



(51) International Patent Classification:

A61K 47/68 (2017.01) A61M 5/14 (2006.01)
A61P 35/02 (2006.01) A61K 51/10 (2006.01)
C07K 16/28 (2006.01)

(21) International Application Number:

PCT/US2020/051324

(22) International Filing Date:

17 September 2020 (17.09.2020)

(25) Filing Language:

English

(26) Publication Language:

English

(30) Priority Data:

62/901,290 17 September 2019 (17.09.2019) US

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(81) Designated States (unless otherwise indicated, for every

kind of national protection available): AE, AG, AL, AM, AO, AT, AU, AZ, BA, BB, BG, BH, BN, BR, BW, BY, BZ, CA, CH, CL, CN, CO, CR, CU, CZ, DE, DJ, DK, DM, DO, DZ, EC, EE, EG, ES, FI, GB, GD, GE, GH, GM, GT, HN, HR, HU, ID, IL, IN, IR, IS, IT, JO, JP, KE, KG, KH, KN, KP, KR, KW, KZ, LA, LC, LK, LR, LS, LU, LY, MA, MD, ME, MG, MK, MN, MW, MX, MY, MZ, NA, NG, NI, NO, NZ, OM, PA, PE, PG, PH, PL, PT, QA, RO, RS, RU, RW, SA, SC, SD, SE, SG, SK, SL, ST, SV, SY, TH, TJ, TM, TN, TR, TT, TZ, UA, UG, US, UZ, VC, VN, WS, ZA, ZM, ZW.

(84) Designated States (unless otherwise indicated, for every

kind of regional protection available): ARIPO (BW, GH,

(54) Title: RADIOLABELING OF ANTI-CD45 IMMUNOGLOBULIN AND METHODS OF USE THEREOF

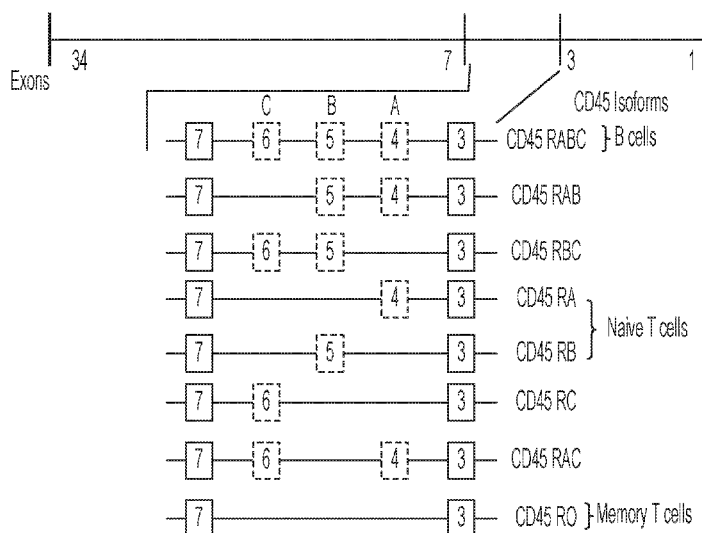


FIG. 1
PRIOR ART

(57) Abstract: Compositions and methods useful for the treatment of hemoglobinopathies and hematological diseases are disclosed herein. The compositions include an actinium-225 labeled anti-CD45 antibody (BC8) formulated as a single patient dose that is wholly deliverable to a patient in a single dose. The actinium-225 labeled anti-CD45 may be administered alone or in combination with additional therapeutic agents, such as other immunotherapeutics or a radiosensitizing agent, or additional therapeutic interventions, such as bone marrow transplant or adoptive cell therapies.



GM, KE, LR, LS, MW, MZ, NA, RW, SD, SL, ST, SZ, TZ, UG, ZM, ZW), Eurasian (AM, AZ, BY, KG, KZ, RU, TJ, TM), European (AL, AT, BE, BG, CH, CY, CZ, DE, DK, EE, ES, FI, FR, GB, GR, HR, HU, IE, IS, IT, LT, LU, LV, MC, MK, MT, NL, NO, PL, PT, RO, RS, SE, SI, SK, SM, TR), OAPI (BF, BJ, CF, CG, CI, CM, GA, GN, GQ, GW, KM, ML, MR, NE, SN, TD, TG).

Published:

- *with international search report (Art. 21(3))*
- *before the expiration of the time limit for amending the claims and to be republished in the event of receipt of amendments (Rule 48.2(h))*
- *with sequence listing part of description (Rule 5.2(a))*

**RADIOLABELING OF ANTI-CD45
IMMUNOGLOBULIN AND METHODS OF USE THEREOF**

CROSS-REFERENCE TO RELATED APPLICATIONS

[0001] This application claims priority to U.S. Provisional Patent Application Serial No. 62/901,290 filed September 17, 2019, which is incorporated herein in its entirety.

REFERENCE TO A SEQUENCE LISTING

[0002] This application contains a sequence listing incorporated herein as a supplemental file submitted via EFS and presented in compliance with 37 CFR §1.52(e)(5) and Rule 13ter.1(a), and which lists sequences identical to the sequences found within this specification.

TECHNICAL FIELD

[0003] The present disclosure relates to methods for radiolabeling monoclonal antibodies against CD45, compositions comprising the radiolabeled monoclonal antibodies against CD45, and methods for use of the radiolabeled anti-CD45 antibodies for the treatment of malignant and non-malignant hematological diseases.

BACKGROUND

[0004] CD45 is a type I transmembrane glycoprotein that is a member of the protein tyrosine phosphatase (PTP) family and plays a key role in T-cell and B-cell receptor signal transduction. CD45 controls activation of the Src family protein-tyrosine kinases Lck and Fyn. CD45 deficiency results in T- and B-lymphocyte dysfunction in the form of severe combined immune deficiency. It is also reported to play a significant role in autoimmune diseases and cancer as well as in infectious diseases including fungal infections (Penninger et al., 2001, *CD45: new jobs for an old acquaintance*, Nat. Immunol., 2(5):389-396), and metabolic disorders. The primary ligands described for CD45 include galectin-1, CD1, CD2, CD3, CD4, TCR, CD22 and Thy-1.

[0005] Also known as leukocyte common antigen (LCA), T200, or Ly-5, CD45 consists of two intracellular phosphatase domains, a transmembrane domain, and an extracellular domain. While both intracellular phosphatase domains are required for appropriate phosphate activity, only one has intrinsic kinase activity (Desai et al., 1994, *The catalytic activity of the CD45 membrane-proximal phosphatase domain is required for TCR signaling and regulation*, EMBO J. 13:4002–4010).

[0006] In general, all cells of hematopoietic origin, with the exception of mature erythrocytes and platelets, express at least one isoform of CD45. High expression of CD45 is seen with most acute lymphoid and myeloid leukemias. Since CD45 is not found on tissues of non-hematopoietic origin, its specific expression in leukemia has made it a good target for developing therapeutics, including immunotherapeutics. For example, CD45 is expressed at a density of approximately 200,000 to 300,000 sites per cell on circulating leukocytes and malignant B cells.

[0007] One particular anti-CD45 antibody, BC8, has been explored as a candidate immunotherapeutic agent alone and in combination with chemotherapy or total body irradiation in the treatment of leukemias. Anti-CD45 antibody-based lymphodepletion is also known (see, e.g., Louis, et al., 2009, *Blood*, 113:2442-2450). However, this approach had shortcomings. For example, in the Louis, et al. study, eight patients were lymphodepleted with anti-CD45 antibody and showed an increase in peripheral blood frequency of desired T-cells after infusion. However, only three patients had clinical benefits, and only one had a complete response.

[0008] CD45 exists as multiple isoforms due to alternative splicing of three of the 34 exons (exons 4, 5, and 6, designated A, B, and C; see **FIG. 1**) in the extracellular domain (Streuli et al., 1987 *Differential usage of three exons generates at least five different mRNAs encoding human leukocyte common antigens*, *J. Exp. Med.* 166:1548–1566; Chang et al., 2016, *Initiation of T cell signaling by CD45 segregation at 'close-contacts'*, *Nat. Immunol.* 17(5):574-582). These three exons encode multiple sites of O-linked glycosylation and are variably modified by sialic acid. As a result, the various isoforms differ substantially in size (391 to 552 amino acids; molecular weight ranging from 180 - 240 kDa), shape, and negative charge. The remaining membrane proximal extracellular domain is heavily N-glycosylated and contains a cysteine-rich spacer region followed by three fibronectin type III repeats.

[0009] While eight isoforms of CD45 are possible, only six are identified in humans: RO (absent all three exons), RA (exon A), RB (exon B), RAB (exons A and B), RBC (exons B and C), and RABC (exons A, B, and C). These different isoforms are differentially expressed on subpopulations of B- and T-cell lymphocytes and are specific to the activation and maturation state of the cell. For example, CD45-RA and CD45-RB are expressed on naïve T-cells, while CD45-RO is expressed on activated T-cells, some B-cell subsets, activated monocytes/macrophages, and granulocytes, and CD45-RABC is preferentially expressed on B-cells (Hermiston et al., 2003, *CD45: A critical regulator of signaling thresholds in immune cells*, *Ann. Rev. Immunol.*, 21:107-137).

[0010] Antibodies that selectively recognize various isoforms of CD45 have been identified. In addition, monoclonal antibodies (mAbs) that bind an epitope common to all the different isoforms have also been identified. For example, the anti-CD45 murine antibody BC8 recognizes all human isoforms of the CD45 antigen.

[0011] While the use of BC8 labelled with Iodine-131 (¹³¹I) for the treatment of subjects needing bone marrow transplant has been explored (*see* International Publication No. WO 2017/155937, incorporated herein by reference in its entirety), there is still a need for compositions and methods of their use for the treatment of malignant and non-malignant hematological diseases. Specifically, there is a need for a therapeutic composition and methods that (i) employ an agent that is more specific than a chemotherapeutic, (ii) is potent enough to be effective at a low dose, and (iii) spares at least some types of hematopoietic stem cells from significant depletion.

SUMMARY OF THE INVENTION

[0012] The present disclosure exploits the pan-specific nature of the BC8 monoclonal antibody to provide compositions and methods useful for depletion, reversible immunosuppression, and/or ablation of specific cell populations, and further, methods for treating certain malignant and non-malignant hematological diseases using these compositions and methods.

[0013] The present disclosure provides compositions and methods of their use for the treatment of various disorders of the hematopoietic system, as well as metabolic disorders, cancers, and autoimmune diseases, among others. The disclosure additionally features methods for conditioning a patient prior to receiving hematopoietic stem cell transplant therapy so as to promote the engraftment of hematopoietic stem cell grafts. The patient may be one that is suffering from one or more blood disorders, such as a hemoglobinopathy or other hematopoietic pathology. The patient may be one that is in need of hematopoietic stem cell transplantation.

[0014] As described herein, hematopoietic stem cells are capable of differentiating into a multitude of cell types in the hematopoietic lineage and can be administered to a patient in order to populate or re-populate a cell type that is deficient in the patient. The present disclosure features methods of treating a patient with a radiolabeled antibody, specifically an actinium-225 (²²⁵Ac) or lutetium-177 (¹⁷⁷Lu) labelled anti-CD45-immunoglobulin, which is capable of targeting hematopoietic cells to (i) directly treat a disease such as a blood disorder, metabolic disease, cancer, or autoimmune disease, among

others described herein, by selectively depleting, reversibly suppressing, or ablating a population of cells that express CD45, such as an aberrant blood cell, cancer cell, or autoimmune cell, and/or (ii) deplete, reversibly suppress, or ablate a population of endogenous hematopoietic stem cells within the patient.

[0015] The former activity enables the direct treatment of a wide range of disorders associated with a cell of the hematopoietic lineage, such as a leukemic cell or a lymphoma cell, of either B or T cell lineage, an autoimmune lymphocyte, such as a T-cell that expresses a T-cell receptor that cross-reacts with a self-antigen, among other cell types. The latter activity, the selective depletion, reversible suppression, or ablation of hematopoietic stem cells, in turn creates a vacancy that can subsequently be filled by transplantation of an exogenous (for instance, an autologous, allogeneic, or syngeneic) hematopoietic stem cell graft.

[0016] The present disclosure thus provides methods for treating a variety of non-cancerous hematopoietic conditions, such as hemoglobinopathies (e.g., sickle cell disease or SCD, and β -thalassemia), congenital immunodeficiencies (e.g., severe combined immunodeficiency or SCID, Fanconi's anemia, Wiskott-Aldrich syndrome, Diamond-Blackfan anemia, and Schwachman-Diamond syndrome, adenosine deaminase deficiency), and viral infections (e.g., HIV infection and acquired immune deficiency syndrome). The present disclosure further provides methods for treating a cancerous disorder, such as a hematological cancer or a solid tumor. Exemplary hematological cancers include acute myeloid leukemia, acute lymphoid leukemia, chronic myeloid leukemia, chronic lymphoid leukemia, multiple myeloma, diffuse large B-cell lymphoma, and non-Hodgkin's lymphoma.

[0017] Accordingly, the present disclosure relates to a stabilized composition comprising an isolated anti-CD45 immunoglobulin (e.g., BC8 mAb clone) in its actinium-225 (^{225}Ac) or lutetium-177 (^{177}Lu) radiolabeled form, and therapeutic uses thereof for the treatment of malignant and non-malignant hematological diseases and disorders. With its ability to bind all isoforms of the CD45 antigen in humans, the BC8 antibody is expected to accumulate a therapeutically high radiation dose specifically and preferentially on high-density CD45 antigen-bearing cells.

[0018] As such, this disclosure also relates to methods for radiolabeling an anti-CD45 immunoglobulin such as the BC8 antibody with radionuclide such as ^{225}Ac or ^{177}Lu . According to certain aspects, the BC8 antibody is conjugated to a chelator such as S-2-(4-Isothiocyanatobenzyl)-1,4,7,10 tetraazacyclododecanetetraacetic acid (p-SCN-Bn-DOTA; referred to as DOTA) to form DOTA-BC8, and radiolabeled with a radionuclide such as

^{225}Ac to form the ^{225}Ac -DOTA-BC8 (i.e., ^{225}Ac -BC8) or ^{177}Lu to provide ^{177}Lu -DOTA-BC8 (i.e., ^{177}Lu -BC8).

[0019] The ^{225}Ac -BC8 or ^{177}Lu -BC8 may be provided as a stabilized formulation comprising one or more pharmaceutically acceptable carriers, salts, or excipients. Certain exemplary carriers or excipients include saline, phosphate buffered saline (e.g., 50 mM PBS buffer, pH 7), and/or 0.5% to 5.0% (w/v) of one or more of ascorbic acid, polyvinylpyrrolidone (PVP), human serum albumin (HSA), a water-soluble salt of HSA, and mixtures thereof.

[0020] The BC8 antibody may comprise a light chain variable domain having the amino acid sequence as set forth in SEQ ID NO:1, or a heavy chain variable domain having an N-terminal amino acid sequence as set forth in SEQ ID NO:9. The BC8 antibody may comprise a light chain variable domain having at least one complementarity determining region (CDR) with the amino acid sequence as set forth in SEQ ID NO:3, SEQ ID NO:4, and SEQ ID NO:5. The BC8 antibody may comprise a light chain having the amino acid sequence as set for the in SEQ ID NO:12 or SEQ ID NO:13.

[0021] The BC8 antibody may comprise a heavy chain variable domain having the amino acid sequence as set forth in SEQ ID NO:2, or a heavy chain variable domain having an N-terminal amino acid sequence as set forth in SEQ ID NO:10. The BC8 antibody may comprise a heavy chain variable domain having at least one complementarity determining region (CDR) with the amino acid sequence as set forth in SEQ ID NO:6, SEQ ID NO:7, and SEQ ID NO:8. The BC8 antibody may comprise a heavy chain having the amino acid sequence as set for the in SEQ ID NO:15 or SEQ ID NO:16.

[0022] According to certain aspects, the BC8 antibody comprises a heavy chain having the amino acid sequence set forth in SEQ ID NO:15 or 16, wherein the amino acid at position 141 (relative to the N-terminal amino acid) is either an ASP or an ASN. A ratio of ASP:ASN at position 141 in a population of BC8 proteins may be within the range 1:99 to 99:1, such as 10:90 to 90:10.

[0023] According to certain aspects, the BC8 antibody comprises a heavy chain variable domain having the amino acid sequence as set forth in SEQ ID NO:2, or a heavy chain variable domain having an N-terminal amino acid sequence as set forth in SEQ ID NO:10, wherein the amino acid at position 141 (relative to the N-terminal amino acid) of the heavy chain is either an ASP or an ASN, with a ratio of ASP:ASN in a population of BC8 proteins within the range 1:99 to 99:1, such as 10:90 to 90:10.

[0024] According to certain aspects, any of the BC8 antibodies indicated above, i.e., those including one or more of SEQ ID NOS:1-10 may be a chimeric or humanized antibody, i.e., BC8c. The BC8c antibody may comprise a human IgG1, IgG2, or IgG4 heavy chain constant region having the amino acid sequence as set forth in SEQ ID NOS:17-19, respectively, a human IgG4 heavy chain constant region having the amino acid sequence as set forth in SEQ ID NO:20 (includes the mutation S228P), and/or a human kappa light chain constant region having the amino acid sequence as set forth in SEQ ID NO:21.

[0025] This disclosure provides methods for directly treating a subject afflicted with a CD45 positive hematological malignancy comprising administering to the subject an effective amount of $^{225}\text{Ac-BC8}$ or $^{177}\text{Lu-BC8}$ as a low dose single therapy agent alone or in combination with other therapies.

[0026] This disclosure provides methods for directly treating a subject afflicted with a CD45 positive hematological malignancy comprising administering to the subject an effective amount of $^{225}\text{Ac-BC8}$ or $^{177}\text{Lu-BC8}$ as a low dose single therapy agent alone or in combination with other therapies with stem cell support.

[0027] The present disclosure relates to methods for depleting, reversibly suppressing, or ablating a subject's hematopoietic stem cells comprising administering to the subject an effective amount of $^{225}\text{Ac-BC8}$ or $^{177}\text{Lu-BC8}$.

[0028] The present disclosure provides methods for depleting or reversibly suppressing circulating tumor cells (e.g., as found in leukemia, lymphoma, myeloma, MDS) by administering to the subject an effective amount of $^{225}\text{Ac-BC8}$ or $^{177}\text{Lu-BC8}$ at a dose that does not myeloablate, and thus does not irreversibly deplete hematopoietic stem cells. Such cells may comprise any of, at least, regulatory T cells, myeloid derived suppressor cells, tumor associated macrophages, activated macrophages secreting IL-1 and/or IL-6, and combinations thereof.

[0029] This disclosure further provides methods for depleting, reversibly suppressing, or ablating a subject's lymphocytes comprising administering to the subject an effective amount of $^{225}\text{Ac-BC8}$ or $^{177}\text{Lu-BC8}$.

[0030] This disclosure also provides methods for treating a subject afflicted with a non-cancerous disorder comprising administering to the subject an amount of $^{225}\text{Ac-BC8}$ or $^{177}\text{Lu-BC8}$ effective to deplete, reversibly suppress, or ablate the subject's hematopoietic stem cells. According to certain aspects, the disorder is treatable via genetically edited cell therapy, and the method further comprises performing the therapy on the subject to treat the subject's disorder after administration of the $^{225}\text{Ac-BC8}$ or $^{177}\text{Lu-BC8}$. According to certain

aspects, the disorder is SCD and the therapy is genetically edited β -globin hematopoietic stem cell therapy. According to certain aspects, the disorder is SCID and the therapy is genetically edited hematopoietic stem cell therapy, wherein the edited gene is the common gamma chain (γ c) gene, the adenosine deaminase (ADA) gene and/or the Janus kinase 3 (JAK3) gene. The stem cell therapy can be allogenic or autologous, for example.

[0031] This disclosure also provides methods for treating a subject afflicted with a cancerous disorder treatable via genetically edited cell therapy comprising (i) administering to the subject an amount of $^{225}\text{Ac-BC8}$ or $^{177}\text{Lu-BC8}$ effective to deplete, reversibly suppress, or ablate the subject's hematopoietic stem cells, and (ii) after a suitable time period, performing the therapy on the subject to treat the subject's disorder. According to certain aspects, the therapy suitable to treat the subject's disorder may be a bone marrow transplant, or an adoptive cell therapy.

[0032] Finally, this disclosure provides an article of manufacture comprising (a) a radiolabeled anti-CD45 antibody such as $^{225}\text{Ac-BC8}$ or $^{177}\text{Lu-BC8}$, and (b) a label instructing the user to administer to a subject an amount of the antibody effective to deplete the subject's hematopoietic stem cells.

BRIEF DESCRIPTION OF THE DRAWINGS

[0033] The patent or application file contains at least one drawing executed in color. Copies of this patent or patent application publication with color drawing(s) will be provided by the Office upon request and payment of the necessary fee.

[0034] **FIG. 1** shows a schematic diagram of exon usage in various isoforms of CD45 produced by differential splicing of the human CD45 gene.

[0035] **FIG. 2** provides the protein sequence of the complementarity determining regions (CDRs), framework regions and variable domain sequences of the light chain (VL) and the heavy chain (VH) of the anti-CD45 mAb BC8. The CDRs are in bold and underlined (SEQ ID NOS: 1 and 2).

[0036] **FIG. 3** provides the CDRs and the N-terminal protein sequences of the light chain and the heavy chain of the anti-CD45 mAb BC8 (SEQ ID NOS: 3 – 10).

[0037] **FIG. 4A** provides the entire nucleotide (SEQ ID NO:11) and amino acid (SEQ ID NO:12) sequence of the light chain of the anti-CD45 mAb BC8.

[0038] **FIG. 4B** provides the amino acid (SEQ ID NO:13) sequence of the light chain of the anti-CD45 mAb BC8 without the leader sequence.

[0039] **FIG. 5A** provides the entire nucleotide (SEQ ID NO:14) and amino acid (SEQ ID NO:15) sequence of the heavy chain of the anti-CD45 mAb BC8, wherein the asparagine at position 141 (from the n terminal of the protein sequence) is found to be deaminated to aspartic acid in at least a portion of the protein population.

[0040] **FIG. 5B** provides the amino acid (SEQ ID NO:16) sequence of the heavy chain of the anti-CD45 mAb BC8 without the leader sequence, wherein the asparagine at position 141 (from the n terminal of the protein sequence) is found to be deaminated to aspartic acid in at least a portion of the protein population.

[0041] **FIGS. 6A-6C** provide the amino acid sequences SEQ ID NOS.17-19 of the human heavy chain constant region, respectively.

[0042] **FIG. 6D** provides the amino acid sequence (SEQ ID NO: 20) of the human heavy chain constant region comprising the mutation S228P.

[0043] **FIG. 6E** provides the amino acid sequence (SEQ ID NO:21) of the human kappa light chain constant region.

[0044] **FIG. 7** depicts a method for lymphodepleting a subject prior to performing an adoptive cell therapy according to certain aspects of the present disclosure.

[0045] **FIG. 8** depicts pharmino-kinetic data demonstrating exemplary clearance and dosing times for a lymphodepletion protocol according to the present disclosure.

[0046] **FIGS. 9A and 9B** provide schematic diagrams of a method for radiolabeling the anti-CD45 mAb BC8 with actinium (^{225}Ac), wherein **FIG. 9A** illustrates attachment of the bifunctional chelator S-2-(4-Isothiocyanatobenzyl)-1,4,7,10 tetraazacyclododecanetetraacetic acid (p-SCN-Bn-DOTA; referred to as DOTA in the figures) to the monoclonal antibody against CD45, and **FIG. 9B** illustrates radiolabeling of the DOTA-anti-CD45 conjugate with ^{225}Ac to provide ^{225}Ac -DOTA-anti-CD45.

[0047] **FIGS. 10A and 10B** provide elution profiles for a BC8 standard and ^{225}Ac -DOTA-BC8 from SEC-HPLC (size exclusion chromatography-high performance liquid chromatography), wherein **FIG. 10A** illustrates elution of the BC8 standard and **FIG. 10B** illustrates elution of the ^{225}Ac -DOTA-BC8 (the peak at 13 minutes is the HSA added to stabilize the conjugated antibody).

[0048] **FIG. 11** provides a graph showing the stability of ^{225}Ac -DOTA-BC8 at various storage dilutions and temperatures as a function of time.

[0049] **FIG. 12** provides a graph showing the ^{225}Ac -DOTA-BC8 immunoreactivity against Ramos cells (CD45 positive cells) and EL4 cells (Cd45 negative cells).

[0050] **FIGS. 13A and 13B** provide graphs showing the binding of various antibody samples to Cytotrol cells measured by flow cytometry, wherein **FIG. 13A** compares binding of the naïve BC8 and naïve 18B7 (control nonspecific) antibodies to the Cytotrol cells, and **FIG. 13B** compares binding of naïve BC8 and DOTA-BC8 antibodies to the Cytotrol cells.

[0051] **FIG. 14A** provides a graph comparing the binding of naïve BC8 and DOTA-BC8 to Cytotrol cells measured by flow cytometry.

[0052] **FIG. 14B** provides a bar graph showing binding of the DOTA-BC8 sample from **FIG. 14A** immediately after labeling with ^{225}Ac (i.e., ^{225}Ac -DOTA-BC8) compared to the binding of ^{225}Ac -18B7 (binding to Cytotrol cells) as measured by fraction radiation retained on the cells after washing.

[0053] **FIGS. 15A and 15B** show graphs comparing the binding of DOTA-BC8 to different human multiple myeloma cell lines, i.e., H929 and U266, measure by flow cytometry (**FIG. 15A**) or after labeling with ^{225}Ac (i.e., ^{225}Ac -DOTA-BC8; **FIG. 15B**) as measured by fraction radiation retained on the cells after washing.

[0054] **FIGS. 16A and 16B** show bar graphs comparing the biodistribution of the ^{225}Ac -DOTA-BC8 (**FIG. 16A**) and ^{225}Ac -DOTA-18B7 (**FIG. 16B**) antibodies in control mice at 1hour, 4 hours, 24 hours, 48 hours, and 96 hours.

[0055] **FIGS. 17A and 17B** show bar graphs comparing the biodistribution of the ^{225}Ac -DOTA-18B7 (control; **FIG. 17A**) and ^{225}Ac -DOTA-BC8 (**FIG. 17B**) antibodies in U266 and H929 SCID-NOD tumor bearing mice at 1hour, 4 hours, 24 hours, 48 hours, and 96 hours.

[0056] **FIG. 18A** shows a graph comparing the tumor volume of H929 multiple myeloma xenograph-bearing SCID-NOD mice after radioimmunotherapy treatment with ^{225}Ac -DOTA-BC8 or ^{225}Ac -DOTA-18B7 (control).

[0057] **FIG. 18B** shows a graph comparing the tumor volume of U266 multiple myeloma xenograph-bearing SCID-NOD mice after radioimmunotherapy treatment with ^{225}Ac -DOTA-BC8 or ^{225}Ac -DOTA-18B7 (control).

[0058] **FIGS. 19A-19D** show histological analysis of tumors excised from U266 and H929 multiple myeloma xenograph-bearing SCID-NOD mice, wherein **FIG. 19A** shows an untreated H929 tumor, **FIG. 19B** shows an ^{225}Ac -DOTA-BC8 treated H929 tumor, **FIG. 19C** shows an untreated U266 tumor, and **FIG. 19D** shows an ^{225}Ac -DOTA-BC8 treated U266 tumor.

[0059] **FIG. 20** shows microSPEC/CT scans of C57Bl/6 mice injected (i.p.) with ¹¹¹Ln-anti-CD45 taken 1 hour, 24 hours, 48hours, 72 hours, 96 hours, and 6 days after injection.

[0060] **FIG. 21** shows bar graphs of the amounts of depletion of various immune cell subpopulations in non-tumor bearing C57Bl/6 mice after treatment with (A) ¹⁷⁷Lu-anti-CD45 or (B) ¹³¹I-anti-CD45.

[0061] **FIG. 22** shows bar graphs of the amounts of depletion of various immune cell populations in the spleens of non-tumor bearing C57Bl/6 mice after treatment with (A) ¹⁷⁷Lu-anti-CD45 or (B) ¹³¹I-anti-CD45.

[0062] **FIG. 23** shows graphs demonstrating that ¹⁷⁷Lu-anti-CD45 and ¹³¹I-anti-CD45 lymphodepletion enable tumor control in an OT I adoptive cell therapy model, wherein (A) demonstrates ¹⁷⁷Lu-anti-CD45 and ¹³¹I-anti-CD45-mediated targeted conditioning prior to adoptively transferred OT I T-cells enabled control of EG.7 tumor growth; (B) shows tumor size for individual mice in each group; and (C) shows survival of control mice that received no conditioning or OT I T-cells, and those conditioned with ¹⁷⁷Lu-anti-CD45 and ¹³¹I-anti-CD45.

BRIEF DESCRIPTION OF THE SEQUENCES

[0063] **SEQ ID NO:1** is the amino acid sequence of the variable domain of the light chain of anti-CD45 murine immunoglobulin BC8.

[0064] **SEQ ID NO:2** is the amino acid sequence of the variable domain of the heavy chain of anti-CD45 murine immunoglobulin BC8.

[0065] **SEQ ID NO:3** is the amino acid sequence of CDR1 of the light chain of anti-CD45 murine immunoglobulin BC8.

[0066] **SEQ ID NO:4** is the amino acid sequence of CDR2 of the light chain of anti-CD45 murine immunoglobulin BC8.

[0067] **SEQ ID NO:5** is the amino acid sequence of CDR3 of the light chain of anti-CD45 murine immunoglobulin BC8.

[0068] **SEQ ID NO:6** is the amino acid sequence of CDR1 of the heavy chain of anti-CD45 murine immunoglobulin BC8.

[0069] **SEQ ID NO:7** is the amino acid sequence of CDR2 of the heavy chain of anti-CD45 murine immunoglobulin BC8.

[0070] **SEQ ID NO:8** is the amino acid sequence of CDR3 of the heavy chain of anti-CD45 murine immunoglobulin BC8.

[0071] **SEQ ID NO:9** is the amino acid sequence of N-terminus of the light chain of anti-CD45 murine immunoglobulin BC8.

[0072] **SEQ ID NO:10** is the amino acid sequence of N-terminus of the heavy chain of anti-CD45 murine immunoglobulin BC8.

[0073] **SEQ ID NO:11** is the nucleotide sequence of the light chain of anti-CD45 murine immunoglobulin BC8.

[0074] **SEQ ID NO:12** is the amino acid sequence of the light chain of anti-CD45 murine immunoglobulin BC8 including a leader sequence.

[0075] **SEQ ID NO:13** is the amino acid sequence of the light chain of anti-CD45 murine immunoglobulin BC8 starting at the protein N-terminal (i.e., absent the leader sequence).

[0076] **SEQ ID NO:14** is the nucleotide sequence of the heavy chain of anti-CD45 murine immunoglobulin BC8.

[0077] **SEQ ID NO:15** is the amino acid sequence of the heavy chain of anti-CD45 murine immunoglobulin BC8 including a leader sequence.

[0078] **SEQ ID NO:16** is the amino acid sequence of the heavy chain of anti-CD45 murine immunoglobulin BC8 starting at the protein N-terminal (i.e., absent the leader sequence).

[0079] **SEQ ID NO:17** is the amino acid sequence of the human IgG1 heavy chain constant region.

[0080] **SEQ ID NO:18** is the amino acid sequence of the human IgG2 heavy chain constant region.

[0081] **SEQ ID NO:19** is the amino acid sequence of the human IgG4 heavy chain constant region.

[0082] **SEQ ID NO:20** is the amino acid sequence of the human IgG4 heavy chain constant region comprising the mutation S228P.

[0083] **SEQ ID NO:21** is the amino acid sequence of the human Kappa light chain constant region.

DEFINITIONS AND ABBREVIATIONS

[0084] Throughout this application, various publications are cited. The disclosure of these publications is hereby incorporated by reference into this application to describe more fully the state of the art to which this disclosure pertains.

[0085] In this application, certain terms are used which shall have the meanings set forth as follows.

[0086] The singular forms “a,” “an,” “the” and the like include plural referents unless the context clearly dictates otherwise. Thus, for example, reference to “an” antibody includes both a single antibody and a plurality of different antibodies.

[0087] The term “about” when used before a numerical designation, e.g., temperature, time, amount, and concentration, including a range, indicates approximations which may vary by $\pm 10\%$, $\pm 5\%$, or $\pm 1\%$.

[0088] As used herein, “administer”, with respect to an antibody, means to deliver the antibody to a subject’s body via any known method suitable for antibody delivery. Specific modes of administration include, without limitation, intravenous, transdermal, subcutaneous, intraperitoneal and intrathecal administration. Exemplary administration methods for antibodies may be as substantially described in International Publication No. WO 2016/187514, incorporated in its entirety herein by reference herein.

