CAR) comprises an extracellular antigen-recognition domain that specifically binds to the antigen and an intracellular signaling domain comprising an ITAM.

129. The combination of embodiment 128, wherein the intracellular signaling domain comprises and intracellular domain of a CD3-zeta (CD3ζ) chain.

130. The combination of embodiment 128 or embodiment 129, wherein the chimeric antigen receptor (CAR) further comprises a costimulatory signaling region.

131. The combination of embodiment 130, wherein the costimulatory signaling region comprises a signaling domain derived from CD28 or 4-1BB, optionally human CD28 or human 4-1BB.

132. The combination of embodiment 130 or embodiment 131, wherein the costimulatory signaling region is a domain derived from 4-1BB, optionally human 4-1BB.

133. The combination of any of embodiments 99-132, wherein the BCMA is human BCMA or the target antigen is a human antigen.

134. The combination of any of embodiments 99-134, wherein the genetically engineered cells comprise T cells or NK cells.

135. The combination of embodiment 134, wherein the genetically engineered cells comprises T cells.

136. The combination of embodiment 134 or embodiment 135, wherein the T cells are CD4+ and/or CD8+.

137. The combination of any of embodiments 134-136, wherein the T cells are primary T cells obtained from a subject.

138. The combination of any of embodiments 99-137, wherein the genetically engineered cells are formulated as a pharmaceutical composition for administration to a subject, optionally wherein the cells are formulated for administration in one or more unit doses for treating a disease or condition.

139. The combination of any of embodiments 99-138, wherein the gamma secretase inhibitor is formulated as a pharmaceutical composition for administration to a subject, optionally wherein the gamma secretase inhibitor is formulated for administration in one or more unit doses.

140. A kit comprising the combination of any of embodiments 99-139 and instructions for administering the genetically engineered cells and/or the gamma secretase inhibitor to a subject having a disease or disorder.

141. The kit of embodiment 140, further comprising a reagent for detecting expression of B cell maturation antigen (BCMA) on the surface of a cell, and instructions for administering the inhibitor to a subject based on the results of use of the reagent for detecting BCMA on the surface of cells of a cancer in the subject, optionally wherein the cells of the cancer express CD138, surface CD38 or a surface plasma cell marker or are derived from plasma cells.

142. The kit of embodiment 141, wherein the instructions specify administering the inhibitor to the subject if the cells comprise low expression of surface BCMA and/or a level of expression of surface BCMA below a threshold level.

143. The kit of embodiment 142, wherein the threshold level of expression of surface BCMA is lower than the average or median expression or level of surface BCMA on plasma cells in a plurality of control subjects, optionally wherein the control subjects are a group of healthy or normal subjects.

144. The kit of embodiment 142 or embodiment 143, wherein;

the low expression of surface BCMA is present when less than or less than about 60%, less than or less than about 50%, less than or less than about 40%, less than or less than about 30%, less than or less than about 20% or less than or less than about 10% of the plasma cells, or cells with a plasma marker or phenotype or the cancer cells, in the subject express surface BCMA; or

the threshold level of surface BCMA is less than or less than about 60%, less than or less than about 50%, less than or less than about 40%, less than or less than about 30%, less than or less than about 20% or less than or less than about 10% of the plasma cells, or cells with a plasma marker or phenotype or the cancer cells, in the subject that express surface BCMA.

145. The kit of any of embodiments 141-144, further comprising genetically engineered cells expressing a recombinant receptor, optionally wherein the genetically engineered cells are formulated for administration of one or more unit doses to a subject having a disease or conditon.

146. The kit of embodiment 145, wherein the recombinant receptor specifically binds to a target antigen associated with the disease or condition.

147. The kit of any of embodiments 140, 145 and 146, wherein the instructions specify administering the gamma secretase inhibitor, or one or more unit doses thereof, to a subject having a disease or disorder prior to, concurrently with or after initiation of administration of a dose of the genetically engineered cells to the subject.

148. The kit of embodiment 147, wherein the instructions specify administering the gamma secretase inhibitor, or one or more unit doses thereof, to a subject having a disease or disorder prior to initiation of administration of a dose of the genetically engineered cells to the subject.

149. The kit of embodiment 148, wherein the instructions specify administering the inhibitor within, or within about, 1 hours, 2 hour, 6 hours, 12 hours, 24 hours, 48 hours, 72 hours, 96 hours or 1 week prior to the initiation of the administration of the cell therapy.

150. The kit of embodiment 147, wherein the instructions specify administering the gamma secretase inhibitor, or one or more unit doses thereof, to a subject having a disease or disorder after initiation of administration of a dose of the genetically engineered cells to the subject.

151. A kit, comprising:

(a) a gamma secretase inhibitor, optionally wherein the gamma secretase inhibitor is formulated in one or more unit doses; and

(b) instructions for administering the gamma secretase inhibitor to a subject after initiation of administration of a cell therapy to a subject, the cell therapy comprising a dose of genetically engineered cells expressing a recombinant receptor.

152. The kit of embodiment 150 or embodiment 151, wherein initiation of administration of the inhibitor is within, or within about, 1 hours, 2 hour, 6 hours, 12 hours, 24 hours, 48 hours, 72 hours, 96 hours,1 week, 14 days, 21 days, 28 days or more after the initiation of the administration of the dose of genetically engineered cells.

153. The kit of any of embodiments 150-152, wherein the instructions specify the inhibitor is for administration at a time in which:

the number of cells of the cell therapy detectable in the blood from the subject is decreased compared to in the subject at a preceding time point after initiation of the administration of the cells;

the number of cells of the cell therapy detectable in the blood is less than or less than about 1.5-fold, 2-fold, 3-fold, 4-fold, 5-fold, 10-fold, 50-fold or 100-fold or less the peak or maximum number of the cells of the cell therapy detectable in the blood of the subject after initiation of administration of the administration of the cells; and/or

at a time after a peak or maximum level of the cells of the cell therapy are detectable in the blood of the subject, the number of cells of or derived from the cells detectable in the blood from the subject is less than less than 10%, less than 5%, less than 1% or less than 0.1% of total peripheral blood mononuclear cells (PBMCs) in the blood of the subject.

154. The kit of any of embodiments150-153, wherein the instructions specify the inhibitor is for administration up to 7 days, 14 days, 21 days, 28 days or more after initiation of the administration of the cells.

155. The kit of any of embodiments 150-154, wherein the recombinant receptor specifically binds to a target antigen associated with the disease or disorder.

156. The kit of embodiment 146 or embodiment 155, wherein the target antigen is selected from among, B cell maturation antigen (BCMA), carbonic anhydrase 9 (CAIX), Her2/neu (receptor tyrosine kinase erbB2), L1-CAM, CD19, CD20, CD22, mesothelin, CEA, and hepatitis B surface antigen, anti-folate receptor, CD23, CD24, CD30, CD33, CD38, CD44, EGFR, epithelial glycoprotein 2 (EPG-2), epithelial glycoprotein 40 (EPG-40), EPHa2, erb-B2, erb-B3, erb-B4, erbB dimers, EGFR vIII, folate binding protein (FBP), FCRL5, FCRH5, fetal acetylcholine receptor, GD2, GD3, HMW-MAA, IL-22R-alpha, IL-13R-alpha2, kinase insert domain receptor (kdr), kappa light chain, Lewis Y, L1-cell adhesion molecule, (L1-CAM), Melanoma-associated antigen (MAGE)-A1, MAGE-A3, MAGE-A6, Preferentially expressed antigen of melanoma (PRAME), survivin, TAG72, B7-H6, IL-13 receptor alpha 2 (IL-13Ra2), CA9, GD3, HMW-MAA, CD171, G250/CAIX, HLA-AI MAGE Al, HLA-A2 NY-ESO-1, PSCA, folate receptor-a, CD44v6, CD44v7/8, avb6 integrin, 8H9, NCAM, VEGF receptors, 5T4, Foetal AchR, NKG2D ligands, CD44v6, dual antigen, a cancer-testes antigen, mesothelin, murine CMV, mucin 1 (MUC1), MUC16, PSCA, NKG2D, NY-ESO-1, MART-1, gp100, oncofetal antigen, ROR1, TAG72, VEGF-R2, carcinoembryonic antigen (CEA), Her2/neu,

estrogen receptor, progesterone receptor, ephrinB2, CD123, c-Met, GD-2, O-acetylated GD2 (OGD2), CE7, Wilms Tumor 1 (WT-1), a cyclin, cyclin A2, CCL-1, CD138, a pathogen-specific antigen and an antigen associated with a universal tag.

157. The kit of embodiment 156, wherein the target antigen is Muc1, optionally human Muc1.

158. The kit of embodiment156, wherein the target antigen is BCMA, optionally human BCMA.

159. The kit of embodiment 158, wherein wherein binding of the recombinant receptor, optionally the CAR, to surface BCMA or a measure indicative of function or activity of the recombinant receptor-expressing cells, optionally CAR-expressing cells, following exposure to cells expressing surface BCMA, is not reduced or blocked or is not substantially reduced or blocked in the presence of a soluble or shed form of the BCMA, optionally at a concentration or amount of the soluble or shed form of the BCMA corresponding to a concentration or amount present in serum or blood or plasma of the subject or of a multiple myeloma patient, or on average in a patient population for the disease or disorder, or at a concentration or amount of the soluble or shed BCMA at which the binding or measure is reduced or blocked, or is substantially reduced or blocked, for cells expressing a reference anti-BCMA recombinant receptor, optionally, anti-BCMA CAR, in the same assay.

160. The kit of any of embodiments 146-159, wherein the recombinant receptor is a transgenic T cell receptor (TCR) or a functional non-T cell receptor.

161. The kit of any of embodiments 146-160, wherein the recombinant receptor is a chimeric receptor, which optionally is a chimeric antigen receptor (CAR).

162. The kit of embodiment 161, wherein the chimeric antigen receptor (CAR) comprises an extracellular antigen-recognition domain that specifically binds to the antigen and an intracellular signaling domain comprising an ITAM.

163. The kit of embodiment 162, wherein the intracellular signaling domain comprises and intracellular domain of a CD3-zeta (CD3ζ) chain.

164. The kit of any of embodiments 161-163, wherein the chimeric antigen receptor (CAR) further comprises a costimulatory signaling region.

165. The kit of embodiment 164, wherein the costimulatory signaling region comprises a signaling domain derived from CD28 or 4-1BB, optionally human CD28 or human 4-1BB.

166. The kit of embodiment 164 or embodiment 165, wherein the costimulatory signaling region is a domain derived from 4-1BB, optionally human 4-1BB.

167. The kit of any of embodiments 146-166, wherein the genetically engineered cells comprise T cells or NK cells.

168. The kit of embodiment 167, wherein the genetically engineered cells comprises T cells.

169. The kit of embodiment 167 or embodiment 168, wherein the T cells are CD4+ and/or CD8+.

170. The kit of any of embodiments 167-169, wherein the T cells are primary T cells obtained from a subject.

171. The kit of any of embodiments 141-170, wherein the gamma secretase inhibitor inhibits cleavage of one or more targets selected from BCMA, Notch 1, Notch 2, Notch 3, Notch 4, Muc1, Ephrin B2, Betaglycan (TGFBR3), CD43, CD44, CSF1R, CXCR1, CXCL16, Delta1, E-cadherein, N-cadherin, HLA-A2, IFNaR2, IL1R1, IL1R2, IL6R or ameloid precursor protein (APP).

172. The kit of embodiment 171, wherein the one or more target of the gamma secretase inhibitor comprises Muc1.

173. The kit of embodiment 171, wherein the one or more target of the gamma secretase inhibitor comprises BCMA.

174. The kit of any of embodiments 141-173, wherein the gamma secretase inhibitor inhibits the target with a half-maximal inhibitory concentration (IC_{50}) of 0.01 nM to 1 μM, 0.01 nM to 100 nM, 0.01 nM to 10 nM, 0.01 nM to 5 nM, 0.01 nM to 1 nM, 0.01 nM to 0.5 nM, 0.01 nM to 0.25 nM, 0.01 nM to 0.1 nM, 0.01 nM to 0.05 nM, 0.05 nM to 1 μM, 0.05 nM to 100 nM, 0.05 nM to 10 nM, 0.05 nM to 5 nM, 0.05 nM to 1 nM, 0.05 nM to 0.5 nM, 0.05 nM to 0.25 nM, 0.05 nM to 0.1 nM, 0.1 nM to 1 μM, 0.1 nM to 100 nM, 0.1 nM to 10 nM, 0.1 nM to 5 nM, 0.1 nM to 1 nM, 0.1 nM to 0.5 nM, 0.1 nM to 0.25 nM, 0.25 nM to 1 μM, 0.25 nM to 100 nM, 0.25 nM to 10 nM, 0.25 nM to 5 nM, 0.25 nM to 1 nM, 0.25 nM to 0.5 nM, 0.5 nM to 1 μM, 0.5 nM to 100 nM, 0.5 nM to 10 nM, 0.5 nM to 5 nM, 0.5 nM to 1 nM, 1 nM to 1 μM, 1

nM to 100 nM, 1 nM to 10 nM, 1 nM to 5 nM, 5 nM to 1 μM, 5 nM to 100 nM, 5 nM to 10 nM, 10 nM to 1 μM, 10 nM to 100 nM or 100 nM to 1 μM.

175. The kit of any of embodiments 141-174, wherein the gamma secretase inhibitor inhibits the target with a half-maximal inhibitory concentration (IC_{50}) of less than or less than about 1 μM, 100 nM, 10 nM, 5 nM, 1 nM, 0.5 nM, 0.35 nM, 0.25 nM, 0.1 nM, 0.05 nM or 0.01 nM or less.

176. The kit of any of embodiments 141-175, wherein the gamma secretase inhibitor is a peptide inhibitor or non-peptide inhibitor.

177. The kit of embodiment 176, wherein the gamma secretase inhibitor is a peptide inhibitor and the peptide inhibitor is selected from among peptide aldehydes derivatives, difluoroketones derivatives, hydroxyethylene dipeptide isotere derivatives, alpha-helical peptide derivatives and dipeptide analogs.

178. The kit of embodiment 177, wherein the gamma secretase inhibitor is a non-peptide inhibitor and the non-peptide inhibitor is a benzodiazepines derivative or a sulfonamides derivative.

179. The kit of any of embodiments 141-178, wherein the gamma secretase inhibitor is a transition state inhibitor or non-transition state inhibitor.

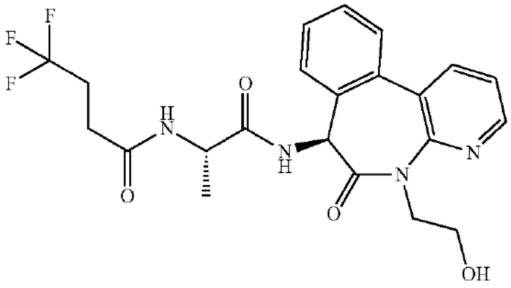
180. The kit of any of embodiments 141-179, wherein the gamma secretase inhibitor is a nonsteroidal anti-inflammatory drug.

181. The kit of any of embodiments 141-180, wherein the gamma secretase inhibitor is selected from LY3039478, secretase inhibitor I (GSI I) Z-Leu-Leu- Norleucine; γ-secretase inhibitor II (GSI II); γ- secretase inhibitor III (GSI III), N- Benzyloxycarbonyl-Leu- leucinal, N-(2-Naphthoyl)-Val- phenylalaninal; γ-secretase inhibitor III (GSI IV); γ-secretase inhibitor III (GSI V), N- Benzyloxycarbonyl-Leu- phenylalaninal; γ- secretase inhibitor III (GSI VI), l-(S)-endo-N- (1,3,3)- Trimethylbicyclo[2.2.1]hept-2-yl)-4- fluorophenyl Sulfonamide; γ-secretase inhibitor III (GSI VII), Menthyloxycarbonyl-LL-CHO; γ- secretase inhibitor III (GSI IX), (DAPT), N- [N-(3,5- Difluorophenacetyl-L- alanyl)]-S- phenylglycine t- Butyl Ester; γ-secretase inhibitor X (GSI X), { 1 S-Benzyl-4R-[1-(1S- carbamoyl- 2- phenethylcarbamoyl)-lS-3-methylbutylcarb-amoyl]-2R- hydroxy-5- phenylpentyl}carbamic Acid tert-butyl Ester; γ-secretase inhibitor XI (GSI XI), 7-Amino-4-chloro-3- methoxyisocoumarin; γ-secretase inhibitor XII (GSI XII), Z-Ile-Leu-CHO; γ-secretase inhibitor XIII (GSI XIII), Z-Tyr-Ile-Leu- CHO; γ-

secretase inhibitor XIV (GSI XIV), Z-Cys(t-Bu)-Ile- Leu-CHO; γ-secretase inhibitor XVI (GSI XVI), N-[N-3,5- Difluorophenacetyl]-L- alanyl-S- phenylglycine Methyl Ester; γ- secretase inhibitor XVII (GSI XVII); γ-secretase inhibitor XIX (GSI XIX), benzo[e][l,4]diazepin-3-yl)-butyramide; γ-secretase inhibitor XX (GSI XX), (S,S)- 2-[2-(3,5- Difluorophenyl)acetylamino]-N-(5-methyl-6-oxo-6,7- dihydro-5H- dibenzo[b,d]azepin-7- yl)propionamide; γ-secretase inhibitor XXI (GSI XXI), (S,S)-2-[2-(3,5- Difluorophenyl)- acetylamino]-N-(l-methyl-2- oxo-5-phenyl-2-,3-dihydro-lH- benzo[e][l,4]diazepin-3- yl)- propionamide; Gamma40 secretase inhibitor I, N-trans-3,5- Dimethoxycinnamoyl-Ile- leucinal; Gamma40 secretase inhibitor II, N-tert-Butyloxycarbonyl- Gly-Val-Valinal Isovaleryl-V V-Sta-A-Sta- OCH3; MK-0752; MRK-003 (Merck); semagacestat/LY450139; RO4929097; PF-03084,014; BMS-708163; MPC-7869 (γ- secretase modifier), YO-01027 (Dibenzazepine), Compound E ([(2S)-2-{[(3,5-Difluorophenyl)acetyl]amino}-N-[(3S)-l-methyl-2-oxo-5-phenyl-2,3-dihydro-lH-l,4-benzodiazepin-3-yl]propanamide], LY411575, L-685,458, BMS-289948 (4-chloro-N-(2,5-difluorophenyl)-N-((lR)- {4-fluoro-2-[3-(lH-imidazol-l-yl)propyl]phenyl}ethyl)benzenesulfonamide hydrochloride) and BMS-299897 (4-[2-((lR)-l-{[(4-chlorophenyl)sulfonyl]-2,5-difluoroanilino}ethyl)-5- fluorophenyljbutanoic acid).

182. The kit of any of embodiments 141-181, wherein the gamma secretase inhibitor is LY3039478 or a compound of the structure:

Compound 1



or a stereoisomer or a pharmaceutically acceptable salt or hydrate of any of the foregoing.

183. The kit of any of embodiments 99-182, wherein the instructions specify administering a dose of genetically engineered cells to a subject having a disease or disorder.

184. The kit of embodiment 183, wherein the disease or disorder is a cancer.

185. The kit of embodiment 184, wherein the cancer is a B cell malignancy.

186. The kit of embodiment 184 or embodiment 185, wherein the cancer is multiple myeloma.

187. The kit of any of embodiments 183-186, wherein the dose comprises from or from about 1 x 10^5 to 5 x 10^8 total recombinant receptor-expressing cells, total T cells, or total peripheral blood mononuclear cells (PBMCs), the cell therapy comprises the administration of from or from about 1 x 10^5 to 1 x 10^8 total recombinant receptor-expressing cells, total T cells, or total peripheral blood mononuclear cells (PBMCs), from or from about 5 x 10^5 to 1 x 10^7 total recombinant receptor-expressing cells, total T cells, or total peripheral blood mononuclear cells (PBMCs) or from or from about 1 x 10^6 to 1 x 10^7 total recombinant receptor-expressing cells, total T cells, or total peripheral blood mononuclear cells (PBMCs), each inclusive, optionally wherein the instructions specify the administration of one or of a plurality of unit doses comprising the dose of cells and/or a volume corresponding to such one or plurality of unit doses comprising the dose of cells.

188. The kit of any of embodiments 183-186, wherein the dose comprises no more than 5 x 10^8 total recombinant receptor-expressing cells, total T cells, or total peripheral blood mononuclear cells (PBMCs), no more than 2.5 x 10^8 total recombinant receptor-expressing cells, total T cells, or total peripheral blood mononuclear cells (PBMCs), no more than 1 x 10^8 total recombinant receptor-expressing cells, total T cells, or total peripheral blood mononuclear cells (PBMCs), no more than 1 x 10^7 total recombinant receptor-expressing cells, total T cells, or total peripheral blood mononuclear cells (PBMCs), no more than 0.5 x 10^7 total recombinant receptor-expressing cells, total T cells, or total peripheral blood mononuclear cells (PBMCs), no more than 1 x 10^6 total recombinant receptor-expressing cells, total T cells, or total peripheral blood mononuclear cells (PBMCs), no more than 0.5 x 10^6 total recombinant receptor-expressing cells, total T cells, or total peripheral blood mononuclear cells (PBMCs), optionally wherein the instructions specify the administration of one or of a plurality of unit doses comprising the dose of cells and/or a volume corresponding to such one or plurality of unit doses comprising the dose of cells.