[0089] In addition, according to aspects of the present disclosure, antibodies can be formulated using one or more routinely used pharmaceutically acceptable carriers. Such carriers are well known to those skilled in the art. For example, injectable drug delivery systems include solutions, suspensions, gels, microspheres and polymeric injectables, and can comprise excipients such as solubility-altering agents (e.g., ethanol, propylene glycol and sucrose) and polymers (e.g., polycaprylactones and PLGA’s).

[0090] As used herein, the term “antibody” includes, without limitation, (a) an immunoglobulin molecule comprising two heavy chains and two light chains, which recognizes an antigen; (b) polyclonal and monoclonal immunoglobulin molecules; (c) monovalent and divalent fragments thereof (e.g., di-Fab); and (d) bi-specific forms thereof. Immunoglobulin molecules may derive from any of the commonly known classes, including but not limited to IgA, secretory IgA, IgG and IgM. IgG subclasses are also well known to those in the art and include, but are not limited to, human IgG1, IgG2, IgG3 and IgG4. Antibodies can be both naturally occurring and non-naturally occurring (e.g., IgG-Fc-silent). Furthermore, antibodies include chimeric antibodies, wholly synthetic antibodies, single chain antibodies, and fragments thereof. Antibodies may be human, humanized or nonhuman.

[0091] A “humanized” antibody refers to an antibody in which some, most or all of the amino acids outside the CDR domains of a non-human antibody are replaced with corresponding amino acids derived from human immunoglobulins. In one embodiment of a

humanized form of an antibody, some, most or all of the amino acids outside the CDR domains have been replaced with amino acids from human immunoglobulins, whereas some, most or all amino acids within one or more CDR regions are unchanged. Small additions, deletions, insertions, substitutions or modifications of amino acids are permissible as long as they do not abrogate the ability of the antibody to bind to a particular antigen. A “humanized” antibody retains an antigenic specificity similar to that of the original antibody.

[0092] A “chimeric antibody” refers to an antibody in which the variable regions are derived from one species and the constant regions are derived from another species, such as an antibody in which the variable regions are derived from a mouse antibody and the constant regions are derived from a human antibody.

[0093] As used herein, “non-cancerous disorders” or “non-malignant disorders” include, without limitation, hemoglobinopathies (e.g., SCD), congenital immunodeficiencies (e.g., SCID), autoimmune disorders (e.g., multiple sclerosis, rheumatoid arthritis, scleroderma, systemic lupus, Type 1 diabetes, myasthenia gravis, sjogren’s disease, polymyositis, etc.), and viral infections (e.g., an HIV infection). Non-cancerous disorders exclude, for example, solid cancers (e.g., tumors) and hematologic malignancies.

[0094] As used herein, “cancer” or “malignant disorder” includes, without limitation, a solid cancer (e.g., a tumor) and a hematologic malignancy. A “hematologic malignancy”, also known as a blood cancer, is a cancer that originates in blood-forming tissue, such as the bone marrow or other cells of the immune system. Hematologic malignancies include, without limitation, leukemias (such as acute myeloid leukemia (AML), acute promyelocytic leukemia, acute lymphoblastic leukemia (ALL), acute mixed lineage leukemia, chronic myeloid leukemia, chronic lymphocytic leukemia (CLL), hairy cell leukemia and large granular lymphocytic leukemia), myelodysplastic syndrome (MDS), myeloproliferative disorders (polycythemia vera, essential thrombocytosis, primary myelofibrosis and chronic myeloid leukemia), lymphomas, multiple myeloma, MGUS and similar disorders, Hodgkin’s lymphoma, non-Hodgkin lymphoma (NHL), primary mediastinal large B-cell lymphoma, diffuse large B-cell lymphoma, follicular lymphoma, transformed follicular lymphoma, splenic marginal zone lymphoma, lymphocytic lymphoma, T-cell lymphoma, and other B-cell malignancies.

[0095] “Solid cancers” include, without limitation, bone cancer, pancreatic cancer, skin cancer, cancer of the head or neck, cutaneous or intraocular malignant melanoma, uterine cancer, ovarian cancer, prostate cancer, rectal cancer, cancer of the anal region, stomach cancer, testicular cancer, uterine cancer, carcinoma of the fallopian tubes, carcinoma

of the endometrium, carcinoma of the cervix, carcinoma of the vagina, carcinoma of the vulva, cancer of the esophagus, cancer of the small intestine, cancer of the endocrine system, cancer of the thyroid gland, cancer of the parathyroid gland, cancer of the adrenal gland, sarcoma of soft tissue, cancer of the urethra, cancer of the penis, pediatric tumors, cancer of the bladder, cancer of the kidney or ureter, carcinoma of the renal pelvis, neoplasm of the central nervous system (CNS), primary CNS lymphoma, tumor angiogenesis, spinal axis tumor, brain stem glioma, pituitary adenoma, Kaposi's sarcoma, epidermoid cancer, squamous cell cancer, environmentally-induced cancers including those induced by asbestos.

[0096] As used herein, the term “burden”, when used in connection with a cancerous cell, means quantity. So, a cancerous cell “burden” means the quantity of cancerous cells. Cancerous cells have a burden with respect to their tissue of origin (i.e., the primary site of disease), such as the “bone marrow blast burden” in the case of AML. Cancerous cells also have a burden with respect to one or more tissues other than those of origin, such as the blast burden in blood, liver and spleen in the case of AML. The term “peripheral burden” relates to such cells. The peripheral burden of cancerous cells, such as blasts in the case of AML, can be measured in different ways with different outcomes. For example, in the case of AML, the “peripheral blast burden” can be measured as the total blast population outside of the bone marrow, or the total blast population of the blood, spleen and liver combined, or simply the blast population of the blood as measured in cells per unit volume. As used herein in connection with AML and other cancers originating in the bone marrow, and unless stated otherwise, the term “peripheral cancerous cell burden” (e.g., peripheral blast burden) refers to the cancerous cell population of the blood as measured in cells per unit volume (e.g., cells/ μ l). This blood-based measurement is a useful proxy for the more cumbersome measurements of spleen and liver burdens, for example.

[0097] Herein, a peripheral cancerous cell burden in a subject is “high” if, when the subject is administered an agent, e.g., a radiolabeled anti-CD45 antibody of the present disclosure, that targets a hematologic malignancy-associated antigen at the maximum safe dose, the agent does not reach the primary site of disease in a sufficient amount to bind to more than 90% of its target antigens at that site. Conversely, a peripheral cancerous cell burden in a subject is “low” if, when the subject is administered the agent at the maximum safe dose, the agent reaches the primary site of disease in a sufficient amount to bind to more than 90% of its target antigens at that site. In the case of AML, examples of low peripheral blast burden are those yielding blood blast burdens at or below 1,000 blast cells/ μ l, at or

below 500 blast cells/ μ l, at or below 400 blast cells/ μ l, at or below 300 blast cells/ μ l, at or below 200 blast cells/ μ l, at or below 100 blast cells/ μ l, and at or below 50 blast cells/ μ l.

[0098] As used herein, a “low dose” of radiolabeled anti-CD45 antibody of the present disclosure is one that is sub-saturating, and as such introduces into the subject's body fewer target antigen-binding sites (i.e., CD45-binding sites on the administered antibody) than there are target antigens (i.e., CD45 molecules). According to certain aspects, a low dose of the radiolabeled anti-CD45 antibody is one where the ratio of CD45-binding sites to CD45 molecules is less than or equal to 9:10, such as less than or equal to 1:2, or less than or equal to 1:5, or less than or equal to 1:10, or less than or equal to 1:20, or less than or equal to 1:100.

[0099] As used herein, the term “subject” or “patient” are interchangeable and include, without limitation, a mammal such as a human, a non-human primate, a dog, a cat, a horse, a sheep, a goat, a cow, a rabbit, a pig, a rat and a mouse. Where the subject is human, the subject can be of any age. According to certain aspects, the subject is an infant. According to further aspects, the subject is one, two, three, four, five, six, seven, eight, nine or 10. According to yet further aspects, the subject is from 10 to 15, or from 15 to 20. According to yet further aspects, the subject is 20 or older, 25 or older, 30 or older, 35 or older, 40 or older, 45 or older, 50 or older, 55 or older, 60 or older, 65 or older, 70 or older, 75 or older, 80 or older, 85 or older, or 90 or older.

[0100] As used herein, “treating” a subject afflicted with a disorder shall include, without limitation, (i) slowing, stopping or reversing the disorder's progression, (ii) slowing, stopping or reversing the progression of the disorder's symptoms, (iii) reducing, and ideally eliminating, the likelihood of the disorder's recurrence, and/or (iv) reducing, and ideally eliminating, the likelihood that the disorder's symptoms will recur. According to certain preferred aspects, treating a subject afflicted with a disorder means (i) reversing the disorder's progression, ideally to the point of eliminating the disorder, and/or (ii) reversing the progression of the disorder's symptoms, ideally to the point of eliminating the symptoms, and/or (iii) reducing or eliminating the likelihood of relapse. Ideally, treating a subject afflicted with a disorder means curing the disorder by removing or otherwise disabling its genetic cause.

[0101] As used herein, “depleting” with respect to a specific cell type of the subject means reducing that cell population within the subject by at least 10%, 20%, 30%, 40%, 50%, 60%, 70%, 80%, 90%, or 95%. As used herein, “ablating” with respect to a specific cell type

of the subject means reducing that cell population within the subject by greater than 95%, such as by at least 96%, or 97%, or 98%, or 99%, or even 100%.

[0102] The specific cell types depleted using the compositions and methods of the present disclosure include at least hematopoietic stem cells (i.e., multipotential hematopoietic stem cells, also referred to as hemocytoblasts), and lymphocytes, such as peripheral blood lymphocytes or bone marrow lymphocytes. Hematopoietic stem cells (“HSCs”) are multipotent, self-renewing progenitor cells from which all differentiated blood cell types arise during the process of hematopoiesis. HSCs are thought to differentiate into two lineage-restricted, lymphoid and myelo-erythroid, oligopotent progenitor cells, although an alternative “myeloid-based” model for blood lineage development describes a novel intermediary myelo-lymphoid progenitor cell, which has the capacity to generate progeny from both lineages.

[0103] As used herein, the term “hematopoietic stem cells” (“HSCs”) refers to immature blood cells having the capacity to self-renew and to differentiate into mature blood cells containing diverse lineages including but not limited to granulocytes (e.g., promyelocytes, neutrophils, eosinophils, basophils), erythrocytes (e.g., reticulocytes, erythrocytes), thrombocytes (e.g., megakaryoblasts, platelet producing megakaryocytes, platelets), monocytes (e.g., monocytes, macrophages), dendritic cells, microglia, osteoclasts, and lymphocytes (e.g., NK cells, B-cells and T-cells). Such cells may include CD34+ cells. CD34+ cells are immature cells that express the CD34 cell surface marker.

[0104] Methods for measuring HSC populations are routine. They include, for example, the use of flow cytometry to detect human HSCs in a bone marrow sample and staining for various cell surface markers (such as Lin, CD34, CD38, CD43, CD45RO, CD45RA, CD59, CD90, CD109, CD117, CD133, CD166, and HLA DR). Reduction of a patient’s immune cells may also be detected in peripheral blood. Methods for measuring peripheral blood lymphocyte populations are routine. They include, for example, flow cytometry on whole blood samples to determine lymphocyte counts based on labeling with a fluorescent antibody directed against a specific a cell surface marker such as CD45, CD3, CD4 or CD8. Methods for measuring peripheral blood neutrophil populations are also routine. They include, for example, flow cytometry on whole blood samples to determine neutrophil counts based on labeling with a fluorescent antibody directed against a specific a cell surface marker such as Ly6G.

[0105] According to certain aspects of the disclosure, a subject’s lymphocyte decrease is determined by measuring the subject’s peripheral blood lymphocyte level. As

used herein, a subject's "peripheral blood lymphocytes" shall mean the mature lymphocytes circulating in the subject's blood. Examples of peripheral blood lymphocytes include, without limitation, peripheral blood T-cells, peripheral blood NK cells and peripheral blood B cells. As such, and by way of example, a subject's lymphocyte population is depleted if the population of at least one type of the subject's peripheral blood lymphocytes is lowered by no more than 95%. For example, a subject's lymphocytes are depleted if the subject's peripheral blood T-cell level is lowered by 50%, the subject's peripheral blood NK cell level is lowered by 40%, and/or the subject's peripheral blood B cell level is lowered by 30%. In this example, the subject's lymphocytes are depleted even if the level of another immune cell type, such as neutrophils, is not lowered. According to certain aspects, depleting a subject's lymphocytes is reflected by a peripheral blood lymphocyte population reduction of at least 10%, 20%, 30%, 40%, 50%, 60%, 70%, 80%, 90% or 95%.

[0106] As used herein, patients that are "in need of" a hematopoietic stem cell transplant include patients that exhibit a defect or deficiency in one or more blood cell types, as well as patients having a stem cell disorder, autoimmune disease, cancer, or other pathology described herein. Hematopoietic stem cells generally exhibit 1) multi-potency, and can thus differentiate into multiple different blood lineages including, but not limited to, granulocytes (e.g., promyelocytes, neutrophils, eosinophils, basophils), erythrocytes (e.g., reticulocytes, erythrocytes), thrombocytes (e.g., megakaryoblasts, platelet producing megakaryocytes, platelets), monocytes (e.g., monocytes, macrophages), dendritic cells, microglia, osteoclasts, and lymphocytes (e.g., NK cells, B-cells and T-cells), 2) self-renewal, and can thus give rise to daughter cells that have equivalent potential as the mother cell, and 3) the ability to be reintroduced into a transplant recipient whereupon they home to the hematopoietic stem cell niche and re-establish productive and sustained hematopoiesis.

[0107] Additionally or alternatively, a patient "in need of" a hematopoietic stem cell transplant may one that is or is not suffering from a pathology, but nonetheless exhibits a reduced level (e.g., as compared to that of an otherwise healthy subject) of one or more endogenous cell types within the hematopoietic lineage, such as megakaryocytes, thrombocytes, platelets, erythrocytes, mast cells, myeloblasts, basophils, neutrophils, eosinophils, microglia, granulocytes, monocytes, osteoclasts, antigen-presenting cells, macrophages, dendritic cells, natural killer cells, T-lymphocytes, and B-lymphocytes.

[0108] *The anti-CD45 antibody*

[0109] As used herein, an "anti-CD45 antibody" or "anti-CD45-immunoglobulin" is an antibody that binds to an epitope of CD45. According to certain aspects, the anti-CD45

antibody may bind to the epitope recognized by the monoclonal antibody "BC8." BC8 is known, as are methods of making it. These methods are described, for example, in International Publication No. WO 2017/155937, incorporated by reference herein in its entirety, and in the examples provided herein.

[0110] The BC8 monoclonal antibody may comprise a light chain having the amino acid sequence set forth in SEQ ID NO:12, which includes the leader sequence (**FIG. 4A**), or SEQ ID NO:13, which excludes the leader sequence (**FIG. 4B**). The BC8 monoclonal antibody may comprise a light chain variable region having the amino acid sequence set forth in SEQ ID NO:1 (**FIG. 2**). The BC8 monoclonal antibody may comprise a light chain having the N-terminal amino acid sequence set forth in SEQ ID NO:9 (**FIG. 3**). According to certain aspects, the light chain comprises at least one complementarity determining region having the amino acid sequence as set forth in SEQ ID NO:3, SEQ ID NO:4, or SEQ ID NO:5 (**FIG. 3**). According to certain aspects, the light chain comprises the N-terminal amino acid sequence set forth in SEQ ID NO:9 and at least one complementarity determining region having the amino acid sequence as set forth in SEQ ID NO:3, SEQ ID NO:4, or SEQ ID NO:5 (**FIG. 3**).

[0111] The BC8 monoclonal antibody may comprise a heavy chain having the amino acid sequence set forth in SEQ ID NO:15, which includes the leader sequence (**FIG. 5A**), or SEQ ID NO:16, which excludes the leader sequence (**FIG. 5B**). The BC8 monoclonal antibody may comprise a heavy chain variable region having the amino acid sequence set forth in SEQ ID NO:2 (**FIG. 2**). The BC8 monoclonal antibody may comprise a heavy chain having the N-terminal amino acid sequence set forth in SEQ ID NO:10 (**FIG. 3**). According to certain aspects, the heavy chain comprises at least one complementarity determining region having the amino acid sequence as set forth in SEQ ID NO:6, SEQ ID NO:7, or SEQ ID NO:8 (**FIG. 3**). According to certain aspects, the heavy chain comprises a heavy chain having the N-terminal amino acid sequence set forth in SEQ ID NO:10 and at least one complementarity determining region having the amino acid sequence as set forth in SEQ ID NO:6, SEQ ID NO:7, or SEQ ID NO:8 (**FIG. 3**).

[0112] According to certain aspects, the BC8 monoclonal antibody comprises a heavy chain having the amino acid sequence set forth in SEQ ID NO:15 or 16, wherein the amino acid at position 141 (relative to the N-terminal amino acid) is either an ASP or an ASN. A ratio of ASP:ASN at position 141 in a population of BC8 proteins may be within the range 1:99 to 99:1, such as 10:90 to 90:10.

[0113] According to certain aspects, the BC8 monoclonal antibody comprises a heavy chain variable region having the amino acid sequence set forth in SEQ ID NO:2, wherein the

amino acid at position 141 in the constant region (relative to the N-terminal amino acid) is either an ASP or an ASN. A ratio of ASP:ASN at position 141 in a population of BC8 proteins may be within the range 1:99 to 99:1, such as 10:90 to 90:10.

[0114] According to certain aspects, the antibody against CD45 (anti-CD45 antibody) may be a chimeric or humanized antibody. For example, the BC8 monoclonal antibody may comprise a humanized or chimeric BC8 antibody (referred to as “BC8c” herein). For example, a humanized BC8c monoclonal antibody may comprise the parent murine variable (V) regions grafted on to human IgG1, IgG2, or IgG4 constant regions for the heavy chain or a human kappa region for the light chain. IgG4 antibodies are capable of exchanging Fab arms by swapping a heavy chain and attached light chain (half molecule) with a heavy-light chain pair from another molecule, resulting in bispecific antibodies. This process, termed “Fab-arm exchange” herein, has been shown to occur under reducing conditions in vitro and in vivo in mice. The ability of IgG4 antibodies to undergo Fab-arm exchange has been accredited to the instable core-hinge sequence in combination with sequence determinants in the IgG4 CH3 domain. Replacement of core-hinge residue Ser228 by Pro (S228P) results in a partial stabilization of an IgG4 molecule in vitro and in vivo. As such, according to certain aspects, the IgG4 can comprise either S or P at position 228, wherein the mutation S228P may help stabilize the Ab and prevent Fab arm exchange.

[0115] Such chimerism, i.e., humanizing BC8 to produce BC8c, may be achieved by methods known in the art, such as by cloning DNA encoding the BC8 murine heavy and light chain V regions and endogenous murine signal sequences in frame into mammalian expression vectors for the heavy and light chains that already contain human heavy chain constant regions (IgG1, IgG2, or IgG4) or human C kappa.

[0116] Thus, according to certain aspects, the BC8 monoclonal antibody may be chimeric BC8, i.e., BC8c, and may comprise a human IgG1 heavy chain constant region having the amino acid sequence as set forth in SEQ ID NO:17, or a human IgG2 heavy chain constant region having the amino acid sequence as set forth in SEQ ID NO:18, or a human IgG4 heavy chain constant region having the amino acid sequence as set forth in SEQ ID NO:19, or a human IgG4 heavy chain constant region having the amino acid sequence as set forth in SEQ ID NO:20, or a human kappa light chain constant region having the amino acid sequence as set forth in SEQ ID NO:21 (**FIG. 6A-6E**).

[0117] According to certain aspects, the BC8 monoclonal antibody may be chimeric (BC8c) comprising a human IgG1, IgG2, or IgG4 heavy chain constant region having the amino acid sequence as set forth in any one of SEQ ID NOS:17-20, and a human kappa light

chain constant region having the amino acid sequence as set forth in SEQ ID NO:21 (**FIG. 6A-6E**).

[0118] According to certain aspects, the chimeric BC8c monoclonal antibody may comprise a light chain variable region having the amino acid sequence set forth in SEQ ID NO:1 (**FIG. 2**). The BC8c monoclonal antibody may comprise a light chain having the N-terminal amino acid sequence set forth in SEQ ID NO:9 (**FIG. 3**). The BC8c monoclonal antibody may comprise a light chain having at least one complementarity determining region having the amino acid sequence as set forth in SEQ ID NO:3, SEQ ID NO:4, or SEQ ID NO:5 (**FIG. 3**).

[0119] According to certain aspects, the chimeric BC8c monoclonal antibody may comprise a heavy chain variable region having the amino acid sequence set forth in SEQ ID NO:2 (**FIG. 2**). The BC8c monoclonal antibody may comprise a heavy chain having the N-terminal amino acid sequence set forth in SEQ ID NO:10 (**FIG. 3**). The BC8c monoclonal antibody may comprise a heavy chain having at least one complementarity determining region having the amino acid sequence as set forth in SEQ ID NO:6, SEQ ID NO:7, or SEQ ID NO:8 (**FIG. 3**).

[0120] According to certain aspects, the heavy chain of the BC8 or BC8c monoclonal antibody comprises a C-terminal lysine, a C-terminal glycine (G) having lost the C-terminal lysine (K), or is lacking both GK. When referring to antibodies comprising a modified heavy chain constant region described herein, the antibody may comprise a provided sequence having the C-terminal GK or K, or alternatively, lacking GK or K.

[0121] *Patient Specific Composition*

[0122] As used herein, a composition comprising ²²⁵Ac-labelled BC8 includes both actinium-225 labeled antibody and non-labeled antibody, with the minority being the actinium-225 labeled antibody. Likewise, for ¹⁷⁷Lu-labelled BC8, the composition will include both labelled and unlabeled antibody populations. The ratio of labeled to non-labeled antibody can be adjusted using known methods. Thus, accordingly to certain aspects of the present disclosure, the anti-CD45 antibody may be provided in a total protein amount of up to 100mg, such as up to 60 mg, such as 5mg to 45mg, or a total protein amount of from 0.001 mg/kg patient weight to 3.0 mg/kg patient weight, such as from 0.005 mg/kg patient weight to 2.0 mg/kg patient weight, or from 0.01 mg/kg patient weight to 1 mg/kg patient weight, or from 0.1 mg/kg patient weight to 0.6 mg/kg patient weight, or 0.3 mg/kg patient weight, or 0.4 mg/kg patient weight, or 0.5 mg/kg patient weight, or 0.6 mg/kg patient weight.

[0123] According to certain aspects of the present disclosure, the radiolabeled anti-CD45 antibody (i.e., ^{225}Ac -labelled BC8 or ^{177}Lu -labelled BC8) may comprise a labeled fraction and an unlabeled fraction, wherein the ratio of labeled : unlabeled may be from about 0.01:10 to 1:10, such as 0.01:5 to 0.1:5, or 0.01:3 to 0.1:3, or 0.01:1 to 0.1:1 labeled : unlabeled. Moreover, the radiolabeled anti-CD45 antibody may be provided as a single dose composition tailored to a specific patient, wherein the amount of labeled and unlabeled anti-CD45 antibody in the composition may depend on at least a patient weight, age, gender, and/or disease state or health status. See for example administration methods disclosed in International Publication No. WO 2016/187514, incorporated herein by reference herein in its entirety. According to certain aspects, the radiolabeled anti-CD45 antibody may be provided in multiple doses, wherein each dose in the regime may comprise a composition tailored to a specific patient, wherein the amount of labeled and unlabeled anti-CD45 antibody in the composition may depend on at least a patient weight, age, gender, and/or disease state or health status.

[0124] This inventive combination of a labeled fraction and an unlabeled fraction of the anti-CD45 antibody allows the composition to be tailored to a specific patient, wherein each of the radiation dose and the protein dose of the monoclonal antibody are personalized to that patient based on at least one patient specific parameter. As such, each vial of the composition may be made for a specific patient, where the entire content of the vial is delivered to that patient in a single dose. When a treatment regime calls for multiple doses, each dose may be formulated as a patient specific dose in a vial to be administered to the patient as a “single dose” (i.e., full contents of the vial administered at one time). The subsequent dose may be formulated in a similar manner, such that each dose in the regime provides a patient specific dose in a single dose container. One of the advantages of the disclosed composition is that there will be no left-over radiation that would need to be discarded or handled by the medical personnel, e.g., no dilution, or other manipulation to obtain a dose for the patient. When provided in a single dose container, the container is simply placed in-line in an infusion tubing set for infusion to the patient. Moreover, the volume can be standardized so that there is a greatly reduced possibility of medical error (i.e., delivery of an incorrect dose, as the entire volume of the composition is to be administered in one infusion).

[0125] *Treatment of hematological diseases*

[0126] The majority of malignancies of hematologic origin, whether myeloid or lymphoid-derived, express CD45 on the surface of tumor cells to varying degrees. This

includes leukemias (such as acute myeloid leukemia (AML), acute promyelocytic leukemia, acute lymphoblastic leukemia (ALL), acute mixed lineage leukemia, chronic myeloid leukemia, chronic lymphocytic leukemia (CLL), hairy cell leukemia and large granular lymphocytic leukemia), myelodysplastic syndrome (MDS), myeloproliferative disorders (polycythemia vera, essential thrombocytosis, primary myelofibrosis and chronic myeloid leukemia), lymphomas, multiple myeloma, MGUS and similar disorders, Hodgkin's lymphoma, non-Hodgkin lymphoma (NHL), primary mediastinal large B-cell lymphoma, diffuse large B-cell lymphoma, follicular lymphoma, transformed follicular lymphoma, splenic marginal zone lymphoma, lymphocytic lymphoma, T-cell lymphoma, and other B-cell malignancies. As such, an amount of a radiolabeled anti-CD45 antibody, when administered to a patient, will be effective as a direct anti-tumor therapy to reduce tumor blast count in the periphery and in the immune cell compartments, such as bone marrow, spleen and lymph nodes.

[0127] Direct therapy with a radiolabeled anti-CD45 antibody may be as a low-dose single agent necessary to reduce tumor blast count, but reversibly spare hematopoietic stem cells, or in combination with other therapeutic agents such as chemotherapy agents or targeted therapy agents (e.g., but not limited to: HDAC inhibitors, BCL2 inhibitors, monoclonal antibodies, or tyrosine kinase receptor inhibitors – TKIs).

[0128] Doses of an ^{225}Ac radiolabeled anti-CD45 antibody effective at controlling tumor growth and reducing blast count without irreversibly depleting hematopoietic stem cells would deliver a radiation exposure to bone marrow below a threshold level. A dose of 2 Gy is considered to be a non-myeloablative dose of radiation. An ideal dose would deliver a dose of at least 2 Gy but high enough to eliminate leukemic or lymphoma tumor cells and provide transient but reversible myelosuppression. Dose levels above 2 Gy, but less than a myeloablative dose are anticipated to be effective at controlling tumor burden in lymphoma and leukemia. Further, combining single low-dose radiolabeled anti-CD45 antibody treatment with another targeted agent, potent anti-tumor activity may be achieved using lower doses of the antibody radioconjugate further sparing depletion of hematopoietic stem cells.

[0129] An exemplary low dose may be a dose of ^{225}Ac -BC8 that is less than 150 μCi , such as from 10 μCi to 100 μCi , or a dose of less than 2 $\mu\text{Ci}/\text{kg}$, such as from 0.01 $\mu\text{Ci}/\text{kg}$ to 1.5 $\mu\text{Ci}/\text{kg}$ or 0.1 $\mu\text{Ci}/\text{kg}$ to 1.0 $\mu\text{Ci}/\text{kg}$.

[0130] *Depletion of circulating tumor and bone marrow blast cells*

[0131] Hematologic malignancies, including without limitation leukemias such as acute myeloid leukemia, acute lymphocytic leukemia, multiple myeloma, etc., pose a unique

set of problems for effective therapy. If killed too quickly, the high burden of circulating tumor cells often associated with leukemias can be toxic to the patient. Cyto-reductive therapy is the process by which the number of circulating blast cells are reduced.

[0132] According to certain aspects, a cyto-reductive therapy may be used to treat a hematological malignancy, and generally includes administration of a low dose of the radiolabeled anti-CD45, such as a dose that depletes the circulating tumor cells (e.g. leukemia, lymphoma, myeloma, MDS) but is not myeloablative and therefore does not irreversibly deplete HSCs. An exemplary low dose may be a dose of $^{225}\text{Ac-BC8}$ that is less than 150 μCi , such as from 10 μCi to 100 μCi , or a dose of less than 2 $\mu\text{Ci/kg}$, such as from 0.01 $\mu\text{Ci/kg}$ to 1.5 $\mu\text{Ci/kg}$ or 0.1 $\mu\text{Ci/kg}$ to 1.0 $\mu\text{Ci/kg}$.

[0133] ***Hematopoietic stem cell therapy***

[0134] The hematopoietic stem cell therapy includes administration of hematopoietic stem cells, such as a bone marrow transplant (BMT). The hematopoietic stem cells may be administered to a patient defective or deficient in one or more cell types of the hematopoietic lineage in order to re-constitute the defective or deficient population of cells in vivo. For example, the patient may be suffering from cancer or from a hemoglobinopathy (e.g., a non-malignant hemoglobinopathy), such as sickle cell anemia, thalassemia, Fanconi anemia, aplastic anemia, and Wiskott-Aldrich syndrome. The subject may be one that is suffering from adenosine deaminase severe combined immunodeficiency (ADA SCID), HIV/AIDS, metachromatic leukodystrophy, Diamond-Blackfan anemia, and Schwachman-Diamond syndrome. The subject may have or be affected by an inherited blood disorder (e.g., sickle cell anemia) or an autoimmune disorder, such as scleroderma, multiple sclerosis, ulcerative colitis, Crohn's disease, Type 1 diabetes, or another autoimmune pathology.

[0135] The cancer may be a neuroblastoma or a hematologic cancer. For instance, the subject may have a leukemia, lymphoma, or myeloma. In some embodiments, the subject has acute myeloid leukemia, acute lymphoid leukemia, chronic myeloid leukemia, chronic lymphoid leukemia, multiple myeloma, diffuse large B-cell lymphoma, or non-Hodgkin's lymphoma. The subject may have myelodysplastic syndrome (MDS).

[0136] ***Gene editing***

[0137] Gene editing technologies have advanced substantially with the advent of site-specific editing methods such as TALEN, CRISPR/cas9 and zinc finger nuclease (ZFN) methods. These methods have therapeutic potential for patients afflicted with malignant and non-malignant hereditary diseases.

[0138] Gene editing precisely and permanently alters a sequence of genomic DNA that remains under endogenous genetic regulation and control for proper and appropriate expression of the modified genetic element. There are presently four major classes of nucleases for human genome gene editing: zinc finger nucleases (ZFNs); transcription activator-like effector nucleases (TALENs); meganucleases (MNs); and clustered regularly interspaced short palindromic repeats (CRISPR/Cas9). Each of these can recognize and bind a specific target sequence of DNA. Depending on the approach, the target DNA can be cleaved on one or both strands. To correct a mutation, a correction template is used for homology-directed repair of the introduced break at the site of the targeted lesion. This technology can also be exploited to silence or ablate a particular gene by incorporating a mutational insertion or deletion. Further, gene-editing technology can also be utilized to functionally replace one gene with another, such as within the T-cell receptor alpha constant locus (*TRAC*), and thereby change the specificity of the T-cells (Eyquem, et. al., 2017, Nature. 543:113-117).

[0139] *Adoptive cell therapy (“ACT”)*

[0140] The adoptive cell therapy may include administration of cells expressing a chimeric antigen receptor (CAR), or a T-cell receptor (TCR), or may include tumor-infiltrating lymphocytes (TIL). The population of cells expressing the CAR/TCR may comprise a population of activated T-cells or natural killer (NK) cells or dendritic cells expressing the CAR/TCR which recognize an antigen. Dendritic cells are capable of antigen presentation, as well as direct killing of tumors. The population of cells expressing the CAR/TCR may comprise a population of gene-edited cells.

[0141] As used herein, the term “gene-edited” CAR T-cell is synonymous with the terms “genetically engineered” CAR T-cell and “engineered” CAR T-cell. A gene-edited CAR T-cell that “fails to properly express” a checkpoint receptor (e.g., PD1, Lag3 or TIM3) does not express the full-length, functional checkpoint receptor. For example, a gene-edited CAR T-cell that fails to properly express PD1 may fail to do so because, without limitation, (i) the cell’s PD1 gene has been ablated, or (ii) the cell’s PD1 gene has been otherwise altered so as not to yield a fully or even partially functional PD1 product. In other words, according to certain aspects, a gene-edited CAR T-cell that fails to properly express PD1 may fail to do so because the cell’s PD1 gene has been altered to diminish PD1 expression. Similarly, a gene-edited CAR T-cell that “fails to properly express” a T-cell receptor does not express the full-length, functional T-cell receptor.

[0142] According to certain aspects, the functional endogenous T-cell receptor is replaced through editing by a “knock-in” to the native TCR locus of an exogenously transduced CAR or recombinant TCR. The gene-edited CAR T-cells may include, without limitation, the following: (i) allogenic gene-edited CAR T-cells that fail to properly express PD1 but do properly express all other checkpoint receptors and T-cell receptors; (ii) allogenic gene-edited CAR T-cells that fail to properly express a particular T-cell receptor but do properly express all checkpoint receptors and all other T-cell receptors; and (iii) allogenic gene-edited CAR T-cells that fail to properly express PD1 and fail to properly express a particular T-cell receptor, but do properly express all other checkpoint receptors and all other T-cell receptors.

[0143] Examples of T-cell gene editing to generate allogeneic, universal CAR T-cells include the work of Eyquem and colleagues (Eyquem, et. al., 2017, Nature. 543:113-117). In that study, the endogenous T-cell receptor alpha constant locus (*TRAC*) was effectively replaced by a recombinant CAR gene construct. By this method, the recombinant CAR was placed effectively under the control of the cell’s native TCR regulatory signals. By this same strategy, CARs or recombinant TCRs may be effectively inserted by knock-in into the T-cell receptor beta constant gene locus (*TRBC*) or into the beta-2 microglobulin (*B2M*) MHC-I-related gene locus, known to be expressed in all T-cells. Another example includes the work of Ren and colleagues (Ren, et. al., 2017, Clin. Cancer Res 23:2255-2266). Recognizing that checkpoint receptors are immune-suppressive and may blunt the stimulation of exogenous autologous or allogeneic CAR T-cells, this group exploited CRISPR/cas9 technology to ablate the endogenous TCR α and β loci (*TRAC* and *TRBC*) and the *B2M* gene, while also silencing the endogenous *PD1* gene. With this approach, the engineered cells did not elicit graft-versus-host disease but did resist immune checkpoint receptor suppression.

[0144] *Lymphodepletion and myeloablation*

[0145] Before administering a dose of HSTs (e.g., bone marrow transplant) or engineered immune cells to a patient, it is common to lymphodeplete the patient. The lymphodepletion process is considered important, indeed essential, to the success of BMT and adoptive cell therapy (ACT) methods. The process creates sufficient space in the immune microenvironment (e.g., bone marrow) to allow the transferred cells to engraft. It also creates a favorable immune homeostatic environment for the successful engraftment, proliferation, and persistence of the transferred cells by eliciting a favorable cytokine profile. It elicits this cytokine profile particularly in the peripheral immune niches (e.g., bone marrow, spleen and

lymph nodes) for the establishment and proliferation of the engineered cells. (*see, e.g.,* Maine, et al., 2002, *J. Clin. Invest.*, 110:157-159; Muranski, et al., 2006, *Nat. Clin. Pract. Oncol.*, 3(12):668-681; Klebanoff, et al., 2005, *Trends Immunol.*, 26(2): 111-117).

[0146] As indicated hereinabove, myeloid and lymphoid-derived cell express CD45. Doses of an ²²⁵Ac radiolabeled anti-CD45 antibody effective at reducing blast count without irreversibly depleting hematopoietic stem cells would deliver a radiation exposure to bone marrow below a threshold level. A dose of 2 Gy is considered to be a non-myeloablative dose of radiation. A dose of at least 2 Gy may provide transient but reversible myelosuppression. Myeloablative doses are variable, such as in the range of 8-18 Gy, but typically include doses delivering more than 10-12 Gy to the bone marrow. Thus, a therapeutic low-dose range that may provide reversible immunosuppression (with or without stem cell support) would be 2 Gy or greater, but below a myeloablative dose, such as at or below 8 Gy. For higher doses, stem cell support may be necessary.

[0147] As used herein, an amount of a radiolabeled anti-CD45 antibody, when administered, is “effective” to deplete a specific targeted cell type if the cell population is reduced by at least 10%, 20%, 30%, 40%, 50%, 60%, 70%, 80%, 90% or 95%. For example, an amount of radiolabeled anti-CD45 antibody, when administered, is “effective” to deplete the subject’s peripheral blood lymphocytes if the peripheral blood lymphocytes are depleted without depletion of the subject’s neutrophils, or with less than 10% or 20% reduction in the subject’s neutrophils. An “effective” amount of radiolabeled anti-CD45 antibody can also be related to an amount that will deplete the subject’s regulatory T cells, myeloid derived suppressor cells, tumor associated macrophages, activated macrophages secreting IL-1 and/or IL-6, and combinations thereof, such as by at least 20%, or 30%, 40%, 50%, 60%, 70%, 80%, 90%, or 95%.

[0148] As used herein, an amount of a radiolabeled anti-CD45 antibody, when administered, is “effective” to reversibly suppress a targeted cell type if the cell population, such as the subject’s HSC level or lymphocyte level, is reduced by greater than 95%, such as by at least 96%, or 97%, or 98%, or even 99%. “Reversible immunosuppression” generally comprises use of a low dose therapeutic or combination thereof, such as the actinium-225 labelled BC8 disclosed herein, to deplete the targeted cells to a greater extent than standard lymphodepletion *without ablating* the target cells, i.e., at a dose that is less than a myeloablation dose (non-myeloablative dose). Moreover, reversible immunosuppression may indicate that the targeted immune population (immune privileged cell population or tissues) is only transiently depleted, while other non-targeted populations are not affected.

[0149] As disclosed hereinbelow, reversible immunosuppression of the present disclosure generally may comprise administration of the actinium-225 labelled BC8 and another agent such as, for example, another immunotherapeutic agent or a radio-sensitizing agent.

[0150] As used herein, an amount of a radiolabeled anti-CD45 antibody, when administered, is “effective” to ablate a targeted cell type if the cell population, such as the subject’s HSC level or lymphocyte level, is reduced by 100% (also referred to as myeloablation).

[0151] According to certain aspects of the present disclosure, the radiolabeled anti-CD45 antibody is actinium-225 labelled BC8 (²²⁵Ac-labelled BC8), and the effective amount of ²²⁵Ac-labelled BC8 is below, for example, 5.0 μCi/kg (i.e., where the amount of ²²⁵Ac-BC8 administered to the subject delivers a radiation dose of below 5.0 μCi per kilogram of subject’s body weight).

[0152] According to aspects of the present disclosure, the effective amount of the ²²⁵Ac-labelled BC8 is below 4.5 μCi/kg, 4.0 μCi/kg, 3.5 μCi/kg, 3.0 μCi/kg, 2.5 μCi/kg, 2.0 μCi/kg, 1.5 μCi/kg, 1.0 μCi/kg, 0.9 μCi/kg, 0.8 μCi/kg, 0.7 μCi/kg, 0.6 μCi/kg, 0.5 μCi/kg, 0.4 μCi/kg, 0.3 μCi/kg, 0.2 μCi/kg, 0.1 μCi/kg, 0.05 μCi/kg, or 0.01 μCi/kg. According to certain aspects, the effective amount of the ²²⁵Ac-labelled BC8 is at least 0.01 μCi/kg, or 0.05 μCi/kg, 0.1 μCi/kg, 0.2 μCi/kg, 0.3 μCi/kg, 0.4 μCi/kg, 0.5 μCi/kg, 0.6 μCi/kg, 0.7 μCi/kg, 0.8 μCi/kg, 0.9 μCi/kg, 1 μCi/kg, 1.5 μCi/kg, 2 μCi/kg, 2.5 μCi/kg, 3 μCi/kg, 3.5 μCi/kg, 4 μCi/kg, or 4.5 μCi/kg. According to certain aspects, the ²²⁵Ac-labeled BC8 may be administered at a dose that includes any combination of upper and lower limits as described herein, such as from at least 0.1 μCi/kg to below 5 μCi/kg, or from at least 0.5 μCi/kg to below 3 μCi/kg.

[0153] According to certain aspects, the effective amount of the ²²⁵Ac-labelled BC8 is below 1.0 mCi, such as below 0.5 mCi (i.e., wherein the ²²⁵Ac is administered to the subject in a non-weight-based dosage). According to certain aspects, the effective dose of the ²²⁵Ac-labelled BC8 may be below 1.0 mCi, such as below 0.9 mCi, 0.8 mCi, 0.7 mCi, 0.6 mCi, 0.5 mCi, 0.45 mCi, 0.4 mCi, 0.35 mCi, 0.3 mCi, 0.25 mCi, 0.2 mCi, 0.1 mCi, 90 μCi, 80 μCi, 70 μCi, 60 μCi, 50 μCi, 40 μCi, 30 μCi, 20 μCi, 10 μCi, or 5 μCi. The effective amount of ²²⁵Ac-labelled BC8 may be at least 2 μCi, such as at least 5 μCi, 10 μCi, 20 μCi, 30 μCi, 40 μCi, 50 μCi, 60 μCi, 70 μCi, 80 μCi, 90 μCi, 100 μCi, 120 μCi, 140 μCi, 160 μCi, 180 μCi, 200 μCi, 300 μCi, 400 μCi, 500 μCi, 600 μCi, 700 μCi, 800 μCi, or 900 μCi. According to certain aspects, the ²²⁵Ac-labelled BC8 may be administered at a dose that

includes any combination of upper and lower limits as described herein, such as from at least 15 μCi to below 120 μCi , or from at least 20 μCi to below 100 μCi , or from 80 μCi to below 500 μCi . According to certain aspects, the ^{225}Ac -labelled BC8 may be administered at a low dose. An exemplary low dose may be a dose of ^{225}Ac -BC8 that is less than 150 μCi , such as from 10 μCi to 100 μCi , or a dose of less than 2 $\mu\text{Ci}/\text{kg}$, such as from 0.01 $\mu\text{Ci}/\text{kg}$ to 1.5 $\mu\text{Ci}/\text{kg}$ or 0.1 $\mu\text{Ci}/\text{kg}$ to 1.0 $\mu\text{Ci}/\text{kg}$.

[0154] According to certain aspects of the present disclosure, the radiolabeled anti-CD45 antibody is lutetium-177 labelled BC8 (^{177}Lu -labelled BC8), and the effective amount of ^{177}Lu -labelled BC8 is below, for example, 500 $\mu\text{Ci}/\text{kg}$ (i.e., where the amount of ^{177}Lu -BC8 administered to the subject delivers a radiation dose of below 500 μCi per kilogram of subject's body weight).

[0155] According to aspects of the present disclosure, the effective amount of the ^{177}Lu -labelled BC8 is below 450 $\mu\text{Ci}/\text{kg}$, 400 $\mu\text{Ci}/\text{kg}$, 350 $\mu\text{Ci}/\text{kg}$, 300 $\mu\text{Ci}/\text{kg}$, 250 $\mu\text{Ci}/\text{kg}$, 200 $\mu\text{Ci}/\text{kg}$, 150 $\mu\text{Ci}/\text{kg}$, 100 $\mu\text{Ci}/\text{kg}$, 90 $\mu\text{Ci}/\text{kg}$, 80 $\mu\text{Ci}/\text{kg}$, 70 $\mu\text{Ci}/\text{kg}$, 60 $\mu\text{Ci}/\text{kg}$, 50 $\mu\text{Ci}/\text{kg}$, 40 $\mu\text{Ci}/\text{kg}$, 30 $\mu\text{Ci}/\text{kg}$, 20 $\mu\text{Ci}/\text{kg}$, 10 $\mu\text{Ci}/\text{kg}$, 5 $\mu\text{Ci}/\text{kg}$, or 1 $\mu\text{Ci}/\text{kg}$. According to certain aspects, the effective amount of the ^{177}Lu -labelled BC8 is at least 1 $\mu\text{Ci}/\text{kg}$, 2.5 $\mu\text{Ci}/\text{kg}$, 5 $\mu\text{Ci}/\text{kg}$, 10 $\mu\text{Ci}/\text{kg}$, 20 $\mu\text{Ci}/\text{kg}$, 30 $\mu\text{Ci}/\text{kg}$, 40 $\mu\text{Ci}/\text{kg}$, 50 $\mu\text{Ci}/\text{kg}$, 60 $\mu\text{Ci}/\text{kg}$, 70 $\mu\text{Ci}/\text{kg}$, 80 $\mu\text{Ci}/\text{kg}$, 90 $\mu\text{Ci}/\text{kg}$, 100 $\mu\text{Ci}/\text{kg}$, 150 $\mu\text{Ci}/\text{kg}$, 200 $\mu\text{Ci}/\text{kg}$, 250 $\mu\text{Ci}/\text{kg}$, 300 $\mu\text{Ci}/\text{kg}$, 350 $\mu\text{Ci}/\text{kg}$, 400 $\mu\text{Ci}/\text{kg}$ or 450 $\mu\text{Ci}/\text{kg}$. According to certain aspects, an ^{177}Lu -labeled BC8 may be administered at a dose that includes any combination of upper and lower limits as described herein, such as from at least 5 $\mu\text{Ci}/\text{kg}$ to below 50 $\mu\text{Ci}/\text{kg}$, or from at least 50 $\mu\text{Ci}/\text{kg}$ to below 500 $\mu\text{Ci}/\text{kg}$.

[0156] According to certain aspects, the effective amount of the ^{177}Lu -labelled BC8 is below 20 mCi, such as below 15 mCi, 10 mCi, 9 mCi, 8 mCi, 7 mCi, 6 mCi, 5 mCi, 3 mCi, 2 mCi, 1 mCi, 800 μCi , 600 μCi , 400 μCi , 200 μCi , 100 μCi , or 50 μCi . The effective amount of ^{177}Lu -labeled BC8 may be at least 10 μCi , such as at least 25 μCi , 50 μCi , 100 μCi , 200 μCi , 300 μCi , 400 μCi , 500 μCi , 600 μCi , 700 μCi , 800 μCi , 900 μCi , 1 mCi, 2 mCi, 3 mCi, 4 mCi, 5 mCi, 10 mCi, or 15 mCi. According to certain aspects, an ^{177}Lu -labeled BC8 may be administered at a dose that includes any combination of upper and lower limits as described herein, such as from at least 10 μCi to below 20 mCi, or from at least 100 μCi to below 3 mCi, or from 3 mCi to below 20 mCi.

[0157] As used herein, a "suitable time period" after administering a radiolabeled anti-CD45 antibody to a subject and before performing an additional therapy on the subject is

a time period sufficient to permit the administered antibody to deplete, reversibly suppress, or ablate the targeted cells of the subject, such as the subject's HSCs and/or lymphocytes. According to certain aspects, the suitable time period is fewer than 15 days, fewer than 14 days, fewer than 13 days, fewer than 12 days, fewer than 11 days, fewer than 10 days, fewer than 9 days, fewer than 8 days, fewer than 7 days, fewer than 6 days, fewer than 5 days, fewer than 4 days, or fewer than 3 days. According to certain aspects, the suitable time period is 1 day, 2 days, 3 days, 4 days, 5 days, 6 days, 7 days, 8 days, 9 days, 10 days, 11 days, 12 days, 13 days, 14 days, 15 days, or more than 15 days.

[0158] According to certain aspects, the suitable time period after administering the radiolabeled anti-CD45 antibody that an ACT procedure may be performed is 3 days, 4 days, 5 days, 6 days, 7 days, 8 days or 9 days, such as preferably 6, 7 or 8 days.

[0159] Throughout this application, various publications are cited. The disclosure of these publications is hereby incorporated by reference into this application to describe more fully the state of the art to which this disclosure pertains. Unless otherwise defined, all technical and scientific terms used herein have the same meaning as commonly understood by one of ordinary skill in the art to which the present disclosure belongs. Although methods and materials similar or equivalent to those described herein can be used in the practice or testing described herein, suitable methods and materials are described below.

DETAILED DESCRIPTION OF THE INVENTION

[0160] This disclosure solves an unmet need in the art by providing an unexpectedly superior way to deplete, reversibly immunosuppress, or ablate specific cells in a subject, such as the subject's hematopoietic stem cells or lymphocytes. Reversible immunosuppression of these cells may be useful in the treatment of a CD45 positive hematological malignancy, and may be achieved using low-dose therapeutics, such as administration of an anti-CD45 antibody at sub-saturating radiation doses.

[0161] Radiolabeled anti-CD45 antibodies have shown clinical potential for targeted myeloablative conditioning prior to bone marrow transplant. CD45 is an attractive target for conditioning as it is highly expressed in all nucleated immune cells including hematopoietic stem cells, lymphoid and myeloid cells. The potent alpha-emitter 225-Actinium (²²⁵Ac) is a promising radionuclide for targeted conditioning, with high linear energy transfer (80-100 keV/ μ m) over a short path length, and a long 9.9-day half-life.

[0162] Moreover, depletion or ablation of these cells using a high dose therapeutic, i.e., higher radiation doses, may be a precursor to a cell-based therapy like a bone marrow

transplant and/or an adoptive cell therapy (e.g., chimeric antigen receptor therapy, CAR T-cell therapy or TCR-cell therapy) or a gene-edited cell-based therapy (e.g., genetically edited β -globin hematopoietic stem cell therapy for sickle cell disease, SCD).

[0163] Accordingly, this disclosure employs a radiolabeled anti-CD45 antibody such as ^{225}Ac -BC8 to deplete, reversibly immunosuppress, or ablate specific cells in a subject. The antibody can safely and effectively deplete or ablate the subject's hematopoietic stem cells or lymphocytes via targeted conditioning. This approach avoids certain adverse effects caused by less specific agents like chemotherapeutics or external beam radiation.

[0164] The present disclosure provides methods of treating a variety of disorders, such as diseases of a cell type in the hematopoietic lineage, cancers, autoimmune diseases, metabolic disorders, and stem cell disorders, among others. The compositions and methods described herein may (i) directly deplete or ablate a population of cells that give rise to a pathology, such as a population of cancer cells (e.g., leukemia cells) and autoimmune cells (e.g., autoreactive T-cells), and/or (ii) deplete or ablate a population of endogenous hematopoietic stem cells so as to promote the engraftment of transplanted hematopoietic stem cells by providing a niche to which the transplanted cells may home.

[0165] The foregoing activities can be achieved by administration of a composition comprising ^{225}Ac -BC8 or ^{177}Lu -BC8. In the case of direct treatment of a disease, this administration can cause a reduction in the quantity of the cells that give rise to the pathology of interest. In the case of preparing a patient for hematopoietic stem cell transplant therapy, this administration can cause the selective depletion, reversible suppression, or ablation of a population of endogenous hematopoietic stem cells, thereby creating a vacancy in the hematopoietic tissue, such as the bone marrow or lymphocytes, that can subsequently be filled by transplanted, exogenous hematopoietic stem cells, i.e. bone marrow transplant or adoptive cell transfer.

[0166] **Radiolabeled Immunotherapeutic**

[0167] According to aspects of the present disclosure, the anti-CD45 immunoglobulin BC8 may comprise ^{225}Ac or ^{177}Lu . According to certain preferred aspects, the anti-CD45 immunoglobulin BC8 may be radiolabeled with the alpha-emitting radionuclide Actinium-225 (^{225}Ac). The ^{225}Ac payload conjugated to the monoclonal antibody BC8 delivers high energy alpha particles directly to the targeted cell(s), generating lethal double strand DNA breaks. Due to its short path length, the range of its high energy alpha particle emission is only a few cell diameters thick, thereby limiting damage to nearby non-malignant or normal

tissues. As such, ^{225}Ac -BC8 may provide a therapeutically effective dose at lower radiation amounts than ^{177}Lu -BC8.

[0168] Furthermore, ^{225}Ac -antibody conjugates offer a crucial advantage over antibody-drug conjugates of the prior art as they are found to be effective even in patients with low target antigen expressing tumors. This is because of the large cytotoxic effects of the ^{225}Ac , which is in striking contrast to antibody-drug conjugates where hundreds of antibody molecules are needed to bind to their respective antigens to exert an effect on a targeted cell or tissue.

[0169] Other advantages of the radioactive payload over drugs or toxins include: 1) the antibody that delivers the radiation does not need to be internalized to kill the cell; 2) not every cell in the targeted tissue or tumor needs to be targeted by the antibody; and 3) In contrast to antibody-drug conjugates, the radioisotope linked to the antibody is unlikely to elicit significant immune responses that would limit subsequent use. Moreover, studies reported herein demonstrate the stability of the ^{225}Ac labelled antibodies, and their highly targeted cytotoxicity.

[0170] According to certain aspects of the present disclosure, the ^{225}Ac may be attached or chelated by a chelating agent that is conjugated to the monoclonal antibody. As detailed in Example 3 below, the anti-CD45-immunoglobulin may be prepared by first forming a chelator conjugated anti-CD45 (“conjugated anti-CD45”), and then chelating the radionuclide with the conjugated anti-CD45 to form the radiolabeled anti-CD45 (i.e., ^{225}Ac -BC8).

[0171] According to the methods of forming the radiolabeled anti-CD45 described herein, the monoclonal antibody against CD45 may be dissolved in a buffered solution comprising a chelant. The pH may be selected to optimize conditions for conjugation of the chelant with the antibody in a conjugation reaction mixture. The conjugation reaction mixture may comprise a bicarbonate buffer or a phosphate buffer. The conjugation reaction mixture may have a pH of about 8.0 to about 9.2. For example, the conjugation reaction mixture may have a pH of about 8.0, about 8.1, about 8.3, about 8.4, about 8.5, about 8.6, about 8.7, about 8.8, about 8.9, about 9.0, about 9.1, or about 9.2. The temperature of the conjugation reaction mixture may be adjusted to promote conjugation of the chelant with the targeting moiety. For example, the conjugation reaction mixture can be incubated at a temperature of about room temperature, or about 37°C . The conjugation reaction mixture may be incubated for any amount of time sufficient to provide conjugation such as, for example, about 1.5 hours.

[0172] The conjugated anti-CD45 may be dissolved in a buffered solution comprising a radionuclide. The pH may be selected to optimize conditions for chelation of the radionuclide with the conjugated anti-CD45 in a chelation reaction mixture. The chelation reaction mixture may have a pH of about 5.5 to about 7.0. For example, the chelation reaction mixture may have a pH of about 5.5, about 5.6, about 5.7, about 5.8, about 5.9, about 6.0, about 6.1, about 6.2, about 6.3, about 6.4, about 6.5, about 6.6, about 6.7, about 6.8, about 6.9 or about 7.0.

[0173] The temperature of the chelation reaction mixture may be adjusted to promote chelation of the radionuclide with the conjugated anti-CD45-immunoglobulin. For example, the chelation reaction mixture may be incubated at a temperature of about 37°C. The chelation reaction mixture may be incubated for about 1.5 hours. After a period of time, the solution may be quenched by the addition of a quenching chelate (e.g. diethylenetriaminepentaacetic acid (DTPA)) and the reaction mixture may be purified. The chelation reaction mixture may be further incubated after addition of the quenching chelate, such as for about 30 minutes at about 37°C after addition of the quenching chelate.

[0174] The chelators useful in the present disclosure are compounds which have the dual functionality of sequestering metal ions plus the ability to covalently bind a biological carrier such as an antibody. Numerous chelators are known in the art. Exemplary chelators suitable for use in the present disclosure include, but are not limited to chelators such as S-2-(4-Isothiocyanatobenzyl)-1,4,7,10 tetraazacyclododecanetetraacetic acid (p-SCN-Bn-DOTA), diethylene triamine pentaacetic acid (DTPA); ethylene diamine tetraacetic acid (EDTA); 1,4,7,10-tetraazacyclododecane-N,N',N'',N'''-tetraacetic acid (DOTA); p-isothiocyanatobenzyl-1,4,7,10-tetraazacyclododecane-1,4,7,10-tetraacetic acid (p-SCN-Bz-DOTA); 1,4,7,10-tetraazacyclododecane-N,N',N''-triacetic acid (DO3A); 1,4,7,10-tetraazacyclododecane-1,4,7,10-tetrakis(2-propionic acid) (DOTMA); 3,6,9-triaza-12-oxa-3,6,9-tricarboxymethylene-10-carboxy-13-phenyl-tridecanoic acid ("B-19036"); 1,4,7-triazacyclononane-N,N',N''-triacetic acid (NOTA); 1,4,8,11-tetraazacyclotetradecane-N,N',N'',N'''-tetraacetic acid (TETA); triethylene tetraamine hexaacetic acid (TTHA); trans-1,2-diaminohexane tetraacetic acid (CYDTA); 1,4,7,10-tetraazacyclododecane-1-(2-hydroxypropyl)-4,7,10-triacetic acid (HP-DO3A); trans-cyclohexane-diamine tetraacetic acid (CDTA); trans(1,2)-cyclohexane diethylene triamine pentaacetic acid (CDTPA); 1-oxa-4,7,10-triazacyclododecane-N,N',N''-triacetic acid (OTTA); 1,4,7,10-tetraazacyclododecane-1,4,7,10-tetrakis{3-(4-carboxyl)-butanoic acid}; 1,4,7,10-tetraazacyclododecane-1,4,7,10-

tetrakis(acetic acid-methyl amide); 1,4,7,10-tetraazacyclo-dodecane-1,4,7,10-tetrakis(methylene phosphonic acid); and derivatives thereof.

[0175] One or more steps may be used to separate the conjugated CD45 from other constituents of the conjugation reaction mixture or the radiolabeled anti-CD45 from other constituents of the chelation reaction mixture. For example, the reaction mixture can be transferred to a filtering device (*e.g.*, a Millipore centrifugal device) having a particular molecular weight cut off such that filtration of the reaction mixture through the filtration device can separate the conjugated anti-CD45 or the radiolabeled anti-CD45 from other constituents of the respective reaction mixture. Filtration can be used to obtain a conjugated anti-CD45 or the radiolabeled anti-CD45 having at least about 60%, at least about 65%, at least about 70%, at least about 75%, at least about 80%, at least about 85%, at least about 90%, %, at least about 95%, at least about 97%, at least about 98%, at least about 99%, or at least about 99.5% purity.

[0176] According to certain aspects of the present disclosure, the yield of the conjugated anti-CD45-immunoglobulin or the radiolabeled anti-CD45-immunoglobulin from the separation (*e.g.*, purification) is at least about 70%, at least about 75%, at least about 80%, at least about 85%, or at least about 90%, at least 95%, at least 96%, at least 97%, at least 98%, or at least 99% of the final product.

[0177] According to one aspect of the present disclosure, the monoclonal antibody may be first conjugated with a p-SCN-Bn-DOTA or DOTA chelating agent to form the conjugated anti-CD45-immunoglobulin, followed by chelation of ²²⁵Ac by the p-SCN-Bn-DOTA or DOTA on the conjugated anti-CD45-immunoglobulin to form the radiolabeled anti-CD45-immunoglobulin. Thus, according to aspects of the present disclosure, only a single step involving ²²⁵Ac is needed to label the anti-CD45-immunoglobulin.

[0178] According to certain aspects of the present disclosure, the radiolabeled anti-CD45-immunoglobulin ²²⁵Ac-BC8 is relatively stable. For example, greater than 85% of the ²²⁵Ac-BC8 may remain intact after storage for 24 hours at 4°C (*see* **FIG. 11**). Moreover, the ²²⁵Ac-BC8 shows specificity toward CD45 expressing cells (*see* **FIG. 12, 15A, 15B**) and tissues (*see* **FIG. 16A**). The labeling efficiency, stability, and immunoreactivity of the ²²⁵Ac-BC8 combine to provide an effective therapeutic agent.

[0179] Radiolabeling with lutetium-177 (¹⁷⁷Lu) to provide ¹⁷⁷Lu -DOTA-BC8 is also possible and within the scope of the present disclosure. Moreover, reference to ²²⁵Ac-BC8 or ¹⁷⁷Lu-BC8 may include reference to either of ²²⁵Ac-DOTA-BC8 or ¹⁷⁷Lu -DOTA-BC8, respectively, unless specifically indicated otherwise.

[0180] Methods for depleting or ablating targeted cells

[0181] This disclosure provides a method for depleting, reversibly suppressing, or ablating a subject's hematopoietic stem cells comprising administering to the subject an effective amount of a radiolabeled anti-CD45-immunoglobulin such as $^{225}\text{Ac-BC8}$. This disclosure provides a method for depleting a subject's lymphocytes comprising administering to the subject an effective amount of a radiolabeled anti-CD45-immunoglobulin such as $^{225}\text{Ac-BC8}$. This disclosure provides a method for depleting, reducing or eliminating a subject's hematopoietic cancer blasts comprising administering to the subject an effective amount of a radiolabeled anti-CD45-immunoglobulin such as $^{225}\text{Ac-BC8}$ alone as a single agent therapy or in combination with other therapies.

[0182] According to certain aspects of this method, the effective amount of $^{225}\text{Ac-BC8}$ is from 0.05 $\mu\text{Ci/kg}$ to 5.0 $\mu\text{Ci/kg}$ of subject's body weight. Examples of effective amounts include, without limitation, from 0.05 $\mu\text{Ci/kg}$ to 5.0 $\mu\text{Ci/kg}$, such as from 0.1 $\mu\text{Ci/kg}$ to 0.2 $\mu\text{Ci/kg}$, from 0.2 $\mu\text{Ci/kg}$ to 0.3 $\mu\text{Ci/kg}$, from 0.3 $\mu\text{Ci/kg}$ to 0.4 $\mu\text{Ci/kg}$, from 0.4 $\mu\text{Ci/kg}$ to 0.5 $\mu\text{Ci/kg}$, from 0.5 $\mu\text{Ci/kg}$ to 0.6 $\mu\text{Ci/kg}$, from 0.6 $\mu\text{Ci/kg}$ to 0.7 $\mu\text{Ci/kg}$, from 0.7 $\mu\text{Ci/kg}$ to 0.8 $\mu\text{Ci/kg}$, from 0.8 $\mu\text{Ci/kg}$ to 0.9 $\mu\text{Ci/kg}$, from 0.9 $\mu\text{Ci/kg}$ to 1.0 $\mu\text{Ci/kg}$, from 1.0 $\mu\text{Ci/kg}$ to 1.5 $\mu\text{Ci/kg}$, from 1.5 $\mu\text{Ci/kg}$ to 2.0 $\mu\text{Ci/kg}$, from 2.0 $\mu\text{Ci/kg}$ to 2.5 $\mu\text{Ci/kg}$, from 2.5 $\mu\text{Ci/kg}$ to 3.0 $\mu\text{Ci/kg}$, from 3.0 $\mu\text{Ci/kg}$ to 3.5 $\mu\text{Ci/kg}$, from 3.5 $\mu\text{Ci/kg}$ to 4.0 $\mu\text{Ci/kg}$, from 4.0 $\mu\text{Ci/kg}$ to 4.5 $\mu\text{Ci/kg}$, or from 4.5 $\mu\text{Ci/kg}$ to 5.0 $\mu\text{Ci/kg}$.

[0183] According to certain aspects of this method, the effective amount of $^{225}\text{Ac-BC8}$ is less than 1 mCi, such as less than 500 μCi . Examples of effective amounts include, without limitation, from 1 μCi to 500 μCi , such as from 10 μCi to 400 μCi , or 10 μCi to 300 μCi , 10 μCi to 200 μCi , 10 μCi to 100 μCi , 15 μCi to 75 μCi , 20 μCi to 75 μCi , 10 μCi to 50 μCi , 50 μCi to 100 μCi , 100 μCi to 150 μCi , 150 μCi to 200 μCi , 200 μCi to 250 μCi , 250 μCi to 300 μCi , 300 μCi to 350 μCi , 350 μCi to 400 μCi , 400 μCi to 450 μCi , or 450 μCi to 500 μCi .

[0184] An exemplary low dose may be a low dose of $^{225}\text{Ac-BC8}$ is less than 120 μCi , such as from 10 μCi to 100 μCi , or a dose of less than 2 $\mu\text{Ci/kg}$, such as from 0.01 $\mu\text{Ci/kg}$ to 1.5 $\mu\text{Ci/kg}$ or 0.1 $\mu\text{Ci/kg}$ to 1.0 $\mu\text{Ci/kg}$.

[0185] An exemplary high dose of $^{225}\text{Ac-BC8}$ may be a moderate to high dose of at least 120 μCi , such as from 120 μCi to 500 μCi , or a dose of at least 2 $\mu\text{Ci/kg}$, such as from 2 $\mu\text{Ci/kg}$ to 5 $\mu\text{Ci/kg}$, or 3 $\mu\text{Ci/kg}$ to 5 $\mu\text{Ci/kg}$.

[0186] According to certain aspects of this method, the effective amount of $^{225}\text{Ac-BC8}$ is an amount effective to deplete a subject's hematopoietic stem cells or lymphocytes,

such as an amount effective to deplete at least 25% of hematopoietic stem cells of the subject, or at least 50% of hematopoietic stem cells of the subject, or at least 70% of hematopoietic stem cells of the subject, or at least 80% of hematopoietic stem cells of the subject, or up to 90% of hematopoietic stem cells of the subject.