189. The kit of any of embodiments 99-188, wherein the instructions specify administering the inhibitor orally, subcutaneously or intravenously.

190. The kit of embodiment 189, wherein the instructions specify administering the inhibitor orally.

191. The kit of any one of embodiments 99-190, wherein the instructions specify the inhibitor is to be administered at least six times daily, five times daily, four times daily, three times daily, twice daily, once daily, every other day, three times a week, at least once a week, or only one time.

192. The kit of any of embodiments 99-191, wherein the instructions specify the inhibitor is administered three times a week.

193. The kit of embodiment 191 or embodiment 192, wherein the instructions specify the administration of the inhibitor is to be carried out in a treatment cycle that is at least or at least about or 14 days, at least or at least about or 21 days or at least or at least about or 28 days.

194. The kit of any of embodiments 99-193, wherein the instructions specify the inhibitor is administered, or each administration of the inhibitor is independently administered, in an amount of 0.5 mg to 500 mg, 0.5 mg to 250 mg, 0.5 mg to 100 mg, 0.5 mg to 50 mg, 0.5 mg to 25 mg, 0.5 mg to 10 mg, 0.5 mg to 5.0 mg, 0.5 mg to 2.5 mg, 0.5 mg to 1.0 mg, 1.0 mg to 500 mg, 1.0 mg to 250 mg, 1.0 mg to 100 mg, 1.0 mg to 50 mg, 1.0 mg to 25 mg, 1.0 mg to 10 mg, 1.0 mg to 5.0 mg, 1.0 mg to 2.5 mg, 2.5 mg to 500 mg, 2.5 mg to 250 mg, 2.5 mg to 100 mg, 2.5 mg to 50 mg, 2.5 mg to 25 mg, 2.5 mg to 10 mg, 2.5 mg to 5.0 mg, 5.0 mg to 500 mg, 5.0 mg to 250 mg, 5.0 mg to 100 mg, 5.0 mg to 50 mg, 5.0 mg to 25 mg, 5.0 mg to 10 mg, 10 mg to 500 mg, 10 mg to 250 mg, 10 mg to 100 mg, 10 mg to 50 mg, 10 mg to 25 mg, 25 mg to 500 mg, 25 mg to 250 mg, 25 mg to 100 mg, 25 mg to 50 mg, 50 mg to 500 mg, 50 mg to 250 mg, 50 mg to 100 mg, 100 mg to 500 mg, 100 mg to 250 mg or 250 mg to 500 mg.

195. The kit of any of embodiments 99-194, wherein the instructions specify the inhibitor is administered, or each administration of the inhibitor is independently administered, in an amount that is at least or at least about or is or is about 0.5 mg, 1.0 mg, 2.5 mg, 5.0 mg, 10.0 mg, 25 mg, 50 mg, 100 mg, 250 mg or 500 mg.

196. An article of manufacture comprising the combination or kit of any of embodiments 99-195.

VII. Examples

[0618] The following examples are included for illustrative purposes only and are not intended to limit the scope of the invention.

Example 1: Assessment of Surface Expression of BCMA on Multiple Myeloma (MM) Cell Lines in the Presence of a Gamma Secretase Inhibitor

[0619] Effects of the exemplary gamma secretase inhibitor (GSI), LY3039478, on surface levels of B cell maturation antigen (BCMA) on multiple myeloma (MM) cell lines were assessed. Different concentrations of LY3039478, including concentrations within the estimated Cmax/Cmin for the inhibitor, were incubated with RPMI 8226 cells ($BCMA^{low}$ human multiple myeloma cell line), MM1.S cells ($BCMA^{med}$ human multiple myeloma cell line) and OPM2 cells ($BCMA^{med}$ human multiple myeloma cell line) for 24 hours at 37ºC. The cell lines also were incubated in the absence of the inhibitor (vehicle control). Surface expression of BCMA on each cell line was assessed by flow cytometry using an-anti-BCMA antibody. As shown in FIG. 1, LY3039478 achieved potent inhibition of BCMA cleavage from the surface of the assessed cell lines, with an IC_{50} of 0.01nM to 0.35 nM.

Example 2: Assessment of anti-BCMA CAR- T cell Function in the Presence of a Gamma Secretase Inhibitor

[0620] Certain functional activities of anti-BCMA CAR-T cells against BCMA-expressing target cells were assessed in the presence or absence of the exemplary GSI LY3039478.

[0621] In this study, to generate anti-BCMA CAR-expressing T cells, CD4+ and CD8+ T cells were isolated by immunoaffinity-based enrichment from leukapheresis samples from two healthy donor subjects and one myeloma patient. Isolated T cells were activated and transduced with a viral vector encoding an anti-BCMA CAR. The CAR contained an scFv antigen-binding domain specific for BCMA having a VH domain and a VL domain with amino acid sequences set forth in SEQ ID NOS: 24 and 25, respectively, a spacer, a spacer having an amino acid sequence set forth in SEQ ID NO:31, a CD28 transmembrane region, a 4-1BB costimulatory signaling region and a CD3-zeta derived intracellular signaling domain. The nucleic acid construct encoding the CAR also included a truncated receptor sequence for use as a transduction marker, separated from the CAR sequence by a self-cleaving T2A sequence.

I. Cytolytic Activity

[0622] Anti-BCMA CAR-T cells generated as described in this Example above were cultured at 37 ºC with either OPM2 or RPMI-8226 target cells at an effector to target (E:T) ratio

of 0.3:1. The target cells were labeled with NucLight Red (NLR), to permit tracking of target cells by microscopy. Cultures were incubated in the absence of LY3039478 or in the presence of LY3039478 at concentrations of 0.1 nM, 1 nM, 10nM, 100nM, 1000nM, 10000nM or 100000 nM (the range covering the estimated human pharmacokinetic range is shown by the boxes in FIGS. 3A-3D). As a control, target cells were incubated with LY3039478 but without culture with CAR-T cells. Cytolytic activity was assessed by measuring the loss of viable target cells after about 150 hours, as determined by red fluorescent signal (using the IncuCyte® Live Cell Analysis System, Essen Bioscience). The area under the curve for killing observed during the culture period (~150 hours) was determined. Percent lysis (% killing) was determined by normalizing the AUC for killing to cultures incubated without the inhibitor.

[0623] Assessment of CAR-T cells from the myeloma patient-derived donor are depicted in FIGS. 2A (OPM2) and 2B (RPMI- 8226). As shown, LY3039478 increased the cytolytic activity of anti-BCMA CAR T cells at all concentrations tested in both cell lines, although the magnitude of the effect across tested inhibitor concentrations was cell-line dependent. No detrimental effects on CAR T cell function in this assay were observed at 100 μM LY3039478, which is 300 times above the calculated Cmax observed in patients. No differences were observed in target survival in the presence of LY3039478 alone without CAR-T cells. Similar results were observed at an E:T ratio of 1:1. CAR-T cells derived from the other donors that were incubated with LY3039478 also showed an increase in CAR-T function, although some donor to donor variability was observed. These results indicate that treatment in the presence of a gamma secretase inhibitor can increase surface BCMA expressiong and anti-BCMA CAR T cell cytolytic function.

[0624] A similar study was carried out to further assess the effects of different concentrations of LY3039478, including concentrations within the estimated Cmax/Cmin for the inhibitor, on the cytolytic function of anti-BCMA CAR-T cells against OPM2 target cells. Results in FIG. 2C from one exemplary donor showed that LY3039478 was capable of increasing anti-BCMA CAR-T killing of BCMA-expressing target cells with an IC50 of 0.1-0.25 nM.

2. Cytokine Production

[0625] The production of cytokines by anti-BCMA CAR-T cells in the presence or absence of LY3039478 was assessed by monitoring cytokine levels in the supernatants of co-cultures of

CAR-T cells with BCMA-expressing target cells. CAR-T cells generated as described in this Example above were cultured with either OPM2 or RPMI-8226 target cells in the absence of LY3039478 or in the presence of LY3039478 at concentrations of 0.1 nM, 1 nM, 10nM, 100nM, 1000nM, 10000nM or 100000 nM (the range covering the estimated human pharmacokinetic range is shown by the boxes in FIGS. 3A-3D). Culture supernatants were harvested after 24 hours, and IFNγ and IL-2 were measured from the culture supernatants.

[0626] The presence of LY3039478 increased the production of IFN-gamma (FIG. 3A and FIG. 3B) and IL-2 (FIG. 3C and 3D) in supernatant of anti-BCMA CAR T cells co-cultured with BCMA-expressing target cells compared to co-cultures incubated in the absence of LY3039478. The magnitude of the effect differed between cell lines and, in some cases, the concentration of the inhibitor. Similar to effects on cytolytic function, the results indicated that combination with a gamma secretase inhibitor increases BCMA antigen availability for the anti-BCMA CAR-T cells, leading to increased effect of the CAR-T cells, including production of cytokines.

[0627] A similar study was carried out to further assess the effects of different concentrations of LY3039478, including concentrations within the estimated Cmax/Cmin for the inhibitor, on the ability of anti-BCMA CAR-T cells generated as described in this Example 2 above to produce cytokines following co-culture with OPM2 target cells. Results in FIG. 3E from one exemplary donor were consistent with an interpretation that LY3039478 was capable of increasing cytokine production (with an IC50 of 0.1-0.25 nM)by anti-BCMA CAR-T cells cultured with BCMA-expressing target cells.

Example 3: Effect of Gamma Secretase Inhibitor on CAR-T Cell Expansion and Cytokine Production after Serial Restimulation

[0628] The ability of CAR T cells to expand ex vivo following repeated rounds of antigen stimulation can correlate with in vivo function and/or capacity of the cells to persist in in vivo (e.g., following administration and initial activation in response to encounter with antigen) (Zhao et al. (2015) Cancer Cell, 28:415-28). To assess the effect of the exemplary gamma secretase inhibitor on CAR-T cell activity after serial restimulation, anti-BCMA CAR+ T cells were generated as described above in Example 2, and irradiated BCMA-expressing target cells (MM1S cells) were added at an effector-to-target (E:T) ratio of 1:2 in the presence of LY3039478 at two different concentrations, either 0.001 μM (a concentration around the IC50)

or a higher concentration of 1.0 μM (approximately 3 times Cmax), or with vehicle only control. The assay was performed on a composition of anti-BCMA CAR+ T cells generated from T cells derived from a healthy donor or from a donor with multiple myeloma. Every 3-4 days, CAR-expressing T cells were harvested, counted and re-plated at the initial seeding density with fresh media containing new irradiated target cells at the same E:T ratio in the presence or absence of the same concentration of LY3039478. A total of 3-5 rounds of stimulation were carried out during a 10-14 day culture period. Effects of the inhibitor on T cell function were assessed by monitoring the cell number (FIGS. 4A and 4B) or cytokine production in harvested supernatant (FIG. 5A and FIG. 5B) at each round of stimulation.

[0629] The number of CAR-T cells was determined at each round prior to re-plating for anti-BCMA CAR+ T cells derived from the healthy donor-derived cells (FIG. 4A) or the patient-derived cells (FIG. 4B). For anti-BCMA CAR+ T cells derived from both donors, increased expansion of anti-BCMA CAR T cells was observed over the period of 10-14 days when CAR-T cells were incubated in the presence of both concentrations of LY3039478 (0.001 μM and 1.0 μM). The results indicated that LY3039478 did not exhibit any detrimental effects on CAR-T cell function, even though it is a known inhibitor of the Notch signaling pathway, which is involved in T cell differentiation. The results were consistent with a conclusion that the exemplary gamma secretase inhibitor LY3039478 can promote continued CAR+ T cell expansion and/or survival following repeated encounter with cognate antigen.

[0630] The production of cytokines by anti-BCMA CAR-T cells upon restimulation with antigen-expressing cells in the presence or absence of LY3039478 was assessed after the first round of re-plating at day 4. Culture supernatants were harvested 24 hours after re-plating, and IFN-gamma, IL-2 and TNF-alpha were measured from the culture supernatants. Increased levels of IFN-gamma, IL-2 and TNF-alpha were observed for both concentration of LY3039478 (0.001 μM and 1.0 μM) in anti-BCMA CAR+ T cells derived from the healthy donor cells (FIG. 5A) or the patient donor cells (FIG. 5B).

Example 4: Assessment of Agents on Blocking Activity of Anti-BCMA CAR Activity

[0631] Polynucleotides encoding exemplary chimeric antigen receptors (CARs), each containing a human anti-BCMA scFv antigen-binding domain, were generated. Among the CARs were those containing anti-BCMA scFvs containing a VH and VL set forth in SEQ ID

NOS: 22 and 23, respectively (referred to in this example as anti-BCMA.3 CAR), a VH and VL set forth in SEQ ID NOS: 24 and 25, respectively (referred to in this example as anti-BCMA.4 CAR), and a VH and VL set forth in SEQ ID NOS: 18 and 19, respectively (referred to in this example as anti-BCMA.1 CAR). The encoded CARS also were generated to contain an exemplary spacer set forth in SEQ ID NO:31, a human CD28 transmembrane domain, a human 4-1BB-derived intracellular co-signaling sequence, and a CD3-zeta derived intracellular signaling domain.

[0632] cDNA clones encoding such CARs, were linked to a downstream ribosomal skip element (such as T2A) followed by a truncated receptor-encoding sequence, and cloned into a lentiviral expression vector.

[0633] To generate anti-BCMA CAR-expressing T cells, T cells were isolated by immunoaffinity-based enrichment from leukapheresis samples from human donor subjects. Isolated T cells were activated and transduced with lentiviral vectors containing the respective polynucleotides encoding the anti-BCMA CARs and expanded; CD4+ and CD8+ T cells were stained and analyzed by flow cytometry to confirm transduction and expression of the anti-BCMA CARs. Various functions of anti-BCMA CAR-expressing cells were assessed following incubation with BCMA-expressing target cells and soluble BCMA.

[0634] To assess cytolytic activity, anti-BCMA CAR-expressing T cells were incubated with OPM2 target cells at an E:T ratio of 5:1 in the presence of soluble BCMA-Fc at 0, 0.3, 3, 30 or 300 ng/mL. Target cells were labeled with NucLight Red (NLR) to permit tracking of target cells by microscopy. Cytolytic activity was assessed by measuring the loss of viable target cells over a period of between 24 and 72 hours, as determined by red fluorescent signal (using the IncuCyte® Live Cell Analysis System, Essen Bioscience). Percent lysis (% Lysis) was normalized to the lysis that occurred in target cells incubated with mock T cells. Cytolytic activity of T cells expressing the anti-BCMA.1-containing CAR or anti-BCMA.3 CAR were substantially reduced in the presence of 3 ng/mL or more BCMA-Fc, however the cytolytic activity of cells expressing the anti-BCMA.4 CAR was not blocked by the presence of up to 300 ng/mL BCMA-Fc.

[0635] To assess cytokine production, anti-BCMA CAR-expressing T cells were incubated with OPM2 target cells at an E:T ratio of 5:1 in the presence of soluble BCMA-Fc at 0, 111, 333 and 1000 ng/mL. T cells not expressing a CAR (mock) also were assessed. Cytokine

accumulation of IFN-γ, TNF-α and IL-2 in supernatant was assessed. Cytokine accumulation in cultures containing T cells expressing the anti-BCMA.1 CAR or anti-BCMA.3 CAR was substantially reduced in the presence of 111 ng/mL or more BCMA-Fc, however less reduction in cytokine accumulation was observed in cultures containing T cells expressing the anti-BCMA.4 CAR in the presence of soluble BCMA-Fc at all concentrations tested.

[0636] In another experiment, activity of the anti-BCMA.4 CAR-expressing T cells in the presence vs. absence of soluble BCMA was assessed. Anti-BCMA.4 CAR-expressing T cells were co-cultured with RPMI-8226 tumor cells, with recombinant BCMA-Fc, or with cell culture supernatant derived from NCI-H929 multiple myeloma cells (BCMA-secreting cell line, the supernatant containing soluble BCMA). Neither tumor-cell lysis nor cytokine production was affected by any of the concentrations of NCI-H929–derived soluble BCMA (up to 1000 ng/mL). Both tumor-cell lysis and cytokine production were only minimally decreased at similarly high physiological levels of recombinant BCMA.

Example 5: Administration of Gamma Secretase Inhibitor Compound LY3039478 and anti-BCMA CAR+ T cells *in vivo* in an Animal Tumor Model

[0637] A pharmacokinetics and pharmacodyamics study was carried out following oral administration of a single dose of LY3039478 (3 mg/kg) in a human multiple myeloma xenograft model. Specifically, NOD.Cg.PrkdcscidIL2rgtm1Wjl/SzJ (NSG) mice were injected subcutaneously (s.c.) with 1E+07 RPMI-8226 cells modified to express GFP and firefly luciferase (RPMI-8226 ffluc) and tumors were allowed to grow. Thirty two (32) days after RPMI-8226 injection (day 0 of study), a single dose of 3 mg/kg LY3039478 was administered orally.

[0638] FIG. 6 depicts analysis of blood samples, taken at 0, 0.5, 2, 6, 18, 24, 36, and 48 hour post-administration, for plasma levels of the drug. FIG. 7 shows plasma BCMA levels over time in plasma samples taken at 0, 0.5, 2, 6, 18, 24, 36, and 48 hours post-administration in mice administered a single oral dose of LY3039478 (3 g/kg) or vehicle in this study. Tumor cells at 2 hours, 24 hours or 48 hours after administration of LY3039478 or vehicle control in the study were assessed for surface expression of BCMA. Mean fluorescence intensity (MFI) of surface BCMA, as assessed by flow cytometry, is shown in FIG. 8. Increased BCMA expression on

tumors was confirmed with a dosing schedule of 0.1 mg/kg or 1 mg/kg administered two oral administrations of LY3039478 in an alternative multiple myeloma tumor cell model.

[0639] Anti-tumor effects following administration of anti-BCMA CAR T cells and a gamma secretase inhibitor compound, LY3039478, each, individually, and in combination, were assessed in a RPMI-8226 human multiple myeloma xenograft mouse model. The anti-BCMA CAR-expressing T cells were generated as described in Example 2.

[0640] NOD.Cg.PrkdcscidIL2rgtm1Wjl/SzJ (NSG) mice were injected subcutaneously (s.c.) with 1E+07 RPMI-8226 cells modified to express GFP and firefly luciferase (RPMI-8226 ffluc), and tumor volume was allowed to increase over the course of 25 days, the 25th day being designated "day 0" in this study. Animals were divided into 8 groups, and each group received one of the following treatments: (1) vehicle only, (2) LY3039478 (3 mg/kg) only, (3) Mock T cells (not expressing the CAR) (3.00E+06 T cells per mouse) and vehicle, (4) Mock T cells (3.00E+06 T cells per mouse) and LY3039478 (3 mg/kg), (5) anti-BCMA CAR+ T cells (1.00E+06 CAR+ cells per mouse) and vehicle, (6) anti-BCMA CAR+ T cells (1.00E+06 CAR+ cells per mouse) and LY3039478 (3 mg/kg), (7) anti-BCMA CAR+ T cells (3.00E+06 CAR+ cells per mouse) and vehicle, or (8) anti-BCMA CAR+ T cells (3.00E+06 CAR+ cells per mouse) and LY3039478 (3 mg/kg). In each group, LY3039478 or vehicle was delivered by mouth (p.o.) at day -1 and at day zero (0) and every other day (e.o.d.) up to day 21. Where applicable, T cells were administered by a single intravenous (i.v.) injection on day 0.

[0641] Tumor volume was assessed twice weekly over the course of the study; results for different groups are shown in FIG. 9A. CAR+ T cells in blood and in tumor were assessed on days 7, 15, 21, and 28 by flow cytometry using antibodies specific for cell surface markers and a reagent specific to the anti-BCMA CAR. Animal survival was monitored throughout the study.

[0642] As shown in FIG. 9A, tumors continued to grow over the course of the study following adoptive transfer of negative control cells (Mock) or in mice not receiving treatment. In mice that had received the higher dose anti-BCMA CAR-expressing T cells, complete regression of tumor growth was observed by about 20 days post-CAR T cell transfer and was sustained throughout the remainder of the study; some reduction in tumor growth was observed in animals having received the lower dose of anti-BCMA CAR-expressing T cells and vehicle. In animals having received administration of LY3039478 in combination with the lower dose of anti-BCMA CAR-expressing cells, there was observed a substantial reduction in tumor growth

compared to mice treated with the lower dose of anti-BCMA CAR-expressing cells and vehicle. LY3039478 administration was well-tolerated throughout the study as indicated by body weight and condition score.

[0643] The percent survival of animals over the course of the study is shown in FIG. 9B. Administration of anti-BCMA CAR-expressing T cells was observed to result in increased survival. A further survival benefit was observed in mice treated with the combination of the lower dose anti-BCMA CAR-expressing T cells and the GSI, LY3039478, as compared to the group treated with the lower dose of CAR-T cells and vehicle.

Table E1. Survival across groups

Group	Median Survival (days)	Number of deaths	Day 65 Survival %	Day 65 tumor free survival
Vehicle	22.5	10	0	NA
LY3039478	22.5	10	0	NA
Mock + Vehicle	21	10	0	NA
Mock + LY3039478	22.5	7	30	2/3
anti-BCMA CAR-expressing T cell 1e6 + Vehicle	60.5	5	50	5/5
anti-BCMA CAR-expressing T cell 1e6 + LY3039478	Undef	2	80	8/8
anti-BCMA CAR-expressing T cell 3e6 + Vehicle	Undef	0	100	10/10
anti-BCMA CAR-expressing T cell 3e6 + LY3039478	Undef	0	100	10/10

[0644] The presence of CAR+ T cells in the blood was monitored over the course of the study (from blood drawn on days 7, 14, 21 and 28 post CAR-T cell administration). White blood cells from blood samples were stained with various reagents to detect various markers on the surface of immune cells, including T cells, and expression of the CAR (including a reagent specific for the CAR and a reagent specific for a transduction marker encoded by the CAR vector). Results are shown in FIGS. 10A and 10B.

[0645] As shown, increased anti-BCMA CAR-expressing CD4+ T cells (FIG. 10A) and anti-BCMA CAR-expressing CD8+ T cells (FIG. 10B) counts were observed in the peripheral blood at Day **15**, **21** and **28** following administration of either the higher or lower dose CAR-expressing T cells in combination with LY3039478 compared to its absence. In the presence of LY3039478, peak expansion was at about day 15 following administration of higher dose anti-BCMA CAR-expressing cells and was about day 21 following administration of lower dose anti-BCMA CAR-expressing T cells. No changes in peripheral blood count were observed for CD4+ or CD8+ T cells that did not express the anti-BCMA CAR when treated with or without LY3039478.

[0646] Tumor digests were also analyzed to determine the number of CD4+ and CD8+ CAR+ T cells in tumors following the various treatment conditions. On days 7 and 15, satellite tumor digests from mice treated with lower dose anti-BCMA CAR+ T cells and LY3039478 showed a higher number of CD4+ and CD8+ CAR+ T cells per 1 x 10^6 cells compared to mice treated with lower dose anti-BCMA CAR+ T cells and vehicle (FIG. 11). Analysis of cells from tumor digests for surface BCMA expression by flow cytometry, as measured by mean fluorescence intensity (MFI), showed that the presence of LY3039478 increased surface BCMA MFI on RPMI-8226 tumor cells (FIG. 12B). Serum BCMA was also assessed according to treatment condition. FIG. 12A shows BCMA serum levels on days 7 and 15 for each treatment condition. These results are consistent with the ability of LY3039478 to reduce or prevent cleavage of surface BCMA.

[0647] These results are consistent with a finding that administration of LY3039478 in combination with the lower dose of anti-BCMA CAR T cells led to increased anti-tumor effects as compared to administration of the lower dose of anti-BCMA CAR-expressing T cells in combination with vehicle alone.

Example 6: <u>Assessment of Anti-BCMA CAR+ T Cells Following Chronic Stimulation and Re-Challenge in the Presence of a Gamma Secretase Inhibitor Compound, LY3039478</u>

[0648] The effect of a gamma secretase inhibitor compound,LY3039478, on the cytotoxicity of anti-BCMA CAR+ T cells after long-term CAR-specific stimulation was assessed. Anti-BCMA CAR+ T cell compositions, generated as described in Example 2 containing anti-BCMA

CAR-expressing CD4+ and CD8+ T cells, were incubated with 50 µg/mL BCMA-Fc conjugated beads for 7 days, under conditions designed to functionally exhaust the CAR T cells.

[0649] The CAR-T cells then were re-challenged with antigen-expressing target cells in the presence or absence of LY3039478. Specifically, anti-BCMA CAR-expressing T cells were co-cultured with multiple myeloma cell lines RPMI-8226, OPM2, or MM1.S in the presence of 1 µM LY3039478 or DMSO vehicle control at an effector to target ratio of 0.3:1. To assess cytolytic activity, the target cells were labeled with NucLight Red (NLR) to permit tracking by fluorescent microscopy. Killing activity was assessed by measuring the loss of viable target cells over time, as determined by loss of fluorescent signal over time by kinetic fluorescence microscopy (using the INCUCYTE® Live Cell Analysis System, Essen Bioscience). Target fluorescence was monitored over time and was normalized to target cell count.

[0650] The results shown in FIG.13A demonstrated that the presence of LY3039478 improved cytolytic activity of anti-BCMA CAR+ T cells that had been chronically stimulated. A killing index was determined as the inverse of the area under the curve (AUC) for target fluorescence over time, and was normalized relative to DMSO vehicle control. As shown in FIG. 13B, the degree of improvement on cell killing correlated to the density of BCMA antigen on the target cells (antigen density: RPMI-8226< OPM2<MM1.S).

[0651] Cell culture supernatant was harvested from co-cultures 24 hours after initiation of culture of anti-BCMA CAR-expressing T cells and target cells in the experiment above, or in a similar experiment but in which anti-BCMA CAR+ T cells and target cells were co-cultured at an E:T ratio of 1:1. The production of TNF-alpha, IFN-gamma and IL-2 in the harvested supernantant was measured using a Luminex Multiplex Assay. At both E:T ratios, the presence of the GSI LY3039478 improved function of the chronically stimulated anti-BCMA CAR-expressing T cells to produce cytokines (FIG. 13C). The degree of improvement in cytokine production also correlated to the density of BCMA antigen on the target cells.

Example 7: <u>Assessment of Cytokine Production by anti-BCMA CAR+ T Cells in the Presence or Absence of a Gamma Secretase Inhibitor</u>

[0652] Various parameters indicative of T cell function were assessed following incubation of anti-BCMA CAR+ T cells with BCMA-containing beads in the presence of different concentrations of the gamma secretase inhibitor, LY3039478, or vehicle (DMSO). Supernatant

was harvested after 24 hours and production of TNF-alpha, IFN-gamma and IL-2 were measured using a Luminex Multiplex Assay. The results are shown in FIG. 14. The results were consistent with a conclusion that the presence of this GSI compound did not impact cytokine production, CAR+ T cell count, or viability.

Example 8: Assessement of BCMA Surface Expression in Multiple Myeloma Cells in Patients Treated with Gamma Secretase Inhibitor

[0653] Three human subjects having multiple myeloma were administered three 25 mg doses of the gamma secretase inhibitor (GSI) LY3039478 by mouth. The absolute B cell maturation antigen (BCMA) antibody binding capacity (ABC) on the surface of multiple myeloma cells was quantitated by flow cytometry performed on bone marrow aspirate specimens obtained from each of the patients prior to administration of the GSI, and immediately after the third oral GSI dose. A mean 66-fold increase in BCMA ABC was observed following GSI administration (FIG. 15A). Percentage of plasma cells in the patient samples that were determind to have measurable surface BCMA expression was observed to increase from 35% (mean) before GSI to 98.4% (mean) following the three oral doses (FIG. 15B).

[0654] The present invention is not intended to be limited in scope to the particular disclosed embodiments, which are provided, for example, to illustrate various aspects of the invention. Various modifications to the compositions and methods described will become apparent from the description and teachings herein. Such variations may be practiced without departing from the true scope and spirit of the disclosure and are intended to fall within the scope of the present disclosure.

SEQUENCES

SEQ ID NO.	SEQUENCE	DESCRIPTION
1	ESKYGPPCPPCP	spacer (IgG4hinge) (aa) Homo sapiens
2	GAATCTAAGTACGGACCGCCCTGCCCCCCTTGCCCT	spacer (IgG4hinge) (nt) Homo sapiens
3	ESKYGPPCPPCPGQPREPQVYTLPPSQEEMTKNQVSLTCLVKGFYPSDIAVEWE SNGQPENNYKTTPPVLDSDGSFFLYSRLTVDKSRWQEGNVFSCSVMHEALHNHY TQKSLSLSLGK	Hinge-CH3 spacer Homo sapiens
4	ESKYGPPCPPCPAPEFLGGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSQEDPE VQFNWYVDGVEVHNAKTKPREEQFNSTYRVVSVLTVLHQDWLNGKEYKCKVSNK GLPSSIEKTISKAKGQPREPQVYTLPPSQEEMTKNQVSLTCLVKGFYPSDIAVE WESNGQPENNYKTTPPVLDSDGSFFLYSRLTVDKSRWQEGNVFSCSVMHEALHN HYTQKSLSLSLGK	Hinge-CH2-CH3 spacer Homo sapiens
5	RWPESPKAQASSVPTAQPQAEGSLAKATTAPATTRNTGRGGEEKKKEKEKEEQE ERETKTPECPSHTQPLGVYLLTPAVQDLWLRDKATFTCFVVGSDLKDAHLTWEV AGKVPTGGVEEGLLERHSNGSQSQHSRLTLPRSLWNAGTSVTCTLNHPSLPPQR LMALREPAAQAPVKLSLNLLASSDPPEAASWLLCEVSGFSPPNILLMWLEDQRE VNTSGFAPARPPPQPGSTTFWAWSVLRVPAPPSPQPATYTCVVSHEDSRTLLNA SRSLEVSYVTDH	IgD-hinge-Fc Homo sapiens
6	LEGGGEGRGSLLTCGDVEENPGPR	T2A artificial
7	RKVCNGIGIGEFKDSLSINATNIKHFKNCTSISGDLHILPVAFRGDSFTHTPPL DPQELDILKTVKEITGFLLIQAWPENRTDLHAFENLEIIRGRTKQHGQFSLAVV SLNITSLGLRSLKEISDGDVIISGNKNLCYANTINWKKLFGTSGQKTKIISNRG ENSCKATGQVCHALCSPEGCWGPEPRDCVSCRNVSRGRECVDKCNLLEGEPREF VENSECIQCHPECLPQAMNITCTGRGPDNCIQCAHYIDGPHCVKTCPAGVMGEN NTLVWKYADAGHVCHLCHPNCTYGCTGPGLEGCPTNGPKIPSIATGMVGALLLL LVVALGIGLFM	tEGFR artificial
8	FWVLVVVGGVLACYSLLVTVAFIIFWV	CD28 (amino acids 153-179 of Accession No. P10747) Homo sapiens
9	IEVMYPPPYLDNEKSNGTIIHVKGKHLCPSPLFPGPSKPFWVLVVVGGVLACYS LLVTVAFIIFWV	CD28 (amino acids 114-179 of Accession No. P10747) Homo sapiens
10	RSKRSRLLHSDYMNMTPRRPGPTRKHYQPYAPPRDFAAYRS	CD28 (amino acids 180-220 of P10747) Homo sapiens
11	RSKRSRGGHSDYMNMTPRRPGPTRKHYQPYAPPRDFAAYRS	CD28 (LL to GG) Homo sapiens
12	KRGRKKLLYIFKQPFMRPVQTTQEEDGCSCRFPEEEEGGCEL	4-1BB (amino acids 214-255 of Q07011.1) Homo sapiens

13	RVKFSRSADAPAYQQGQNQLYNELNLGRREEYDVLDKRRGRDPEMGGKPRRKNP QEGLYNELQKDKMAEAYSEIGMKGERRRGKGHDGLYQGLSTATKDTYDALHMQA LPPR	CD3 zeta Homo sapiens
14	RVKFSRSAEPPAYQQGQNQLYNELNLGRREEYDVLDKRRGRDPEMGGKPRRKNP QEGLYNELQKDKMAEAYSEIGMKGERRRGKGHDGLYQGLSTATKDTYDALHMQA LPPR	CD3 zeta Homo sapiens
15	RVKFSRSADAPAYKQGQNQLYNELNLGRREEYDVLDKRRGRDPEMGGKPRRKNP QEGLYNELQKDKMAEAYSEIGMKGERRRGKGHDGLYQGLSTATKDTYDALHMQA LPPR	CD3 zeta Homo sapiens
16	PGGG-(SGGGG)5-P- wherein P is proline, G is glycine and S is serine	linker
17	GSADDAKKDAAKKDGKS	Linker
18	QIQLVQSGPELKKPGETVKISCKASGYTFTDYSINWVKRAPGKGLKWMGWINTE TREPAYAYDFRGRFAFSLETSASTAYLQINNLKYEDTATYFCALDYSYAMDYWG QGTSVTVSS	Variable heavy (VH) Anti-BCMA
19	DIVLTQSPPSLAMSLGKRATISCRASESVTILGSHLIHWYQQKPGQPPTLLIQL ASNVQTGVPARFSGSGSRTDFTLTIDPVEEDDVAVYYCLQSRTIPRTFGGGTKL EIK	Variable light (VL) Anti-BCMA
20	QIQLVQSGPDLKKPGETVKLSCKASGYTFTNFGMNWVKQAPGKGFKWMAWINTY TGESYFADDFKGRFAFSVETSATTAYLQINNLKTEDTATYFCARGEIYYGYDGG FAYWGQGTLVTVSA	Variable heavy (VH) Anti-BCMA
21	DVVMTQSHRFMSTSVGDRVSITCRASQDVNTAVSWYQQKPGQSPKLLIFSASYR YTGVPDRFTGSGSGADFTLTISSVQAEDLAVYYCQQHYSTPWTFGGGTKLDIK	Variable light (VL) Anti-BCMA
22	EVQLVQSGAEVKKPGESLKISCKGSGYSFTSYWIGWVRQMPGKGLEWMGIIYPG DSDTRYSPSFQGHVTISADKSISTAYLQWSSLKASDTAMYYCARYSGSFDNWGQ GTLVTVSS	Variable heavy (VH) Anti-BCMA
23	SYELTQPPSASGTPGQRVTMSCSGTSSNIGSHSVNWYQQLPGTAPKLLIYTNNQ RPSGVPDRFSGSKSGTSASLAISGLQSEDEADYYCAAWDGSLNGLVFGGGTKLT VLG	Variable light (VL) Anti-BCMA
24	EVQLVQSGAEMKKPGASLKLSCKASGYTFIDYYVYWMRQAPGQGLESMGWINPN SGGTNYAQKFQGRVTMTRDTSISTAYMELSRLRSDDTAMYYCARSQRDGYMDYW GQGTLVTVSS	Variable heavy (VH) Anti-BCMA
25	QSALTQPASVSASPGQSIAISCTGTSSDVGWYQQHPGKAPKLMIYEDSKRPSGV SNRFSGSKSGNTASLTISGLQAEDEADYYCSSNTRSSTLVFGGGTKLTVLG	Variable light (VL) Anti-BCMA
26	GGGGS	Linker
27	GGGS	Linker
28	GGGGSGGGGSGGGGS	Linker
29	GSTSGSGKPGSGEGSTKG	Linker
30	SRGGGGSGGGGSGGGGSLEMA	Linker
31	ESKYGPPCPPCPAPPVAGPSVFLFPPKPKDTLMISRTPEVTCVVVDVSQEDPEV QFNWYVDGVEVHNAKTKPREEQFQSTYRVVSVLTVLHQDWLNGKEYKCKVSNKG LPSSIEKTISKAKGQPREPQVYTLPPSQEEMTKNQVSLTCLVKGFYPSDIAVEW ESNGQPENNYKTTPPVLDSDGSFFLYSRLTVDKSRWQEGNVFSCSVMHEALHNH YTQKSLSLSLGK	Hinge-CH2-CH3 spacer Homo sapiens
32	EVQLVQSGAEVKKPGSSVKVSCKASGGTFSSYAISWVRQAPGQGLEWMGRIIPI LGIANYAQKFQGRVTMTEDTSTDTAYMELSSLRSEDTAVYYCARSGYSKSIVSY MDYWGQGTLVTVSS	Variable heavy (VH) Anti-BCMA
33	LPVLTQPPSTSGTPGQRVTVSCSGSSSNIGSNVVFWYQQLPGTAPKLVIYRNNQ RPSGVPDRFSVSKSGTSASLAISGLRSEDEADYYCAAWDDSLSGYVFGTGTKVT VLG	Variable light (VL) Anti-BCMA
34	QVQLVQSGAEVKKPGSSVKVSCKASGGTFSSYAISWVRQAPGQGLEWMGRIIPI LGTANYAQKFQGRVTITADESTSTAYMELSSLRSEDTAVYYCARSGYGSYRWED SWGQGTLVTVSS	Variable heavy (VH) Anti-BCMA

35	QAVLTQPPSASGTPGQRVTISCSGSSSNIGSNYVFWYQQLPGTAPKLLIYSNNQRPSGVPDRFSGSKSGTSASLAISGLRSEDEADYYCAAWDDSLSASYVFGTGTKVTVLG	Variable light (VL) Anti-BCMA
36	QVQLVQSGAEVKKPGASVKVSCKASGYTFTDYYMHWVRQAPGQRLEWMGWINPNSGGTNYAQKFQDRITVTRDTSSNTGYMELTRLRSDDTAVYYCARSPYSGVLDKWGQGTLVTVSS	Variable heavy (VH) Anti-BCMA
37	QSVLTQPPSVSGAPGQRVTISCTGSSSNIGAGFDVHWYQQLPGTAPKLLIYGNSNRPSGVPDRFSGSKSGTSASLAITGLQAEDEADYYCQSYDSSLSGYVFGTGTKVTVLG	Variable light (VL) Anti-BCMA
38	ASGGGGSGGRASGGGGS	Linker
39	MALPVTALLLPLALLLHAARP	CD8a signal peptide
40	METDTLLLWVLLLWVPGSTG	signal peptide
41	EVQLLESGGGLVQPGGSLRLSCAASGFTFSSYAMSWVRQAPGKGLEWVSAISGSGGSTYYADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCARAEMGAVFDIWGQGTMVTVSS	Variable heavy (VH) Anti-BCMA
42	EIVLTQSPATLSLSPGERATLSCRASQSVSRYLAWYQQKPGQAPRLLIYDASNRATGIPARFSGSGSGTDFTLTISSLEPEDFAVYYCQQRISWPFTFGGGTKVEIK	Variable light (VL) Anti-BCMA
43	QVQLVESGGGVVQPGRSLRLSCAASGFTFSSYGMHWVRQAPGKGLEWVAVISYDGSNKYYADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCARDGTYLGGLWYFDLWGRGTLVTVSS	Variable heavy (VH) Anti-BCMA
44	DIVMTQSPLSLPVTPGEPASISCRSSQSLLHSNGYNYLDWYLQKPGQSPQLLIYLGSNRASGVPDRFSGSGSGTDFTLKISRVEAEDVGVYYCMQGLGLPLTFGGGTKVEIK	Variable light (VL) Anti-BCMA
45	QVQLVQSGAEVKKPGASVKVSCKASGYTFTSYYMHWVRQAPGQGLEWMGIINPGGGSTSYAQKFQGRVTMTRDTSTSTVYMELSSLRSEDTAVYYCARESWPMDVWGQGTTVTVSS	Variable heavy (VH) Anti-BCMA
46	EIVMTQSPATLSVSPGERATLSCRASQSVSSNLAWYQQKPGQAPRLLIYGASTRATGIPARFSGSGSGTEFTLTISSLQSEDFAVYYCQQYAAYPTFGGGTKVEIK	Variable light (VL) Anti-BCMA
47	QLQLQESGPGLVKPSETLSLTCTVSGGSISSSSYYWGWIRQPPGKGLEWIGSISYSGSTYYNPSLKSRVTISVDTSKNQFSLKLSSVTAADTAVYYCARGRGYATSLAFDIWGQGTMVTVSS	Variable heavy (VH) Anti-BCMA
48	EIVLTQSPATLSLSPGERATLSCRASQSVSSYLAWYQQKPGQAPRLLIYDASNRATGIPARFSGSGSGTDFTLTISSLEPEDFAVYYCQQRHVWPPTFGGGTKVEIK	Variable light (VL) Anti-BCMA
49	EVQLVESGGGLVQPGGSLRLSCAASGFTFSSYSMNWVRQAPGKGLEWVSTISSSSSTIYYADSVKGRFTISRDNAKNSLYLQMNSLRAEDTAVYYCARGSQEHLIFDYWGQGTLVTVSS	Variable heavy (VH) Anti-BCMA
50	EIVLTQSPATLSLSPGERATLSCRASQSVSRYLAWYQQKPGQAPRLLIYDASNRATGIPARFSGSGSGTDFTLTISSLEPEDFAVYYCQQRFYYPWTFGGGTKVEIK	Variable light (VL) Anti-BCMA
51	QVQLVESGGGVVQPGRSLRLSCAASGFTFSSYGMHWVRQAPGKGLEWVAVISYDGSNKYYADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCARTDFWSGSPPGLDYWGQGTLVTVSS	Variable heavy (VH) Anti-BCMA
52	DIQLTQSPSSVSASVGDRVTITCRASQGISSWLAWYQQKPGKAPKLLIYGASSLQSGVPSRFSGSGSGTDFTLTISSLQPEDFATYYCQQIYTFPFTFGGGTKVEIK	Variable light (VL) Anti-BCMA
53	QVQLVQSGAEVKKPGSSVKVSCKASGGTFSSYAISWVRQAPGQGLEWMGGIIPIFGTANYAQKFQGRVTITADESTSTAYMELSSLRSEDTAVYYCARTPEYSSSIWHYYYGMDVWGQGTTVTVSS	Variable heavy (VH) Anti-BCMA
54	DIVMTQSPDSLAVSLGERATINCKSSQSVLYSSNNKNYLAWYQQKPGQPPKLLIYWASTRESGVPDRFSGSGSGTDFTLTISSLQAEDVAVYYCQQFAHTPFTFGGGTKVEIK	Variable light (VL) Anti-BCMA
55	QVQLVESGGGVVQPGRSLRLSCAASGFTFSSYGMHWVRQAPGKGLEWVAVISYDGSNKYYADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCVKGPLQEPPYDYGMDVWGQGTTVTVSS	Variable heavy (VH) Anti-BCMA