[0187] According to certain aspects of this method, the effective amount of $^{225}\text{Ac-BC8}$ is an amount effective to reversibly immunosuppress a subject's hematopoietic stem cells or lymphocytes, such as at least 90% of hematopoietic stem cells of the subject, or at least 92%, or at least 94%, or at least 96%, or up to 98% of hematopoietic stem cells of the subject without completely ablating the hematopoietic stem cells or lymphocytes of the subject.

[0188] According to certain aspects of this method, the effective amount of $^{225}\text{Ac-BC8}$ is an amount effective to deplete a subject's circulating tumor cells, such as hematopoietic stem cells or lymphocytes, such as an amount effective to deplete at least 25% of the hematopoietic stem cells or lymphocytes of the subject, or at least 50% of hematopoietic stem cells or lymphocytes of the subject, or at least 70% of hematopoietic stem cells or lymphocytes of the subject, or at least 80% of hematopoietic stem cells or lymphocytes of the subject, or up to 90% of hematopoietic stem cells or lymphocytes of the subject, without myeloablation. According to certain aspects, this amount may be a low dose of the $^{225}\text{Ac-BC8}$, such as less than 150 μCi or 120 μCi , such as from 10 μCi to 100 μCi , or a dose of less than 2 $\mu\text{Ci/kg}$, such as from 0.01 $\mu\text{Ci/kg}$ to 1.5 $\mu\text{Ci/kg}$ or 0.1 $\mu\text{Ci/kg}$ to 1.0 $\mu\text{Ci/kg}$. this dose may be administered alone or with further additional therapeutics, as disclosed hereinbelow.

[0189] According to certain aspects of this method, the effective amount of $^{225}\text{Ac-BC8}$ is an amount effective to deplete, reduce or eliminate a patient's hematopoietic cancer blasts by 25%, 50%, or 100% as a single agent or in combination with other therapies. A dose that is effective at depleting, reducing or eliminating a patient's cancer blasts may require stem cell support depending on the dose administered.

[0190] Stem cell support may be offered to the patient when such cells have been depleted or reversibly suppressed, or after complete ablation thereof. For example, treatment of a patient with a high cancer cell burden may require higher doses of the radiolabeled anti-CD45 antibody, and as such may result in depletion or suppression of a significant percent of the hematopoietic stem cells of the patient. In such a case, stem cell support may be needed to effect repopulation of those cells. According to certain aspects, stem cell support may not

be needed, and a sufficient quantity of hematopoietic stem cells may be present and capable of repopulating.

[0191] According to certain aspects of this method, the effective amount of $^{225}\text{Ac-BC8}$ is an amount effective to ablate 100% of the hematopoietic stem cells of the subject (also referred to as myeloablation). Exemplary doses include at least those indicated herein as high doses.

[0192] The method generally comprises administering to the subject an effective amount of the $^{225}\text{Ac-BC8}$ in a single dose, such as a single patient specific dose. The amount of reduction of the lymphocytes or hematopoietic stem cells may be determined by any of the methods disclosed herein above. The dose of $^{225}\text{Ac-BC8}$ may depend on the amount of depletion or immunosuppression desired. For example, depletion of the hematopoietic stem cells may be achieved at low doses, such as less than 2Gy of the $^{225}\text{Ac-BC8}$. Reversible immunosuppression of the hematopoietic stem cells may be achieved at doses of less than 8 Gy of the $^{225}\text{Ac-BC8}$, such as doses of 2 Gy to 8 Gy, and ablation may be achieved at high doses such as 8 Gy or greater of the $^{225}\text{Ac-BC8}$, such as about 10 – 12 Gy.

[0193] The method may further comprise administering to the subject an effective amount of the $^{225}\text{Ac-BC8}$ in a fractionated dose, such as multiple administrations of portions of a single patient specific dose, or administration of multiple single patient specific doses. When administered with a second agent, the dose of $^{225}\text{Ac-BC8}$ may depend on the amount of depletion or immunosuppression desired.

[0194] **Methods for treating non-malignant hematological disorders**

[0195] This depletion method (also referred to herein as a conditioning method) may be useful, for example, for improving the outcome of a subsequent gene-edited cell-based therapy where the depletion of hematopoietic stem cells is desirable. According to certain aspects of this method, the subject is afflicted with a non-cancerous disorder treatable via genetically edited cell therapy and is about to undergo such therapy to treat the disorder. The present disclosure also provides a method for treating a subject afflicted with a non-cancerous disorder treatable via genetically edited cell therapy comprising (i) administering to the subject an amount of a radiolabeled anti-CD45 antibody effective to deplete the subject's hematopoietic stem cells, and (ii) after a suitable time period, performing the therapy on the subject to treat the subject's disorder.

[0196] Examples of non-cancerous disorders include, without limitation, hemoglobinopathies (e.g., SCD and β -thalassemia), congenital immunodeficiencies (e.g., SCID and Fanconi's anemia) and viral infections (e.g., HIV infection). According to certain

aspects, the disorder is SCD and the therapy is genetically edited β -globin hematopoietic stem cell therapy. The stem cell therapy can be allogenic or autologous, for example. According to certain aspects, the disorder is SCID and the therapy is genetically edited hematopoietic stem cell therapy, wherein the edited gene is the common gamma chain (γ c) gene, the adenosine deaminase (ADA) gene and/or the Janus kinase 3 (JAK3) gene. The stem cell therapy can be allogenic or autologous, for example.

[0197] According to certain preferred aspects of the subject method, the radiolabeled anti-CD45 antibody is radiolabeled BC8, such as $^{225}\text{Ac-BC8}$. The effective amount of the $^{225}\text{Ac-BC8}$ is an amount effective to deplete, reversibly suppress, or ablate the hematopoietic stem cells of the subject, and can be, for example, from 0.01 $\mu\text{Ci/kg}$ to 1.0 $\mu\text{Ci/kg}$, from 1.0 $\mu\text{Ci/kg}$ to 3.0 $\mu\text{Ci/kg}$, from 3.0 $\mu\text{Ci/kg}$ to 5.0 $\mu\text{Ci/kg}$, or from 0.1 $\mu\text{Ci/kg}$ to 5.0 $\mu\text{Ci/kg}$ of the subject's weight. According to certain aspects, the method comprises (i) administering to the subject from 0.1 $\mu\text{Ci/kg}$ to 5.0 $\mu\text{Ci/kg}$ of $^{225}\text{Ac-BC8}$, and (ii) after 6, 7 or 8 days, performing the therapy on the subject to treat the subject's disorder. According to certain other aspects, the method comprises (i) administering to the subject from 0.1 $\mu\text{Ci/kg}$ to 1.0 $\mu\text{Ci/kg}$ of $^{225}\text{Ac-BC8}$, and (ii) after 6, 7 or 8 days, performing the therapy on the subject to treat the subject's disorder. According to yet further aspects, the method comprises (i) administering to the subject from 1.0 $\mu\text{Ci/kg}$ to 3.0 $\mu\text{Ci/kg}$ of $^{225}\text{Ac-BC8}$, and (ii) after 6, 7 or 8 days, performing the therapy on the subject to treat the subject's disorder. According to yet further aspects, the method comprises (i) administering to the subject from 3.0 $\mu\text{Ci/kg}$ to 5.0 $\mu\text{Ci/kg}$ of $^{225}\text{Ac-BC8}$, and (ii) after 6, 7 or 8 days, performing the therapy on the subject to treat the subject's disorder.

[0198] **Methods for treating malignant hematological disorders**

[0199] The present disclosure also provides a method for treating a subject afflicted with a malignant disease or disorder such as a cancer. The method may generally comprise administering to the subject an amount of a radiolabeled anti-CD45 antibody effective to deplete, reversibly suppress, or ablate the subject's HSC or lymphocytes or hematopoietic cancer blasts. According to at least one exemplary method, a low dose of a radiolabeled anti-CD45 antibody is administered to reduce or deplete the number of hematopoietic stem cells or lymphocytes, and/or circulating tumor cells, without myeloablation. The dose may be a low dose as defined herein.

[0200] According to certain aspects, the method may further comprise administering stem cell support. According to certain aspects, the method may further comprise performing a conditioning therapy after a suitable time period. The conditioning therapy may be a

hematopoietic stem cell transplant, such as a bone marrow transplant, or an adoptive cell therapy on the subject to treat the subject's cancer.

[0201] According to certain aspects of this method, the subject is afflicted with cancer and is about to undergo adoptive cell therapy to treat the cancer. Adoptive cell therapy is known, and includes, for example, CAR T-cell therapy (e.g., autologous cell therapy and allogeneic cell therapy). Adoptive cell therapies provide a method of promoting regression of a cancer in a subject, and generally comprise (i) collecting autologous T-cells (leukapheresis); (ii) expanding the T-cells (culturing); (iii) administering to the subject nonmyeloablative lymphodepleting chemotherapy; and (iv) after administering nonmyeloablative lymphodepleting chemotherapy, administering to the subject the expanded T-cells. The methods of the present disclosure include using a radiolabeled anti-CD45 antibody in lieu of the lymphodepleting chemotherapy, and/or after administration of the expanded cells (e.g., T-cell, NK-cells, dendritic cells, etc.). This later administration of the anti-CD45 antibody (i.e., after administration of the expanded cells) may be used in preparation for transplantation of autologous stem cells (HSCT), or administration of a second effective amount or number of expanded cells.

[0202] Accordingly, the present disclosure provides methods for the treatment of a proliferative disease, such as a hematological malignancy, which include administration of a radiolabeled anti-CD45 antibody and an adoptive cell therapy. The adoptive cell therapy may generally include apheresis of autologous cells which may be gene edited prior to reinfusion (adoptive cell therapy such as CAR T-cell therapy) after lymphodepletion by the radiolabeled anti-CD45 antibody. Alternatively, allogeneic cells may be reinfused after lymphodepletion to provide the adoptive cell therapy. According to methods of the present disclosure, the radiolabeled anti-CD45 antibody may be provided as a single dose 3 to 9 days, such as 6 to 8 days, prior to the adoptive cell therapy.

[0203] According to certain aspects of this method, the radiolabeled anti-CD45 antibody is radiolabeled BC8 as described hereinabove, provided at the doses as described hereinabove, wherein the dose generally depends on the specific radionuclide label (e.g., ²²⁵Ac-BC8). According to certain aspects of this method, the suitable time period after administering the radiolabeled anti-CD45 antibody is 3 days, 4 days, 5 days, 6 days, 7 days, 8 days or 9 days, such as preferably 6, 7 or 8 days.

[0204] According to certain aspects, the method for treating a subject afflicted with cancer consists of (i) administering to the subject a single dose of a radiolabeled anti-CD45 antibody effective to deplete the subject's lymphocytes, and (ii) after a suitable time period

(e.g., 6, 7 or 8 days), performing adoptive cell therapy on the subject to treat the subject's cancer. According to certain aspects, the method for treating a subject afflicted with cancer consists of (i) administering to the subject a single dose of a radiolabeled anti-CD45 antibody effective to deplete the subject's lymphocytes, and (ii) after a suitable time period (e.g., 6, 7 or 8 days), performing adoptive cell therapy on the subject to treat the subject's cancer.

[0205] According to certain aspects of this method, the effective amount of ^{225}Ac -BC8 is from 0.01 $\mu\text{Ci}/\text{kg}$ to 5.0 $\mu\text{Ci}/\text{kg}$ of subject's body weight.

[0206] According to certain aspects, the method for treating a subject afflicted with cancer consists of (i) administering to the subject a single dose of a radiolabeled anti-CD45 antibody effective to myelosuppress the subject, and (ii) after a suitable time period (e.g., 4, 5, 6, 7 or 8 days), performing a bone marrow transplant on the subject to treat the subject's cancer. According to certain aspects, the method for treating a subject afflicted with cancer consists of (i) administering to the subject a single dose of a radiolabeled anti-CD45 antibody effective to deplete the subject's myelocytes, and (ii) after a suitable time period (e.g., 4, 5, 6, 7 or 8 days), performing a bone marrow transplant on the subject to treat the subject's cancer.

[0207] According to certain aspects of this method, the effective amount of ^{225}Ac -BC8 is a low dose, such as a dose of less than 120 μCi , such as from 10 μCi to 100 μCi , or a dose of less than 2 $\mu\text{Ci}/\text{kg}$, such as from 0.01 $\mu\text{Ci}/\text{kg}$ to 1.5 $\mu\text{Ci}/\text{kg}$ or 0.1 $\mu\text{Ci}/\text{kg}$ to 1.0 $\mu\text{Ci}/\text{kg}$.

[0208] Additional therapeutic agents

[0209] The presently disclosed compositions and methods may be used in combination with certain additional therapeutic agents. For example, additional immunotherapeutic agents may be administered in combination with the anti-CD45 antibody compositions disclosed herein. Exemplary additional immunotherapeutic agents include at least antibodies against CD33 and/or CD38 (see for example International Publication No. WO 2019/094931, incorporated herein by reference herein in its entirety), and/or antibodies against CD34, CD117, and/or CD135 (see for example US. Provisional Patent application Nos. 62/838,646 and 62/838,589. Incorporated herein by reference in their entirety).

[0210] The presently disclosed compositions and methods may be used in combination with a radiosensitizer, which may enhance the efficacy of the disclosed radioimmunotherapy (e.g., ^{225}Ac -BC8). For example, Bcl-2 inhibitors may be active against a number of cancer cell lines in combination with radiation, such as provided by the ^{225}Ac -BC8. Additionally, small molecule inhibitors of Bcl-2 proteins display synergy with other

anticancer agents, including, but not limited to etoposide, doxorubicin, cisplatin, paclitaxel, and radiation.

[0211] Inhibiting apoptosis is widely accepted as a necessary step in the transition from normal to cancer cells, and several cancer therapies exert their effects by reversing this process. Commitment to apoptosis is caused by permeabilization of the outer mitochondrial membrane—a process regulated by the binding between different members of the Bcl-2 family. Furthermore, Bcl-2 family members also bind to the endoplasmic reticulum, where they modify processes such as the unfolded-protein response and autophagy that also cause or modify different types of cell death.

[0212] Bcl-2 overexpression was initially described in follicular lymphomas as a consequence of a t(14; 18) translocation, and as a poor prognostic marker in acute myelogenous leukemia (AML) and non-Hodgkin's lymphomas. Overexpression of Bcl-2 was subsequently described in prostate, breast and colon carcinomas, as well as glioblastomas. Overexpression of Mcl-1, another anti-apoptotic, Bcl-2-related protein, was identified in relapsed AML, and was associated with poor prognosis. Other changes in Bcl-2-related protein expression identified in cancer cells include different mutations in the Bax gene, and changes in the proapoptotic to antiapoptotic Bcl-2 protein ratio. The inability of cancer cells to execute an apoptotic program due to defects in the normal apoptotic machinery is thus often associated with an increase in resistance to radiation and/or immunotherapy-induced apoptosis.

[0213] Accordingly, the presently disclosed methods may include addition of a radiosensitizer such as a Bcl-2 inhibitor that may act to directly or indirectly induce apoptosis in cancer cells in a manner that is synergistic with the radiolabeled-anti-CD45 antibody. Bcl-2 inhibitors include small molecule and antisense oligonucleotide drugs, such as AT-101 ((-)-gossypol), GENASENSE® (G3139 or oblimersen; Bcl-2-targeting antisense oligonucleotide), IPI-194, IPI-565, ABT-737, ABT-263, GX-070 (obatoclax) and the like.

[0214] According to preferred aspects, the Bcl-2 inhibitor may be venetoclax, a drug that has been approved for treating chronic lymphocytic leukemia ("CLL"). Venetoclax binds to the BH3-binding groove of BCL-2, displacing pro-apoptotic proteins like BIM to initiate mitochondrial outer membrane permeabilization ("MOMP"), the release of cytochrome c, and caspase activation, ultimately resulting in programmed cancer cell death (i.e., apoptosis). Ideally, by changing the balance between pro-apoptotic and anti-apoptotic stimuli, venetoclax would facilitate programmed cell death of cancer cells and thus improve patient outcomes.

[0215] According to certain aspects, the presently disclosed radioimmunotherapy (e.g., $^{225}\text{Ac-BC8}$), may be used in combination with a BCL-2 inhibitor such as venetoclax to provide a method for treating a subject afflicted with cancer, comprising administering to the subject (i) a BCL-2 inhibitor in conjunction with (ii) the radiolabeled anti-CD45 antibody (e.g., $^{225}\text{Ac-BC8}$), wherein the amounts of the BCL-2 inhibitor and radiolabeled anti-CD45 antibody, when administered in conjunction with one another, are therapeutically effective to treat the cancer.

[0216] This disclosure also provides a method for treating a human subject afflicted with a hematological disease or disorder, comprising administering to the subject (i) a BCL-2 inhibitor such as venetoclax in conjunction with (ii) $^{225}\text{Ac-BC8}$, wherein the amounts of venetoclax and $^{225}\text{Ac-BC8}$, when administered in conjunction with one another, are therapeutically effective to treat the acute myeloid leukemia.

[0217] As used herein, administering to a subject a BCL-2 inhibitor "in conjunction with" a radiolabeled anti-CD45 antibody such as $^{225}\text{Ac-BC8}$ means administering the BCL-2 inhibitor before, during or after administration of the $^{225}\text{Ac-BC8}$. This administration includes, without limitation, the following scenarios: (i) the BCL-2 inhibitor is administered first (e.g., orally once per day for 21 days, 28 days, 35 days, 42 days, 49 days, or a longer period during which the cancer being treated does not progress and during which the BCL-2 inhibitor does not cause unacceptable toxicity), and the $^{225}\text{Ac-BC8}$ is administered second (e.g., intravenously in a single dose or a plurality of doses over a period of weeks); (ii) the BCL-2 inhibitor is administered concurrently with the $^{225}\text{Ac-BC8}$ (e.g., the BCL-2 inhibitor is administered orally once per day for n days, and the $^{225}\text{Ac-BC8}$ is administered intravenously in a single dose on one of days 2 through n-1 of the BCL-2 inhibitor regimen); (iii) the BCL-2 inhibitor is administered concurrently with the $^{225}\text{Ac-BC8}$ (e.g., the BCL-2 inhibitor is administered orally for a duration of greater than one month (e.g., orally once per day for 35 days, 42 days, 49 days, or a longer period during which the cancer being treated does not progress and during which the BCL-2 inhibitor does not cause unacceptable toxicity), and the $^{225}\text{Ac-BC8}$ is administered intravenously in a single dose on a day within the first month of the BCL-2 inhibitor regimen); and (iv) the $^{225}\text{Ac-BC8}$ is administered first (e.g., intravenously in a single dose or a plurality of doses over a period of weeks), and the BCL-2 inhibitor is administered second (e.g., orally once per day for 21 days, 28 days, 35 days, 42 days, 49 days, or a longer period during which the cancer being treated does not progress and during which the BCL-2 inhibitor does not cause unacceptable toxicity).

[0218] The amount of the radiolabeled anti-CD45 antibody administered may be sufficient to deplete, reversibly immunosuppress, or ablate the hematological stem cells in the patient. In general, the dose of the radiolabeled anti-CD45 antibody is a sub-saturating dose that may reversibly immunosuppress the hematological stem cells.

[0219] Additional radiosensitizing agents include, for example, histone deacetylase inhibitors (HDACi) such as vorinostat, belinostat, and romidepsin; metronidazole, misonidazole, intra-arterial Budr, intravenous iododeoxyuridine (IudR), nitroimidazole, 5-substituted-4-nitroimidazoles, 2H-isoindolediones, [[(2-bromoethyl)-amino]methyl]-nitro-1H-imidazole-1-ethanol, nitroaniline derivatives, DNA-affinic hypoxia selective cytotoxins, halogenated DNA ligand, 1,2,4 benzotriazine oxides, 2-nitroimidazole derivatives, fluorine-containing nitroazole derivatives, benzamide, nicotinamide, acridine-intercalator, 5-thiotretazole derivative, 3-nitro-1,2,4-triazole, 4,5-dinitroimidazole derivative, hydroxylated texaphrins, cisplatin, mitomycin, tiripazamine, nitrosourea, mercaptopurine, methotrexate, fluorouracil, bleomycin, vincristine, carboplatin, epirubicin, doxorubicin, cyclophosphamide, vindesine, etoposide, paclitaxel, heat (hyperthermia), and the like.

[0220] The term “histone deacetylase inhibitor” or “HDACi” refers to histone deacetylase inhibitors that can be grouped in four classes: hydroxamates (panobinostat (LBH-589), trichostatin-A (TSA), vorinostat (SAHA), belinostat (PXD101), NVP-LAQ824 and givinostat (ITF2357)), cyclic peptide (romidepsin (depsipeptide)), aliphatic acids (valproic acid (VPA) and sodium phenylbutyrate) and benzamides (MS-275, MGCD0103). HDACi are characterized as class I-specific HDACs inhibitors (MGCD0103, romidepsin and MS-275) or as pan-HDAC inhibitors, denoting activity against both classes I and II HDACs (TSA, panobinostat, vorinostat and belinostat).

[0221] Histone deacetylase inhibitors are recognized to exert multiple cytotoxic actions in cancer cells, often through acetylation of non-histone proteins. Some well-recognized mechanisms of HDACi lethality include, in addition to relaxation of DNA and de-repression of gene transcription, interference with chaperone protein function, free radical generation, induction of DNA damage, up-regulation of endogenous inhibitors of cell cycle progression, and promotion of apoptosis. Intriguingly, this class of agents is relatively selective for transformed cells where they have been found to cause DNA repair to be halted after chemotherapy, and to promote the efficacy of chemotherapy.

[0222] According to certain aspects, the presently disclosed radioimmunotherapy (e.g., ²²⁵Ac-BC8), may be used in combination with an HDACi such as vorinostat, belinostat, or romidepsin to provide a method for treating a subject afflicted with cancer, comprising

administering to the subject (i) an HDACi in conjunction with (ii) the radiolabeled anti-CD45 antibody (e.g., $^{225}\text{Ac-BC8}$), wherein the amounts of the HDACi and radiolabeled anti-CD45 antibody, when administered in conjunction with one another, are therapeutically effective to treat the cancer.

[0223] This disclosure also provides a method for treating a human subject afflicted with a hematological disease or disorder, comprising administering to the subject (i) an HDACi such as vorinostat, belinostat, or romidepsin in conjunction with (ii) $^{225}\text{Ac-BC8}$, wherein the amounts of the HDACi and $^{225}\text{Ac-BC8}$, when administered in conjunction with one another, are therapeutically effective to treat the acute myeloid leukemia.

[0224] As with the BCL-2 inhibitors, administering to a subject an HDACi "in conjunction with" a radiolabeled anti-CD45 antibody such as $^{225}\text{Ac-BC8}$ means administering the HDACi before, during or after administration of the $^{225}\text{Ac-BC8}$. This administration includes, without limitation, the following scenarios: (i) the HDACi is administered first (e.g., orally once per day for 21 days, 28 days, 35 days, 42 days, 49 days, or a longer period during which the cancer being treated does not progress and during which the HDACi does not cause unacceptable toxicity, or intravenously on days 1, 8, 15, of a 28 day cycle), and the $^{225}\text{Ac-BC8}$ is administered second (e.g., intravenously in a single dose or a plurality of doses over a period of weeks); (ii) the HDACi is administered concurrently with the $^{225}\text{Ac-BC8}$ (e.g., the HDACi is administered orally once per day for n days, or intravenously for n days, and the $^{225}\text{Ac-BC8}$ is administered intravenously in a single dose on one of days 2 through n-1 of the HDACi regimen); (iii) the HDACi is administered concurrently with the $^{225}\text{Ac-BC8}$ (e.g., the HDACi is administered orally for a duration of greater than one month as described herein, and the $^{225}\text{Ac-BC8}$ is administered intravenously in a single dose on a day within the first month of the HDACi regimen); and (iv) the $^{225}\text{Ac-BC8}$ is administered first (e.g., intravenously in a single dose or a plurality of doses over a period of weeks), and the HDACi is administered second (as described herein).

[0225] Article of Manufacture

[0226] The present disclosure further provides an article of manufacture comprising (a) a radiolabeled anti-CD45-immunoglobulin, and (b) a label instructing the user to administer to a subject an amount of the immunoglobulin effective to deplete the subject's hematopoietic stem cells.

[0227] According to certain aspects of the subject article, the radiolabeled anti-CD45-immunoglobulin is $^{225}\text{Ac-BC8}$, wherein the effective amount can be, for example, from 0.01 $\mu\text{Ci/kg}$ to 5.0 $\mu\text{Ci/kg}$, or from 0.01 $\mu\text{Ci/kg}$ to 1.0 $\mu\text{Ci/kg}$ of the $^{225}\text{Ac-BC8}$, or from 1.0 $\mu\text{Ci/kg}$

to 3.0 $\mu\text{Ci}/\text{kg}$ of the $^{225}\text{Ac-BC8}$, or from 3.0 $\mu\text{Ci}/\text{kg}$ to 5.0 $\mu\text{Ci}/\text{kg}$ of the $^{225}\text{Ac-BC8}$, or from 10 μCi to 120 μCi of the $^{225}\text{Ac-BC8}$, or from 100 μCi to 250 μCi of the $^{225}\text{Ac-BC8}$, or from 200 μCi to 500 μCi of the $^{225}\text{Ac-BC8}$, or from 5 μCi to 80 μCi of the $^{225}\text{Ac-BC8}$.

[0228] According to certain aspects of the subject article, the radiolabeled anti-CD45-immunoglobulin is $^{177}\text{Lu-BC8}$, wherein the effective amount can be, for example, from 1 $\mu\text{Ci}/\text{kg}$ to 500 $\mu\text{Ci}/\text{kg}$, or from 1 $\mu\text{Ci}/\text{kg}$ to 100 $\mu\text{Ci}/\text{kg}$ of the $^{177}\text{Lu-BC8}$, or from 100 $\mu\text{Ci}/\text{kg}$ to 300 $\mu\text{Ci}/\text{kg}$ of the $^{177}\text{Lu-BC8}$, or from 300 $\mu\text{Ci}/\text{kg}$ to 500 $\mu\text{Ci}/\text{kg}$ of the $^{177}\text{Lu-BC8}$.

[0229] This disclosure will be better understood by reference to the examples which follow, but those skilled in the art will readily appreciate that the specific examples detailed are only illustrative of the disclosure as described more fully in the claims which follow thereafter.

[0230] **Aspects of the Invention**

[0231] The following aspects are disclosed in this application:

[0232] Aspect 1. A method for depleting a subject's hematopoietic stem cells comprising administering to the subject an effective amount of a radiolabeled anti-CD45-immunoglobulin.

[0233] Aspect 2. The method of aspect 1, wherein the effective amount of the radiolabeled anti-CD45-immunoglobulin depletes at least 25% of hematopoietic stem cells of the subject, or 50% of hematopoietic stem cells of the subject, or at least 70% of hematopoietic stem cells of the subject, or at least 80% of hematopoietic stem cells of the subject, or at least 90% of hematopoietic stem cells of the subject, or at least 95% of hematopoietic stem cells of the subject, or not more than 90% of hematopoietic stem cells of the subject, or not more than 95% of hematopoietic stem cells of the subject.

[0234] Aspect 3. The method of aspect 1, wherein the effective amount of the radiolabeled anti-CD45-immunoglobulin depletes at least 90% of hematopoietic stem cells of the subject, or at least 95% of hematopoietic stem cells of the subject, or at least 98% of hematopoietic stem cells of the subject, or at least 99% of hematopoietic stem cells of the subject, or not more than 98% of hematopoietic stem cells of the subject, or not more than 99% of hematopoietic stem cells of the subject.

[0235] Aspect 4. The method according to any one of aspects 1 to 3, further comprising: administering an effective amount of a second therapeutic agent comprising one or more of an immunotherapeutic agent, a radiosensitizer, or a chemotherapeutic agent.

[0236] Aspect 5. The method of aspect 4, wherein the immunotherapeutic agent comprises one or more antibodies against CD33, CD34, CD38, CD119, and CD135.

[0237] Aspect 6. The method of aspect 4, wherein the radiosensitizer comprises a Bcl-2 inhibitor, or an HDAC inhibitor (HDACi).

[0238] Aspect 7. The method of aspect 1, wherein the effective amount of the radiolabeled anti-CD45-immunoglobulin depletes 100% of hematopoietic stem cells of the subject (i.e., ablates the hematopoietic stem cells).

[0239] Aspect 8. A method for depleting a subject's lymphocytes comprising administering to the subject an effective amount of a radiolabeled anti-CD45-immunoglobulin.

[0240] Aspect 9. The method of aspect 8, wherein the effective amount of the radiolabeled anti-CD45-immunoglobulin depletes at least 25% of lymphocytes of the subject, or 50% of lymphocytes of the subject, or at least 70% of lymphocytes of the subject, or at least 80% of lymphocytes of the subject, or at least 90% of lymphocytes of the subject, or at least 95% of lymphocytes of the subject, or at least 98% of lymphocytes of the subject, or at least 99% of lymphocytes of the subject, or not more than 90% of lymphocytes of the subject, or not more than 95% of lymphocytes of the subject, or not more than 98% of lymphocytes of the subject, or not more than 99% of lymphocytes of the subject.

[0241] Aspect 10. The method of aspect 8, wherein the effective amount of the radiolabeled anti-CD45-immunoglobulin depletes at least 25% of hematopoietic cancer blasts of the subject, or 50% of hematopoietic cancer blasts of the subject, or at least 70% of hematopoietic cancer blasts of the subject, or at least 80% of hematopoietic cancer blasts of the subject, or at least 90% of hematopoietic cancer blasts of the subject, or at least 95% of hematopoietic cancer blasts of the subject, or at least 98% of hematopoietic cancer blasts of the subject, or at least 99% of hematopoietic cancer blasts of the subject, or not more than 90% of hematopoietic cancer blasts of the subject, or not more than 95% of hematopoietic cancer blasts of the subject, or not more than 98% of hematopoietic cancer blasts of the subject, or not more than 99% of hematopoietic cancer blasts of the subject.

[0242] Aspect 11. The method of aspect 8, wherein the effective amount of the radiolabeled anti-CD45-immunoglobulin depletes 100% of lymphocytes of the subject (i.e., ablates the lymphocytes), or 100% of the hematopoietic cancer blasts of the subject.

[0243] Aspect 12. The method according to any one of aspects 1 to 11, wherein the subject is afflicted with a non-cancerous disorder treatable via genetically edited cell therapy and is about to undergo such therapy to treat the disorder, and the effective amount of the radiolabeled anti-CD4-immunoglobulin is administered as a single dose.

[0244] Aspect 13. A method for treating a subject afflicted with a non-cancerous disorder treatable via genetically edited cell therapy comprising (i) administering to the subject an amount of a radiolabeled anti-CD45-immunoglobulin effective to deplete the subject's hematopoietic stem cells, and (ii) after a suitable time period, performing the therapy on the subject to treat the subject's disorder.

[0245] Aspect 14. The method according to aspect 12 or 13, wherein the disorder is selected from the group consisting of a hemoglobinopathy, a congenital immunodeficiency, and a viral infection.

[0246] Aspect 15. The method according to aspect 14, wherein the disorder is selected from the group consisting of sickle cell disease (SCD), severe combined immunodeficiency disease (SCID), β -thalassemia and Fanconi's anemia.

[0247] Aspect 16. The method of aspect 14, wherein the disorder is SCD and the therapy is genetically edited β -globin hematopoietic stem cell therapy.

[0248] Aspect 17. The method of aspect 14, wherein the disorder is SCID and the therapy is genetically edited hematopoietic stem cell therapy, wherein the edited gene is selected from the group consisting of the common gamma chain (γ c) gene, the adenosine deaminase (ADA) gene and the Janus kinase 3 (JAK3) gene.

[0249] Aspect 18. The method according to any one of aspects 1 to 7, wherein the subject is afflicted with a cancerous disorder and is about to undergo a hematopoietic stem cell transplant such as a bone marrow transplant to treat the disorder, and wherein the effective amount of the radiolabeled anti-CD4-immunoglobulin is administered as a single dose.

[0250] Aspect 19. The method according to any one of aspects 8 to 11, wherein the subject is afflicted with a cancerous disorder treatable via genetically edited cell therapy and is about to undergo such therapy to treat the disorder, and the effective amount of the radiolabeled anti-CD4-immunoglobulin is administered as a single dose.

[0251] Aspect 20: A method for treating a subject afflicted with a cancerous disorder treatable via genetically edited cell therapy comprising (i) administering to the subject an amount of a radiolabeled anti-CD4-immunoglobulin effective to deplete the subject's lymphocytes, and (ii) after a suitable time period, performing the therapy on the subject to treat the subject's disorder.

[0252] Aspect 21: The method according to any one of aspects 18 to 20, wherein the cancerous disorder is acute myeloid leukemia, acute lymphoid leukemia, chronic myeloid

leukemia, chronic lymphoid leukemia, multiple myeloma, diffuse large B-cell lymphoma, of non-Hodgkin's lymphoma.