56	EIVMTQSPATLSVSPGERATLSCRASQSVSSNLAWYQQKPGQAPRLLIYSASTR ATGIPARFSGSGSGTEFTLTISSLQSEDFAVYYCQQHHVWPLTFGGGTKVEIK	Variable light (VL) Anti-BCMA
57	QVQLVQSGAEVKKPGSSVKVSCKASGGTFSSYAISWVRQAPGQGLEWMGRIIPI LGIANYAQKFQGRVTITADKSTSTAYMELSSLRSEDTAVYYCARGGYYSHDMWS EDWGQGTLVTVSS	Variable heavy (VH) Anti-BCMA
58	LPVLTQPPSASGTPGQRVTISCSGRSSNIGSNSVNWYRQLPGAAPKLLIYSNNQ RPPGVPVRFSGSKSGTSASLAISGLQSEDEATYYCATWDDNLNVHYVFGTGTKV TVLG	Variable light (VL) Anti-BCMA
59	QVQLVQSGSELKKPGASVKVSCKASGYTFTDYSINWVRQAPGQGLEWMGWINTE TREPAYAYDFRGRFVFSLDTSVSTAYLQISSLKAEDTAVYYCARDYSYAMDYWG QGTLVTVSS	Variable heavy (VH) Anti-BCMA
60	DIVLTQSPASLAVSLGERATINCRASESVSVIGAHLIHWYQQKPGQPPKLLIYL ASNLETGVPARFSGSGSGTDFTLTISSLQAEDAAIYYCLQSRIFPRTFGQGTKL EIK	Variable light (VL) Anti-BCMA
61	EVQLVESGGGLVQPGGSLRLSCAVSGFALSNHGMSWVRRAPGKGLEWVSGIVYS GSTYYAASVKGRFTISRDNSRNTLYLQMNSLRPEDTAIYYCSAHGGESDVWGQG TTVTVSS	Variable heavy (VH) Anti-BCMA
62	DIQLTQSPSSLSASVGDRVTITCRASQSISSYLNWYQQKPGKAPKLLIYAASSL QSGVPSRFSGSGSGTDFTLTISSLQPEDFATYYCQQSYSTPYTFGQGTKVEIK	Variable light (VL) Anti-BCMA
63	QVQLVESGGGLVQPGRSLRLSCAASGFTFSNYAMSWVRQAPGKGLGWVSGISRS GENTYYADSVKGRFTISRDNSKNTLYLQMNSLRDEDTAVYYCARSPAHYYGGMD VWGQGTTVTVSS	Variable heavy (VH) Anti-BCMA
64	DIVLTQSPGTLSLSPGERATLSCRASQSISSSFLAWYQQKPGQAPRLLIYGASR RATGIPDRFSGSGSGTDFTLTISRLEPEDSAVYYCQQYHSSPSWTFGQGTKLEI K	Variable light (VL) Anti-BCMA
65	QVQLVESGGGLVQPGGSLRLSCAVSGFALSNHGMSWVRRAPGKGLEWVSGIVYS GSTYYAASVKGRFTISRDNSRNTLYLQMNSLRPEDTAIYYCSAHGGESDVWGQG TTVTVSS	Variable heavy (VH) Anti-BCMA
66	DIRLTQSPSPLSASVGDRVTITCQASEDINKFLNWYHQTPGKAPKLLIYDASTL QTGVPSRFSGSGSGTDFTLTINSLQPEDIGTYYCQQYESLPLTFGGGTKVEIK	Variable light (VL) Anti-BCMA
67	EVQLVESGGGLVQPGGSLRLSCAVSGFALSNHGMSWVRRAPGKGLEWVSGIVYS GSTYYAASVKGRFTISRDNSRNTLYLQMNSLRPEDTAIYYCSAHGGESDVWGQG TTVTVSS	Variable heavy (VH) Anti-BCMA
68	EIVLTQSPGTLSLSPGERATLSCRASQSIGSSSLAWYQQKPGQAPRLLMYGASS RASGIPDRFSGSGSGTDFTLTISRLEPEDFAVYYCQQYAGSPPFTFGQGTKVEI K	Variable light (VL) Anti-BCMA
69	QIQLVQSGPELKKPGETVKISCKASGYTFRHYSMNWVKQAPGKGLKWMGRINTE SGVPIYADDFKGRFAFSVETSASTAYLVINNLKDEDTASYFCSNDYLYSLDFWG QGTALTVSS	Variable heavy (VH) Anti-BCMA
70	DIVLTQSPPSLAMSLGKRATISCRASESVTILGSHLIYWYQQKPGQPPTLLIQL ASNVQTGVPARFSGSGSRTDFTLTIDPVEEDDVAVYYCLQSRTIPRTFGGGTKL EIK	Variable light (VL) Anti-BCMA
71	QIQLVQSGPELKKPGETVKISCKASGYTFTHYSMNWVKQAPGKGLKWMGRINTE TGEPLYADDFKGRFAFSLETSASTAYLVINNLKNEDTATFFCSNDYLYSCDYWG QGTTLTVSS	Variable heavy (VH) Anti-BCMA
72	DIVLTQSPASLAMSLGKRATISCRASESVSVIGAHLIHWYQQKPGQPPKLLIYL ASNLETGVPARFSGSGSGTDFTLTIDPVEEDDVAIYSCLQSRIFPRTFGGGTKL EIK	Variable light (VL) Anti-BCMA
73	QVQLVQSGAEVKKPGASVKVSCKASGYSFPDYYINWVRQAPGQGLEWMGWIYFA SGNSEYNQKFTGRVTMTRDTSINTAYMELSSLTSEDTAVYFCASLYDYDWYFDV WGQGTMVTVSS	Variable heavy (VH) Anti-BCMA
74	DIVMTQTPLSLSVTPGQPASISCKSSQSLVHSNGNTYLHWYLQKPGQSPQLLIY KVSNRFSGVPDRFSGSGSGTDFTLKISRVEAEDVGIYYCSQSSIYPWTFGQGTK LEIK	Variable light (VL) Anti-BCMA

75	QVQLVQSGAEVKKPGASVKVSCKASGYSFPDYYINWVRQAPGQGLEWMGWIYFASGNSEYNQKFTGRVTMTRDTSSSTAYMELSSLRSEDTAVYFCASLYDYDWYFDVWGQGTMVTVSS	Variable heavy (VH) Anti-BCMA
76	DIVMTQTPLSLSVTPGEPASISCKSSQSLVHSNGNTYLHWYLQKPGQSPQLLIYKVSNRFSGVPDRFSGSGSGADFTLKISRVEAEDVGVYYCAETSHVPWTFGQGTKLEIK	Variable light (VL) Anti-BCMA
77	QVQLVESGGGLVQPGGSLRLSCEASGFTLDYYAIGWFRQAPGKEREGVICISRSDGSTYYADSVKGRFTISRDNAKKTVYLQMISLKPEDTAAYYCAAGADCSGYLRDYEFRGQGTQVTVSS	Anti-BCMA sdAb
78	IEVMYPPPYLDNEKSNGTIIHVKGKHLCPSPLFPGPSKP	CD28 spacer
79	IYIWAPLAGTCGVLLLSLVITLYCN	CD8a TM
80	LDNEKSNGTIIHVKGKHLCPSPLFPGPSKP	CD28 spacer (truncated)
81	PTTTPAPRPPTPAPTIASQPLSLRPEACRPAAGGAVHTRGLDFACD	CD8a hinge
82	TTTPAPRPPTPAPTIASQPLSLRPEACRPAAGGAVHTRGLDFACD	CD8a hinge
83	FVPVFLPAKPTTTPAPRPPTPAPTIASQPLSLRPEACRPAAGGAVHTRGLDFACD	CD8a hinge
84	DTGLYICKVELMYPPPYYLGIGNGTQIYVIDPEPCPDSD	CTLA4 hinge
85	FLLWILAAVSSGLFFYSFLLTAVS	CTLA4 TM
86	QIKESLRAELRVTERRAEVPTAHPSPSPRPAGQFQTLV	PD-1 hinge
87	VGVVGGLLGSLVLLVWVLAVI	PD-1 TM
88	GLAVSTISSFFPPGYQ	Fc(gamma)RIIIa hinge
89	EPKSPDKTHTCPPCPAPPVAGPSVFLFPPKPKDTLMIARTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYNSTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSRDELTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSKLTVDKSRWQQGNVFSCSVMHEALHNHYTQKSLSLSPGK	IgG1 hinge
90	EVQLLESGGGLVQPGGSLRLSCAASGFTFSSYAMSWVRQAPGKGLEWVSAISGSGGSTYYADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCARAEMGAVFDIWGQGTMVTVSSGSTSGSGKPGSGEGSTKGEIVLTQSPATLSLSPGERATLSCRASQSVSRYLAWYQQKPGQAPRLLIYDASNRATGIPARFSGSGSGTDFTLTISSLEPEDFAVYYCQQRISWPFTFGGGTKVEIKRAAALDNEKSNGTIIHVKGKHLCPSPLFPGPSKPFWVLVVVGGVLACYSLLVTVAFIIFWVRSKRSRLLHSDYMNMTPRRPGPTRKHYQPYAPPRDFAAYRSRVKFSRSADAPAYQQGQNQLYNELNLGRREEYDVLDKRRGRDPEMGGKPRRKNPQEGLYNELQKDKMAEAYSEIGMKGERRRGKGHDGLYQGLSTATKDTYDALHMQALPPR	anti-BCMA CAR
91	EIVLTQSPATLSLSPGERATLSCRASQSVSRYLAWYQQKPGQAPRLLIYDASNRATGIPARFSGSGSGTDFTLTISSLEPEDFAVYYCQQRISWPFTFGGGTKVEIKRGSTSGSGKPGSGEGSTKGEVQLLESGGGLVQPGGSLRLSCAASGFTFSSYAMSWVRQAPGKGLEWVSAISGSGGSTYYADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCARAEMGAVFDIWGQGTMVTVSSAAALDNEKSNGTIIHVKGKHLCPSPLFPGPSKPFWVLVVVGGVLACYSLLVTVAFIIFWVRSKRSRLLHSDYMNMTPRRPGPTRKHYQPYAPPRDFAAYRSRVKFSRSADAPAYQQGQNQLYNELNLGRREEYDVLDKRRGRDPEMGGKPRRKNPQEGLYNELQKDKMAEAYSEIGMKGERRRGKGHDGLYQGLSTATKDTYDALHMQALPPR	anti-BCMA CAR
92	QVQLVESGGGVVQPGRSLRLSCAASGFTFSSYGMHWVRQAPGKGLEWVAVISYDGSNKYYADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCARDGTYLGGLWYFDLWGRGTLVTVSSGSTSGSGKPGSGEGSTKGDIVMTQSPLSLPVTPGEPASISCRSSQSLLHSNGYNYLDWYLQKPGQSPQLLIYLGSNRASGVPDRFSGSGSGTDFTLKISRVEAEDVGVYYCMQGLGLPLTFGGGTKVEIKRAAALDNEKSNGTIIHVKGKHLCPSPLFPGPSKPFWVLVVVGGVLACYSLLVTVAFIIFWVRSKRSRLLHSDYMNMTPRRPGPTRKHYQPYAPPRDFAAYRSRVKFSRSADAPAYQQGQNQLYNELNLGRREEYDVLDKRRGRDPEMGGKPRRKNPQEGLYNELQKDKMAEAYSEIGMKGERRRGKGHDGLYQGLSTATKDTYDALHMQALPPR	anti-BCMA CAR

93	DIVMTQSPLSLPVTPGEPASISCRSSQSLLHSNGYNYLDWYLQKPGQSPQLLIY LGSNRASGVPDRFSGSGSGTDFTLKISRVEAEDVGVYYCMQGLGLPLTFGGGTK VEIKRGSTSGSGKPGSGEGSTKGQVQLVESGGGVVQPGRSLRLSCAASGFTFSS YGMHWVRQAPGKGLEWVAVISYDGSNKYYADSVKGRFTISRDNSKNTLYLQMNS LRAEDTAVYYCARDGTYLGGLWYFDLWGRGTLVTVSSAAALDNEKSNGTIIHVK GKHLCPSPLFPGPSKPFWVLVVVGGVLACYSLLVTVAFIIFWVRSKRSRLLHSD YMNMTPRRPGPTRKHYQPYAPPRDFAAYRSRVKFSRSADAPAYQQGQNQLYNEL NLGRREEYDVLDKRRGRDPEMGGKPRRKNPQEGLYNELQKDKMAEAYSEIGMKG ERRRGKGHDGLYQGLSTATKDTYDALHMQALPPR	anti-BCMA CAR
94	QVQLVQSGAEVKKPGASVKVSCKASGYTFTSYYMHWVRQAPGQGLEWMGIINPG GGSTSYAQKFQGRVTMTRDTSTSTVYMELSSLRSEDTAVYYCARESWPMDVWGQ GTTVTVSSGSTSGSGKPGSGEGSTKGEIVMTQSPATLSVSPGERATLSCRASQS VSSNLAWYQQKPGQAPRLLIYGASTRATGIPARFSGSGSGTEFTLTISSLQSED FAVYYCQQYAAYPTFGGGTKVEIKRAAALDNEKSNGTIIHVKGKHLCPSPLFPG PSKPFWVLVVVGGVLACYSLLVTVAFIIFWVRSKRSRLLHSDYMNMTPRRPGPT RKHYQPYAPPRDFAAYRSRVKFSRSADAPAYQQGQNQLYNELNLGRREEYDVLD KRRGRDPEMGGKPRRKNPQEGLYNELQKDKMAEAYSEIGMKGERRRGKGHDGLY QGLSTATKDTYDALHMQALPPR	anti-BCMA CAR
95	EIVMTQSPATLSVSPGERATLSCRASQSVSSNLAWYQQKPGQAPRLLIYGASTR ATGIPARFSGSGSGTEFTLTISSLQSEDFAVYYCQQYAAYPTFGGGTKVEIKRG STSGSGKPGSGEGSTKGQVQLVQSGAEVKKPGASVKVSCKASGYTFTSYYMHWV RQAPGQGLEWMGIINPGGGSTSYAQKFQGRVTMTRDTSTSTVYMELSSLRSEDT AVYYCARESWPMDVWGQGTTVTVSSAAALDNEKSNGTIIHVKGKHLCPSPLFPG PSKPFWVLVVVGGVLACYSLLVTVAFIIFWVRSKRSRLLHSDYMNMTPRRPGPT RKHYQPYAPPRDFAAYRSRVKFSRSADAPAYQQGQNQLYNELNLGRREEYDVLD KRRGRDPEMGGKPRRKNPQEGLYNELQKDKMAEAYSEIGMKGERRRGKGHDGLY QGLSTATKDTYDALHMQALPPR	anti-BCMA CAR
96	QLQLQESGPGLVKPSETLSLTCTVSGGSISSSSYYWGWIRQPPGKGLEWIGSIS YSGSTYYNPSLKSRVTISVDTSKNQFSLKLSSVTAADTAVYYCARGRGYATSLA FDIWGQGTMVTVSSGSTSGSGKPGSGEGSTKGEIVLTQSPATLSLSPGERATLS CRASQSVSSYLAWYQQKPGQAPRLLIYDASNRATGIPARFSGSGSGTDFTLTIS SLEPEDFAVYYCQQRHVWPPTFGGGTKVEIKRAAALDNEKSNGTIIHVKGKHLC PSPLFPGPSKPFWVLVVVGGVLACYSLLVTVAFIIFWVRSKRSRLLHSDYMNMT PRRPGPTRKHYQPYAPPRDFAAYRSRVKFSRSADAPAYQQGQNQLYNELNLGRR EEYDVLDKRRGRDPEMGGKPRRKNPQEGLYNELQKDKMAEAYSEIGMKGERRRG KGHDGLYQGLSTATKDTYDALHMQALPPR	anti-BCMA CAR
97	EIVLTQSPATLSLSPGERATLSCRASQSVSSYLAWYQQKPGQAPRLLIYDASNR ATGIPARFSGSGSGTDFTLTISSLEPEDFAVYYCQQRHVWPPTFGGGTKVEIKR GSTSGSGKPGSGEGSTKGQLQLQESGPGLVKPSETLSLTCTVSGGSISSSSYYW GWIRQPPGKGLEWIGSISYSGSTYYNPSLKSRVTISVDTSKNQFSLKLSSVTAA DTAVYYCARGRGYATSLAFDIWGQGTMVTVSSAAALDNEKSNGTIIHVKGKHLC PSPLFPGPSKPFWVLVVVGGVLACYSLLVTVAFIIFWVRSKRSRLLHSDYMNMT PRRPGPTRKHYQPYAPPRDFAAYRSRVKFSRSADAPAYQQGQNQLYNELNLGRR EEYDVLDKRRGRDPEMGGKPRRKNPQEGLYNELQKDKMAEAYSEIGMKGERRRG KGHDGLYQGLSTATKDTYDALHMQALPPR	anti-BCMA CAR
98	EVQLVESGGGLVQPGGSLRLSCAASGFTFSSYSMNWVRQAPGKGLEWVSTISSS SSTIYYADSVKGRFTISRDNAKNSLYLQMNSLRAEDTAVYYCARGSQEHLIFDY WGQGTLVTVSSGSTSGSGKPGSGEGSTKGEIVLTQSPATLSLSPGERATLSCRA SQSVSRYLAWYQQKPGQAPRLLIYDASNRATGIPARFSGSGSGTDFTLTISSLE PEDFAVYYCQQRFYYPWTFGGGTKVEIKRAAALDNEKSNGTIIHVKGKHLCPSP LFPGPSKPFWVLVVVGGVLACYSLLVTVAFIIFWVRSKRSRLLHSDYMNMTPRR PGPTRKHYQPYAPPRDFAAYRSRVKFSRSADAPAYQQGQNQLYNELNLGRREEY DVLDKRRGRDPEMGGKPRRKNPQEGLYNELQKDKMAEAYSEIGMKGERRRGKGH DGLYQGLSTATKDTYDALHMQALPPR	anti-BCMA CAR
99	EIVLTQSPATLSLSPGERATLSCRASQSVSRYLAWYQQKPGQAPRLLIYDASNR ATGIPARFSGSGSGTDFTLTISSLEPEDFAVYYCQQRFYYPWTFGGGTKVEIKR GSTSGSGKPGSGEGSTKGEVQLVESGGGLVQPGGSLRLSCAASGFTFSSYSMNW	anti-BCMA CAR