[0253] Aspect 22: The method according to aspects 19 or 20, wherein the genetically edited cell therapy is an adoptive cell therapy to treat the cancerous disorder.

[0254] Aspect 23: The method according to aspect 22, wherein the adoptive cell therapy is CAR T-cell therapy, wherein the CAR T-cell therapy comprises the administration of gene-edited CAR T-cells, and wherein the gene-edited CAR T-cells fail to properly express at least one checkpoint receptor and/or at least one T-cell receptor.

[0255] Aspect 24: The method according to aspect 23, wherein the CAR T-cell therapy is autologous cell therapy.

[0256] Aspect 25: The method according to aspect 23, wherein the CAR T-cell therapy is allogeneic cell therapy.

[0257] Aspect 26: The method according to any one of aspects 1 to 25, wherein the radiolabeled anti-CD45 antibody is $^{225}\text{Ac-BC8}$ or $^{177}\text{Lu-BC8}$.

[0258] Aspect 27. The method according to aspect 26, wherein the effective amount of $^{225}\text{Ac-BC8}$ is from 0.01 $\mu\text{Ci/kg}$ to 5.0 $\mu\text{Ci/kg}$ subject weight, or from 0.01 $\mu\text{Ci/kg}$ to 1.0 $\mu\text{Ci/kg}$ subject weight, or from 1.0 $\mu\text{Ci/kg}$ to 3.0 $\mu\text{Ci/kg}$ subject weight, or from 3.0 $\mu\text{Ci/kg}$ to 5.0 $\mu\text{Ci/kg}$ subject weight; OR wherein the effective amount of $^{225}\text{Ac-BC8}$ is from 2 μCi to below 0.5 mCi, or from at least 2 μCi to below 120 μCi , or from 10 μCi to below 120 μCi , or from 50 μCi to below 250 μCi ; OR wherein the effective amount of $^{177}\text{Lu-BC8}$ is from 1 $\mu\text{Ci/kg}$ to 500 $\mu\text{Ci/kg}$ subject weight, or from 1 $\mu\text{Ci/kg}$ to 100 $\mu\text{Ci/kg}$ subject weight, or from 100 $\mu\text{Ci/kg}$ to 300 $\mu\text{Ci/kg}$ subject weight, or from 300 $\mu\text{Ci/kg}$ to 500 $\mu\text{Ci/kg}$ subject weight; OR wherein the effective amount of $^{177}\text{Lu-BC8}$ is from 10 μCi to 20 mCi, or from 100 μCi to 3 mCi, or from 3 mCi to 20 mCi.

[0259] Aspect 28: The method according to any one of aspects 1 to 27, wherein the anti-CD45-immunoglobulin comprises BC8, wherein the BC8 comprises a light chain having the amino acid sequence as set forth in SEQ ID NO:1, or a light chain N-terminal amino acid sequence as set forth in SEQ ID NO: 9.

[0260] Aspect 29: The method according to any one of aspects 1 to 28, wherein the anti-CD45-immunoglobulin comprises BC8, wherein the light chain of the BC8 comprises at least one complementarity determining region having the amino acid sequence as set forth in SEQ ID NO:3, SEQ ID NO:4, or SEQ ID NO:5.

[0261] Aspect 30: The method according to any one of aspects 1 to 29, wherein the anti-CD45-immunoglobulin comprises BC8, wherein the BC8 comprises a light chain having the amino acid sequence set forth in SEQ ID NO:12 or SEQ ID NO:13.

[0262] Aspect 31: The method according to any one of aspects 1 to 30, wherein the anti-CD45-immunoglobulin comprises BC8, wherein the BC8 comprises a heavy chain having the amino acid sequence set forth in SEQ ID NO:2, or a heavy chain N-terminal amino acid sequence as set forth in SEQ ID NO: 10.

[0263] Aspect 32: The method according to any one of aspects 1 to 31, wherein the anti-CD45-immunoglobulin comprises BC8, wherein the heavy chain of the BC8 comprises at least one complementarity determining region having the amino acid sequence as set forth in SEQ ID NO:6, SEQ ID NO:7, or SEQ ID NO:8.

[0264] Aspect 33: The method according to any one of aspects 1 to 32, wherein the anti-CD45-immunoglobulin comprises BC8, wherein the BC8 comprises a heavy chain having the amino acid sequence set forth in SEQ ID NO:15 or SEQ ID NO:16.

[0265] Aspect 34: The method according to any one of aspects 1 to 33, wherein the anti-CD45-immunoglobulin comprises BC8, and the heavy chain of the BC8 comprises the amino acid ASP or ASN at position 141 (relative to the N-terminal amino acid).

[0266] Aspect 35: The method according to aspect 34, wherein a ratio of ASP:ASN in a population of BC8 proteins within the range 1:99 to 99:1, such as 10:90 to 90:10.

[0267] Aspect 36: The method according to any one of aspects 1 to 35, wherein the anti-CD45-immunoglobulin comprises BC8 that has been modified to comprise a heavy chain constant region from human IgG1, IgG2, or IgG4, i.e., the amino acid sequence as set forth in one of SEQ ID NOS:17-19.

[0268] Aspect 37: The method according to any one of aspects 1 to 36, wherein the anti-CD45-immunoglobulin comprises BC8 that has been modified to comprise a heavy chain constant region from human IgG4 comprising the mutation S228P, having the amino acid sequence as set forth in SEQ ID NO:20.

[0269] Aspect 38: The method according to any one of aspects 1 to 37, wherein the anti-CD45-immunoglobulin comprises BC8 that has been modified to comprise a light chain kappa constant region from human having the amino acid sequence as set forth in SEQ ID NO:21.

[0270] Aspect 39. An article of manufacture comprising (a) a radiolabeled anti-CD45 antibody, and (b) a label instructing the user to administer to a subject an amount of the

antibody effective to deplete or ablate the subject's hematopoietic stem cells or the subject's lymphocytes.

[0271] Aspect 40. The article of aspect 39, wherein the radiolabeled BC8 is ^{225}Ac -BC8, and the effective amount of ^{225}Ac -BC8 is from 0.01 $\mu\text{Ci}/\text{kg}$ to 5.0 $\mu\text{Ci}/\text{kg}$, or from 0.01 $\mu\text{Ci}/\text{kg}$ to 1.0 $\mu\text{Ci}/\text{kg}$ subject weight, or from 1.0 $\mu\text{Ci}/\text{kg}$ to 3.0 $\mu\text{Ci}/\text{kg}$ subject weight, or from 3.0 $\mu\text{Ci}/\text{kg}$ to 5.0 $\mu\text{Ci}/\text{kg}$ subject weight; OR wherein the effective amount of ^{225}Ac -BC8 is from 2 μCi to below 0.5 mCi, or from 2 μCi to 250 μCi , or from 75 μCi to 400 μCi

EXAMPLES

[0272] Example 1 – Production of the anti-CD45 Immunoglobulin BC8

[0273] The murine anti-CD45 mAb BC8 was prepared from a hybridoma (ATCC No. HB-10507) that was initially developed by fusing mouse myeloma NS1 cells with spleen cells from a BALB/C mouse hyperimmunized with human phytohemagglutinin (PHA)-stimulated mononuclear cells. The original fused cells, after screening for microbial contaminations, were cultured using the JRH-Biosciences EXCell 300 medium supplemented with 1-2% Fetal Bovine Serum (FBS).

[0274] The hybridoma cell line was adapted for culture in a serum-free culture medium. Briefly, the cells in culture were slowly and gradually weaned off the serum albumin using the combo medium supplemented with glutamine, cholesterol, insulin and transferrin. The cells were then grown in up to 500L scale to a density of $>1 \times 10^6$ cells/ml. The medium was harvested and processed for the purification of the anti-CD45 antibody using a combination of cation exchange chromatography, protein-A chromatography, and anion exchange membrane separation. The purified antibody was concentrated by nanofiltration (30kD cutoff). The concentration of the purified product was measured at 5.2 mg/ml and was stored at 2-8°C.

[0275] The purified antibody was characterized by SDS-PAGE, IEF, and SEC-HPLC techniques. A single product peak (99.4%) was recorded with SEC-HPLC with about 0.6% aggregates. The non-reducing SDS-PAGE showed a single band for the antibody. The SDS-PAGE under reduced conditions confirmed the presence of the light and the heavy chains (99.9% together).

[0276] Example 2 – Sequencing of the Anti-CD45-Immunoglobulin BC8

[0277] Total RNA was isolated from the hybridoma cells following the technical manual of Trizol® Reagent. The total RNA was analyzed by agarose gel electrophoresis and was reverse transcribed into cDNA using isotype-specific anti-sense primers or universal primers following the technical manual of PrimeScript™ 1st Strand cDNA Synthesis Kit.

The antibody fragments of VH, VL, CH and CL were amplified and were separately cloned into a standard cloning vector using standard molecular cloning procedures. Colony PCR screening was performed to identify clones with inserts of correct sizes. More than five single colonies with inserts of correct sizes were sequenced for each antibody fragment. The complete nucleotide sequence of the light and the heavy chains are shown in **FIGS. 4A, 4B, 5A, and 5B**.

[0278] The anti-CD45-immunoglobulin (i.e., BC8 antibody) was sequenced using the mass spectrometry peptide mapping approach. The BC8 antibody was de-glycosylated, reduced and digested with individual enzymes; trypsin, Lys-C and chymotrypsin. The peptide fragments were then analyzed by the LC-coupled mass spectrometry technique using the MS/MS fragmentation analysis approach. Protein sequencing of the heavy and light chains of the BC8 antibody showed that the actual amino acid sequence differs from that predicted by the DNA sequence by only a single amino acid in the heavy chain. As highlighted in **FIGS. 5A and 5B**, the codon that codes for the amino acid at position 141 predicts an ASN-141 and not the actual ASP-141 found by protein sequencing. Moreover, sequencing of various batches of the protein indicated differing amounts of the ASP and ASN at position 141, i.e., the protein was found to comprise both ASN-141 and ASP-141 in ratios of from 1:99 to 99:1, such as 10:90 to 90:10 (ASN-141 : ASP-141). See **Table 1**.

Table 1:

LCMS/MS analysis of peptide fragment 124-151 from two batches of isolated BC8 mAb showing the presences of both "N" and "D" at position-141 in difference abundance.

BC8 mAb Lot	Peptide Fragment Sequence (124-151 fragment)	Observed [M+H] ⁺ (Da)	Theoretical [M+H] ⁺ (Da)	% Abundance
GBI Lot	TTPPSVYPLAPGSAAQT <u>N</u> SM VTLGCLVK	2860.4663	2860.4583	84.4%
	TTPPSVYPLAPGSAAQT <u>D</u> SM VTLGCLVK	2861.4470	2861.4423	15.6%
Lot#2014	TTPPSVYPLAPGSAAQT <u>N</u> SM VTLGCLVK	2860.4627	2860.4583	43.6%
	TTPPSVYPLAPGSAAQT <u>D</u> SM VTLGCLVK	2861.4479	2861.4423	56.4%

[0279] This type of post-translational modification, i.e., deamination, may depend on the cellular environment and, in some cases, has been postulated to be related to protein age (e.g., may provide a signal for protein degradation). The fact that other deaminated amino acids were not identified, however, may be indicative of an important and specific role for

ASP-141. At the very least, ASP-141 may be in an exposed or accessible region on the folded protein. That is, ASN-141 may be solvent accessible and reside within a conformationally flexible region of the antibody. The effect of deamination on the biological activity of the BC8 antibody may be determined from the results of human clinical trials.

[0280] Example 3 – BC8: Labeling and purification to form ²²⁵Ac-BC8

[0281] Conjugation of BC8 and irrelevant control mAb 18B7 (mouse IgG1) with DOTA

[0282] The antibodies against CD45 (i.e., BC8 antibody) and a control (i.e., mouse monoclonal reactive against the fungal polysaccharide glucuronoxylomannan; 2mg each) were equilibrated with conjugation buffer (Na carbonate buffer with 1 mM EDTA, pH=8.5-9.0) by four ultrafiltration spins using either a Centricon filter with MW cutoff of 50,000, or a Vivaspin ultrafiltration tube with MW cutoff of 50,000; 1.5 milliliters (ml) of conjugation buffer per spin was used. For each spin, the antibodies were spun at 53,000 RPM for 5-20 minutes at 4°C in a Thermo IEC Centra CL3R centrifuge with a fixed angle rotor to a residual retentate volume of 100-200 microliters (μl). Spin times vary for different antibodies and different protein concentrations. The antibodies were incubated at 4°C for 30 minutes following the second and third spins to allow time for equilibration.

Table 2

DOTA/Protein Molar Ratio		Radiolabeling Yield
Initial	Final	
5	1.2	70
5	1.2	68
5	1.2	67
5	1.2	75
7.5	1.5	82
7.5	1.5	83
7.5	1.5	81
7.5	1.5	83
15	1.4	80
15	1.4	82
15	1.4	80
15	1.4	83

[0283] For the conjugation, a solution of DOTA-pSCN (MW=678) at 3 mg/ml in 0.15M NH₄OAc was prepared by dissolved by vortexing. DOTA-pSCN and the antibodies (at >5 mg/ml) and were mixed together at 5, 7.5 and 15 molar ratios in Eppendorf tubes and incubated for 15 hours at room temperature (see FIG. 9A). For purification of the DOTA-antibody conjugates the unreacted DOTA-pSCN was removed by 7 rounds of ultrafiltration

as described above, washing each time with 1.5ml of 0.15 M NH₄OAc buffer, pH=6.5 down a volume of approximately 100µl. After the final wash, 0.15 M NH₄OAc buffer was added to bring each sample to a final concentration of ~1 mg/ml.

[0284] The final concentration of the DOTA antibody conjugates was measured by the simplified Lowry method. The number of DOTA molecules conjugated to the antibodies (DOTA to protein molar ratio) was determined as described in Dadachova et al., 1999, Spectrophotometric method for determination of bifunctional macrocyclic ligands in macrocyclic ligand-protein conjugates, *Nuclear Medicine & Biology*, 26:977-982. The results of the DOTA to protein molar ratio determination are shown in **Table 2**.

[0285] Radiolabeling of DOTA-Antibody Conjugates with ²²⁵Ac

[0286] A reaction comprising 15µl 0.15M NH₄OAc buffer, pH=6.5 and 2µL (10µg) DOTA-BC8 (5 mg/ml) was mixed in an Eppendorf reaction tube, and 4µL ²²⁵Ac (10 µCi) in 0.05 M HCl was subsequently added (*see FIG. 9B*). The contents of the tube were mixed with a pipette tip and the reaction mixture was incubated at 37°C for 90 min with shaking at 100 rpm. At the end of the incubation period, 3 µL of a 1mM DTPA solution was added to the reaction mixture and it was incubated at room temperature for 20 min to bind the unreacted ²²⁵Ac into the ²²⁵Ac-DTPA complex.

[0287] Instant thin layer chromatography (ITLC) with 10cm silica gel strip and 10mM EDTA/normal saline mobile phase was used to determine the radiochemical purity of ²²⁵Ac-DOTA-BC8 through separating ²²⁵Ac-labeled BC8 (²²⁵Ac-DOTA-BC8) from free ²²⁵Ac (²²⁵Ac-DTPA). In this system the radiolabeled antibody stays at the point of application and ²²⁵Ac-DTPA moves with the solvent front. The strips were cut in halves and counted in the gamma counter equipped with the multichannel analyzer using channels 72-110 for ²²⁵Ac to exclude its daughters. Select radiolabeling results are presented in **Table 2**, which demonstrate that the conjugate formed at an initial molar ratio of DOTA to BC8 of 7.5 provided the highest conjugation ratio (DOTA to BC8 protein) and was chosen for all follow-up experiments described below (batch A).

[0288] Purification of ²²⁵Ac-DOTA-BC8 and HPLC of the purified ²²⁵Ac-DOTA-BC8

[0289] The ²²⁵Ac-DOTA-BC8 samples were purified either on PD10 columns pre-blocked with 1% HSA or on the Vivaspin centrifugal concentrators with 50 kDa MW cut-off with 2 x 1.5 mL washes, 3 min per spin. The HPLC analyses of the ²²⁵Ac-DOTA-BC8 after purification were conducted using a Waters HPLC system equipped with flow-through Waters UV and Bioscan Radiation detectors. The injected samples size was 30 µL. The

elution was carried out on a TSK3000SW XL column using PBS at pH=7.4 as an eluent and a flow rate of 1ml/min. Example chromatograms are provided in **FIGS. 10A** and **10B**, which show SEC-HPLC (size exclusion chromatography-HPLC) of ^{225}Ac -DOTA-BC8, wherein **FIG. 10A** shows the BC8 standard, and **FIG. 10B** shows the ^{225}Ac -DOTA-BC8 (the peak at 13 min is HSA added to stabilize the final formulation).

[0290] Example 4 – ^{225}Ac -BC8: Stability

[0291] ^{225}Ac -DOTA-BC8 stability determination

[0292] The DOTA-BC8 (batch A) was used in all immunoreactivity experiments and was radiolabeled with ^{225}Ac as described in the procedure above at $1\mu\text{Ci}/\mu\text{g}$ specific activity. For the stability determination, the ^{225}Ac -DOTA-BC8 was tested either in the original volume of $20\mu\text{L}$, or diluted to $40\mu\text{L}$ or $60\mu\text{L}$ with the working buffer ($0.15\text{ M NH}_4\text{OAc}$) and incubated at room temperature (rt) for 48 hours or at 4°C for 96 hours and tested by ITLC. All samples were analyzed in duplicate and the experiment was performed three times.

[0293] The results shown in **FIG. 11** demonstrate that the actinium-225 labeled BC8 (^{225}Ac -DOTA-BC8) was stable at 4°C for up to 96 hours.

[0294] Example 5 – ^{225}Ac -BC8: Immunoreactivity

[0295] ^{225}Ac -DOTA-BC8 immunoreactivity (IR) determination using cell lines

[0296] The DOTA-BC8 (batch A) was used in all immunoreactivity experiments and was radiolabeled with ^{225}Ac as described in the procedure above at $1\mu\text{Ci}/\mu\text{g}$ specific activity. The Ramos CD45 positive cells and control CD45 negative EL4 cells were used in the amounts of 1.0-7.5 million cells per sample, in duplicate. The experiment was performed twice. The results presented in **FIG. 12** demonstrate that ^{225}Ac -DOTA-BC8 bound specifically to Ramos cells with 50% of the radiolabeled antibody binding to these cells versus only around 10% binding to control EL4 cells. However, continuously growing two cell lines in the laboratory for QC is not cost effective, and simpler assays were desired for IR determination. As controls the following conditions were used: Ramos cells pre-blocked with 1% BSA; EL4 cells; EL4 cells pre-blocked with 1% BSA.

[0297] ^{225}Ac -DOTA-BC8 immunoreactivity (IR) determination using Cytotrol cells

[0298] Cytotrol cells (Beckman Coulter) were initially used to determine the binding of naïve BC8 antibody to those cells versus control 18B7 antibody (nonspecific control antibody against the fungal polysaccharide glucuronoxylomannan) by flow cytometry (**FIG. 13A**). The cells were taken up in RPMI medium, and the secondary antibody was PE labeled rat anti-mouse IgG1 from Biolegend. CytoTrol cells are lyophilized human lymphocytes

isolated from peripheral blood that exhibit CD45 surface antigen and were selected based on their commercial availability (Beckman Coulter) and consistency.

[0299] The binding of naïve BC8 to Cytotrol cells was compared to that of DOTA-BC8 (**FIG. 13B**). Naïve BC8 showed strong binding to Cytotrol cells, while control 18B7 mAb bound only at the background level (**FIG. 13B**). The attachment of DOTA to BC8 reduced its IR to approximately 70% of the naïve BC8 IR (**FIG. 13B**).

Table 3
IR determination using Cytotrol cells for six samples of ^{225}Ac -DOTA-BC8.

Counts in the washes and bound to the cells	BC8 1	BC8 2	BC8 3	BC8 4	BC8 5	BC8 6
S1	6,456	6,000	6,234	7,011	7,534	6,589
S2	2,443	1,830	1,900	1,300	890	1,711
S3	780	650	521	543	499	623
cells	15,354	17,023	16,743	16,985	15,999	15,235
% bound	61.3	66.5	65.9	65.7	66.4	63.01
Mean \pm SD for 6 samples	64.8 \pm 2.14%					

[0300] Subsequently we performed the IR determination for ^{225}Ac -DOTA-BC8. To measure binding of the radiolabeled antibodies to the Cytotrol cells, we used 3 tubes of Cytotrol cells (lot 729154) for each sample, and measured binding for duplicate samples. Using 0.5ml reconstitution buffer, washed from vial to vial, we pooled the cells. The vials were washed with two more aliquots of 0.5ml, and the washes were pooled. The cells were collected by centrifugation for 4 minutes at 4000 rpm, blocked with 1ml of RPMI containing 1% Bovine Serum Albumin (BSA), respun, and resuspended in 0.5ml RPMI/BSA. Around 25,000 CPM of labeled antibody were added/vial. The vials were incubated for 1-hour at 37°C, shaking at 150 RPM, spun 4 minutes at 4000 RPM, collect three washes, count washes and cells. Table 3 shows the IR determination for 6 samples of ^{225}Ac -DOTA-BC8. The mean IR was 64.8 \pm 2.14%.

[0301] Finally, we performed the side by side comparison of the binding of DOTA-BC8 sample to Cytotrol cells by flow cytometry (**FIG. 14B**), followed by immediate radiolabeling of the same sample with ^{225}Ac and binding of the radiolabeled sample to Cytotrol cells (**FIG. 14A**). The binding of the DOTA-BC8 to the cells by flow cytometry (around 60% of the naïve BC8 binding) matched that of radiolabeled ^{225}Ac -DOTA-BC8 to Cytotrol cells. Accordingly, the Cytotrol assay was found to be a convenient and cost-effective way to evaluate the IR of ^{225}Ac -DOTA-BC8 which routinely binds to the cells at 64.8 \pm 2.14%.

[0302] Example 6 – ²²⁵Ac-BC8: Radioimmunotherapy for Multiple Myeloma

[0303] *Evaluation of the suitability of human H929 and U266 multiple myeloma cells as model cell lines for radioimmunotherapy (RIT) of multiple myeloma with ²²⁵Ac-DOTA-BC8*

[0304] The multiple myeloma (MM) cell lines H929 and U266 were purchased from the American Type Tissue Collection ATTC and grown according to the ATCC instructions. Binding of the unlabeled DOTA-BC8 to both cell lines was measured by flow cytometry (**FIG. 15A**), followed by the binding of ²²⁵Ac-DOTA-BC8 to the cells (**Fig. 15B**).

[0305] Subsequently, the H929 and U266 cells were used for an in vitro killing assay with ²²⁵Ac-DOTA-BC8. Two doses of ²²⁵Ac-DOTA-BC8 (20 pCi/ml and 250 pCi/ml) were used. The incubation of the cells with the radiolabeled antibodies was done in 96 well plates in a 200ul total volume. The same two doses of the control antibody ²²⁵Ac-DOTA-18B7 were used. The cells were washed from unbound radioactivity at 4- and 12-hour time points, and their survival was evaluated with a Trypan blue assay 3 days later (**Table 4**). The killing of both cell lines was antibody-specific and dose dependent. That is, both cell lines expressed a sufficient amount of CD45 on their surface for specific targeting by ²²⁵Ac-DOTA-BC8 and could be used for the subsequent in vivo experiments.

Table 4

Combined data for two experiments on treatment of H929 and U266 cells with ²²⁵Ac-DOTA-BC8 and tested for survival 3 days later

Treatment group	Surviving cells, %	
	4 hours	12 hours
U266, untreated	92±10	83±8
U266, ²²⁵ AcBC8, 20 pCi/mL	24±7	14±2
U266, ²²⁵ AcBC8, 250 pCi/ML	2±1	5±1
U266, ²²⁵ Ac18B7, 20 pCi/ML	75±8	78±10
U266, ²²⁵ Ac18B7, 250 pCi/ML	50±9	57±7
H929, untreated	97±12	92±10
H929, ²²⁵ AcBC8, 20 pCi/mL	33±7	22±6
H929, ²²⁵ AcBC8, 250 pCi/mL	12±3	6±1
H929, ²²⁵ Ac18B7, 20 pCi/mL	72±12	80±13
H929, ²²⁵ Ac18B7, 250 pCi/mL	60±11	55±14

[0306] Example 7 – ²²⁵Ac-BC8: Biodistribution

[0307] *Biodistribution of ²²⁵Ac-DOTA-BC8 in a naïve mouse model*

[0308] The purpose of this study was to evaluate the pharmacokinetic biodistribution of ²²⁵Ac-DOTA-BC8 versus a control ²²⁵Ac-DOTA-18B7 antibody in a naïve mouse model to ascertain baseline biodistribution and clearance in the absence of disease. The DOTA

conjugated BC8 antibody (batch A) and 18B7 antibody (produced at the same molar ratio as batch A; 7.5 moles DOTA to Ab) were radiolabeled with ^{225}Ac as detailed above. The antibodies were radiolabeled with the specific activity of $0.4 \mu\text{Ci}/\mu\text{g}$. The ^{225}Ac -DOTA-BC8 immunoreactivity was tested with the Cytotrol cells and demonstrated 55% binding, thus meeting the 50% minimal binding requirement.

[0309] Fifty (50) healthy CD-1 mice female mice were randomly assigned to two groups and injected intraperitoneally with either ^{225}Ac -DOTA-BC8 or control ^{225}Ac -DOTA-18B7. Each mouse received $5\mu\text{g}$ ($2\mu\text{Ci}$) of the radiolabeled antibody in $100\mu\text{l}$ of $0.15 \text{ M NH}_4\text{OAc}$ buffer with ascorbic acid. The intraperitoneal route is preferential to tail vein injections in the case of the long-lived radionuclides such as ^{225}Ac to avoid contamination of the personnel, animals, and the facility (i.e., due to the possible back pressure splash from the tails). According to our own and other groups data, intraperitoneally injected antibodies completely leave peritoneum within one-hour post injection. The mice were euthanized at 1, 4, 24, 48, and 96 hours ($n = 5$ mice per construct per time-point). Tissue samples (brain, muscle, bone (femur with the bone marrow), heart, lung, liver, spleen, kidneys, stomach, intestine, and blood) were collected from each mouse, weighed, and the accumulated activity per tissue counted in the gamma counter using ^{225}Ac energy window.

[0310] The percentages of injected dose per gram (ID/g, %) are shown in **FIGS. 16A** and **16B**. The results show that the patterns of the biodistribution and pharmacokinetic clearance of the two antibodies were very close to each other, which attests to the overall stability of the one-step labeled antibodies *in vivo*. The clearance from the blood and blood-rich organs and uptake of the control ^{225}Ac -DOTA-18B7 was somewhat lower, which is explained by the lack of homology between murine proteins and the 18B7 antigen (the fungal polysaccharide glucuronoxylomannan).

[0311] We also performed calculations to determine the ^{225}Ac -DOTA-BC8 and ^{225}Ac -DOTA-18B7 antibody half-lives in the blood and blood-rich organs (lungs and heart) using the data in **FIGS. 16A** and **16B**, and Prizm 5.0 software (GraphPad, San Diego, CA). The results are presented in **Table 5**, which show the half-life of ^{225}Ac -DOTA-BC8 to be around 100 hours (4.2 days), which is typical for a full-size murine IgG1 to a mammalian antigen, and the half-life of ^{225}Ac -DOTA-18B7 (also a murine IgG1) to be only 30 hours (1.25 days), likely because of the foreign nature of its respective antigen (the fungal polysaccharide). Thus, it appears that the radiolabeled antibodies are stable *in vivo*, clear fast from the blood and blood rich organs, and are suitable for use in subsequent pharmacokinetic experiments in CD45-positive tumor-bearing mice.

Table 5
 ^{225}Ac -DOTA-BC8 and ^{225}Ac -DOTA-18B7
 half-lives in the blood and blood-rich organs

Organ	Half-life, hours	
	^{225}Ac -BC8	^{225}Ac -18B7
blood	100 ± 5	30 ± 3
lungs	90 ± 3	20 ± 2
heart	104 ± 4	22 ± 4

[0312] Biodistribution of ^{225}Ac -DOTA-BC8 mAb in myeloma tumor-bearing SCID mice

[0313] The purpose of this study was to understand the biodistribution of ^{225}Ac -DOTA-BC8 versus a control ^{225}Ac -DOTA-18B7 antibody in a multiple myeloma SCID mouse model. The DOTA-conjugated BC8 (batch A) and 18B7 (as above) were radiolabeled with ^{225}Ac as above with the specific activity of 0.4 $\mu\text{Ci}/\mu\text{g}$. Their immunoreactivity was tested with the Cytotrol cells and demonstrated 61% binding, thus meeting the 50% minimal binding requirement.

[0314] Fifty (50) SCID-NOD (severe combined immunodeficiency non-obese diabetic) female 4-5 weeks old mice (Charles River Laboratories) were injected subcutaneously with 10^7 human multiple myeloma H929 cells (ATCC) into the right flank and with 10^7 human multiple myeloma U266 cells (ATCC) into the left flank. In approximately 20 days, when the tumors reached 3-4 mm in diameter, the mice were randomized into two 2 groups of 25 mice and injected intraorbitally with either ^{225}Ac -DOTA-BC8 or control ^{225}Ac -DOTA-18B7 mAb. Each mouse then received 0.4 μCi (1 μg) of the radiolabeled antibody in 50 μl of 0.15M NH_4OAc buffer with ascorbic acid. As indicated above, the intraorbital route is preferential to tail vein injections to avoid possible back pressure splash from the tails. The mice were euthanized at 1, 4, 24, 48, and 96 hours (n = 5 mice per construct per time-point). The tumors and tissue samples (brain, muscle, femur, bone marrow, heart, lung, liver, spleen, kidneys, stomach, intestine, and blood) were collected from each mouse, weighed, and the accumulated activity per tissue counted in the gamma counter using ^{225}Ac energy window.

[0315] The results are presented in **FIGS. 17A** and **17B** as a percentages of injected dose per gram (ID/g, %). The uptake of ^{225}Ac -DOTA-BC8 in the H929 and U266 tumors was significantly (P=0.01) higher than that of ^{225}Ac -DOTA-18B7. Both antibodies cleared quickly from the blood and blood rich organs. Importantly, there was no uptake of ^{225}Ac -

DOTA-BC8 in bone marrow, attesting to the lack of any homology to human CD45 in mouse bone marrow. These results show that ^{225}Ac -DOTA-BC8 localizes specifically in H929 and U266 tumors and thus could be used for further radioimmunotherapy (RIT) experiments.

[0316] Example 8 – ^{225}Ac -BC8: Radioimmunotherapy of tumors in mice

[0317] *Radioimmunotherapy (RIT) of H929 and U266 tumors in SCID-NOD mice with ^{225}Ac -DOTA-BC8*

[0318] The therapeutic potential of ^{225}Ac -DOTA-BC8 for the treatment of multiple myeloma xenografts in a mouse model was evaluated using forty (40) SCID-NOD female 4-5 weeks old mice. The mice were injected subcutaneously with 10^7 H929 (right flank) and U266 (left flank) human multiple myeloma cells as in the biodistribution experiments. In approximately 19 days, when the tumors reached 3-4 mm in diameter, the mice were randomized into the groups of 8 and treated intraorbitally with: 0.3 μCi ^{225}Ac -DOTA-BC8; 0.3 μCi ^{225}Ac -DOTA-18B7 control mAb; a matching amount of unlabeled BC8; or left untreated. The size of the tumors was measured on the day of treatment and every three days thereafter with the electronic calipers. The mice were monitored for their tumor size and well-being for 30 days.

[0319] The results of the RIT study are shown in **FIGS. 18A** and **18B**. There was a pronounced therapeutic effect of ^{225}Ac -DOTA-BC8 on both H929 and U266 tumors. The matching activity of the control ^{225}Ac -DOTA-18B7 had some effect on the tumor size, however, it was significantly ($P=0.02$) less than that of ^{225}Ac -DOTA-BC8. Thus, it is clear that the RIT of mice bearing multiple myeloma xenografts was effective in almost completely abrogating the tumor growth and not having any undesirable side effects.

[0320] *Histological analysis of the tumors post RIT*

[0321] At the completion of the RIT experiment, the mice were sacrificed, their tumors excised, placed in ethanol followed by buffered formaline, parafinized, cut into 5 μm sections and stained with H&E. **FIGS. 19A-19D** show the H929 and U266 tumors from the untreated and ^{225}Ac -DOTA-BC8 treated mice. The untreated tumors are much more coherent than the RIT treated tumors which show lack of coherence and necrosis.

[0322] Example 9 – ^{225}Ac -30F11: Marrow ablative effect of anti-CD45 surrogate

[0323] In this study we have evaluated the tolerability and myeloablative effects of ^{225}Ac -labelled anti-mouse pan-CD45 antibody clone 30F11 in mice for targeted conditioning prior to BMT. Thus, the dose-dependent myeloablative effects of ^{225}Ac -anti-CD45 antibody (30F11) on B6-Ly5^a mice was evaluated. Further, the study evaluated the extent of

engraftment and donor chimerism following congenic bone marrow transplant with B6-Ly5^b (CD45 allotype difference for monitoring chimerism).

[0324] *Experimental Methods:*

[0325] (1) *Conjugation and labeling of 30F11.*

[0326] The anti-CD45 antibody 30F11 was conjugated with the chelator DOTA as described above. To test if the DOTA-conjugated 30F11 retains immunoreactivity, cells shown to be CD45 positive were incubated with naked 30F11 and DOTA-30F11 and the amount of bound Ab was determined by flow cytometry using anti-ratIgG2b^{PE} to detect bound antibodies.

[0327] The DOTA-30F11 was radiolabeled with ¹¹¹In or ²²⁵Ac as described hereinabove to a specific activity of 5μCi/1μg (1:1) or 1μCi/1μg (1:1) antibody, respectively, and radiochemical purity of 99±1.

[0328] (1) *Biodistribution of anti-CD45 antibody 30F11 in C57Bl/6 mice.*

[0329] C57Bl/6 mice were injected i.v. with 60μg ¹¹¹In-30F11 with a specific activity of 5 μCi/μg. From 1 to 240 h after injection, the spleen was found to have the highest uptake of ¹¹¹In-labeled-30F11, followed by bone marrow and liver. Kidneys, ovaries, lungs and blood showed minimal uptake. The biodistribution for each organ was fitted to a time-activity curve to calculate the accumulated activity for each organ. The equilibrium dose constants of ²²⁵Ac was then applied to obtain the dose to organ per administered activity, reported in **Table 6**.

Table 6
²²⁵Ac-CD45 antibody
Absorbed dose to Organs

Organ	cGy/uCi
Blood	24.3
Spleen	8691.8
Liver	2310.6
Ovary	97.5
Kidnet	578.0
Lung	140.8
Bone Marrow	3963.5

[0330] (2) *Dose dependent tolerability of ²²⁵Ac-30F11 in B6-Ly5^a mice.*

[0331] In order to determine tolerability of ²²⁵Ac-CD45 antibody radio-conjugate, C57Bl/6 (5 per cohort) were treated with escalating doses of ²²⁵Ac-30F11 on day zero. Three

ascending dose levels (100nCi, 250nCi, and 500nCi) were administered in a total of 10ug (ca. 0.5 mg/kg) of 30F11 antibody (volume 100 to 200ul) injected into the tail vein (IV). Five untreated mice served as control for this study. Immediately prior to conditioning, pre-treatment blood samples were drawn from the control mice for baseline blood cell count measurements. Each of RBC and WBC were measured at weeks 1 and 2, and the mice euthanize at week 4. Blood was collected from euthanized mice & analyzed for liver & kidney toxicity i.e., blood urea nitrogen (BUN), creatine, alanine transaminase (ALT), and aspartate aminotransferase (AST) measured. A Kaplan-Meier graph showed that each of the 100nCi and 250nCi doses were well tolerated, while the 500nCi dose showed a decreased probability of survival after 1 week.

[0332] (3) *Safety profile of ²²⁵Ac-30F11 in B6-Ly5^a mice.*

[0333] In order to determine the safety profile of ²²⁵Ac-CD45 bone marrow engraftment, C57Bl/6 mice were treated with ²²⁵Ac-30F11 and reconstituted with donor bone marrow (CD45.1) as follows: 100nCi or 250nCi of ²²⁵Ac-30F11 was injected on Day 0 (as above). Four days after conditioning, half of the cohorts received congenic bone marrow (BMT) harvested from C57Bl/6-Ly5^b mice at a target density of marrow of 10⁷ nucleated cells injected via tail vein. Mice were regularly monitored for body weight and overt signs of changes in health and behavior.

[0334] Engraftment and donor chimerism was evaluated by blood collection, and mice were euthanized at 12 weeks. Engraftment was assessed by total WBC, RBC, HSC, neutrophil, and platelet counts, and BUN, creatinine, ALT, and AST.

[0335] *Results:* 500 nCi ²²⁵Ac-30F11 was found to be a maximum tolerated dose for this myeloablation modality. Mice treated with 250 and 100nCi of ²²⁵Ac-30F11 demonstrated effective myeloablative conditioning and donor BM engraftment in a dose dependent manner without any long-term hematological toxicity.

[0336] *Conclusion:* The pan-CD45-targeting antibody 30F11 armed with ²²⁵Ac appears to be a safe and potent targeted conditioning approach for BMT. This data supports the development of CD45 targeted ablation prior to BMT using ²²⁵Ac-armed antibodies.

[0337] **Example 10 – Lymphodepletion with ¹⁷⁷Lu-anti-CD45 and ¹³¹I-anti-CD45**

[0338] Prior to a patient receiving a dose of an adoptive cell transfer such as engineered autologous or allogeneic CAR-T cells, it is common to perform a lymphodepletion step often using high dose chemotherapy. This process is considered important to create sufficient space in the immune microenvironment, e.g. bone marrow, to allow the transferred cells to engraft. Further, it appears to elicit a favorable cytokine profile

for establishment and proliferation of the donor lymphocytes. In this study, use of the beta emitter ^{177}Lu (6.6 day half-life; 1.5 mm path length) for mediating effective lymphodepletion in mouse models is tested. Preclinical studies using a ^{177}Lu -labeled and ^{131}I -labeled surrogate anti-mouse pan-CD45 antibody (30F11) were performed to investigate in a mouse model the response of targeted RIT lymphodepletion on particular immune cell types and resulting changes in immune cytokine expression.

[0339] Following single dose administration of non-myeloablative doses of ^{177}Lu -CD45-RIT, peripheral blood, bone marrow and spleen samples were collected from 8-12 week C57Bl/6 mice at 96 hours and 10 days post-treatment for immunophenotyping to evaluate lymphoid and myeloid subsets for lymphodepletion, and serum for cytokine profiling. ^{177}Lu -CD45-RIT was shown to effectively lymphodeplete both lymphocyte and myeloid cells, inclusive of immune suppressive T regs and MDSCs. Studies evaluating this targeted lymphodepletion regimen in E.G7 lymphoma tumor bearing mice prior to adoptive cell transfer with OVA-specific CD8+ T cells will also be presented.

[0340] *Methods and materials*

[0341] The anti-mouse pan-CD45 antibody 30F11 was labeled with Lutetium-177 (^{177}Lu -CD45) and Iodine-131 (^{131}I -CD45) and used as a surrogate for radiolabeled pan-human BC8 to perform targeted lymphodepletion in mice. Immunoreactivity was confirmed in CD45+ cell-based binding assay to be > 95%.

[0342] For lymphodepletion studies in mice: Female adolescent C57Bl/6 mice were treated with 20ug of 30F11 labelled with 20 or 40 μCi of ^{177}Lu or 50 or 100 μCi of ^{131}I to determine the ability to selectively deplete immune cell subsets. Immune cell subset quantitation was measured by flow cytometry.

[0343] For lymphodepletion studies in OT I mouse model: Female adolescent C57Bl/6 CD45.1 mice were injected subcutaneously with OVA expressing CD45+ E.G7-OVA lymphoma tumor cells until 100mm³ tumor volume reached. Approximately 7 days post-tumor cell injection, mice were treated with ^{177}Lu -CD45 (40 μCi), ^{131}I -CD45 (100 μCi), or received no lymphodepletion treatment. Four days post-lymphodepletion, isolated CD8+ T cells isolated from CD45.2 OT I mice were administered to mice. Tumor volume and body weight were monitored, and mice were sacrificed when tumor volume exceeded 4000 mm³ or became necrotic.

[0344] *Results*

[0345] Anti-CD45 antibody was conjugated to DOTA at a ratio 20:1 and then labeled with ^{111}In at a ratio of 5:1. C57Bl/6 mice were injected i.p. with 60 μg ^{111}In -labeled anti-CD45

antibody with a specific activity of 5 $\mu\text{Ci}/\mu\text{g}$ and antibody distribution was monitored by microSPECT/CT at indicated time points. CD45 antibody homed to immune system organs: lymph nodes, spleen, and bone marrow (see **FIG. 20**).

[0346] The radiolabeled anti-CD45 antibodies ^{177}Lu -CD45 and ^{131}I -CD45 were found to transiently deplete CD45+ immune cell subsets without affecting platelets, red blood cells, or bone marrow cells. As shown in **FIG. 21**, treatment of non-tumor bearing C57B/6 mice with (A) 20 or 40 μCi ^{177}Lu -CD45 or (B) 50 or 100 μCi ^{131}I CD45 antibody was similarly effective in transiently lymphodepleting various immune cell populations without affecting bone marrow cells, red blood cells, or platelets.

[0347] Moreover, the ^{177}Lu radiolabeled anti-CD45 antibodies were found to transiently deplete CD45-expressing immune cell subsets in the spleen. As shown in **FIG. 22**, treatment of non-tumor bearing C57B/6 mice with 40 μCi ^{177}Lu -CD45 antibody was effective in transiently depleting various immune populations in the spleen including regulatory T cells (T-regs). This lymphodepletion enabled tumor control in an OT 1 adoptive cell therapy model.

[0348] As shown in **FIG. 23**, Following E.G7 tumor engraftment, mice either received no conditioning (Untreated and OT I) or were conditioned with 40 μCi ^{177}Lu -CD45 or 100 μCi of ^{131}I -CD45 on Day 0 and then received 1×10^6 OT I CD8+ CD45.2 OVA reactive T cells on day 4. Panel A shows results from ^{177}Lu -CD45 and ^{131}I -CD45-mediated targeted conditioning prior to adoptively transferred OT I T cells enabled control of EG.7 tumor growth. Panel B shows the tumor size for individual mice in each group. The OT1 T cell persistence and expansion was confirmed in mice at the time of sacrifice. Panel C shows the overall survival of control mice (i.e., received no conditioning or OT I T cells), mice who received OT 1 T cells, and those that also received the ^{177}Lu -CD45 and ^{131}I -CD45-mediated targeted conditioning.

[0349] *Conclusions*

[0350] These studies demonstrate the feasibility of using a low dose of ^{177}Lu -CD45 or ^{131}I -CD45 radioimmunotherapy as a transient non-myeloablative targeted lymphodepletion regimen prior to adoptive cell therapy. ^{111}In -CD45 imaging demonstrated that CD45 targeting delivers radiation selectively to immune privileged tissues. Studies determined that 40 μCi ^{177}Lu -CD45 or 100 μCi ^{131}I -CD45 could effectively deplete various immune cell subsets in mice but spare bone marrow cells, red blood cells, and platelets. In a model of adoptive cell therapy using CD45.1 OT1 mice bearing EG.7-OVA tumors, mice that received RIT-mediated lymphodepletion demonstrated enhanced tumor control over mice that did not

receive lymphodepletion. This data supports CD45 targeted lymphodepletion prior to adoptive cell therapy using a non-myeloablative dose of ^{131}I -CD45 or ^{177}Lu -CD45 RIT.

[0351] Example 11 – ^{225}Ac -BC8: Sickle Cell Disease (SCD)

[0352] This example describes HSC ablation (i.e., 100% depletion) preceding transplant with gene-edited HSCs in patients with SCD.

[0353] SCD is the most common hemoglobinopathy worldwide. The incidence of SCD among African Americans is approximately 1 in 500. It is estimated that 100,000 individuals are afflicted in the United States. SCD is caused by a single nucleotide mutation in the β -globin gene that produces sickle hemoglobin. SCD patients may exhibit anemia, vaso-occlusive crises (VOCs), hemolysis, chronic organ dysfunction, and early mortality. The mortality rate among children with SCD is 0.5 per 100,000. However, the mortality rate in adults is more than 2.5 per 100,000, and median life expectancy is less than 50 years of age for both men and women with SCD.

[0354] Currently, the only curative treatment for SCD is a hematopoietic stem cell transplant (HSCT). Unfortunately, HSCTs for SCD are not without problems. According to the Center for International Blood and Marrow Transplant Research, only 1,089 patients with SCD underwent HSCTs from 1991 to April 2017. Risks associated with HSCTs include complications (such as graft-versus-host disease) arising from the use of allogeneic donor stem cells.

[0355] With the advent of gene editing technologies, there is now an opportunity to cure SCD patients using autologous stem cells in which the mutation in the β -globin gene responsible for SCD has been corrected. ZFN, TALEN, CRISPR/cas9 and other nuclease-mediated editing approaches could be used to repair, or remove and replace, stem cells from an SCD patient. For example, Sun and Zhao (Biotech. And Bioeng., 2014, 111(5)) demonstrated the successful repair of the human β -globin gene mutation in patient pluripotent HSCs using TALENs. In addition, Dever, et al., (Nature, 2016, 539:384-389) demonstrated efficient repair of the Glu6Val mutation responsible for SCD in patient HSCs using CRISPR/cas9. Clinical trials using this approach for SCD are now starting.

[0356] Unfortunately, standard myeloablative conditioning regimens (i.e., 100% HSC-depleting regimens) using high dose chemotherapy or total body irradiation are currently used for transplants, including for autologous gene-edited cell transplants. There is a need for a safer and more effective conditioning method for these patients. Radiolabeled BC8 (e.g., ^{225}Ac -BC8) would be more sparing of a patient's normal tissues. Notably, older patients with SCD may already have organ damage as a result of their disease, and exposure

to non-specific radiation or chemotherapy as a myeloablative conditioning regimen could make performing a stem cell transplant even riskier. A radiolabeled BC8 approach presents a better option for these patients.

[0357] Further, due to the hereditary nature of the disease, correcting the disease through transplantation of gene-edited HSCs is preferred as early in life as possible, as complications of the disease may be irreversible and have a negative impact on long-term survival for the patient. As such, treating infants or young children afflicted with SCD using gene-edited HSCs is envisioned. To this end, radiolabeled BC8, particularly BC8 labeled with an alpha-emitting radionuclide such as ^{225}Ac , would be ideal. The use of an alpha-emitting radionuclide such as ^{225}Ac , with its very short, high energy radiation path length, would focus the radiation on CD45-positive cells and allow for effective myeloablation without the need to isolate treated patients (as would be required for conditioning with a myeloablative dose of a ^{131}I -BC8).

[0358] Example 12 – ^{225}Ac -BC8: Severe combined immunodeficiency (SCID)

[0359] This example describes HSC ablation preceding transplantation with gene-edited HSCs in patients with Severe combined immunodeficiency (SCID).

[0360] SCID is a germline genetic disorder in which afflicted patients present with severe T cell defects, with or without accompanying B cell defects. SCID involves a defective adaptive immune response that prevents patients from mounting an effective antibody response to pathogens. SCID is the most severe form of primary immunodeficiencies, and there are at least nine different known genes where mutations lead to SCID. Because SCID patients are incapable of mounting an adaptive immune response, they are susceptible to infection, and early mortality is high. SCID is also known as the “bubble boy” disease because patients must be kept in a sterile environment to avoid life-threatening infections.

[0361] The most frequent genetic defect in SCID is in the common gamma chain (γc), which is a protein that is shared by the receptors for interleukins IL-2, IL-4, IL-7, IL-9, IL-15 and IL-21. Other mutated genes that can lead to SCID are ADA and JAK3. As with SCD, only treatment with a stem cell transplant is potentially curative for SCID. However, delayed immune recovery and GVHD are significant risks for these patients. Also, as with SCD patients, SCID patients are young and therefore need effective and safe methods for treatment, including a better conditioning regimen prior to transplant.

[0362] Gene editing technology may precisely repair the defect in a SCID patient’s own HSCs. Once returned to the body, these engineered HSCs can produce normal

lymphocytes and establish a working adaptive immune response to protect against infection. Recently, Chang et al (Cell Reports, 2015, 12:1668-1677) reported effectively restoring normal lymphocyte development via CRISPR/cas9-mediated repair of a mutation in the JAK3 gene in mice. Further Alzubi, et al., (Nature, Scientific Reports, 2017, 7:12475) recently demonstrated using TALEN technology to precisely repair in mice a genetic defect in the IL2RG (common gamma chain), the gene responsible for X-SCID.

[0363] It is important that safer and more effective methods for conditioning human SCID patients are developed. Alpha-emitter CD45 radioimmunotherapy, such as with ²²⁵Ac-BC8, is needed to safely condition these predominantly young patients.

[0364] Example 13 – ²²⁵Ac-BC8: Treatment Synopsis for non-malignant disorders

[0365] **Table 7** summarizes selected treatment regimens using gene-edited stem cell administration preceded by HSC depletion via administration of an actinium radiolabeled BC8 antibody (i.e., conditioning agent; ²²⁵Ac-BC8).

Table 7

Disease	Therapy with Gene-edited HSCs or Pleuripotent Stem Cells Genes repaired include:
SCD	b-globin (HBB)
SCID	JAK3, Janus Family Kinase, ADA, adenosine deaminase, IL2RG, common gamma chain gene
β -Thalassemia	b-globin (HBB), BCL11A
Fanconi's Anemia	FANCC
Wiskott-Aldrich Syndrome	WAS
AIDS	CCR5 and CXCR4

CLAIMS

What is claimed is:

1. A method for treating a hemoglobinopathy or a hematological disease or disorder comprising:
administering a composition comprising an effective amount of a radiolabeled anti-CD45 antibody, wherein the effective amount is administered as a single dose, and wherein the radiolabeled anti-CD45 antibody comprises an actinium-225 labeled (²²⁵Ac) antibody or a lutetium-177 labeled (¹⁷⁷Lu) antibody.
2. The method of Claim 1, wherein the radiolabeled anti-CD45 antibody comprises ²²⁵Ac-BC8 having a light chain with the amino acid sequence set forth in SEQ ID NO:1, or a light chain with the N-terminal amino acid sequence as set forth in SEQ ID NO: 9.
3. The method of Claim 1, wherein the radiolabeled anti-CD45 antibody comprises ²²⁵Ac-BC8 having a light chain with at least one complementarity determining region having the amino acid sequence as set forth in SEQ ID NO:3, SEQ ID NO:4, or SEQ ID NO:5.
4. The method of Claim 1, wherein the radiolabeled anti-CD45 antibody comprises ²²⁵Ac-BC8 having a light chain with the amino acid sequence as set forth in SEQ ID NO:12 or SEQ ID NO:13.
5. The method of Claim 1, wherein the radiolabeled anti-CD45 antibody comprises ²²⁵Ac-BC8 having a heavy chain with the amino acid sequence set forth in SEQ ID NO:2, or a heavy chain with the N-terminal amino acid sequence as set forth in SEQ ID NO: 10.
6. The method of Claim 1, wherein the radiolabeled anti-CD45 antibody comprises ²²⁵Ac-BC8 having a heavy chain with at least one complementarity determining region having the amino acid sequence as set forth in SEQ ID NO:6, SEQ ID NO:7, or SEQ ID NO:8.
7. The method of Claim 1, wherein the radiolabeled anti-CD45 antibody comprises ²²⁵Ac-BC8 having a heavy chain with the amino acid sequence as set forth in SEQ ID NO:15 or SEQ ID NO:16.

8. The method of Claim 1, wherein the radiolabeled anti-CD45 antibody comprises ²²⁵Ac-BC8 with the amino acid ASP or ASN at position 141 (relative to the N-terminal amino acid).
9. The method of Claim 8, wherein a ratio of ASP:ASN in a population of BC8 proteins is 1:99 to 99:1, such as 10:90 to 90:10.
10. The method of Claim 1, further comprising:
administering an effective amount of a secondary agent.
11. The method of Claim 12, wherein the secondary agent comprises a radiosensitizer.
12. The method of Claim 15, wherein the radiosensitizer comprises a Bcl-2 inhibitor, an HDAC inhibitor, or a combination thereof.
13. The method of Claim 1, wherein the composition comprises an actinium-225 labeled BC8 antibody (²²⁵Ac-BC8) and an unlabeled anti-CD45 antibody, wherein a dose of each of the labelled BC8 and unlabeled anti-CD45 antibody are selected based on a patient specific characteristic selected from a patient weight, a patient age, a patient gender, and/or a patient health status.
14. The method of Claim 1, wherein the effective amount of the radiolabeled anti-CD45 antibody depletes at least 50% of lymphocytes of the subject but does not induce myeloablation in the subject.
15. The method of Claim 14, wherein the effective amount of the radiolabeled anti-CD45 antibody depletes circulating tumor cells of hematopoietic origin.
16. The method of Claim 14, wherein the effective amount of the radiolabeled anti-CD45 provides a radiation dose of 2 Gy or less to the bone marrow, or from at least 2 Gy to less than 8 Gy to the bone marrow.
17. The method of Claim 15, wherein the radiolabeled anti-CD45 comprises ²²⁵Ac-BC8, and the effective amount comprises a dose of 0.1 uCi/kg – 1.0 uCi/kg or a dose of 10 uCi – 150 uCi.

18. The method according to Claim 26, wherein the effective amount of the actinium-225 anti-CD45 depletes CD45+ circulating tumor cells but does not
19. The method of Claim 14, wherein the patient is in need of a bone marrow transplant.
20. The method of Claim 19, wherein the method further comprises performing the bone marrow transplant 4, 5, 6, 7, 8, or 9 days after the administration of the radiolabeled anti-CD45 antibody.
21. The method of Claim 1, wherein the effective amount of the radiolabeled anti-CD45 depletes regulatory T cells, myeloid derived suppressor cells, tumor associated macrophages, activated macrophages secreting IL-1 and/or IL-6, and combinations thereof.
22. The method of Claim 21, wherein the effective amount of the radiolabeled anti-CD45 provides a radiation dose of greater than 8 Gy to the bone marrow.
23. The method of Claim 16 or 22, further comprising:
administering to the subject an effective amount of a population of cells expressing a chimeric antigen receptor or a T-cell receptor (CAR/TCR) 6, 7, or 8 days after administration of the radiolabeled anti-CD45 antibody.
24. The method of claim 1, wherein the subject is afflicted with a non-cancerous disorder treatable via genetically edited cell therapy and is about to undergo such therapy to treat the disorder, and the effective amount of the radiolabeled anti-CD45 antibody is administered as a single dose.
25. The method of claim 24, wherein the disorder is selected from the group consisting of a hemoglobinopathy, a congenital immunodeficiency, and a viral infection.
26. The method of claim 24, wherein the disorder is selected from the group consisting of sickle cell disease (SCD), severe combined immunodeficiency disease (SCID), β -thalassemia and Fanconi's anemia.
27. The method of claim 26, wherein the disorder is SCD and the therapy is genetically edited β -globin hematopoietic stem cell therapy.

28. The method of claim 27, wherein the disorder is SCID and the therapy is genetically edited hematopoietic stem cell therapy, wherein the edited gene is selected from the group consisting of the common gamma chain (γ c) gene, the adenosine deaminase (ADA) gene and the Janus kinase 3 (JAK3) gene.
29. The method of Claim 1, wherein the hematological disease is myelodysplastic syndrome (MDS), multiple myeloma (MM), acute myeloid leukemia (AML), myeloproliferative neoplasm, or a combination thereof.

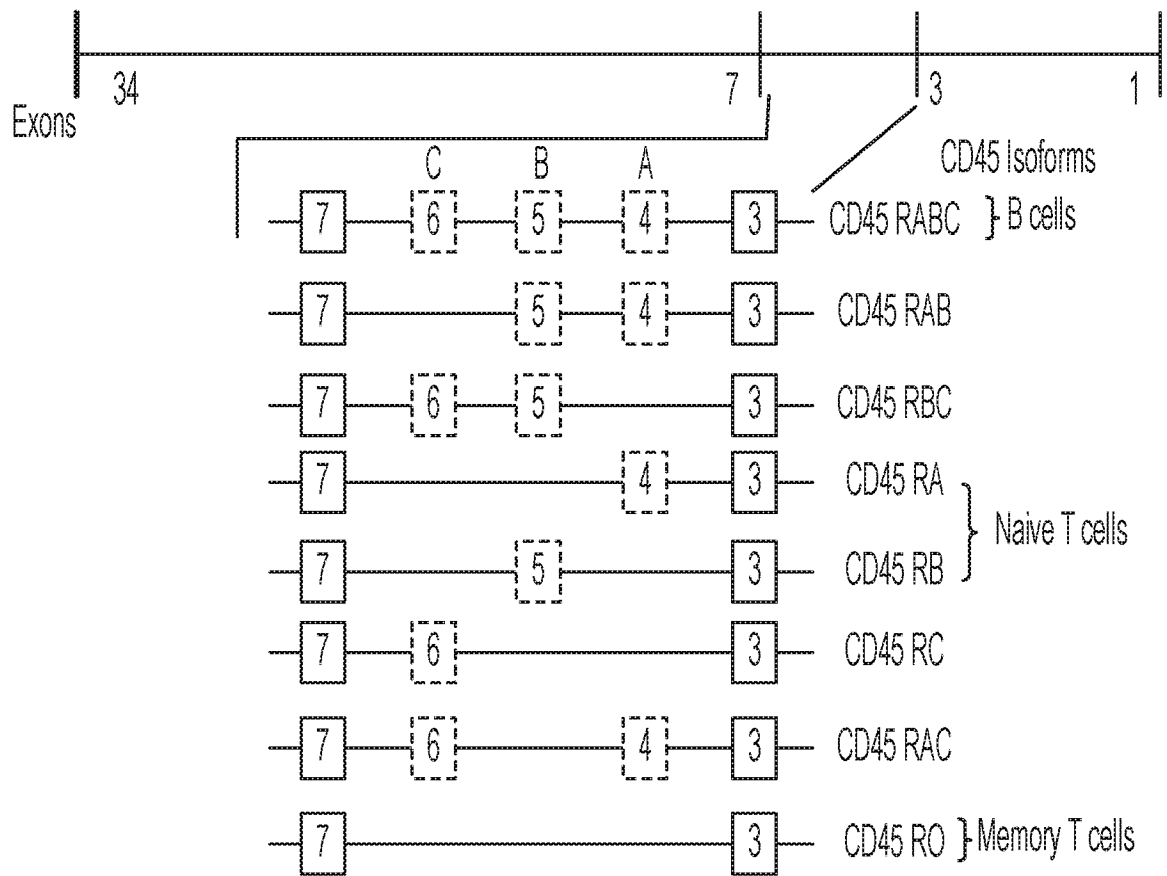


FIG. 1
PRIOR ART

FIG. 2

Variable Light Chain (VL)	DIALTQSPASLAVSLGQRATISCRASKSVSTSGYSYLHWYQOKPGQPPKLLIYLAS <u>NLESGV</u> PARFSGSGSGTDFTLNIHPVEEEDAATYYC <u>QHSRELPFT</u> FGSGTKLEIK (Sequence ID No: 1)
Variable Heavy Chain (VH)	EVKLLLESGGGLVQPGGSLKLSCAASGFDFSRYWMSWVRQAPGKGLEWIGEINPTSS <u>TINFTPSLKD</u> KVFI SRDNAKNTLYLQMSKVRSEDTALYYCARG <u>GNYYRYGDAMDY</u> WG QGTSVTVSSAK (Sequence ID No: 2)

FIG. 3

The Light Chain CDRs:

LC-CDR-1: RASKSVSTSGYSYLH (Sequence ID NO:3)
 LC-CDR-2: LASNLES (Sequence ID NO:4)
 LC-CDR-3: QHSRELPFT (Sequence ID NO:5)

The Heavy Chain CDRs:

HC-CDR-1: GFDFSRYWMS (Sequence ID NO:6)
 HC-CDR-2: EINPTSSTINFTPSLKD (Sequence ID NO:7)
 HC-CDR-3: GNYYRYGDAMDY (Sequence ID NO:8)

The Light Chain N-term: DIALTQS (Sequence ID NO:9)

The Heavy Chain N-term: EVKLLLES (Sequence ID NO:10)

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FIG. 4A

	Leader Sequence														
1	M	E	T	D	T	L	L	L	W	V	L	L	L	W	V
	ATG	GAG	ACA	GAC	ACA	CTC	CTG	TTA	TGG	GTA	CTG	CTG	CTC	TGG	GTT
	Leader Sequence							FR1							
46	F	G	S	T	G	D	I	A	L	T	Q	S	P	A	S
	CCA	GGT	TCC	ACT	GGT	GAC	ATT	GCG	CTG	ACA	CAG	TCT	CCT	GCT	TCC
	FR1											CDR1			
91	L	A	V	S	L	G	Q	R	A	T	I	S	C	R	A
	TTA	GCT	GTA	TCT	CTG	GGA	CAG	AGG	GCC	ACC	ATC	TCA	TGC	AGG	GCC
	CDR1											FR2			
136	S	K	S	V	S	T	S	G	Y	S	Y	L	H	W	Y
	AGC	AAA	AGT	GTC	AGT	ACA	TCT	GGC	TAT	AGT	TAT	CTG	CAC	TGG	TAC
	FR2											CDR2			
181	Q	Q	K	P	G	Q	P	P	K	L	L	I	Y	L	A
	CAA	CAG	AAA	CCA	GGA	CAG	CCA	CCC	AAA	CTC	CTC	ATC	TAT	CTT	GCA
	CDR2							FR3							
226	S	N	L	E	S	G	V	P	A	R	F	S	G	S	G
	TCC	AAC	CTA	GAA	TCT	GGG	GTC	CCT	GCC	AGG	TTC	AGT	GGC	AGT	GGG
	FR3											CDR3			
271	S	G	T	D	F	T	L	N	I	H	P	V	E	E	E
	TCT	GGG	ACA	GAC	TTC	ACC	CTC	AAC	ATC	CAT	CCT	GTG	GAG	GAG	GAG
	FR3							CDR3							
316	D	A	A	T	Y	Y	C	Q	H	S	R	E	L	P	F
	GAT	GCT	GCA	ACC	TAT	TAC	TGT	CAG	CAC	AGT	AGG	GAG	CTT	CCA	TTC
	CDR3			FR4							C kappa				
361	T	F	G	S	G	T	K	L	E	I	K	R	A	D	A
	ACG	TTC	GGC	TCG	GGG	ACA	AAG	TTG	GAA	ATA	AAA	CGG	GCT	GAT	GCT
	C kappa														
406	A	P	T	V	S	I	F	P	P	S	S	E	Q	L	T
	GCA	CCA	ACT	GTA	TCC	ATC	TTC	CCA	CCA	TCC	AGT	GAG	CAG	TTA	ACA
	C kappa														
451	S	G	G	A	S	V	V	C	F	L	N	N	F	Y	P
	TCT	GGA	GGT	GCC	TCA	GTC	GTG	TGC	TTC	TTG	AAC	AAC	TTC	TAC	CCC
	C kappa														
496	K	D	I	N	V	K	W	K	I	D	G	S	E	R	Q
	AAA	GAC	ATC	AAT	GTC	AAG	TGG	AAG	ATT	GAT	GGC	AGT	GAA	CGA	CAA
	C kappa														
541	N	G	V	L	N	S	W	T	D	Q	D	S	K	D	S
	AAT	GGC	GTC	CTG	AAC	AGT	TGG	ACT	GAT	CAG	GAC	AGC	AAA	GAC	AGC

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FIG. 4A (continued)

		<i>C kappa</i>													
586	T ACC	Y TAC	S AGC	M ATG	S AGC	S AGC	T ACC	L CTC	T ACG	L TTG	T ACC	K AAG	D GAC	E GAG	Y TAT
		<i>C kappa</i>													
631	E GAA	R CGA	H CAT	N AAC	S AGC	Y TAT	T ACC	C TGT	E GAG	A GCC	T ACT	H CAC	K AAG	T ACA	S TCA
		<i>C kappa</i>													<i>Stop</i>
676	T ACT	S TCA	P CCC	I ATT	V GTC	K AAG	S AGC	F TTC	N AAC	R AGG	N AAT	E GAG	C TGT	*	TAG