	VRQAPGKGLEWVSTISSSSSTIYYADSVKGRFTISRDNAKNSLYLQMNSLRAED TAVYYCARGSQEHLIFDYWGQGTLVTVSSAAALDNEKSNGTIIHVKGKHLCPSP LFPGPSKPFWVLVVVGGVLACYSLLVTVAFIIFWVRSKRSRLLHSDYMNMTPRR PGPTRKHYQPYAPPRDFAAYRSRVKFSRSADAPAYQQGQNQLYNELNLGRREEY DVLDKRRGRDPEMGGKPRRKNPQEGLYNELQKDKMAEAYSEIGMKGERRRGKGH DGLYQGLSTATKDTYDALHMQALPPR	
100	QVQLVESGGGVVQPGRSLRLSCAASGFTFSSYGMHWVRQAPGKGLEWVAVISYD GSNKYYADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCARTDFWSGSPPG LDYWGQGTLVTVSSGSTSGSGKPGSGEGSTKGDIQLTQSPSSVSASVGDRVTIT CRASQGISSWLAWYQQKPGKAPKLLIYGASSLQSGVPSRFSGSGSGTDFTLTIS SLQPEDFATYYCQQIYTFPFTFGGGTKVEIKRAAALDNEKSNGTIIHVKGKHLC PSPLFPGPSKPFWVLVVVGGVLACYSLLVTVAFIIFWVRSKRSRLLHSDYMNMT PRRPGPTRKHYQPYAPPRDFAAYRSRVKFSRSADAPAYQQGQNQLYNELNLGRR EEYDVLDKRRGRDPEMGGKPRRKNPQEGLYNELQKDKMAEAYSEIGMKGERRRG KGHDGLYQGLSTATKDTYDALHMQALPPR	anti-BCMA CAR
101	DIQLTQSPSSVSASVGDRVTITCRASQGISSWLAWYQQKPGKAPKLLIYGASSL QSGVPSRFSGSGSGTDFTLTISSLQPEDFATYYCQQIYTFPFTFGGGTKVEIKR GSTSGSGKPGSGEGSTKGQVQLVESGGGVVQPGRSLRLSCAASGFTFSSYGMHW VRQAPGKGLEWVAVISYDGSNKYYADSVKGRFTISRDNSKNTLYLQMNSLRAED TAVYYCARTDFWSGSPPGLDYWGQGTLVTVSSAAALDNEKSNGTIIHVKGKHLC PSPLFPGPSKPFWVLVVVGGVLACYSLLVTVAFIIFWVRSKRSRLLHSDYMNMT PRRPGPTRKHYQPYAPPRDFAAYRSRVKFSRSADAPAYQQGQNQLYNELNLGRR EEYDVLDKRRGRDPEMGGKPRRKNPQEGLYNELQKDKMAEAYSEIGMKGERRRG KGHDGLYQGLSTATKDTYDALHMQALPPR	anti-BCMA CAR
102	QVQLVQSGAEVKKPGSSVKVSCKASGGTFSSYAISWVRQAPGQGLEWMGGIIPI FGTANYAQKFQGRVTITADESTSTAYMELSSLRSEDTAVYYCARTPEYSSSIWH YYYGMDVWGQGTTVTVSSGSTSGSGKPGSGEGSTKGDIVMTQSPDSLAVSLGER ATINCKSSQSVLYSSNNKNYLAWYQQKPGQPPKLLIYWASTRESGVPDRFSGSG SGTDFTLTISSLQAEDVAVYYCQQFAHTPFTFGGGTKVEIKRAAALDNEKSNGT IIHVKGKHLCPSPLFPGPSKPFWVLVVVGGVLACYSLLVTVAFIIFWVRSKRSR LLHSDYMNMTPRRPGPTRKHYQPYAPPRDFAAYRSRVKFSRSADAPAYQQGQNQ LYNELNLGRREEYDVLDKRRGRDPEMGGKPRRKNPQEGLYNELQKDKMAEAYSE IGMKGERRRGKGHDGLYQGLSTATKDTYDALHMQALPPR	anti-BCMA CAR
103	DIVMTQSPDSLAVSLGERATINCKSSQSVLYSSNNKNYLAWYQQKPGQPPKLLI YWASTRESGVPDRFSGSGSGTDFTLTISSLQAEDVAVYYCQQFAHTPFTFGGGT KVEIKRGSTSGSGKPGSGEGSTKGQVQLVQSGAEVKKPGSSVKVSCKASGGTFS SYAISWVRQAPGQGLEWMGGIIPIFGTANYAQKFQGRVTITADESTSTAYMELS SLRSEDTAVYYCARTPEYSSSIWHYYYGMDVWGQGTTVTVSSAAALDNEKSNGT IIHVKGKHLCPSPLFPGPSKPFWVLVVVGGVLACYSLLVTVAFIIFWVRSKRSR LLHSDYMNMTPRRPGPTRKHYQPYAPPRDFAAYRSRVKFSRSADAPAYQQGQNQ LYNELNLGRREEYDVLDKRRGRDPEMGGKPRRKNPQEGLYNELQKDKMAEAYSE IGMKGERRRGKGHDGLYQGLSTATKDTYDALHMQALPPR	anti-BCMA CAR
104	QVQLVESGGGVVQPGRSLRLSCAASGFTFSSYGMHWVRQAPGKGLEWVAVISYD GSNKYYADSVKGRFTISRDNSKNTLYLQMNSLRAEDTAVYYCVKGPLQEPPYDY GMDVWGQGTTVTVSSGSTSGSGKPGSGEGSTKGEIVMTQSPATLSVSPGERATL SCRASQSVSSNLAWYQQKPGQAPRLLIYSASTRATGIPARFSGSGSGTEFTLTI SSLQSEDFAVYYCQQHHVWPLTFGGGTKVEIKRAAALDNEKSNGTIIHVKGKHL CPSPLFPGPSKPFWVLVVVGGVLACYSLLVTVAFIIFWVRSKRSRLLHSDYMNM TPRRPGPTRKHYQPYAPPRDFAAYRSRVKFSRSADAPAYQQGQNQLYNELNLGR REEYDVLDKRRGRDPEMGGKPRRKNPQEGLYNELQKDKMAEAYSEIGMKGERRR GKGHDGLYQGLSTATKDTYDALHMQALPPR	anti-BCMA CAR
105	EIVMTQSPATLSVSPGERATLSCRASQSVSSNLAWYQQKPGQAPRLLIYSASTR ATGIPARFSGSGSGTEFTLTISSLQSEDFAVYYCQQHHVWPLTFGGGTKVEIKR GSTSGSGKPGSGEGSTKGQVQLVESGGGVVQPGRSLRLSCAASGFTFSSYGMHW VRQAPGKGLEWVAVISYDGSNKYYADSVKGRFTISRDNSKNTLYLQMNSLRAED TAVYYCVKGPLQEPPYDYGMDVWGQGTTVTVSSAAALDNEKSNGTIIHVKGKHL CPSPLFPGPSKPFWVLVVVGGVLACYSLLVTVAFIIFWVRSKRSRLLHSDYMNM	anti-BCMA CAR

	TPRRPGPTRKHYQPYAPPRDFAAYRSRVKFSRSADAPAYQQGQNQLYNELNLGR REEYDVLDKRRGRDPEMGGKPRRKNPQEGLYNELQKDKMAEAYSEIGMKGERRR GKGHDGLYQGLSTATKDTYDALHMQALPPR	
106	QSALTQPASVSASPGQSIAISCTGTSSDVGWYQQHPGKAPKLMIYEDSKRPSGV SNRFSGSKSGNTASLTISGLQAEDEADYYCSSNTRSSTLVFGGGTKLTVLGSRG GGGSGGGGSGGGGSLEMAEVQLVQSGAEMKKPGASLKLSCKASGYTFIDYYVYW MRQAPGQGLESMGWINPNSGGTNYAQKFQGRVTMTRDTSISTAYMELSRLRSDD TAMYYCARSQRDGYMDYWGQGTLVTVSSAAAIEVMYPPPYLDNEKSNGTIIHVK GKHLCPSPLFPGPSKPFWVLVVVGGVLACYSLLVTVAFIIFWVRSKRSRLLHSD YMNMTPRRPGPTRKHYQPYAPPRDFAAYRSRVKFSRSADAPAYQQGQNQLYNEL NLGRREEYDVLDKRRGRDPEMGGKPRRKNPQEGLYNELQKDKMAEAYSEIGMKG ERRRGKGHDGLYQGLSTATKDTYDALHMQALPPR	anti-BCMA CAR
107	QSVLTQPPSVSGAPGQRVTISCTGSSSNIGAGFDVHWYQQLPGTAPKLLIYGNS NRPSGVPDRFSGSKSGTSASLAITGLQAEDEADYYCQSYDSSLSGYVFGTGTKV TVLGSRGGGGSGGGGSGGGGSLEMAQVQLVQSGAEVKKPGASVKVSCKASGYTF TDYYMHWVRQAPGQRLEWMGWINPNSGGTNYAQKFQDRITVTRDTSSNTGYMEL TRLRSDDTAVYYCARSPYSGVLDKWGQGTLVTVSSAAAIEVMYPPPYLDNEKSN GTIIHVKGKHLCPSPLFPGPSKPFWVLVVVGGVLACYSLLVTVAFIIFWVRSKR SRLLHSDYMNMTPRRPGPTRKHYQPYAPPRDFAAYRSRVKFSRSADAPAYQQGQ NQLYNELNLGRREEYDVLDKRRGRDPEMGGKPRRKNPQEGLYNELQKDKMAEAY SEIGMKGERRRGKGHDGLYQGLSTATKDTYDALHMQALPPR	anti-BCMA CAR
108	SYELTQPPSASGTPGQRVTMSCSGTSSNIGSHSVNWYQQLPGTAPKLLIYTNNQ RPSGVPDRFSGSKSGTSASLAISGLQSEDEADYYCAAWDGSLNGLVFGGGTKLT VLGSRGGGGSGGGGSGGGGSLEMAEVQLVQSGAEVKKPGESLKISCKGSGYSFT SYWIGWVRQMPGKGLEWMGIIYPGDSDTRYSPSFQGHVTISADKSISTAYLQWS SLKASDTAMYYCARYSGSFDNWGQGTLVTVSSAAAIEVMYPPPYLDNEKSNGTI IHVKGKHLCPSPLFPGPSKPFWVLVVVGGVLACYSLLVTVAFIIFWVRSKRSRL LHSDYMNMTPRRPGPTRKHYQPYAPPRDFAAYRSRVKFSRSADAPAYQQGQNQL YNELNLGRREEYDVLDKRRGRDPEMGGKPRRKNPQEGLYNELQKDKMAEAYSEI GMKGERRRGKGHDGLYQGLSTATKDTYDALHMQALPPR	anti-BCMA CAR
109	LPVLTQPPSASGTPGQRVTISCSGRSSNIGSNSVNWYRQLPGAAPKLLIYSNNQ RPPGVPVRFSGSKSGTSASLAISGLQSEDEATYYCATWDDNLNVHYVFGTGTKV TVLGSRGGGGSGGGGSGGGGSLEMAQVQLVQSGAEVKKPGSSVKVSCKASGGTF SSYAISWVRQAPGQGLEWMGRIIPILGIANYAQKFQGRVTITADKSTSTAYMEL SSLRSEDTAVYYCARGGYYSHDMWSEDWGQGTLVTVSSAAAIEVMYPPPYLDNE KSNGTIIHVKGKHLCPSPLFPGPSKPFWVLVVVGGVLACYSLLVTVAFIIFWVR SKRSRLLHSDYMNMTPRRPGPTRKHYQPYAPPRDFAAYRSRVKFSRSADAPAYQ QGQNQLYNELNLGRREEYDVLDKRRGRDPEMGGKPRRKNPQEGLYNELQKDKMA EAYSEIGMKGERRRGKGHDGLYQGLSTATKDTYDALHMQALPPR	anti-BCMA CAR
110	QAVLTQPPSASGTPGQRVTISCSGSSSNIGSNYVFWYQQLPGTAPKLLIYSNNQ RPSGVPDRFSGSKSGTSASLAISGLRSEDEADYYCAAWDDSLSASYVFGTGTKV TVLGSRGGGGSGGGGSGGGGSLEMAQVQLVQSGAEVKKPGSSVKVSCKASGGTF SSYAISWVRQAPGQGLEWMGRIIPILGTANYAQKFQGRVTITADESTSTAYMEL SSLRSEDTAVYYCARSGYGSYRWEDSWGQGTLVTVSSAAAIEVMYPPPYLDNEK SNGTIIHVKGKHLCPSPLFPGPSKPFWVLVVVGGVLACYSLLVTVAFIIFWVRS KRSRLLHSDYMNMTPRRPGPTRKHYQPYAPPRDFAAYRSRVKFSRSADAPAYQQ GQNQLYNELNLGRREEYDVLDKRRGRDPEMGGKPRRKNPQEGLYNELQKDKMAE AYSEIGMKGERRRGKGHDGLYQGLSTATKDTYDALHMQALPPR	anti-BCMA CAR
111	LPVLTQPPSASGTPGQRVTISCSGRSSNIGSNSVNWYRQLPGAAPKLLIYSNNQ RPPGVPVRFSGSKSGTSASLAISGLQSEDEATYYCATWDDNLNVHYVFGTGTKV TVLGSRGGGGSGGGGSGGGGSLEMAQVQLVQSGAEVKKPGSSVKVSCKASGGTF SSYAISWVRQAPGQGLEWMGRIIPILGIANYAQKFQGRVTITADKSTSTAYMEL SSLRSEDTAVYYCARGGYYSHDMWSEDWGQGTLVTVSSAAAPTTTPAPRPPTPA PTIASQPLSLRPEACRPAAGGAVHTRGLDFACDIYIWAPLAGTCGVLLLSLVIT LYCNKRGRKKLLYIFKQPFMRPVQTTQEEDGCSCRFPEEEEGGCELRVKFSRSA EPPAYQQGQNQLYNELNLGRREEYDVLDKRRGRDPEMGGKPRRKNPQEGLYNEL QKDKMAEAYSEIGMKGERRRGKGHDGLYQGLSTATKDTYDALHMQALPPR	anti-BCMA CAR

112	SYELTQPPSASGTPGQRVTMSCSGTSSNIGSHSVNWYQQLPGTAPKLLIYTNNQ RPSGVPDRFSGSKSGTSASLAISGLQSEDEADYYCAAWDGSLNGLVFGGGTKLT VLGSRGGGGSGGGGSGGGGSLEMAEVQLVQSGAEVKKPGESLKISCKGSGYSFT SYWIGWVRQMPGKGLEWMGIIYPGDSDTRYSPSFQGHVTISADKSISTAYLQWS SLKASDTAMYYCARYSGSFDNWGQGTLVTVSSAAAPTTTPAPRPPTPAPTIASQ PLSLRPEACRPAAGGAVHTRGLDFACDIYIWAPLAGTCGVLLLSLVITLYCNKR GRKKLLYIFKQPFMRPVQTTQEEDGCSCRFPEEEEGGCELRVKFSRSAEPPAYQ QGQNQLYNELNLGRREEYDVLDKRRGRDPEMGGKPRRKNPQEGLYNELQKDKMA EAYSEIGMKGERRRGKGHDGLYQGLSTATKDTYDALHMQALPPR	anti-BCMA CAR
113	QAVLTQPPSASGTPGQRVTISCSGSSSNIGSNYVFWYQQLPGTAPKLLIYSNNQ RPSGVPDRFSGSKSGTSASLAISGLRSEDEADYYCAAWDDSLSASYVFGTGTKV TVLGSRGGGGSGGGGSGGGGSLEMAQVQLVQSGAEVKKPGSSVKVSCKASGGTF SSYAISWVRQAPGQGLEWMGRIIPILGTANYAQKFQGRVTITADESTSTAYMEL SSLRSEDTAVYYCARSGYGSYRWEDSWGQGTLVTVSSAAAPTTTPAPRPPTPAP TIASQPLSLRPEACRPAAGGAVHTRGLDFACDIYIWAPLAGTCGVLLLSLVITL YCNKRGRKKLLYIFKQPFMRPVQTTQEEDGCSCRFPEEEEGGCELRVKFSRSAE PPAYQQGQNQLYNELNLGRREEYDVLDKRRGRDPEMGGKPRRKNPQEGLYNELQ KDKMAEAYSEIGMKGERRRGKGHDGLYQGLSTATKDTYDALHMQALPPR	anti-BCMA CAR
114	QSVLTQPPSVSGAPGQRVTISCTGSSSNIGAGFDVHWYQQLPGTAPKLLIYGNS NRPSGVPDRFSGSKSGTSASLAITGLQAEDEADYYCQSYDSSLSGYVFGTGTKV TVLGSRGGGGSGGGGSGGGGSLEMAQVQLVQSGAEVKKPGASVKVSCKASGYTF TDYYMHWVRQAPGQRLEWMGWINPNSGGTNYAQKFQDRITVTRDTSSNTGYMEL TRLRSDDTAVYYCARSPYSGVLDKWGQGTLVTVSSAAAPTTTPAPRPPTPAPTI ASQPLSLRPEACRPAAGGAVHTRGLDFACDIYIWAPLAGTCGVLLLSLVITLYC NKRGRKKLLYIFKQPFMRPVQTTQEEDGCSCRFPEEEEGGCELRVKFSRSAEPP AYQQGQNQLYNELNLGRREEYDVLDKRRGRDPEMGGKPRRKNPQEGLYNELQKD KMAEAYSEIGMKGERRRGKGHDGLYQGLSTATKDTYDALHMQALPPR	anti-BCMA CAR
115	QSALTQPASVSASPGQSIAISCTGTSSDVGWYQQHPGKAPKLMIYEDSKRPSGV SNRFSGSKSGNTASLTISGLQAEDEADYYCSSNTRSSTLVFGGGTKLTVLGSRG GGGSGGGGSGGGGSLEMAEVQLVQSGAEMKKPGASLKLSCKASGYTFIDYYVYW MRQAPGQGLESMGWINPNSGGTNYAQKFQGRVTMTRDTSISTAYMELSRLRSDD TAMYYCARSQRDGYMDYWGQGTLVTVSSAAAPTTTPAPRPPTPAPTIASQPLSL RPEACRPAAGGAVHTRGLDFACDIYIWAPLAGTCGVLLLSLVITLYCNKRGRKK LLYIFKQPFMRPVQTTQEEDGCSCRFPEEEEGGCELRVKFSRSAEPPAYQQGQN QLYNELNLGRREEYDVLDKRRGRDPEMGGKPRRKNPQEGLYNELQKDKMAEAYS EIGMKGERRRGKGHDGLYQGLSTATKDTYDALHMQALPPR	anti-BCMA CAR
116	DIVLTQSPPSLAMSLGKRATISCRASESVTILGSHLIHWYQQKPGQPPTLLIQL ASNVQTGVPARFSGSGSRTDFTLTIDPVEEDDVAVYYCLQSRTIPRTFGGGTKL EIKGSTSGSGKPGSGEGSTKGQIQLVQSGPELKKPGETVKISCKASGYTFTDYS INWVKRAPGKGLKWMGWINTETREPAYAYDFRGRFAFSLETSASTAYLQINNLK YEDTATYFCALDYSYAMDYWGQGTSVTVSSAAATTTPAPRPPTPAPTIASQPLS LRPEACRPAAGGAVHTRGLDFACDIYIWAPLAGTCGVLLLSLVITLYCKRGRKK LLYIFKQPFMRPVQTTQEEDGCSCRFPEEEEGGCELRVKFSRSADAPAYQQGQN QLYNELNLGRREEYDVLDKRRGRDPEMGGKPRRKNPQEGLYNELQKDKMAEAYS EIGMKGERRRGKGHDGLYQGLSTATKDTYDALHMQALPPR	anti-BCMA CAR
117	DIVLTQSPASLAVSLGERATINCRASESVSVIGAHLIHWYQQKPGQPPKLLIYL ASNLETGVPARFSGSGSGTDFTLTISSLQAEDAAIYYCLQSRIFPRTFGQGTKL EIKGSTSGSGKPGSGEGSTKGQVQLVQSGSELKKPGASVKVSCKASGYTFTDYS INWVRQAPGQGLEWMGWINTETREPAYAYDFRGRFVFSLDTSVSTAYLQISSLK AEDTAVYYCARDYSYAMDYWGQGTLVTVSSAAATTTPAPRPPTPAPTIASQPLS LRPEACRPAAGGAVHTRGLDFACDIYIWAPLAGTCGVLLLSLVITLYCKRGRKK LLYIFKQPFMRPVQTTQEEDGCSCRFPEEEEGGCELRVKFSRSADAPAYQQGQN QLYNELNLGRREEYDVLDKRRGRDPEMGGKPRRKNPQEGLYNELQKDKMAEAYS EIGMKGERRRGKGHDGLYQGLSTATKDTYDALHMQALPPR	anti-BCMA CAR
118	DIVLTQSPASLAVSLGERATINCRASESVSVIGAHLIHWYQQKPGQPPKLLIYL ASNLETGVPARFSGSGSGTDFTLTISSLQAEDAAIYYCLQSRIFPRTFGQGTKL EIKGSTSGSGKPGSGEGSTKGQVQLVQSGSELKKPGASVKVSCKASGYTFTDYS	anti-BCMA CAR

	INWVRQAPGQGLEWMGWINTETREPAYAYDFRGRFVFSLDTSVSTAYLQISSLK AEDTAVYYCARDYSYAMDYWGQGTLVTVSSAAADTGLYICKVELMYPPPYYLGI GNGTQIYVIDPEPCPDSDFLLWILAAVSSGLFFYSFLLTAVSKRGRKKLLYIFK QPFMRPVQTTQEEDGCSCRFPEEEEGGCELRVKFSRSADAPAYQQGQNQLYNEL NLGRREEYDVLDKRRGRDPEMGGKPRRKNPQEGLYNELQKDKMAEAYSEIGMKG ERRRGKGHDGLYQGLSTATKDTYDALHMQALPPR	
119	DIVLTQSPASLAVSLGERATINCRASESVSVIGAHLIHWYQQKPGQPPKLLIYL ASNLETGVPARFSGSGSGTDFTLTISSLQAEDAAIYYCLQSRIFPRTFGQGTKL EIKGSTSGSGKPGSGEGSTKGQVQLVQSGSELKKPGASVKVSCKASGYTFTDYS INWVRQAPGQGLEWMGWINTETREPAYAYDFRGRFVFSLDTSVSTAYLQISSLK AEDTAVYYCARDYSYAMDYWGQGTLVTVSSAAAQIKESLRAELRVTERRAEVPT AHPSPSPRPAGQFQTLVVGVVGGLLGSLVLLVWVLAVICSKRGRKKLLYIFKQP FMRPVQTTQEEDGCSCRFPEEEEGGCELRVKFSRSADAPAYQQGQNQLYNELNL GRREEYDVLDKRRGRDPEMGGKPRRKNPQEGLYNELQKDKMAEAYSEIGMKGER RRGKGHDGLYQGLSTATKDTYDALHMQALPPR	anti-BCMA CAR
120	EVQLVESGGGLVKPGGSLRLSCAASGFTFSDYYMSWIRQAPGKGLEWVSYISSS GSTIYYADSVKGRFTISRDNAKNSLYLQMNSLRAEDTAVYYCAKVDGDYTEDYW GQGTLVTVSSGGGGSGGGGSGGGGSQSALTQPASVSGSPGQSITISCTGSSSDV GKYNLVSWYQQPPGKAPKLIIYDVNKRPSGVSNRFSGSKSGNTATLTISGLQGD DEADYYCSSYGGSRSYVFGTGTKVTVLESKYGPPCPPCPAPPVAGPSVFLFPPK PKDTLMISRTPEVTCVVVDVSQEDPEVQFNWYVDGVEVHNAKTKPREEQFQSTY RVVSVLTVLHQDWLNGKEYKCKVSNKGLPSSIEKTISKAKGQPREPQVYTLPPS QEEMTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYS RLTVDKSRWQEGNVFSCSVMHEALHNHYTQKSLSLSLGKMFWVLVVVGGVLACY SLLVTVAFIIFWVKRGRKKLLYIFKQPFMRPVQTTQEEDGCSCRFPEEEEGGCE LRVKFSRSADAPAYQQGQNQLYNELNLGRREEYDVLDKRRGRDPEMGGKPRRKN PQEGLYNELQKDKMAEAYSEIGMKGERRRGKGHDGLYQGLSTATKDTYDALHMQ ALPPR	anti-BCMA CAR
121	EVQLVQSGGGLVQPGRSLRLSCTASGFTFGDYAMSWFKQAPGKGLEWVGFIRSK AYGGTTEYAASVKGRFTISRDDSKSIAYLQMNSLKTEDTAVYYCAAWSAPTDYW GQGTLVTVSSGGGGSGGGGSGGGGSDIQMTQSPAFLSASVGDRVTVTCRASQGI SNYLAWYQQKPGNAPRLLIYSASTLQSGVPSRFRGTGYGTEFSLTIDSLQPEDF ATYYCQQSYTSRQTFGPGTRLDIKESKYGPPCPPCPAPPVAGPSVFLFPPKPKD TLMISRTPEVTCVVVDVSQEDPEVQFNWYVDGVEVHNAKTKPREEQFQSTYRVV SVLTVLHQDWLNGKEYKCKVSNKGLPSSIEKTISKAKGQPREPQVYTLPPSQEE MTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSRLT VDKSRWQEGNVFSCSVMHEALHNHYTQKSLSLSLGKMFWVLVVVGGVLACYSLL VTVAFIIFWVKRGRKKLLYIFKQPFMRPVQTTQEEDGCSCRFPEEEEGGCELRV KFSRSADAPAYQQGQNQLYNELNLGRREEYDVLDKRRGRDPEMGGKPRRKNPQE GLYNELQKDKMAEAYSEIGMKGERRRGKGHDGLYQGLSTATKDTYDALHMQALP PR	anti-BCMA CAR
122	EVQLVESGGGLVKPGGSLRLSCAASGFTFSDYYMSWIRQAPGKGLEWVSYISSS GSTIYYADSVKGRFTISRDNAKNSLYLQMNSLRAEDTAVYYCAKVDGPPSFDIW GQGTMVTVSSGGGGSGGGGSGGGGSSYVLTQPPSVSVAPGQTARITCGANNIGS KSVHWYQQKPGQAPMLVVYDDDDRPSGIPERFSGSNSGNTATLTISGVEAGDEA DYFCHLWDRSRDHYVFGTGTKLTVLESKYGPPCPPCPAPPVAGPSVFLFPPKPK DTLMISRTPEVTCVVVDVSQEDPEVQFNWYVDGVEVHNAKTKPREEQFQSTYRV VSVLTVLHQDWLNGKEYKCKVSNKGLPSSIEKTISKAKGQPREPQVYTLPPSQE EMTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLYSRL TVDKSRWQEGNVFSCSVMHEALHNHYTQKSLSLSLGKMFWVLVVVGGVLACYSL LVTVAFIIFWVKRGRKKLLYIFKQPFMRPVQTTQEEDGCSCRFPEEEEGGCELR VKFSRSADAPAYQQGQNQLYNELNLGRREEYDVLDKRRGRDPEMGGKPRRKNPQ EGLYNELQKDKMAEAYSEIGMKGERRRGKGHDGLYQGLSTATKDTYDALHMQAL PPR	anti-BCMA CAR
123	SYELTQPPSASGTPGQRVTMSCSGTSSNIGSHSVNWYQQLPGTAPKLLIYTNNQ RPSGVPDRFSGSKSGTSASLAISGLQSEDEADYYCAAWDGSLNGLVFGGGTKLT VLGSRGGGGSGGGGSGGGGSLEMAEVQLVQSGAEVKKPGESLKISCKGSGYSFT	anti-BCMA CAR

	SYWIGWVRQMPGKGLEWMGIIYPGDSDTRYSPSFQGHVTISADKSISTAYLQWS SLKASDTAMYYCARYSGSFDNWGQGTLVTVSSESKYGPPCPPCPAPPVAGPSVF LFPPKPKDTLMISRTPEVTCVVVDVSQEDPEVQFNWYVDGVEVHNAKTKPREEQ FQSTYRVVSVLTVLHQDWLNGKEYKCKVSNKGLPSSIEKTISKAKGQPREPQVY TLPPSQEEMTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGS FFLYSRLTVDKSRWQEGNVFSCSVMHEALHNHYTQKSLSLSLGKMFWVLVVVGG VLACYSLLVTVAFIIFWVKRGRKKLLYIFKQPFMRPVQTTQEEDGCSCRFPEEE EGGCELRVKFSRSADAPAYQQGQNQLYNELNLGRREEYDVLDKRRGRDPEMGGK PRRKNPQEGLYNELQKDKMAEAYSEIGMKGERRRGKGHDGLYQGLSTATKDTYD ALHMQALPPR	
124	QSALTQPASVSASPGQSIAISCTGTSSDVGWYQQHPGKAPKLMIYEDSKRPSGV SNRFSGSKSGNTASLTISGLQAEDEADYYCSSNTRSSTLVFGGGTKLTVLGSRG GGGSGGGGSGGGGSLEMAEVQLVQSGAEMKKPGASLKLSCKASGYTFIDYYVYW MRQAPGQGLESMGWINPNSGGTNYAQKFQGRVTMTRDTSISTAYMELSRLRSDD TAMYYCARSQRDGYMDYWGQGTLVTVSSESKYGPPCPPCPAPPVAGPSVFLFPP KPKDTLMISRTPEVTCVVVDVSQEDPEVQFNWYVDGVEVHNAKTKPREEQFQST YRVVSVLTVLHQDWLNGKEYKCKVSNKGLPSSIEKTISKAKGQPREPQVYTLPP SQEEMTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLY SRLTVDKSRWQEGNVFSCSVMHEALHNHYTQKSLSLSLGKMFWVLVVVGGVLAC YSLLVTVAFIIFWVKRGRKKLLYIFKQPFMRPVQTTQEEDGCSCRFPEEEEGGC ELRVKFSRSADAPAYQQGQNQLYNELNLGRREEYDVLDKRRGRDPEMGGKPRRK NPQEGLYNELQKDKMAEAYSEIGMKGERRRGKGHDGLYQGLSTATKDTYDALHM QALPPR	anti-BCMA CAR
125	QSALTQPASVSASPGQSIAISCTGTSSDVGWYQQHPGKAPKLMIYEDSKRPSGV SNRFSGSKSGNTASLTISGLQAEDEADYYCSSNTRSSTLVFGGGTKLTVLGSRG GGGSGGGGSGGGGSLEMAEVQLVQSGAEMKKPGASLKLSCKASGYTFIDYYVYW MRQAPGQGLESMGWINPNSGGTNYAQKFQGRVTMTRDTSISTAYMELSRLRSDD TAMYYCARSQRDGYMDYWGQGTLVTVSSESKYGPPCPPCPAPPVAGPSVFLFPP KPKDTLMISRTPEVTCVVVDVSQEDPEVQFNWYVDGVEVHNAKTKPREEQFQST YRVVSVLTVLHQDWLNGKEYKCKVSNKGLPSSIEKTISKAKGQPREPQVYTLPP SQEEMTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDGSFFLY SRLTVDKSRWQEGNVFSCSVMHEALHNHYTQKSLSLSLGKMFWVLVVVGGVLAC YSLLVTVAFIIFWVRSKRSRLLHSDYMNMTPRRPGPTRKHYQPYAPPRDFAAYR SRVKFSRSADAPAYQQGQNQLYNELNLGRREEYDVLDKRRGRDPEMGGKPRRKN PQEGLYNELQKDKMAEAYSEIGMKGERRRGKGHDGLYQGLSTATKDTYDALHMQ ALPPR	anti-BCMA CAR
126	EVQLVESGGGLVQPGGSLRLSCAVSGFALSNHGMSWVRRAPGKGLEWVSGIVYS GSTYYAASVKGRFTISRDNSRNTLYLQMNSLRPEDTAIYYCSAHGGESDVWGQG TTVTVSSASGGGGSGGRASGGGGSDIQLTQSPSSLSASVGDRVTITCRASQSIS SYLNWYQQKPGKAPKLLIYAASSLQSGVPSRFSGSGSGTDFTLTISSLQPEDFA TYYCQQSYSTPYTFGQGTKVEIKTTTPAPRPPTPAPTIASQPLSLRPEACRPAA GGAVHTRGLDFACDIYIWAPLAGTCGVLLLSLVITLYCKRGRKKLLYIFKQPFM RPVQTTQEEDGCSCRFPEEEEGGCELRVKFSRSADAPAYKQGQNQLYNELNLGR REEYDVLDKRRGRDPEMGGKPRRKNPQEGLYNELQKDKMAEAYSEIGMKGERRR GKGHDGLYQGLSTATKDTYDALHMQALPPR	anti-BCMA CAR
127	QVQLVESGGGLVQPGRSLRLSCAASGFTFSNYAMSWVRQAPGKGLGWVSGISRS GENTYYADSVKGRFTISRDNSKNTLYLQMNSLRDEDTAVYYCARSPAHYYGGMD VWGQGTTVTVSSASGGGGSGGRASGGGGSDIVLTQSPGTLSLSPGERATLSCRA SQSISSSFLAWYQQKPGQAPRLLIYGASRRATGIPDRFSGSGSGTDFTLTISRL EPEDSAVYYCQQYHSSPSWTFGQGTKLEIKTTTPAPRPPTPAPTIASQPLSLRP EACRPAAGGAVHTRGLDFACDIYIWAPLAGTCGVLLLSLVITLYCKRGRKKLLY IFKQPFMRPVQTTQEEDGCSCRFPEEEEGGCELRVKFSRSADAPAYKQGQNQLY NELNLGRREEYDVLDKRRGRDPEMGGKPRRKNPQEGLYNELQKDKMAEAYSEIG MKGERRRGKGHDGLYQGLSTATKDTYDALHMQALPPR	anti-BCMA CAR
128	QVQLVESGGGLVQPGGSLRLSCAVSGFALSNHGMSWVRRAPGKGLEWVSGIVYS GSTYYAASVKGRFTISRDNSRNTLYLQMNSLRPEDTAIYYCSAHGGESDVWGQG TTVTVSSASGGGGSGGRASGGGGSDIRLTQSPSPLSASVGDRVTITCQASEDIN	anti-BCMA CAR

	KFLNWYHQTPGKAPKLLIYDASTLQTGVPSRFSGSGSGTDFTLTINSLQPEDIG TYYCQQYESLPLTFGGGTKVEIKTTTPAPRPPTPAPTIASQPLSLRPEACRPAA GGAVHTRGLDFACDIYIWAPLAGTCGVLLLSLVITLYCKRGRKKLLYIFKQPFM RPVQTTQEEDGCSCRFPEEEEGGCELRVKFSRSADAPAYKQGQNQLYNELNLGR REEYDVLDKRRGRDPEMGGKPRRKNPQEGLYNELQKDKMAEAYSEIGMKGERRR GKGHDGLYQGLSTATKDTYDALHMQALPPR	
129	EVQLVESGGGLVQPGGSLRLSCAVSGFALSNHGMSWVRRAPGKGLEWVSGIVYS GSTYYAASVKGRFTISRDNSRNTLYLQMNSLRPEDTAIYYCSAHGGESDVWGQG TTVTVSSASGGGGSGGRASGGGGSEIVLTQSPGTLSLSPGERATLSCRASQSIG SSSLAWYQQKPGQAPRLLMYGASSRASGIPDRFSGSGSGTDFTLTISRLEPEDF AVYYCQQYAGSPPFTFGQGTKVEIKTTTPAPRPPTPAPTIASQPLSLRPEACRP AAGGAVHTRGLDFACDIYIWAPLAGTCGVLLLSLVITLYCKRGRKKLLYIFKQP FMRPVQTTQEEDGCSCRFPEEEEGGCELRVKFSRSADAPAYKQGQNQLYNELNL GRREEYDVLDKRRGRDPEMGGKPRRKNPQEGLYNELQKDKMAEAYSEIGMKGER RRGKGHDGLYQGLSTATKDTYDALHMQALPPR	anti-BCMA CAR
130	QIQLVQSGPDLKKPGETVKLSCKASGYTFTNFGMNWVKQAPGKGFKWMAWINTY TGESYFADDFKGRFAFSVETSATTAYLQINNLKTEDTATYFCARGEIYYGYDGG FAYWGQGTLVTVSAGGGGSGGGGSGGGGSDVVMTQSHRFMSTSVGDRVSITCRA SQDVNTAVSWYQQKPGQSPKLLIFSASYRYTGVPDRFTGSGSGADFTLTISSVQ AEDLAVYYCQQHYSTPWTFGGGTKLDIKTTTPAPRPPTPAPTIASQPLSLRPEA CRPAAGGAVHTRGLDFACDIYIWAPLAGTCGVLLLSLVITLYCKRGRKKLLYIF KQPFMRPVQTTQEEDGCSCRFPEEEEGGCELRVKFSRSADAPAYKQGQNQLYNE LNLGRREEYDVLDKRRGRDPEMGGKPRRKNPQEGLYNELQKDKMAEAYSEIGMK GERRRGKGHDGLYQGLSTATKDTYDALHMQALPPR	anti-BCMA CAR
131	QIQLVQSGPELKKPGETVKISCKASGYTFTDYSINWVKRAPGKGLKWMGWINTE TREPAYAYDFRGRFAFSLETSASTAYLQINNLKYEDTATYFCALDYSYAMDYWG QGTSVTVSSGGGGSGGGGSGGGGSQIQLVQSGPELKKPGETVKISCKASGYTFT DYSINWVKRAPGKGLKWMGWINTETREPAYAYDFRGRFAFSLETSASTAYLQIN NLKYEDTATYFCALDYSYAMDYWGQGTSVTVSSTTTPAPRPPTPAPTIASQPLS LRPEACRPAAGGAVHTRGLDFACDIYIWAPLAGTCGVLLLSLVITLYCKRGRKK LLYIFKQPFMRPVQTTQEEDGCSCRFPEEEEGGCELRVKFSRSADAPAYKQGQN QLYNELNLGRREEYDVLDKRRGRDPEMGGKPRRKNPQEGLYNELQKDKMAEAYS EIGMKGERRRGKGHDGLYQGLSTATKDTYDALHMQALPPR	anti-BCMA CAR
132	QIQLVQSGPELKKPGETVKISCKASGYTFTDYSINWVKRAPGKGLKWMGWINTE TREPAYAYDFRGRFAFSLETSASTAYLQINNLKYEDTATYFCALDYSYAMDYWG QGTSVTVSSGGGGSGGGGSGGGGSDIVLTQSPASLAMSLGKRATISCRASESVS VIGAHLIHWYQQKPGQPPKLLIYLASNLETGVPARFSGSGSGTDFTLTIDPVEE DDVAIYSCLQSRIFPRTFGGGTKLEIKTTTPAPRPPTPAPTIASQPLSLRPEAC RPAAGGAVHTRGLDFACDIYIWAPLAGTCGVLLLSLVITLYCKRGRKKLLYIFK QPFMRPVQTTQEEDGCSCRFPEEEEGGCELRVKFSRSADAPAYKQGQNQLYNEL NLGRREEYDVLDKRRGRDPEMGGKPRRKNPQEGLYNELQKDKMAEAYSEIGMKG ERRRGKGHDGLYQGLSTATKDTYDALHMQALPPR	anti-BCMA CAR
133	QIQLVQSGPELKKPGETVKISCKASGYTFRHYSMNWVKQAPGKGLKWMGRINTE SGVPIYADDFKGRFAFSVETSASTAYLVINNLKDEDTASYFCSNDYLYSLDFWG QGTALTVSSGGGGSGGGGSGGGGSDIVLTQSPPSLAMSLGKRATISCRASESVT ILGSHLIYWYQQKPGQPPTLLIQLASNVQTGVPARFSGSGSRTDFTLTIDPVEE DDVAVYYCLQSRTIPRTFGGGTKLEIKTTTPAPRPPTPAPTIASQPLSLRPEAC RPAAGGAVHTRGLDFACDIYIWAPLAGTCGVLLLSLVITLYCKRGRKKLLYIFK QPFMRPVQTTQEEDGCSCRFPEEEEGGCELRVKFSRSADAPAYKQGQNQLYNEL NLGRREEYDVLDKRRGRDPEMGGKPRRKNPQEGLYNELQKDKMAEAYSEIGMKG ERRRGKGHDGLYQGLSTATKDTYDALHMQALPPR	anti-BCMA CAR
134	QIQLVQSGPELKKPGETVKISCKASGYTFTHYSMNWVKQAPGKGLKWMGRINTE TGEPLYADDFKGRFAFSLETSASTAYLVINNLKNEDTATFFCSNDYLYSCDYWG QGTTLTVSSGGGGSGGGGSGGGGSDIVLTQSPPSLAMSLGKRATISCRASESVT ILGSHLIYWYQQKPGQPPTLLIQLASNVQTGVPARFSGSGSRTDFTLTIDPVEE DDVAVYYCLQSRTIPRTFGGGTKLEIKTTTPAPRPPTPAPTIASQPLSLRPEAC RPAAGGAVHTRGLDFACDIYIWAPLAGTCGVLLLSLVITLYCKRGRKKLLYIFK	anti-BCMA CAR