SEQ ID NO:11 Nucleotide Sequence

SEQ ID NO:12 Amino Acid Sequence

FIG. 4B

N-Terminal

```

Asp Ile Ala Leu Thr Gln Ser Pro Ala Ser Leu Ala Val Ser Leu Gly Gln Arg Ala Thr
1           5           10           15           20

Ile Ser Cys Arg Ala Ser Lys Ser Val Ser Thr Ser Gly Tyr Ser Tyr Leu His Trp Tyr
21          25          30          35          40
          | CDR1

Gln Gln Lys Pro Gly Gln Pro Pro Lys Leu Leu Ile Tyr Leu Ala Ser Asn Leu Glu Ser
41          45          50          55          60
          | CDR2

Gly Val Pro Ala Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Asn Ile His
61          65          70          75          80

Pro Val Glu Glu Glu Asp Ala Ala Thr Tyr Tyr Cys Gln His Ser Arg Glu Leu Pro Phe
81          85          90          95          100
          | CDR3

Thr Phe Gly Ser Gly Thr Lys Leu Glu Ile Lys Arg Ala Asp Ala Ala Pro Thr Val Ser
101         105         110         115         120

Ile Phe Pro Pro Ser Ser Glu Gln Leu Thr Ser Gly Gly Ala Ser Val Val Cys Phe Leu
121         125         130         135         140

Asn Asn Phe Tyr Pro Lys Asp Ile Asn Val Lys Trp Lys Ile Asp Gly Ser Glu Arg Gln
141         145         150         155         160

Asn Gly Val Leu Asn Ser Trp Thr Asp Gln Asp Ser Lys Asp Ser Thr Tyr Ser Met Ser
161         165         170         175         180

Ser Thr Leu Thr Leu Thr Lys Asp Glu Tyr Glu Arg His Asn Ser Tyr Thr Cys Glu Ala
181         185         190         195         200

                                     C-Terminal
Thr His Lys Thr Ser Thr Ser Pro Ile Val Lys Ser Phe Asn Arg Asn Glu Cys
201         205         210         215         218
    
```

SEQ ID NO:13 Amino Acid Sequence

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FIG. 5A

	<i>Leader Sequence</i>														
1	M	D	F	G	L	I	F	F	I	V	A	L	L	K	G
	ATG	GAT	TTT	GGG	CTG	ATT	TTT	TTT	ATT	GTT	GCT	CTT	TTA	AAA	GGG
	<i>Leader Seq.</i>					<i>FR1</i>									
46	V	Q	C	E	V	K	L	L	E	S	G	G	G	L	V
	GTC	CAG	TGT	GAG	GTG	AAG	CTT	CTC	GAG	TCT	GGA	GGT	GGC	CTG	GTG
	<i>FR1</i>														
91	Q	P	G	G	S	L	K	L	S	C	A	A	S	G	F
	CAG	CCT	GGA	GGA	TCC	CTG	AAA	CTC	TCC	TGT	GCA	GCC	TCA	GGA	TTC
	<i>FR1</i>			<i>CDR1</i>					<i>FR2</i>						
136	D	F	S	R	Y	W	M	S	W	V	R	Q	A	P	G
	GAT	TTC	AGT	AGA	TAC	TGG	ATG	AGT	TGG	GTC	CGG	CAG	GCT	CCA	GGG
	<i>FR2</i>							<i>CDR2</i>							
181	K	G	L	E	W	I	G	E	I	N	P	T	S	S	T
	AAA	GGG	CTA	GAA	TGG	ATT	GGA	GAG	ATT	AAT	CCA	ACT	AGC	AGT	ACG
	<i>CDR2</i>							<i>FR3</i>							
226	I	N	F	T	P	S	L	K	D	K	V	F	I	S	R
	ATA	AAC	TTT	ACG	CCA	TCT	CTA	AAG	GAT	AAA	GTC	TTC	ATC	TCC	AGA
	<i>FR3</i>														
271	D	N	A	K	N	T	L	Y	L	Q	M	S	K	V	R
	GAC	AAC	GCC	AAA	AAT	ACG	CTG	TAC	CTG	CAA	ATG	AGC	AAA	GTG	AGA
	<i>FR3</i>										<i>CDR3</i>				
316	S	E	D	T	A	L	Y	Y	C	A	R	G	N	Y	Y
	TCT	GAG	GAC	ACA	GCC	CTT	TAT	TAC	TGT	GCA	AGA	GGG	AAC	TAC	TAT
	<i>CDR3</i>								<i>FR4</i>						
361	R	Y	G	D	A	M	D	Y	W	G	Q	G	T	S	V
	AGG	TAC	GGA	GAT	GCT	ATG	GAC	TAC	TGG	GGT	CAA	GGA	ACC	TCA	GTC
	<i>FR4</i>					<i>Constant region</i>									
406	T	V	S	S	A	K	T	T	P	P	S	V	Y	P	L
	ACC	GTC	TCC	TCA	GCC	AAA	ACG	ACA	CCC	CCA	TCT	GTC	TAT	CCA	CTG
	<i>Constant region</i>														
451	A	P	G	S	A	A	Q	T	D141	S	M	V	T	L	G
	GCC	CCT	GGA	TCT	GCT	GCC	CAA	ACT	AAC	TCC	ATG	GTG	ACC	CTG	GGA
	<i>Constant region</i>														
496	C	L	V	K	G	Y	F	P	E	P	V	T	V	T	W
	TGC	CTG	GTC	AAG	GGC	TAT	TTC	CCT	GAG	CCA	GTG	ACA	GTG	ACC	TGG
	<i>Constant region</i>														
541	N	S	G	S	L	S	S	G	V	H	T	F	P	A	V
	AAC	TCT	GGA	TCC	CTG	TCC	AGC	GGT	GTG	CAC	ACC	TTC	CCA	GCT	GTC

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FIG. 5A (continued)

	<i>Constant region</i>														
586	L	Q	S	D	L	Y	T	L	S	S	S	V	T	V	P
	CTG	CAG	TCT	GAC	CTC	TAC	ACT	CTG	AGC	AGC	TCA	GTG	ACT	GTC	CCC
	<i>Constant region</i>														
631	S	S	T	W	P	S	E	T	V	T	C	N	V	A	H
	TCC	AGC	ACC	TGG	CCC	AGC	GAG	ACC	GTC	ACC	TGC	AAC	GTT	GCC	CAC
	<i>Constant region</i>														
676	P	A	S	S	T	K	V	D	K	K	I	V	P	R	D
	CCG	GCC	AGC	AGC	ACC	AAG	GTG	GAC	AAG	AAA	ATT	GTG	CCC	AGG	GAT
	<i>Constant region</i>														
721	C	G	C	K	P	C	I	C	T	V	P	E	V	S	S
	TGT	GGT	TGT	AAG	CCT	TGC	ATA	TGT	ACA	GTC	CCA	GAA	GTA	TCA	TCT
	<i>Constant region</i>														
766	V	F	I	F	P	P	K	P	K	D	V	L	T	I	T
	GTC	TTC	ATC	TTC	CCC	CCA	AAG	CCC	AAG	GAT	GTG	CTC	ACC	ATT	ACT
	<i>Constant region</i>														
811	L	T	P	K	V	T	C	V	V	V	D	I	S	K	D
	CTG	ACT	CCT	AAG	GTC	ACG	TGT	GTT	GTG	GTA	GAC	ATC	AGC	AAG	GAT
	<i>Constant region</i>														
856	D	P	E	V	Q	F	S	W	F	V	D	D	V	E	V
	GAT	CCC	GAG	GTC	CAG	TTC	AGC	TGG	TTT	GTA	GAT	GAT	GTG	GAG	GTG
	<i>Constant region</i>														
901	H	T	A	Q	T	Q	P	R	E	E	Q	F	N	S	T
	CAC	ACA	GCT	CAG	ACG	CAA	CCC	CGG	GAG	GAG	CAG	TTC	AAC	AGC	ACT
	<i>Constant region</i>														
946	F	R	S	V	S	E	L	P	I	M	H	Q	D	W	L
	TTC	CGC	TCA	GTC	AGT	GAA	CTT	CCC	ATC	ATG	CAC	CAG	GAC	TGG	CTC
	<i>Constant region</i>														
991	N	G	K	E	F	K	C	R	V	N	S	A	A	F	P
	AAT	GGC	AAG	GAG	TTC	AAA	TGC	AGG	GTC	AAC	AGT	GCA	GCT	TTC	CCT
	<i>Constant region</i>														
1036	A	P	I	E	K	T	I	S	K	T	K	G	R	P	K
	GCC	CCC	ATC	GAG	AAA	ACC	ATC	TCC	AAA	ACC	AAA	GGC	AGA	CCG	AAG
	<i>Constant region</i>														
1081	A	P	Q	V	Y	T	I	P	P	P	K	E	Q	M	A
	GCT	CCA	CAG	GTG	TAC	ACC	ATT	CCA	CCT	CCC	AAG	GAG	CAG	ATG	GCC
	<i>Constant region</i>														
1126	K	D	K	V	S	L	T	C	M	I	T	D	F	F	P
	AAG	GAT	AAA	GTC	AGT	CTG	ACC	TGC	ATG	ATA	ACA	GAC	TTC	TTC	CCT

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FIG. 5A (continued)

	<i>Constant region</i>														
1171	E	D	I	T	V	E	W	Q	W	N	G	Q	P	A	E
	GAA	GAC	ATT	ACT	GTG	GAG	TGG	CAG	TGG	AAT	GGG	CAG	CCA	GCG	GAG
	<i>Constant region</i>														
1216	N	Y	K	N	T	Q	P	I	M	D	T	D	G	S	Y
	AAC	TAC	AAG	AAC	ACT	CAG	CCC	ATC	ATG	GAC	ACA	GAT	GGC	TCT	TAC
	<i>Constant region</i>														
1261	F	V	Y	S	K	L	N	V	Q	K	S	N	W	E	A
	TTC	GTC	TAC	AGC	AAG	CTC	AAT	GTG	CAG	AAG	AGC	AAC	TGG	GAG	GCA
	<i>Constant region</i>														
1306	G	N	T	F	T	C	S	V	L	H	E	G	L	H	N
	GGA	AAT	ACT	TTC	ACC	TGC	TCT	GTG	TTA	CAT	GAG	GGC	CTG	CAC	AAC
	<i>Constant region</i>													<i>Stop</i>	
1351	H	H	T	E	K	S	L	S	H	S	P	G	K	*	
	CAC	CAT	ACT	GAG	AAG	AGC	CTC	TCC	CAC	TCT	CCT	GGT	AAA	TGA	

SEQ ID NO:14 Nucleotide Sequence

SEQ ID NO:15 Amino Acid Sequence

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FIG. 5B

N-Terminal

Glu	Val	Lys	Leu	Leu	Glu	Ser	Gly	Gly	Gly	Leu	Val	Gln	Pro	Gly	Gly	Ser	Leu	Lys	Leu
1				5					10					15					20
										CDR1									
Ser	Cys	Ala	Ala	Ser	Gly	Phe	Asp	Phe	Ser	<u>Arg</u>	<u>Tyr</u>	<u>Trp</u>	<u>Met</u>	<u>Ser</u>	Trp	Val	Arg	Gln	Ala
21				25					30					35					40
										CDR2									
Pro	Gly	Lys	Gly	Leu	Glu	Trp	Ile	Gly	<u>Glu</u>	<u>Ile</u>	<u>Asn</u>	<u>Pro</u>	<u>Thr</u>	<u>Ser</u>	<u>Ser</u>	<u>Thr</u>	<u>Ile</u>	<u>Asn</u>	<u>Phe</u>
41				45					50					55					60
<u>Thr</u>	<u>Pro</u>	<u>Ser</u>	<u>Leu</u>	<u>Lys</u>	<u>Asp</u>	Lys	Val	Phe	Ile	Ser	Arg	Asp	Asn	Ala	Lys	Asn	Thr	Leu	Tyr
61				65					70					75					80
										CDR3									
Leu	Gln	Met	Ser	Lys	Val	Arg	Ser	Glu	Asp	Thr	Ala	Leu	Tyr	Tyr	Cys	Ala	Arg	<u>Gly</u>	<u>Asn</u>
81				85					90					95					100
<u>Tyr</u>	<u>Tyr</u>	<u>Arg</u>	<u>Tyr</u>	<u>Gly</u>	<u>Asp</u>	<u>Ala</u>	<u>Met</u>	<u>Asp</u>	<u>Tyr</u>	Trp	Gly	Gln	Gly	Thr	Ser	Val	Thr	Val	Ser
101				105					110					115					120
Ser	Ala	Lys	Thr	Thr	Pro	Pro	Ser	Val	Tyr	Pro	Leu	Ala	Pro	Gly	Ser	Ala	Ala	Gln	Thr
121				125					130					135					140
Asp	Ser	Met	Val	Thr	Leu	Gly	Cys	Leu	Val	Lys	Gly	Tyr	Phe	Pro	Glu	Pro	Val	Thr	Val
141				145					150					155					160
Thr	Trp	Asn	Ser	Gly	Ser	Leu	Ser	Ser	Gly	Val	His	Thr	Phe	Pro	Ala	Val	Leu	Gln	Ser
161				165					170					175					180
Asp	Leu	Tyr	Thr	Leu	Ser	Ser	Ser	Val	Thr	Val	Pro	Ser	Ser	Thr	Trp	Pro	Ser	Glu	Thr
181				185					190					195					200
Val	Thr	Cys	Asn	Val	Ala	His	Pro	Ala	Ser	Ser	Thr	Lys	Val	Asp	Lys	Lys	Ile	Val	Pro
201				205					210					215					220
Arg	Asp	Cys	Gly	Cys	Lys	Pro	Cys	Ile	Cys	Thr	Val	Pro	Glu	Val	Ser	Ser	Val	Phe	Ile
221				225					230					235					240
Phe	Pro	Pro	Lys	Pro	Lys	Asp	Val	Leu	Thr	Ile	Thr	Leu	Thr	Pro	Lys	Val	Thr	Cys	Val
241				245					250					255					260

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FIG. 6A

Human IgG1	ASTKGPSVFPLAPSSKSTSGGTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAVLQSS GLYSLSSVWVTPSSSLGTQTYICNVNHKPSNTKVDKKVEPKSCDKTHTCPPCPAPELLGG PSVFLFPPKPKDTLMISRTPEVTCVVDVSHEDPEVKFNWYVDGVEVHNAKTKPREEQYN STYRVVSVLTVLHQDWLNGKEYKCKVSNKALPAPIEKTISKAKGQPREPQVYTLPPSRDE LTKNQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTTPVLDSDGSFFLYSKLTVDKSRW QQGNVVFSCSVMHREALHNHYTQKSLSLSPGK
------------	---

(Sequence ID No: 17)

FIG. 6B

Human IgG2	ASTKGPSVFPLAPCSRSTSESTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAVLQSS GLYSLSSVWVTPSSNFGTQTYTCNVDHKPSNTKVDKTVKCCVECPAPPVAGPSVF LFPPKPKDTLMISRTPEVTCVVDVSHEDPEVQFNWYVDGVEVHNAKTKPREEQFNSTFR VVSVLTVLHQDWLNGKEYKCKVSNKGLPAPIEKTISKAKGQPREPQVYTLPPSREEMTKN QVSLTCLVKGFYPSDISVEWESNGQPENNYKTTTPMLDSDGSFFLYSKLTVDKSRWQQGN VFSCSVMHREALHNHYTQKSLSLSPGK
------------	---

(Sequence ID No: 18)

FIG. 6C

Human IgG4	ASTKGPSVFPLAPCSRSTSESTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAVLQSS GLYSLSSVWVTPSSSLGKTYTCNVDHKPSNTKVDKRVESKYGPPCPAPEFLGGPSV FLFPPKPKDTLMISRTPEVTCVVDVSDQEDPEVQFNWYVDGVEVHNAKTKPREEQFNSTY RVVSVLTVLHQDWLNGKEYKCKVSNKGLPSSIEKTISKAKGQPREPQVYTLPPSQQEEMTK NQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTTPVLDSDGSFFLYSRLTVDKSRWQEG NVFSCSVMHREALHNHYTQKSLSLSLGGK
------------	--

(Sequence ID No: 19)

FIG. 6D

Human IgG4 with S228P	ASTKGPSVFPLAPCSRSTSESTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAVLQSS GLYSLSSVWVTPSSSLGKTYTCNVDHKPSNTKVDKRVESKYGPPCPAPEFLGGPSV FLFPPKPKDTLMISRTPEVTCVVDVSDQEDPEVQFNWYVDGVEVHNAKTKPREEQFNSTY RVVSVLTVLHQDWLNGKEYKCKVSNKGLPSSIEKTISKAKGQPREPQVYTLPPSQQEEMTK NQVSLTCLVKGFYPSDIAVEWESNGQPENNYKTTTPVLDSDGSFFLYSRLTVDKSRWQEG NVFSCSVMHREALHNHYTQKSLSLSLGGK
--------------------------	--

(Sequence ID No: 20)

FIG. 6E

Human Kappa Light Chain Constant Region	RTVAAPSVFIFPPSDEQLKSGTASVVCLLNNFYPREAKVQWKVDNALQSGNSQESVTEQD SKDSTYLSSTLTLSKADYEEKHKVYACEVTHQGLSSPVTKSFNRGEC
---	---

(Sequence ID No: 21)

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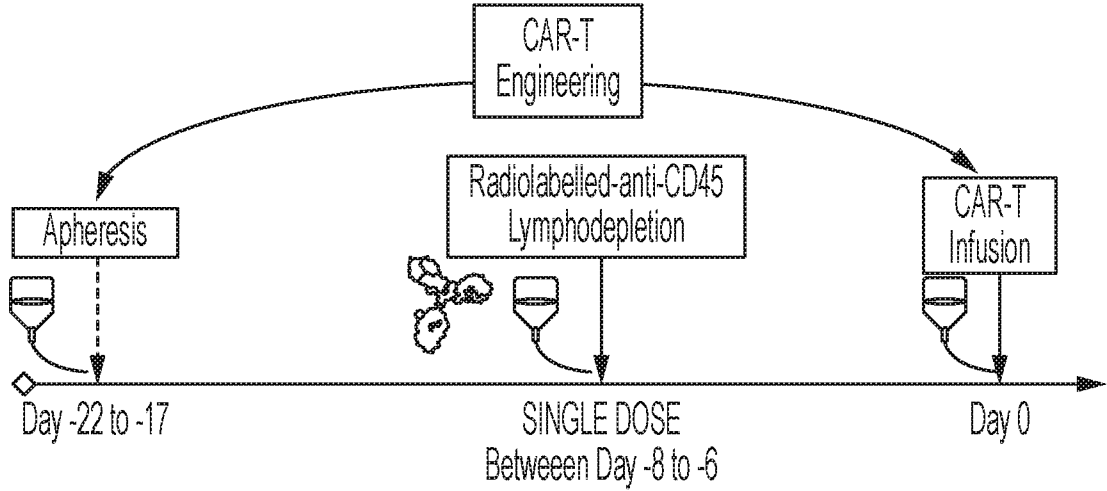


FIG. 7

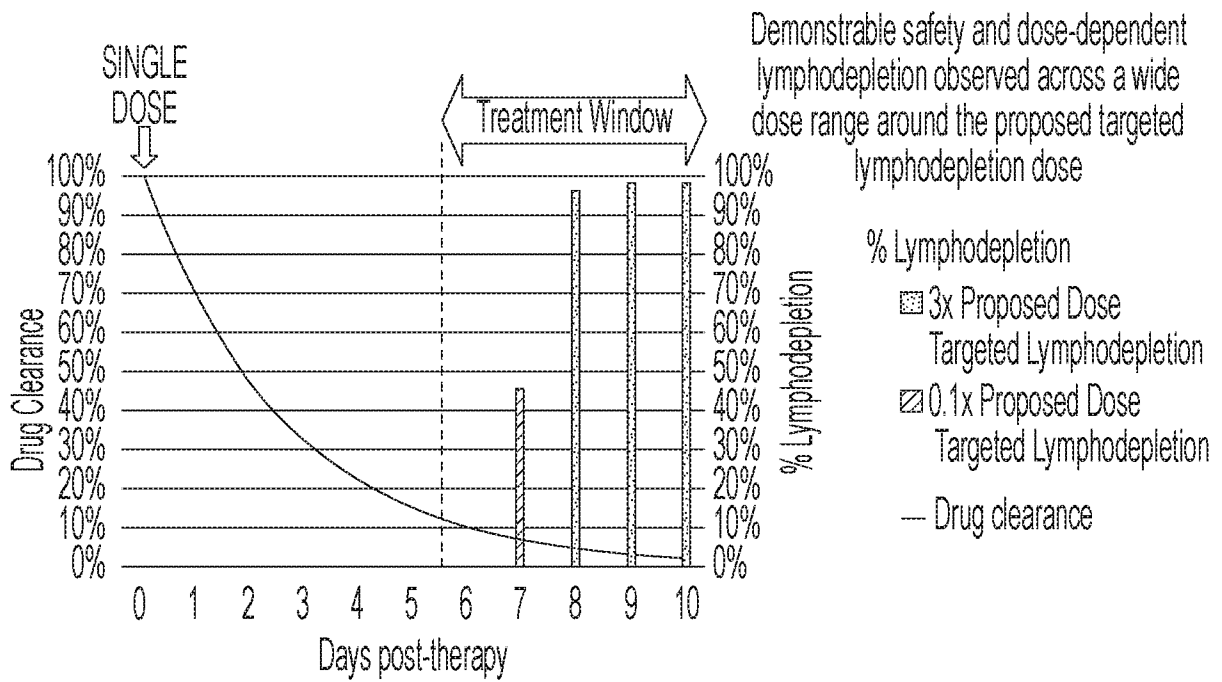


FIG. 8

FIG. 9A

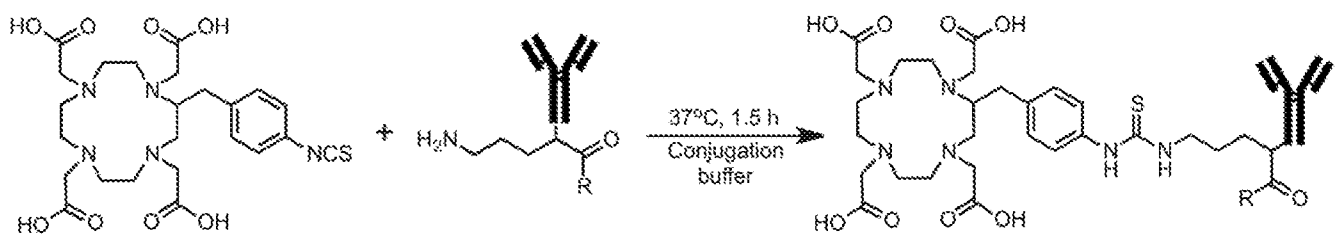
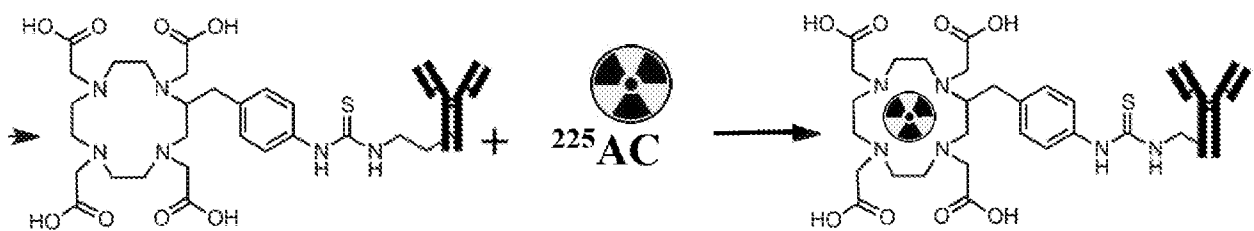


FIG. 9B



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FIG. 10A

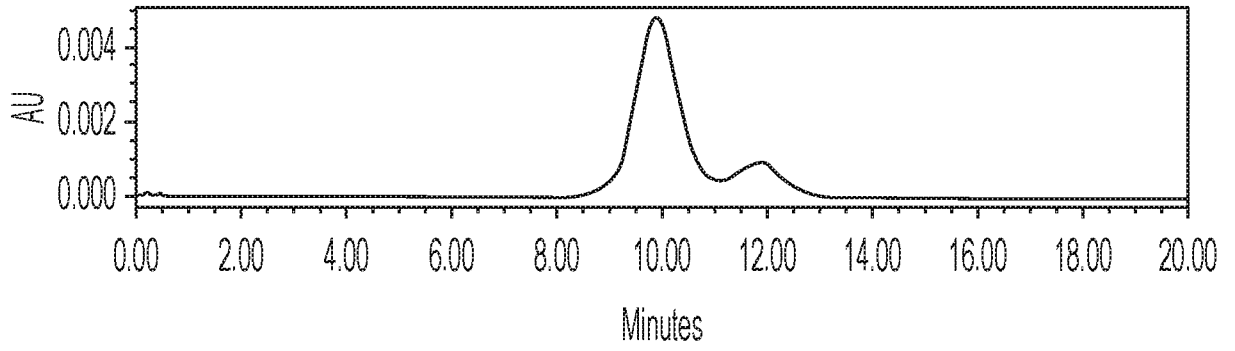
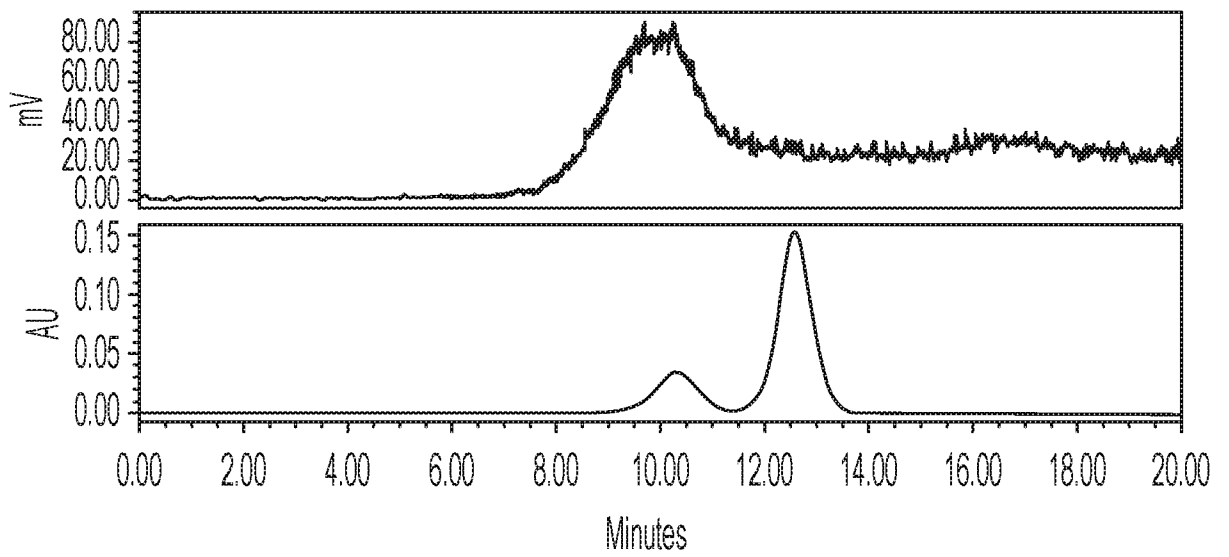