	QPFMRPVQTTQEEDGCSCRFPEEEEGGCELRVKFSRSADAPAYKQGQNQLYNEL NLGRREEYDVLDKRRGRDPEMGGKPRRKNPQEGLYNELQKDKMAEAYSEIGMKG ERRRGKGHDGLYQGLSTATKDTYDALHMQALPPR	
135	DIVLTQSPPSLAMSLGKRATISCRASESVTILGSHLIHWYQQKPGQPPTLLIQL ASNVQTGVPARFSGSGSRTDFTLTIDPVEEDDVAVYYCLQSRTIPRTFGGGTKL EIKGSTSGSGKPGSGEGSTKGQIQLVQSGPELKKPGETVKISCKASGYTFTDYS INWVKRAPGKGLKWMGWINTETREPAYAYDFRGRFAFSLETSASTAYLQINNLK YEDTATYFCALDYSYAMDYWGQGTSVTVSSFVPVFLPAKPTTTPAPRPPTPAPT IASQPLSLRPEACRPAAGGAVHTRGLDFACDIYIWAPLAGTCGVLLLSLVITLY CNHRNRSKRSRLLHSDYMNMTPRRPGPTRKHYQPYAPPRDFAAYRSRVKFSRSA DAPAYQQGQNQLYNELNLGRREEYDVLDKRRGRDPEMGGKPRRKNPQEGLYNEL QKDKMAEAYSEIGMKGERRRGKGHDGLYQGLSTATKDTYDALHMQALPPR	anti-BCMA CAR
136	QVQLVQSGAEVKKPGASVKVSCKASGYSFPDYYINWVRQAPGQGLEWMGWIYFA SGNSEYNQKFTGRVTMTRDTSINTAYMELSSLTSEDTAVYFCASLYDYDWYFDV WGQGTMVTVSSGGGGSGGGGSGGGGSDIVMTQTPLSLSVTPGQPASISCKSSQS LVHSNGNTYLHWYLQKPGQSPQLLIYKVSNRFSGVPDRFSGSGSGTDFTLKISR VEAEDVGIYYCSQSSIYPWTFGQGTKLEIKGLAVSTISSFFPPGYQIYIWAPLA GTCGVLLLSLVITLYCKRGRKKLLYIFKQPFMRPVQTTQEEDGCSCRFPEEEEG GCELRVKFSRSADAPAYQQGQNQLYNELNLGRREEYDVLDKRRGRDPEMGGKPR RKNPQEGLYNELQKDKMAEAYSEIGMKGERRRGKGHDGLYQGLSTATKDTYDAL HMQALPPR	anti-BCMA CAR
137	QVQLVQSGAEVKKPGASVKVSCKASGYSFPDYYINWVRQAPGQGLEWMGWIYFA SGNSEYNQKFTGRVTMTRDTSINTAYMELSSLTSEDTAVYFCASLYDYDWYFDV WGQGTMVTVSSGGGGSGGGGSGGGGSDIVMTQTPLSLSVTPGQPASISCKSSQS LVHSNGNTYLHWYLQKPGQSPQLLIYKVSNRFSGVPDRFSGSGSGTDFTLKISR VEAEDVGIYYCSQSSIYPWTFGQGTKLEIKTTTPAPRPPTPAPTIASQPLSLRP EACRPAAGGAVHTRGLDFACDIYIWAPLAGTCGVLLLSLVITLYCKRGRKKLLY IFKQPFMRPVQTTQEEDGCSCRFPEEEEGGCELRVKFSRSADAPAYQQGQNQLY NELNLGRREEYDVLDKRRGRDPEMGGKPRRKNPQEGLYNELQKDKMAEAYSEIG MKGERRRGKGHDGLYQGLSTATKDTYDALHMQALPPR	anti-BCMA CAR
138	QVQLVQSGAEVKKPGASVKVSCKASGYSFPDYYINWVRQAPGQGLEWMGWIYFA SGNSEYNQKFTGRVTMTRDTSINTAYMELSSLTSEDTAVYFCASLYDYDWYFDV WGQGTMVTVSSGGGGSGGGGSGGGGSDIVMTQTPLSLSVTPGQPASISCKSSQS LVHSNGNTYLHWYLQKPGQSPQLLIYKVSNRFSGVPDRFSGSGSGTDFTLKISR VEAEDVGIYYCSQSSIYPWTFGQGTKLEIKEPKSPDKTHTCPPCPAPPVAGPSV FLFPPKPKDTLMIARTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREE QYNSTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQV YTLPPSRDELTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDG SFFLYSKLTVDKSRWQQGNVFSCSVMHEALHNHYTQKSLSLSPGKIYIWAPLAG TCGVLLLSLVITLYCKRGRKKLLYIFKQPFMRPVQTTQEEDGCSCRFPEEEEGG CELRVKFSRSADAPAYQQGQNQLYNELNLGRREEYDVLDKRRGRDPEMGGKPRR KNPQEGLYNELQKDKMAEAYSEIGMKGERRRGKGHDGLYQGLSTATKDTYDALH MQALPPR	anti-BCMA CAR
139	QVQLVQSGAEVKKPGASVKVSCKASGYSFPDYYINWVRQAPGQGLEWMGWIYFA SGNSEYNQKFTGRVTMTRDTSSSTAYMELSSLRSEDTAVYFCASLYDYDWYFDV WGQGTMVTVSSGGGGSGGGGSGGGGSDIVMTQTPLSLSVTPGEPASISCKSSQS LVHSNGNTYLHWYLQKPGQSPQLLIYKVSNRFSGVPDRFSGSGSGADFTLKISR VEAEDVGVYYCAETSHVPWTFGQGTKLEIKGLAVSTISSFFPPGYQIYIWAPLA GTCGVLLLSLVITLYCKRGRKKLLYIFKQPFMRPVQTTQEEDGCSCRFPEEEEG GCELRVKFSRSADAPAYQQGQNQLYNELNLGRREEYDVLDKRRGRDPEMGGKPR RKNPQEGLYNELQKDKMAEAYSEIGMKGERRRGKGHDGLYQGLSTATKDTYDAL HMQALPPR	anti-BCMA CAR
140	QVQLVQSGAEVKKPGASVKVSCKASGYSFPDYYINWVRQAPGQGLEWMGWIYFA SGNSEYNQKFTGRVTMTRDTSSSTAYMELSSLRSEDTAVYFCASLYDYDWYFDV WGQGTMVTVSSGGGGSGGGGSGGGGSDIVMTQTPLSLSVTPGEPASISCKSSQS LVHSNGNTYLHWYLQKPGQSPQLLIYKVSNRFSGVPDRFSGSGSGADFTLKISR VEAEDVGVYYCAETSHVPWTFGQGTKLEIKTTTPAPRPPTPAPTIASQPLSLRP	anti-BCMA CAR

	EACRPAAGGAVHTRGLDFACDIYIWAPLAGTCGVLLLSLVITLYCKRGRKKLLY IFKQPFMRPVQTTQEEDGCSCRFPEEEEGGCELRVKFSRSADAPAYQQGQNQLY NELNLGRREEYDVLDKRRGRDPEMGGKPRRKNPQEGLYNELQKDKMAEAYSEIG MKGERRRGKGHDGLYQGLSTATKDTYDALHMQALPPR	
141	QVQLVQSGAEVKKPGASVKVSCKASGYSFPDYYINWVRQAPGQGLEWMGWIYFA SGNSEYNQKFTGRVTMTRDTSSSTAYMELSSLRSEDTAVYFCASLYDYDWYFDV WGQGTMVTVSSGGGGSGGGGSGGGGSDIVMTQTPLSLSVTPGEPASISCKSSQS LVHSNGNTYLHWYLQKPGQSPQLLIYKVSNRFSGVPDRFSGSGSGADFTLKISR VEAEDVGVYYCAETSHVPWTFGQGTKLEIKEPKSPDKTHTCPPCPAPPVAGPSV FLFPPKPKDTLMIARTPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREE QYNSTYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQV YTLPPSRDELTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDG SFFLYSKLTVDKSRWQQGNVFSCSVMHEALHNHYTQKSLSLSPGKIYIWAPLAG TCGVLLLSLVITLYCKRGRKKLLYIFKQPFMRPVQTTQEEDGCSCRFPEEEEGG CELRVKFSRSADAPAYQQGQNQLYNELNLGRREEYDVLDKRRGRDPEMGGKPRR KNPQEGLYNELQKDKMAEAYSEIGMKGERRRGKGHDGLYQGLSTATKDTYDALH MQALPPR	anti-BCMA CAR
142	IYIWAPLAGTCGVLLLSLVITLYCNHRN	CD8a TM
143	IYIWAPLAGTCGVLLLSLVIT	CD8a TM
144	RAAA	linking peptide
145	EVQLVESGGGLVKPGGSLRLSCAASGFTFSDYYMSWIRQAPGKGLEWVSYISSS GSTIYYADSVKGRFTISRDNAKNSLYLQMNSLRAEDTAVYYCAKVDGDYTEDYW GQGTLVTVSS	Variable heavy (VH) Anti-BCMA
146	QSALTQPASVSGSPGQSITISCTGSSSDVGKYNLVSWYQQPPGKAPKLIIYDVN KRPSGVSNRFSGSKSGNTATLTISGLQGDDEADYYCSSYGGSRSYVFGTGTKVT VL	Variable light (VL) Anti-BCMA
147	EVQLVQSGGGLVQPGRSLRLSCTASGFTFGDYAMSWFRQAPGKGLEWVGFIRSK AYGGTTEYAASVKGRFTISRDDSKSIAYLQMNSLKTEDTAVYYCAAWSAPTDYW GQGTLVTVSS	Variable heavy (VH) Anti-BCMA
148	DIQMTQSPAFLSASVGDRVTVTCRASQGISNYLAWYQQKPGNAPRLLIYSASTL QSGVPSRFRGTGYGTEFSLTIDSLQPEDFATYYCQQSYTSRQTFGPGTRLDIK	Variable light (VL) Anti-BCMA
149	EVQLVESGGGLVKPGGSLRLSCAASGFTFSDYYMSWIRQAPGKGLEWVSYISSS GSTIYYADSVKGRFTISRDNAKNSLYLQMNSLRAEDTAVYYCAKVDGPPSFDIW GQGTMVTVSS	Variable heavy (VH) Anti-BCMA
150	SYVLTQPPSVSVAPGQTARITCGANNIGSKSVHWYQQKPGQAPMLVVYDDDDRP SGIPERFSGSNSGNTATLTISGVEAGDEADYFCHLWDRSRDHYVFGTGTKLTVL	Variable light (VL) Anti-BCMA
151	EVQLVQSGAEVKKPGSSVKVSCKASGGTFSSYAISWVRQAPGQGLEWMGRIIPI LGIANYAQKFQGRVTMTEDTSTDTAYMELSSLRSEDTAVYYCARSGYSKSIVSY MDYWGQGTLVTVSS	Variable heavy (VH) Anti-BCMA
152	LPVLTQPPSTSGTPGQRVTVSCSGSSSNIGSNVVFWYQQLPGTAPKLVIYRNNQ RPSGVPDRFSVSKSGTSASLAISGLRSEDEADYYCAAWDDSLSGYVFGTGTKVT VLG	Variable light (VL) Anti-BCMA
153	MPLLLLLPLLWAGALA	CD33 Signal peptide
154	MALPVTALLLPLALLLHA	CD8 alpha signal peptide
155	atgcttctcctggtgacaagccttctgctctgtgagttaccacacccagcattc ctcctgatccca	GMCSFR alpha chain signal sequence
156	MLLLVTSLLLCELPHPAFLLIP	GMCSFR alpha chain signal sequence

157	Glu Val Val Val Lys Tyr Gly Pro Pro Cys Pro Pro Cys Pro	Exemplary IgG Hinge
158	X1PPX2P X1 is glycine, cysteine or arginine X2 is cysteine or threonine	Exemplary IgG Hinge
159	Glu Pro Lys Ser Cys Asp Lys Thr His Thr Cys Pro Pro Cys Pro	Exemplary IgG Hinge
160	Glu Arg Lys Cys Cys Val Glu Cys Pro Pro Cys Pro	Exemplary IgG Hinge
161	ELKTPLGDTHTCPRCPEPKSCDTPPPCPRCPEPKSCDTPPPCPRCPEPK SCDTPPPCPRCP	Exemplary IgG Hinge
162	Glu Ser Lys Tyr Gly Pro Pro Cys Pro Ser Cys Pro	Exemplary IgG Hinge
163	Glu Ser Lys Tyr Gly Pro Pro Cys Pro Pro Cys Pro	Exemplary IgG Hinge
164	Tyr Gly Pro Pro Cys Pro Pro Cys Pro	Exemplary IgG Hinge
165	Lys Tyr Gly Pro Pro Cys Pro Pro Cys Pro	Exemplary IgG Hinge
166	MLLLVTSLLLCELPHPAFLLIPRKVCNGIGIGEFKDSLSINATNIKHFK NCTSISGDLHILPVAFRGDSFTHTPPLDPQELDILKTVKEITGFLLIQA WPENRTDLHAFENLEIIRGRTKQHGQFSLAVVSLNITSLGLRSLKEISD GDVIISGNKNLCYANTINWKKLFGTSGQKTKIISNRGENSCKATGQVCH ALCSPEGCWGPEPRDCVSCRNVSRGRECVDKCNLLEGEPREFVENSECI QCHPECLPQAMNITCTGRGPDNCIQCAHYIDGPHCVKTCPAGVMGENNT LVWKYADAGHVCHLCHPNCTYGCTGPGLEGCPTNGPKIPSIATGMVGAL LLLLVVALGIGLFM	tEGFR artificial
167	EGRGSLLTCGDVEENPGP	T2A artificial
168	GSGATNFSLLKQAGDVEENPGP	P2A
169	ATNFSLLKQAGDVEENPGP	P2A
170	QCTNYALLKLAGDVESNPGP	E2A
171	VKQTLNFDLLKLAGDVESNPGP	F2A
172	MLQMAGQCSQNEYFDSLLHACIPCQLRCSSNTPPLTCQRYCNASVTNSV KGTNAGGGGSPKSSDKTHTCPPCPAPEAEGAPSVFLFPPKPKDTLMISR TPEVTCVVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYNSTYRVVS VLTVLHQDWLNGKEYKCKVSNKALPSSIEKTISKAKGQPREPQVYTLPP SRDELTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTPPVLDSDG SFFLYSKLTVDKSRWQQGNVFSCSVMHEALHNHYTQKSLSLSPGK	BCMA-Fc fusion polypeptide
173	DYYVY	CDR-H1
174	WINPNSGGTNYAQKFQG	CDR-H2
175	SQRDGYMDY	CDR-H3
176	GYTFIDY	CDR-H1
177	NPNSGG	CDR-H2
178	GYTFIDYYVY	CDR-H1

179	WINPNSGGTN	CDR-H2
180	GYTFIDYY	CDR-H1
181	INPNSGGT	CDR-H2
182	ARSQRDGYMDY	CDR-H3
183	TGTSSDVG	CDR-L1
184	EDSKRPS	CDR-L2
185	SSNTRSSTLV	CDR-L3
186	ISCTGTSSD	CDR-L1
187	EDS	CDR-L2
188	QSALTQPASVSASPGQSIAISCTGTSSDVGWYQQHPGKAPKLMIYEDSK RPSGVSNRFSGSKSGNTASLTISGLQAEDEADYYCSSNTRSSTLVFGGG TKLTVLGSRGGGGSGGGGSGGGGSLEMAEVQLVQSGAEMKKPGASLKLS CKASGYTFIDYYVYWMRQAPGQGLESMGWINPNSGGTNYAQKFQGRVTM TRDTSISTAYMELSRLRSDDTAMYYCARSQRDGYMDYWGQGTLVTVSS	anti-BCMA scFv

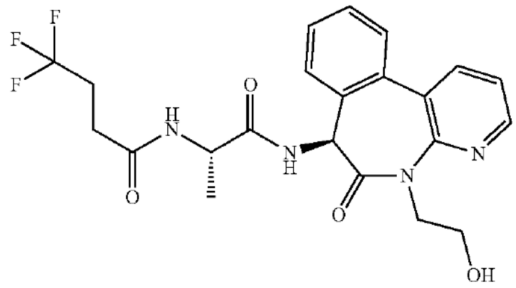
CLAIMS

~~1. Use of combination comprising a cell therapy and a compound, in the manufacture of a medicament for the treatment of a B cell maturation antigen (BCMA) expressing cancer in a subject, wherein~~

~~(a) the cell therapy is to be administered to the subject having the BCMA-expressing cancer said cell therapy comprising a dose of T cells expressing a recombinant receptor, wherein the recombinant receptor specifically binds to BCMA; and~~

~~(b) the compound is to be administered to the subject, wherein the compound is of the structure:~~

Compound 1

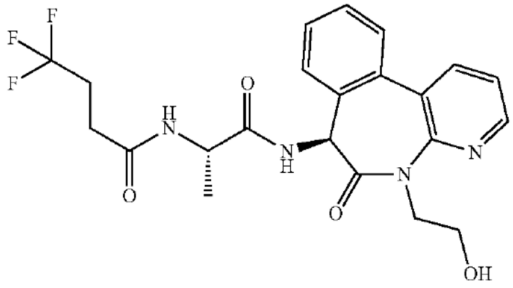


~~or a stereoisomer thereof, or a pharmaceutically acceptable salt or hydrate of either of the foregoing.~~

~~2. The use of claim 1, wherein the compound is to be administered subsequently to initiation of administration of the cell therapy.~~

~~3. Use of a compound in the manufacture of a medicament for the treatment of a B cell maturation antigen (BCMA) expressing cancer in a subject, wherein the compound is to be administered to the subject having the BCMA-expressing cancer, said compound having the structure:~~

Compound 1

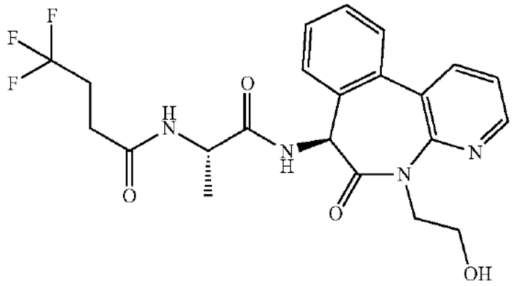


~~or a stereoisomer thereof, or a pharmaceutically acceptable salt or hydrate of either of the foregoing,~~

~~wherein, at the time of initiation of the administration of the compound, the subject has been previously administered, and/or is undergoing treatment with, a cell therapy, said cell therapy comprising a dose of T cells expressing a recombinant receptor, wherein the recombinant receptor specifically binds to BCMA.~~

~~4. Use of a cell therapy for preparation of a medicament for treating a B cell maturation antigen (BCMA)-expressing cancer in a subject, wherein the cell therapy is to be administered to a subject having the BCMA-expressing cancer, said cell therapy comprising a dose of T cells expressing a recombinant receptor, wherein the recombinant receptor specifically binds to BCMA, and wherein, at the time of initiation of the administration of the cell therapy, the subject has been previously administered, and/or is undergoing treatment with, a compound having the structure:~~

Compound 1



~~or a stereoisomer thereof, or a pharmaceutically acceptable salt or hydrate of either of the foregoing.~~

5. The use of claim 1, claim 3 or claim 4, wherein the initiation of administration of the compound is prior to, concurrently with or subsequently to initiation of administration of the cell therapy.

6. The use of claim 5, wherein the compound is to be administered prior to initiation of administration of the cell therapy.

7. The use of any of claims 1-6, wherein the compound is capable of inhibiting cleavage of the target antigen and/or wherein the target antigen is cleaved by a gamma secretase.

8. The use of any of claims 1 to 7, wherein the target antigen is BCMA.

9. The use of any one of claims 1-8, wherein the recombinant receptor is a chimeric antigen receptor.

10. The use of any one of claims 1-9, wherein the compound is capable of inhibiting, or inhibiting an activity or function of, a gamma secretase.

11. The use of any of claims 1-10, wherein the compound, stereoisomer, pharmaceutically acceptable salt or hydrate is to be administered orally.

12. The use of any one of claims 1-11, wherein:

the compound, stereoisomer, pharmaceutically acceptable salt or hydrate is to be administered at least or is to be administered six times daily, five times daily, four times daily, three times daily, twice daily, once daily, every other day, three times a week, at least once a week, twice a week or only one time.

13. The use of any of claims 1-12, wherein the compound, stereoisomer, pharmaceutically acceptable salt or hydrate is to be administered once every other day.

~~14. The use of any of claims 1-12, wherein the compound, stereoisomer, pharmaceutically acceptable salt or hydrate is to be administered no more than once per day.~~

~~15. The use of any of claims 1-12, wherein the compound, stereoisomer, pharmaceutically acceptable salt or hydrate is to be administered once three times per week.~~

~~16. The use of any one of claims 12-15, wherein the compound, stereoisomer, pharmaceutically acceptable salt or hydrate is to be administered on days 2, 4, 7, 9, 11, 14, 16 and 18 after initiation of the administration of the cell therapy.~~

~~17. The use of any of claims 1-16, wherein the compound, stereoisomer, pharmaceutically acceptable salt or hydrate is to be administered at an amount of 1.0 mg to 100 mg.~~

~~18. The use of any of claims 1-17, wherein the compound, stereoisomer, pharmaceutically acceptable salt or hydrate to be administered is or is about 25 mg.~~

~~19. The use of any of claims 1-18, wherein the compound, stereoisomer, pharmaceutically acceptable salt or hydrate is to be administered at an amount of at or about 50 mg.~~

~~20. The use of any of claims 1-19, wherein the cancer is a B cell malignancy.~~

~~21. The use of any of claims 1-20, wherein the cancer is multiple myeloma, plasmacytoma, or a cancer of plasma cell origin.~~

~~22. The use of any of claims 1-21, wherein the cancer is a multiple myeloma.~~

~~23. The use of any of claims 1-22, wherein the cancer is relapsed or treatment refractory multiple myeloma.~~

24. The use of any of claims 1-23, wherein prior to said administration of said cell therapy, a subject is to be selected for administration that has a cancer that, or the cells of which, (i) express CD138, surface CD38 or a surface plasma cell marker or are derived from plasma cells.

25. The use of claim 24, wherein the subject further comprises low expression of surface B-cell maturation (BCMA) and/or a level of expression of surface BCMA below the threshold level.

26. The use of any of claims 1-25, wherein the recombinant receptor is a chimeric antigen receptor (CAR) comprising an antigen-binding domain that binds BCMA, wherein the antigen-binding domain comprises:

(a) a VH set forth in SEQ ID NO: 18 and a VL set forth in SEQ ID NO: 19;
(b) a VH set forth in SEQ ID NO: 20 and a VL set forth in SEQ ID NO: 21;
(c) a VH set forth in SEQ ID NO: 22 and a VL set forth in SEQ ID NO: 23;
(d) a VH set forth in SEQ ID NO: 24 and a VL set forth in SEQ ID NO: 25;
(e) a VH set forth in SEQ ID NO: 32 and a VL set forth in SEQ ID NO: 33;
(f) a VH set forth in SEQ ID NO: 34 and a VL set forth in SEQ ID NO: 35;
(g) a VH set forth in SEQ ID NO: 36 and a VL set forth in SEQ ID NO: 37;
(h) a VH set forth in SEQ ID NO: 41 and a VL set forth in SEQ ID NO: 42;
(i) a VH set forth in SEQ ID NO: 43 and a VL set forth in SEQ ID NO: 44;
(j) a VH set forth in SEQ ID NO: 45 and a VL set forth in SEQ ID NO: 46;
(k) a VH set forth in SEQ ID NO: 47 and a VL set forth in SEQ ID NO: 48;
(l) a VH set forth in SEQ ID NO: 49 and a VL set forth in SEQ ID NO: 50;
(m) a VH set forth in SEQ ID NO: 51 and a VL set forth in SEQ ID NO: 52;
(n) a VH set forth in SEQ ID NO: 53 and a VL set forth in SEQ ID NO: 54;
(o) a VH set forth in SEQ ID NO: 55 and a VL set forth in SEQ ID NO: 56;
(p) a VH set forth in SEQ ID NO: 57 and a VL set forth in SEQ ID NO: 58;
(q) a VH set forth in SEQ ID NO: 59 and a VL set forth in SEQ ID NO: 60;
(r) a VH set forth in SEQ ID NO: 61 and a VL set forth in SEQ ID NO: 62;
(s) a VH set forth in SEQ ID NO: 63 and a VL set forth in SEQ ID NO: 64;

(t) a VH set forth in SEQ ID NO: 65 and a VL set forth in SEQ ID NO: 66;

(u) a VH set forth in SEQ ID NO: 67 and a VL set forth in SEQ ID NO: 68;

(v) a VH set forth in SEQ ID NO: 69 and a VL set forth in SEQ ID NO: 70;

(w) a VH set forth in SEQ ID NO: 71 and a VL set forth in SEQ ID NO: 72;

(x) a VH set forth in SEQ ID NO: 73 and a VL set forth in SEQ ID NO: 74;

(y) a VH set forth in SEQ ID NO: 75 and a VL set forth in SEQ ID NO: 76;

(z) a VH set forth in SEQ ID NO: 145 and a VL set forth in SEQ ID NO: 146;

(aa) a VH set forth in SEQ ID NO: 147 and a VL set forth in SEQ ID NO: 148;

(bb) a VH set forth in SEQ ID NO: 149 and a VL set forth in SEQ ID NO: 150; or

(cc) a VH set forth in SEQ ID NO: 151 and a VL set forth in SEQ ID NO: 152.

27. The use of any of claims 1-26, wherein the recombinant receptor is a chimeric antigen receptor comprising an antigen-binding domain that binds BCMA, wherein the antigen-binding domain comprises:

a V_H region comprising a CDRH1, a CDRH2 and a CDRH3 comprising the amino acid sequence of SEQ ID NOS: 173, 174 and 175, respectively and a V_L region comprising a CDRL1, a CDRL2 and a CDRL3 comprising the amino acid sequence of SEQ ID NOS: 183, 184 and 185, respectively;

a V_H region comprising a CDRH1, a CDRH2 and a CDRH3 comprising the amino acid sequence of SEQ ID NOS: 176, 177 and 175, respectively and a V_L region comprising a CDRL1, a CDRL2 and a CDRL3 comprising the amino acid sequence of SEQ ID NOS: 183, 184 and 185, respectively;

a V_H region comprising a CDRH1, a CDRH2 and a CDRH3 comprising the amino acid sequence of SEQ ID NOS: 178, 179 and 175, respectively and a V_L region comprising a CDRL1, a CDRL2 and a CDRL3 comprising the amino acid sequence of SEQ ID NOS: 183, 184 and 185, respectively; or

a V_H region comprising a CDRH1, a CDRH2 and a CDRH3 comprising the amino acid sequence of SEQ ID NOS: 180, 181 and 182, respectively and a V_L region comprising a CDRL1, a CDRL2 and a CDRL3 comprising the amino acid sequence of SEQ ID NOS: 186, 187 and 185, respectively.

28. The use of claim 27, wherein antigen-binding domain comprises a V_H region comprising an amino acid sequence having at least at or about 90% sequence identity to SEQ ID NO: 24; and a V_L region comprising an amino acid sequence having at least at or about 90% sequence identity to SEQ ID NO: 25.

29. The use of claim 27 or claim 28 wherein the antigen-binding domain comprises a V_H region comprising the amino acid sequence set forth in SEQ ID NO: 24; and a V_L region comprising the amino acid sequence set forth in SEQ ID NO: 25.

30. The use of any of claims 27-29, wherein the antigen-binding domain is an scFv comprising an amino acid sequence having at least at or about 90% sequence identity to SEQ ID NO:188.

31. The use of claim 30, wherein the antigen-binding domain is an scFv comprising an amino acid sequence set forth in SEQ ID NO:188.

32. The use of any of claims 1-26, wherein the recombinant receptor is a chimeric antigen receptor (CAR) comprising an antigen-binding domain that binds BCMA, wherein the antigen-binding domain comprises the VH region set forth in SEQ ID NO:18 and the VL region set forth in SEQ ID NO:19.

33. The use of any of claims 27-32, wherein the intracellular signaling domain comprises an intracellular domain of a CD3-zeta (CD3ζ) chain.

34. The use of any of claims 27-33, wherein the chimeric antigen receptor (CAR) further comprises a costimulatory signaling region.

35. The use of claim 34, wherein the costimulatory signaling region comprises a signaling domain derived from CD28 or 4-1BB, optionally human CD28 or human 4-1BB.

36. The use of any of claims 1-35, wherein the CAR comprises a sequence as set forth in any one of SEQ ID NOS: 90-141 or a sequence of amino acids that exhibits at least 90%, at least 91%, at least 92%, at least 93%, at least 94%, at least 95%, at least 96%, at least 97%, at least 98%, or at least 99% sequence identity to any one of SEQ ID NOS: 90-141.

37. The use of any of claims 1-36, wherein the dose of T cells to be administered comprises between at or about 2.5×10^7 CAR-expressing T cells and 1.2×10^9 CAR-expressing T cells, inclusive.

38. The use of any of claims 1-37, wherein a lymphodepleting chemotherapy is to be administered prior to administration of the cell therapy and/or wherein the subject has received a lymphodepleting chemotherapy prior to administration of the cells.

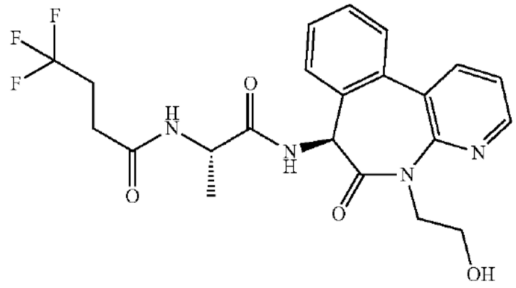
39. The use of claim 38, wherein the lymphodepleting chemotherapy that is to be administered comprises fludarabine and/or cyclophosphamide to the subject.

40. A combination, comprising:

(a) T cells expressing a recombinant receptor that specifically binds to B cell maturation antigen (BCMA); and

(b) a compound having the structure:

Compound 1



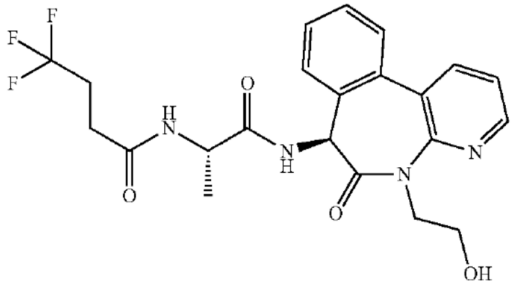
or a stereoisomer thereof, or a pharmaceutically acceptable salt or hydrate of any of the foregoing.

~~41. The combination of claim 40, wherein the compound inhibits or reduces or is capable of inhibiting or reducing intramembrane cleavage of B cell maturation antigen (BCMA).~~

~~42. The combination of claim 40 or claim 41, wherein the recombinant receptor is a chimeric antigen receptor (CAR).~~

~~43. The combination of any of claims 40-42, comprising the compound of the structure:~~

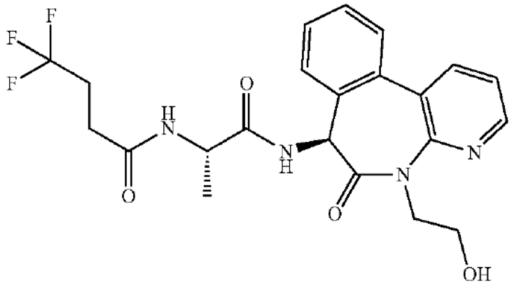
Compound 1



1. Use of a cell therapy and a compound in the manufacture of a medicament for combination therapy of a B cell maturation antigen (BCMA)-expressing cancer in a subject, wherein

(a) the cell therapy comprises a dose of T cells expressing a recombinant receptor, wherein the recombinant receptor specifically binds to BCMA; and

Compound 1



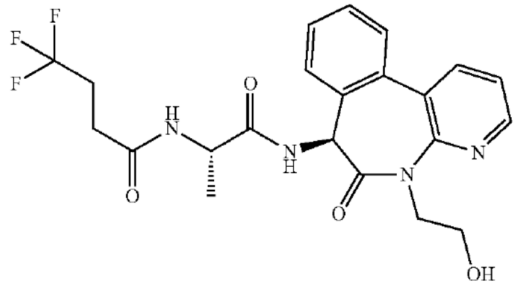
(b) the compound is _____

or a stereoisomer thereof, or a pharmaceutically acceptable salt or hydrate of either of the foregoing.

2. The use of claim 1, wherein the compound is to be administered subsequently to initiation of administration of the cell therapy.

3. Use of a compound in the manufacture of a medicament for the treatment of a B cell maturation antigen (BCMA)-expressing cancer in a subject, wherein the compound is to be administered to the subject having the BCMA-expressing cancer, said compound is:

Compound 1

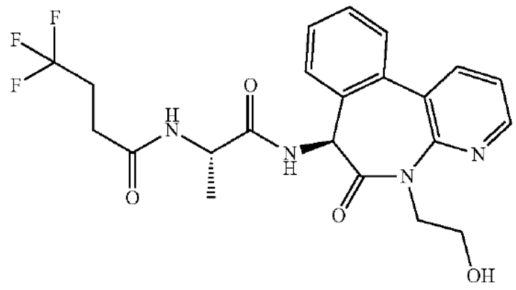


or a stereoisomer thereof, or a pharmaceutically acceptable salt or hydrate of either of the foregoing,

wherein, at the time of initiation of the administration of the compound, the subject has been previously administered, and/or is undergoing treatment with, a cell therapy, said cell therapy comprising a dose of T cells expressing a recombinant receptor, wherein the recombinant receptor specifically binds to BCMA.

4. Use of a cell therapy for preparation of a medicament for treating a B cell maturation antigen (BCMA)-expressing cancer in a subject, wherein the cell therapy is to be administered to a subject having the BCMA-expressing cancer, said cell therapy comprising a dose of T cells expressing a recombinant receptor, wherein the recombinant receptor specifically binds to BCMA, and wherein, at the time of initiation of the administration of the cell therapy, the subject has been previously administered, and/or is undergoing treatment with, a compound having the structure:

Compound 1



or a stereoisomer thereof, or a pharmaceutically acceptable salt or hydrate of either of the foregoing.

5. The use of claim 1 wherein the initiation of administration of the compound is prior to, concurrently with or subsequently to initiation of administration of the cell therapy.

6. The use of claim 5, wherein the compound is to be administered prior to initiation of administration of the cell therapy.

7. The use of any of claims 1-6, wherein the compound is capable of inhibiting cleavage of BCMA and/or wherein the BCMA is cleaved by a gamma secretase.

8. The use of any one of claims 1-7, wherein the recombinant receptor is a chimeric antigen receptor.

9. The use of any one of claims 1-8, wherein the compound is capable of inhibiting, or inhibiting an activity or function of, a gamma secretase.

10. The use of any of claims 1-9, wherein the compound, stereoisomer, pharmaceutically acceptable salt or hydrate is to be administered orally.

11. The use of any one of claims 1-10, wherein:

the compound, stereoisomer, pharmaceutically acceptable salt or hydrate is to be administered at least or is to be administered six times daily, five times daily, four times daily,

three times daily, twice daily, once daily, every other day, three times a week, at least once a week, twice a week or only one time.

12. The use of any of claims 1-11, wherein the compound, stereoisomer, pharmaceutically acceptable salt or hydrate is to be administered once every other day.

13. The use of any of claims 1-11, wherein the compound, stereoisomer, pharmaceutically acceptable salt or hydrate is to be administered no more than once per day.

14. The use of any of claims 1-11, wherein the compound, stereoisomer, pharmaceutically acceptable salt or hydrate is to be administered once three times per week.

15. The use of any one of claims 10-14, wherein the compound, stereoisomer, pharmaceutically acceptable salt or hydrate is to be administered on days 2, 4, 7, 9, 11, 14, 16 and 18 after initiation of the administration of the cell therapy.

16. The use of any of claims 1-15, wherein the compound, stereoisomer, pharmaceutically acceptable salt or hydrate is to be administered at an amount of 1.0 mg to 100 mg.

17. The use of any of claims 1-16, wherein the compound, stereoisomer, pharmaceutically acceptable salt or hydrate to be administered is or is about 25 mg.

18. The use of any of claims 1-17, wherein the compound, stereoisomer, pharmaceutically acceptable salt or hydrate is to be administered at an amount of at or about 50 mg.

19. The use of any of claims 1-18, wherein the cancer is a B cell malignancy.

20. The use of any of claims 1-19, wherein the cancer is multiple myeloma, plasmacytoma, or a cancer of plasma cell origin.

21. The use of any of claims 1-20, wherein the cancer is a multiple myeloma.

22. The use of any of claims 1-21, wherein the cancer is relapsed or treatment refractory multiple myeloma.

23. The use of any of claims 1-22, wherein the subject has been identified, prior to said administration of said cell therapy, to have a cancer in which cells of the cancer express CD138, surface CD38 or a surface plasma cell marker or are derived from plasma cells.

24. The use of any of claims 1-23, wherein the subject has been identified, prior to said administration of said cell therapy, to have B cell maturation antigen (BCMA)-expressing cancer cells comprising low expression of surface B cell maturation (BCMA) and/or a level of expression of surface BCMA below a threshold level.

25. The use of any of claims 1-24, wherein the recombinant receptor is a chimeric antigen receptor (CAR) comprising an antigen-binding domain that binds BCMA, wherein the antigen-binding domain comprises:

(a) a VH set forth in SEQ ID NO: 18 and a VL set forth in SEQ ID NO: 19;

(b) a VH set forth in SEQ ID NO: 20 and a VL set forth in SEQ ID NO: 21;

(c) a VH set forth in SEQ ID NO: 22 and a VL set forth in SEQ ID NO: 23;

(d) a VH set forth in SEQ ID NO: 24 and a VL set forth in SEQ ID NO: 25;

(e) a VH set forth in SEQ ID NO: 32 and a VL set forth in SEQ ID NO: 33;

(f) a VH set forth in SEQ ID NO: 34 and a VL set forth in SEQ ID NO: 35;

(g) a VH set forth in SEQ ID NO: 36 and a VL set forth in SEQ ID NO: 37;

(h) a VH set forth in SEQ ID NO: 41 and a VL set forth in SEQ ID NO: 42;

(i) a VH set forth in SEQ ID NO: 43 and a VL set forth in SEQ ID NO: 44;

(j) a VH set forth in SEQ ID NO: 45 and a VL set forth in SEQ ID NO: 46;

(k) a VH set forth in SEQ ID NO: 47 and a VL set forth in SEQ ID NO: 48;

(l) a VH set forth in SEQ ID NO: 49 and a VL set forth in SEQ ID NO: 50;

(m) a VH set forth in SEQ ID NO: 51 and a VL set forth in SEQ ID NO: 52;

(n) a VH set forth in SEQ ID NO: 53 and a VL set forth in SEQ ID NO: 54;

(o) a VH set forth in SEQ ID NO: 55 and a VL set forth in SEQ ID NO: 56;

(p) a VH set forth in SEQ ID NO: 57 and a VL set forth in SEQ ID NO: 58;

(q) a VH set forth in SEQ ID NO: 59 and a VL set forth in SEQ ID NO: 60;

(r) a VH set forth in SEQ ID NO: 61 and a VL set forth in SEQ ID NO: 62;

(s) a VH set forth in SEQ ID NO: 63 and a VL set forth in SEQ ID NO: 64;

(t) a VH set forth in SEQ ID NO: 65 and a VL set forth in SEQ ID NO: 66;

(u) a VH set forth in SEQ ID NO: 67 and a VL set forth in SEQ ID NO: 68;

(v) a VH set forth in SEQ ID NO: 69 and a VL set forth in SEQ ID NO: 70;

(w) a VH set forth in SEQ ID NO: 71 and a VL set forth in SEQ ID NO: 72;

(x) a VH set forth in SEQ ID NO: 73 and a VL set forth in SEQ ID NO: 74;

(y) a VH set forth in SEQ ID NO: 75 and a VL set forth in SEQ ID NO: 76;

(z) a VH set forth in SEQ ID NO: 145 and a VL set forth in SEQ ID NO: 146;

(aa) a VH set forth in SEQ ID NO: 147 and a VL set forth in SEQ ID NO: 148;

(bb) a VH set forth in SEQ ID NO: 149 and a VL set forth in SEQ ID NO: 150; or

(cc) a VH set forth in SEQ ID NO: 151 and a VL set forth in SEQ ID NO: 152.

26. The use of any of claims 1-25, wherein the recombinant receptor is a chimeric antigen receptor comprising an antigen-binding domain that binds BCMA, wherein the antigen-binding domain comprises:

a V_H region comprising a CDRH1, a CDRH2 and a CDRH3 comprising the amino acid sequence of SEQ ID NOS: 173, 174 and 175, respectively and a V_L region comprising a CDRL1, a CDRL2 and a CDRL3 comprising the amino acid sequence of SEQ ID NOS: 183, 184 and 185, respectively;

a V_H region comprising a CDRH1, a CDRH2 and a CDRH3 comprising the amino acid sequence of SEQ ID NOS: 176, 177 and 175, respectively and a V_L region comprising a CDRL1, a CDRL2 and a CDRL3 comprising the amino acid sequence of SEQ ID NOS: 183, 184 and 185, respectively;

a V_H region comprising a CDRH1, a CDRH2 and a CDRH3 comprising the amino acid sequence of SEQ ID NOS: 178, 179 and 175, respectively and a V_L region comprising a

CDRL1, a CDRL2 and a CDRL3 comprising the amino acid sequence of SEQ ID NOS: 183, 184 and 185, respectively; or

a V_H region comprising a CDRH1, a CDRH2 and a CDRH3 comprising the amino acid sequence of SEQ ID NOS: 180, 181 and 182, respectively and a V_L region comprising a CDRL1, a CDRL2 and a CDRL3 comprising the amino acid sequence of SEQ ID NOS: 186, 187 and 185, respectively.

27. The use of claim 26, wherein antigen-binding domain comprises a V_H region comprising an amino acid sequence having at least at or about 90% sequence identity to SEQ ID NO: 24; and a V_L region comprising an amino acid sequence having at least at or about 90% sequence identity to SEQ ID NO: 25.

28. The use of claim 26 or claim 27 wherein the antigen-binding domain comprises a V_H region comprising the amino acid sequence set forth in SEQ ID NO: 24; and a V_L region comprising the amino acid sequence set forth in SEQ ID NO: 25.

29. The use of any of claims 26-28, wherein the antigen-binding domain is an scFv comprising an amino acid sequence having at least at or about 90% sequence identity to SEQ ID NO:188.

30. The use of claim 29, wherein the antigen-binding domain is an scFv comprising an amino acid sequence set forth in SEQ ID NO:188.

31. The use of any of claims 1-25, wherein the recombinant receptor is a chimeric antigen receptor (CAR) comprising an antigen-binding domain that binds BCMA, wherein the antigen-binding domain comprises the VH region set forth in SEQ ID NO:18 and the VL region set forth in SEQ ID NO:19.

32. The use of any of claims 26-31, wherein the intracellular signaling domain comprises an intracellular domain of a CD3-zeta (CD3ζ) chain.

33. The use of any of claims 26-32, wherein the chimeric antigen receptor (CAR) further comprises a costimulatory signaling region.

34. The use of claim 33, wherein the costimulatory signaling region comprises a signaling domain derived from CD28 or 4-1BB, optionally human CD28 or human 4-1BB.

35. The use of any of claims 1-34, wherein the CAR comprises a sequence as set forth in any one of SEQ ID NOS: 90-141 or a sequence of amino acids that exhibits at least 90%, at least 91%, at least 92%, at least 93%, at least 94%, at least 95%, at least 96%, at least 97%, at least 98%, or at least 99% sequence identity to any one of SEQ ID NOS: 90-141.

36. The use of any of claims 1-35, wherein the dose of T cells to be administered comprises between at or about 2.5×10^7 CAR-expressing T cells and 1.2×10^9 CAR-expressing T cells, inclusive.

37. The use of any of claims 1-36, wherein a lymphodepleting chemotherapy is to be administered prior to administration of the cell therapy and/or wherein the subject has received a lymphodepleting chemotherapy prior to administration of the cells.

38. The use of claim 37, wherein the lymphodepleting chemotherapy that is to be administered comprises fludarabine and/or cyclophosphamide to the subject.