FIG. 10B



Stability of labeling in different volumes and at different temperatures

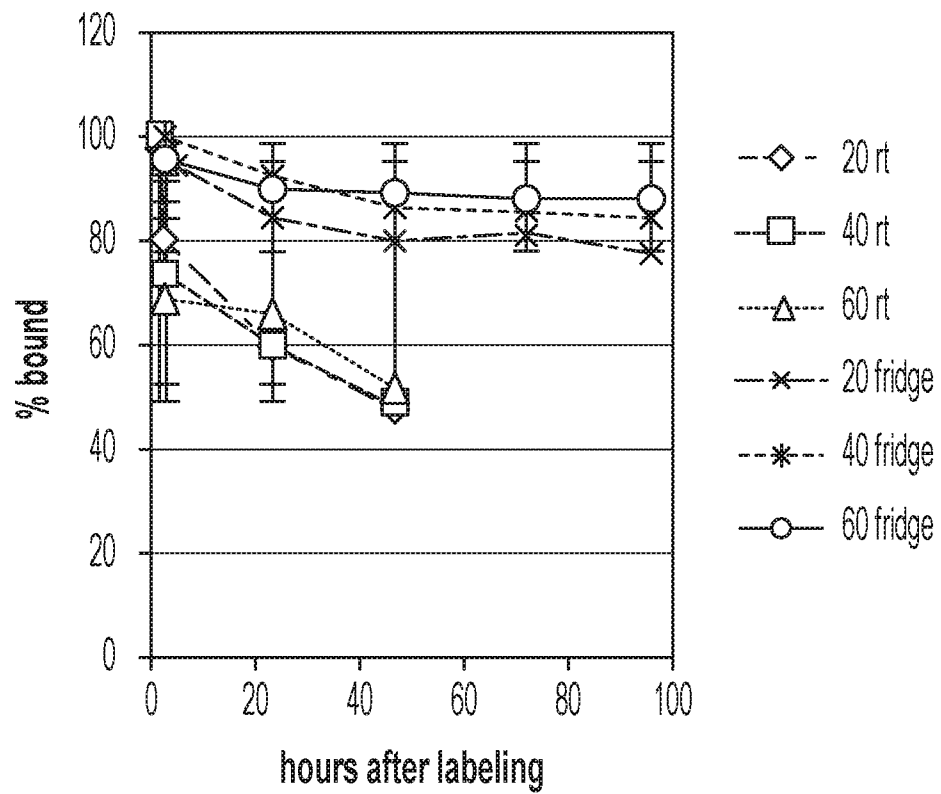


FIG. 11

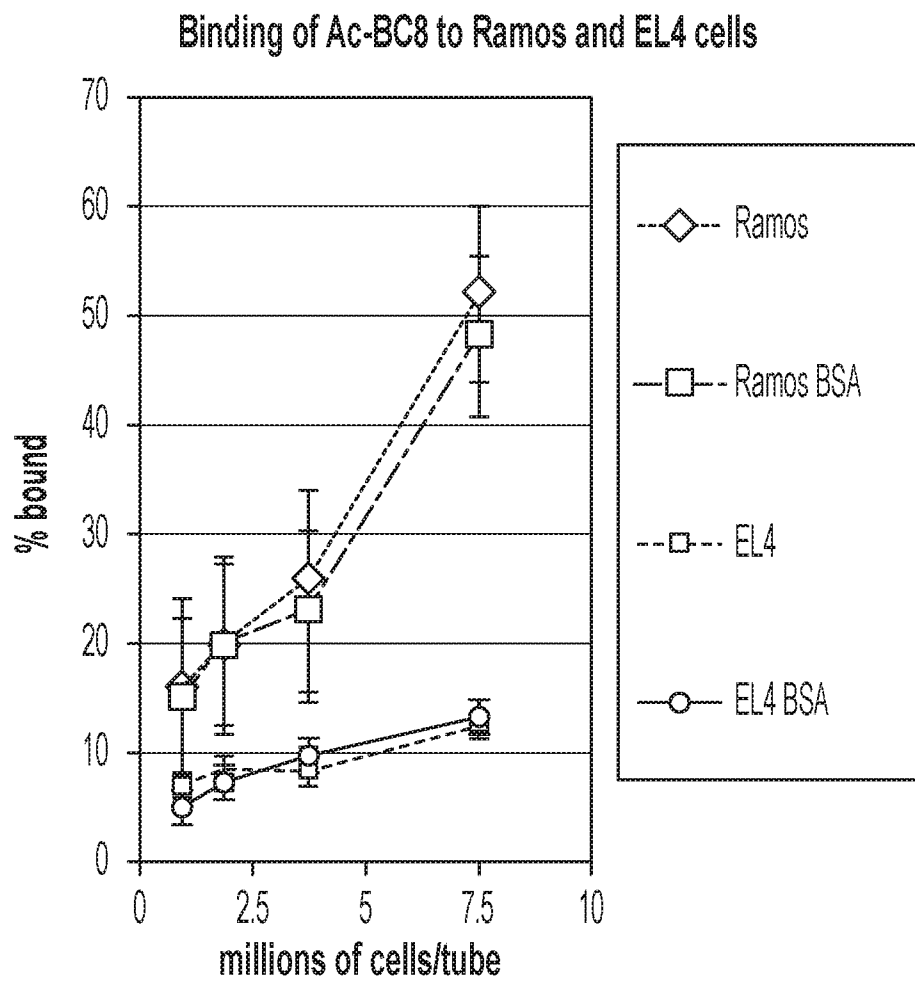


FIG. 12

FIG. 13A

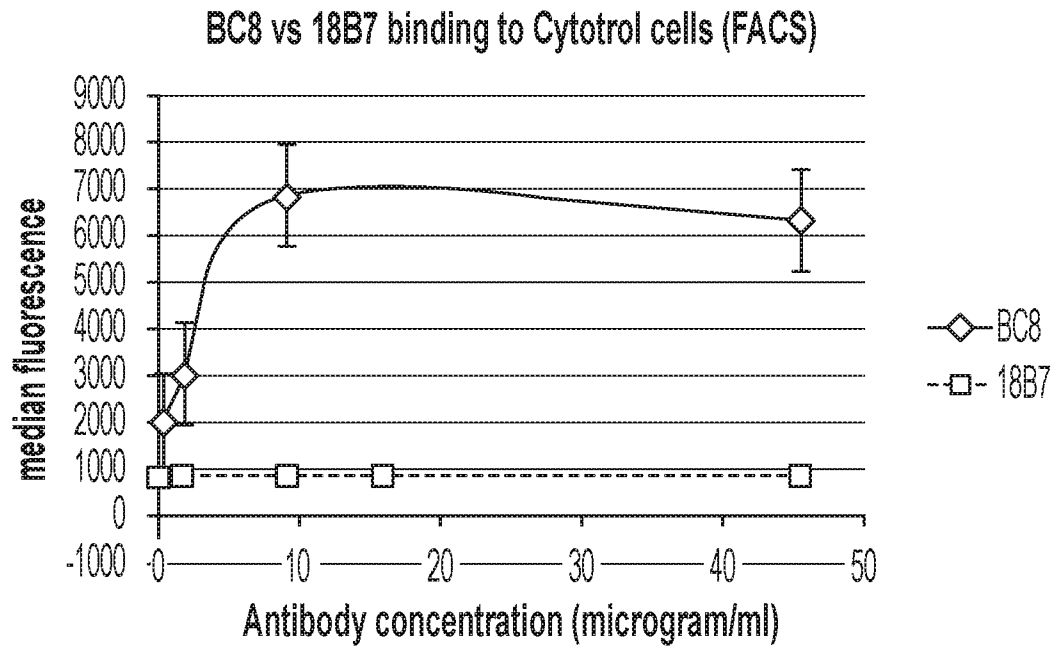


FIG. 13B

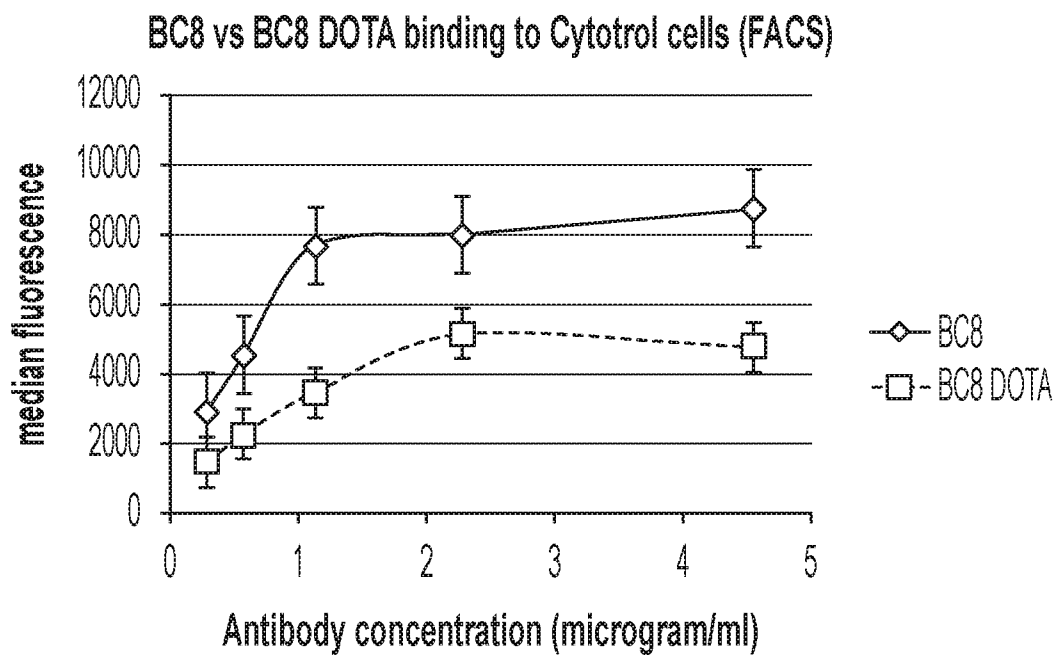


FIG. 14A

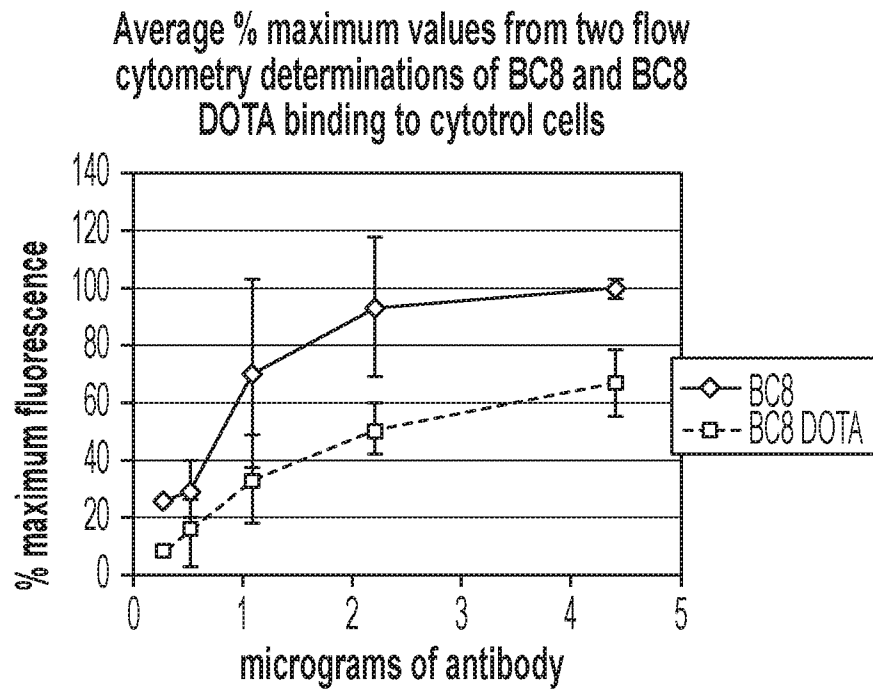


FIG. 14B

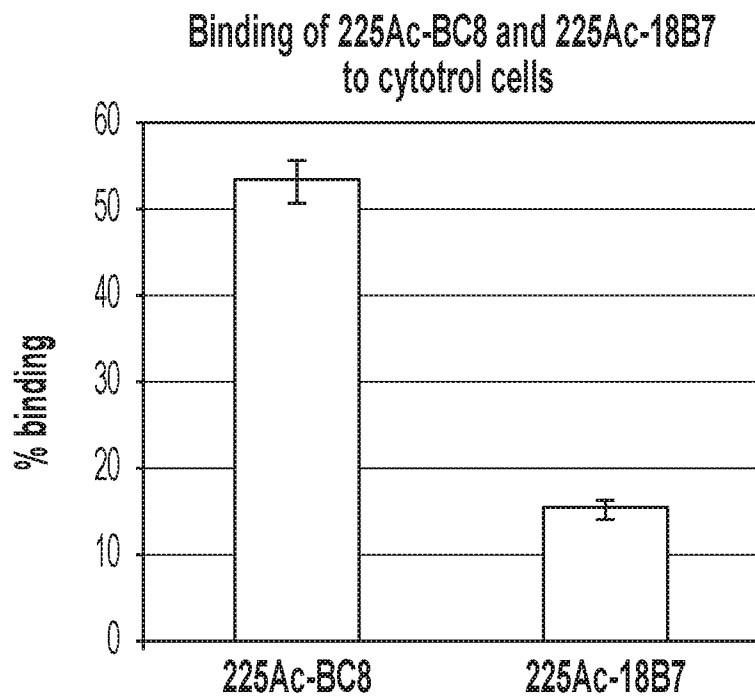


FIG. 15A

Binding of BC8-DOTA to H929 and U266 cells by Flow Cytometry

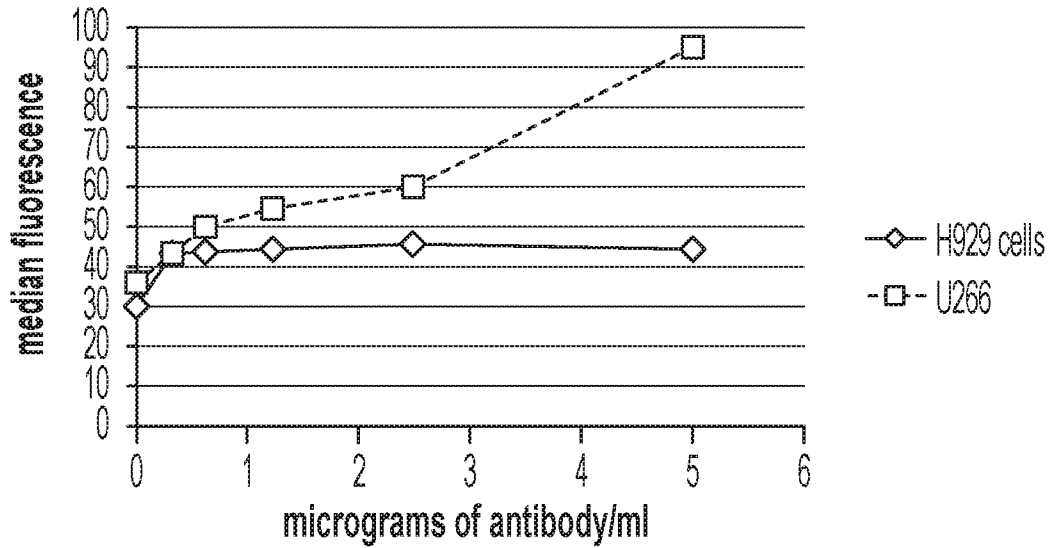
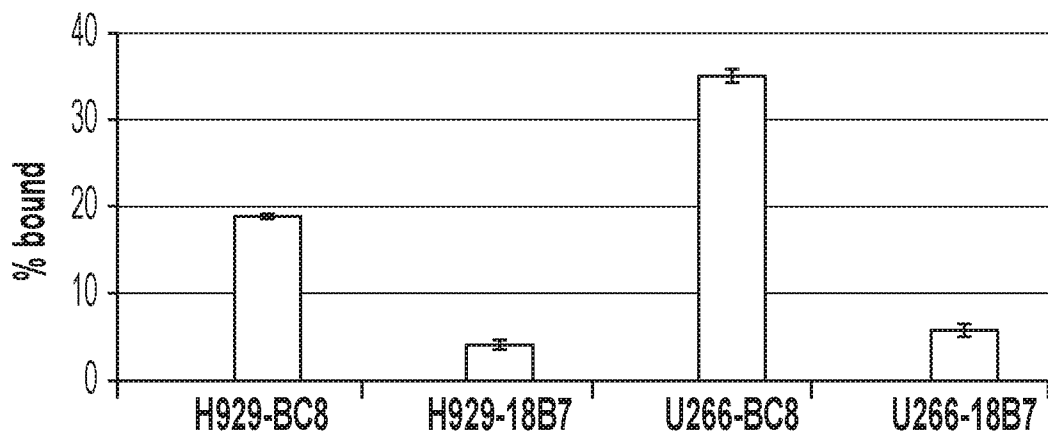


FIG. 15B

Binding of Ac225 labeled antibodies to H929 and U266 cells



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FIG. 16A

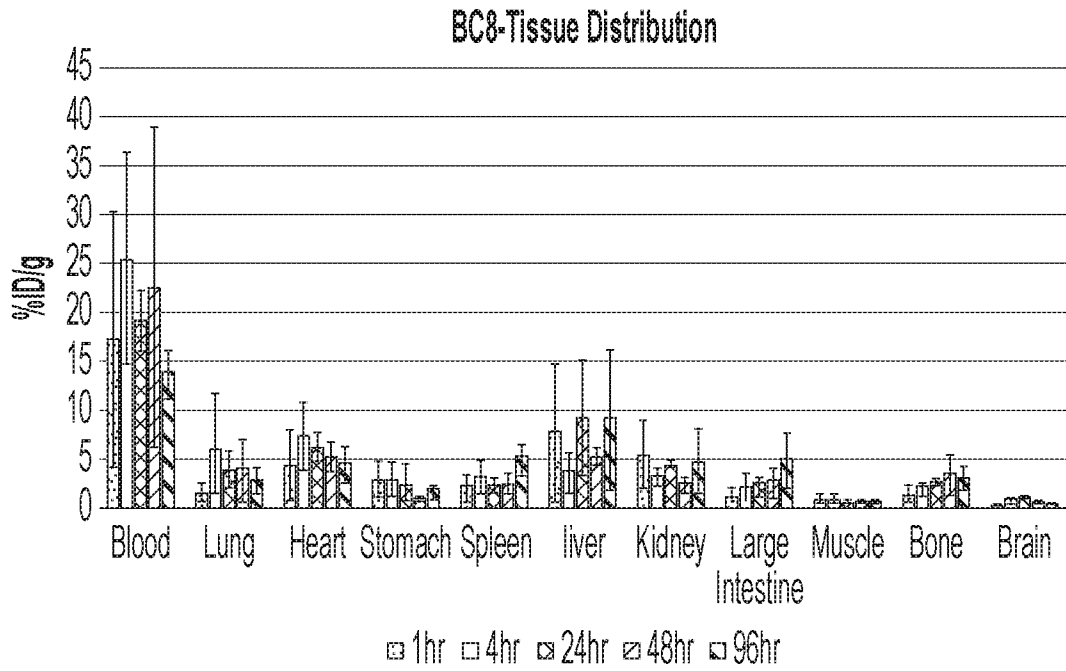


FIG. 16B

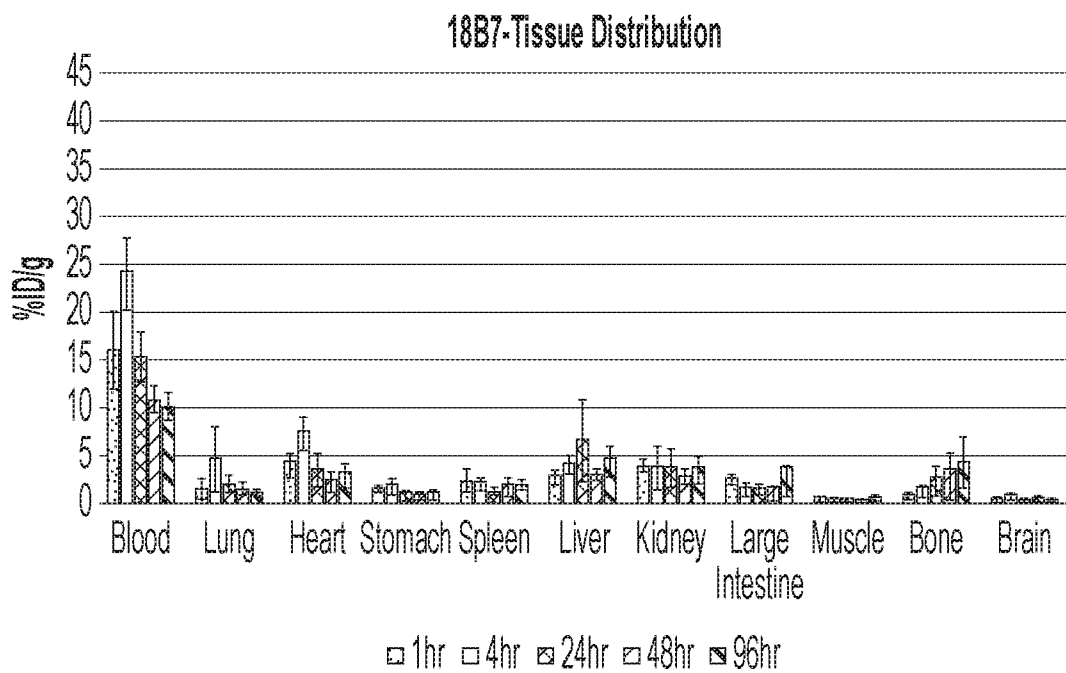


FIG. 17A

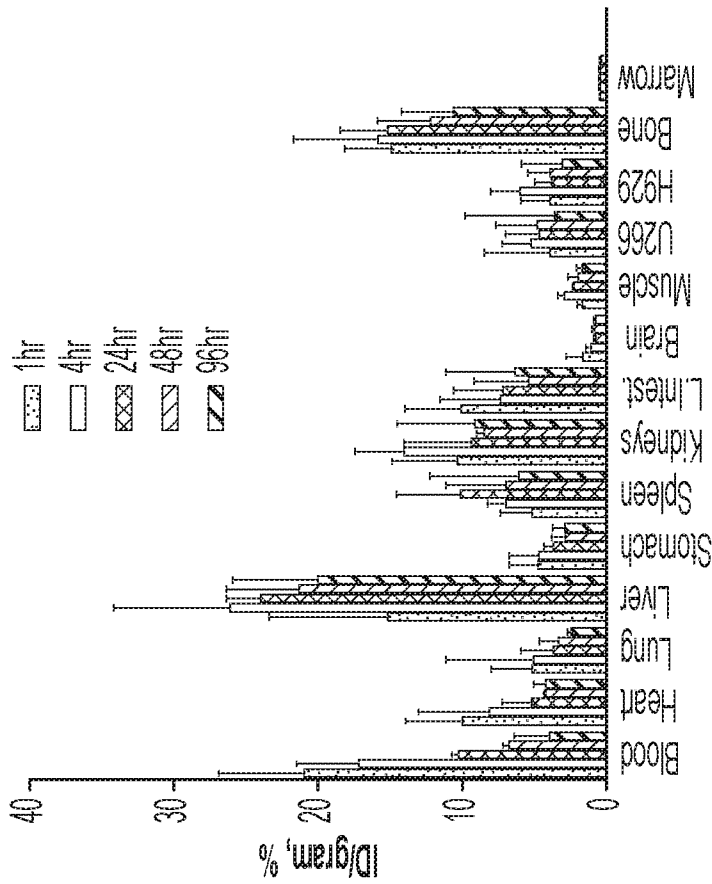
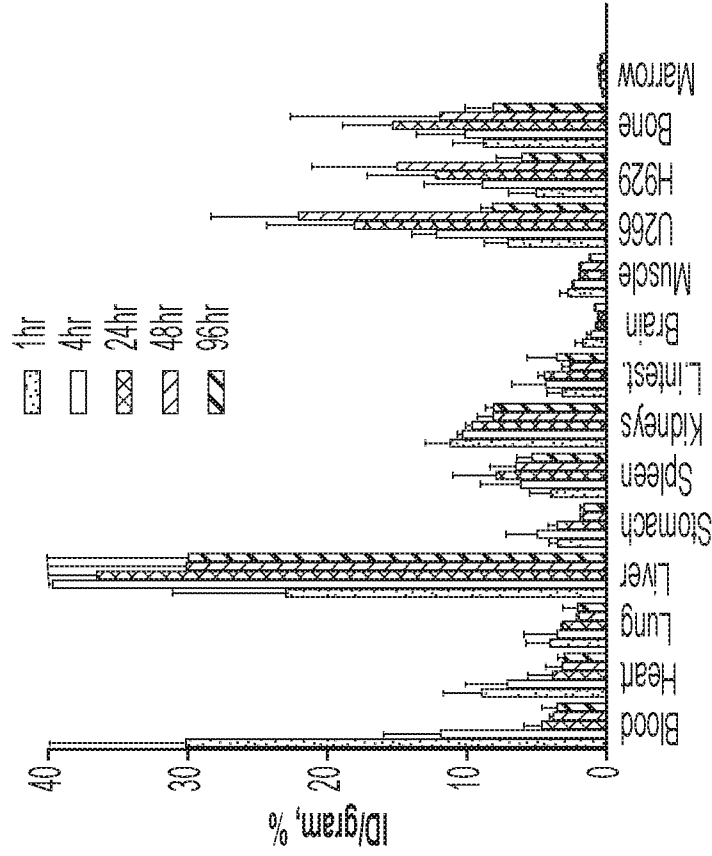


FIG. 17B



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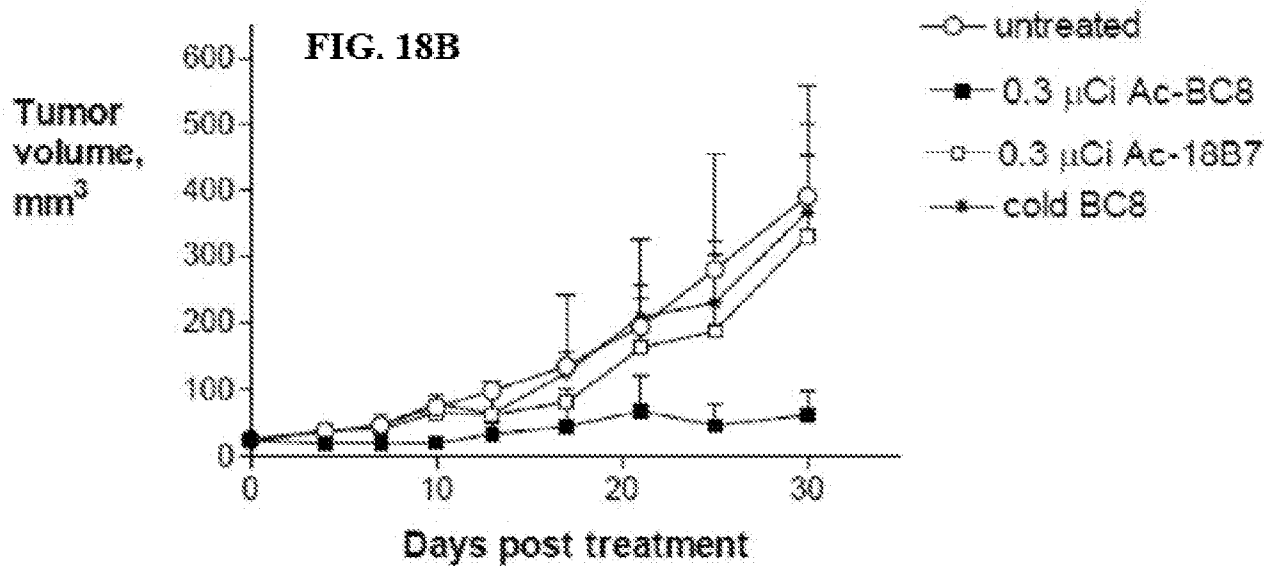
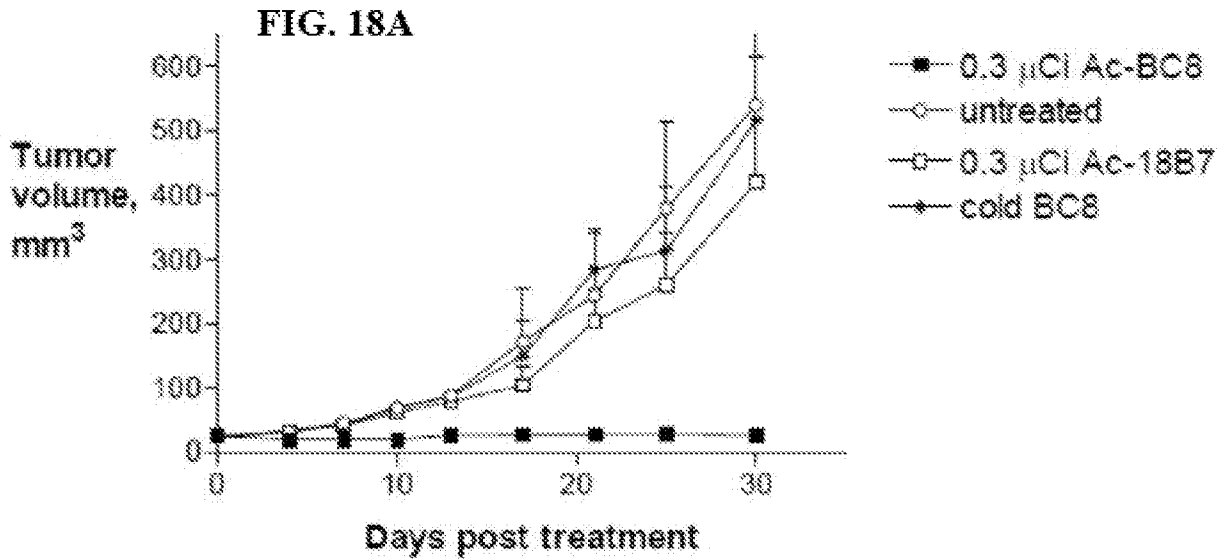


FIG. 19B

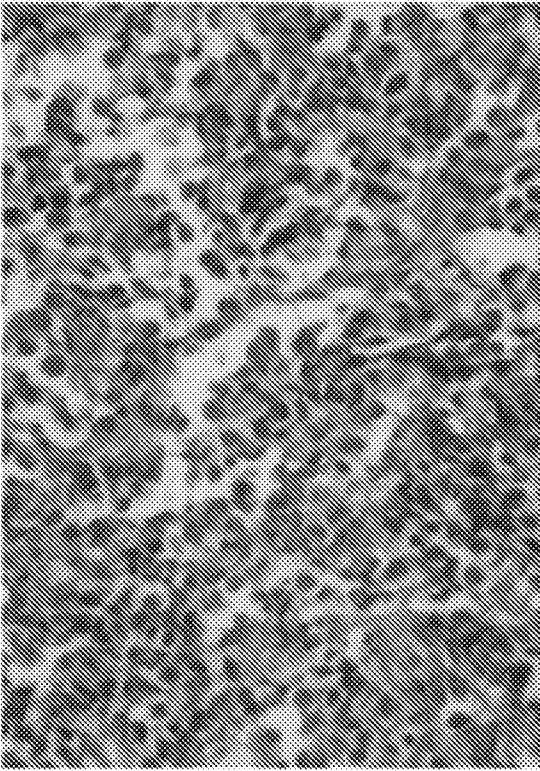


FIG. 19D

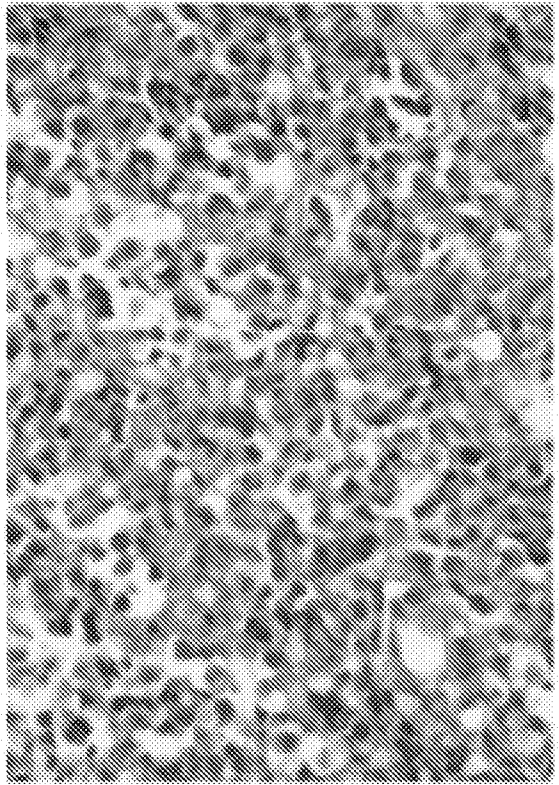


FIG. 19A

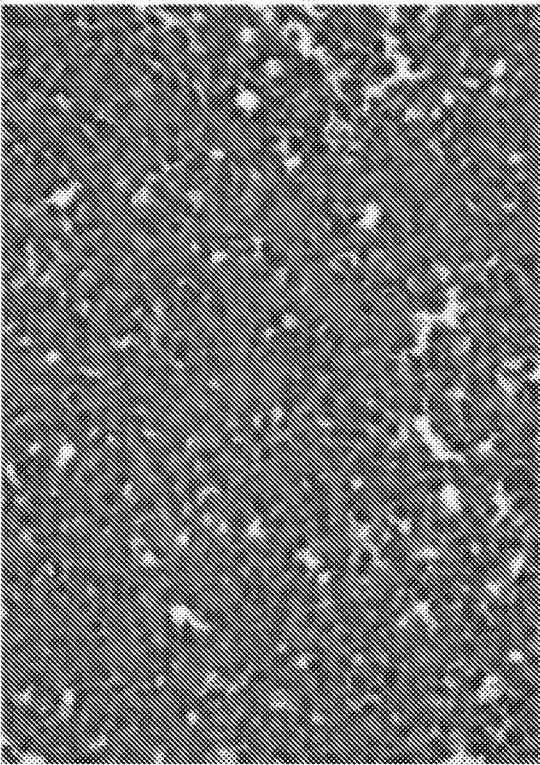
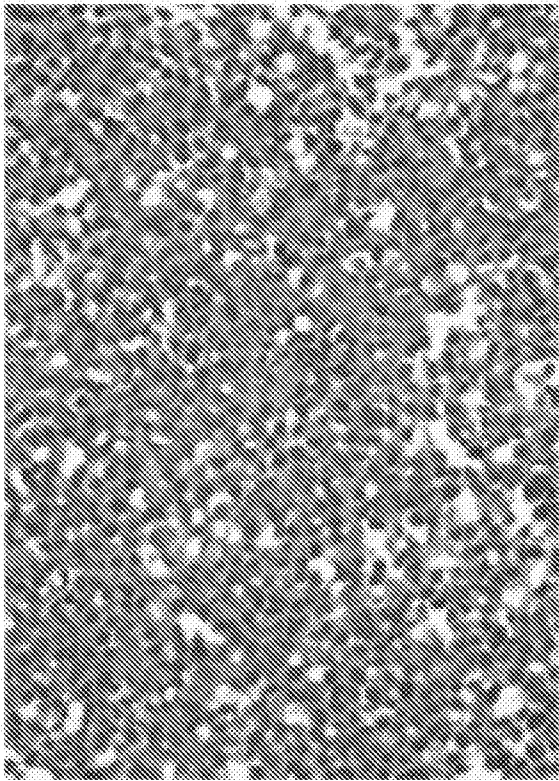


FIG. 19C



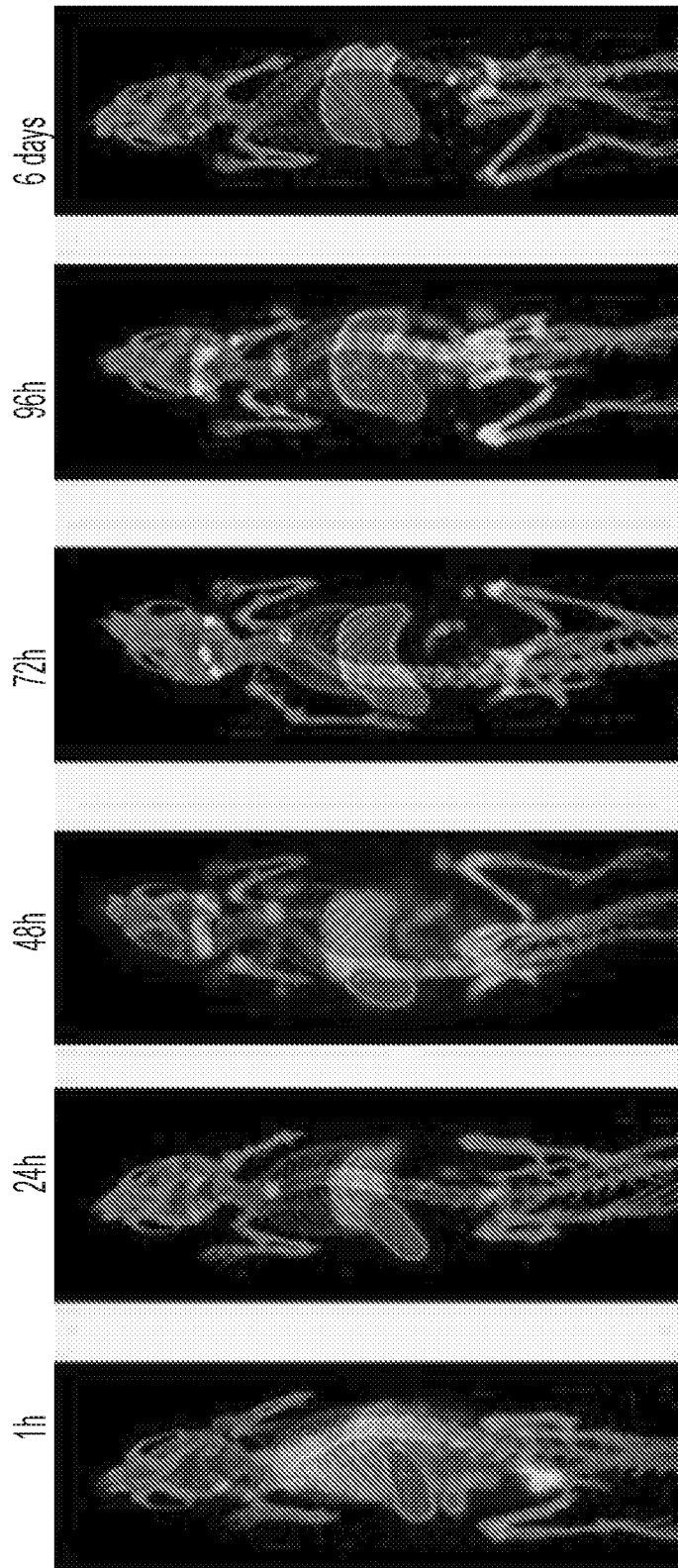


FIG. 20

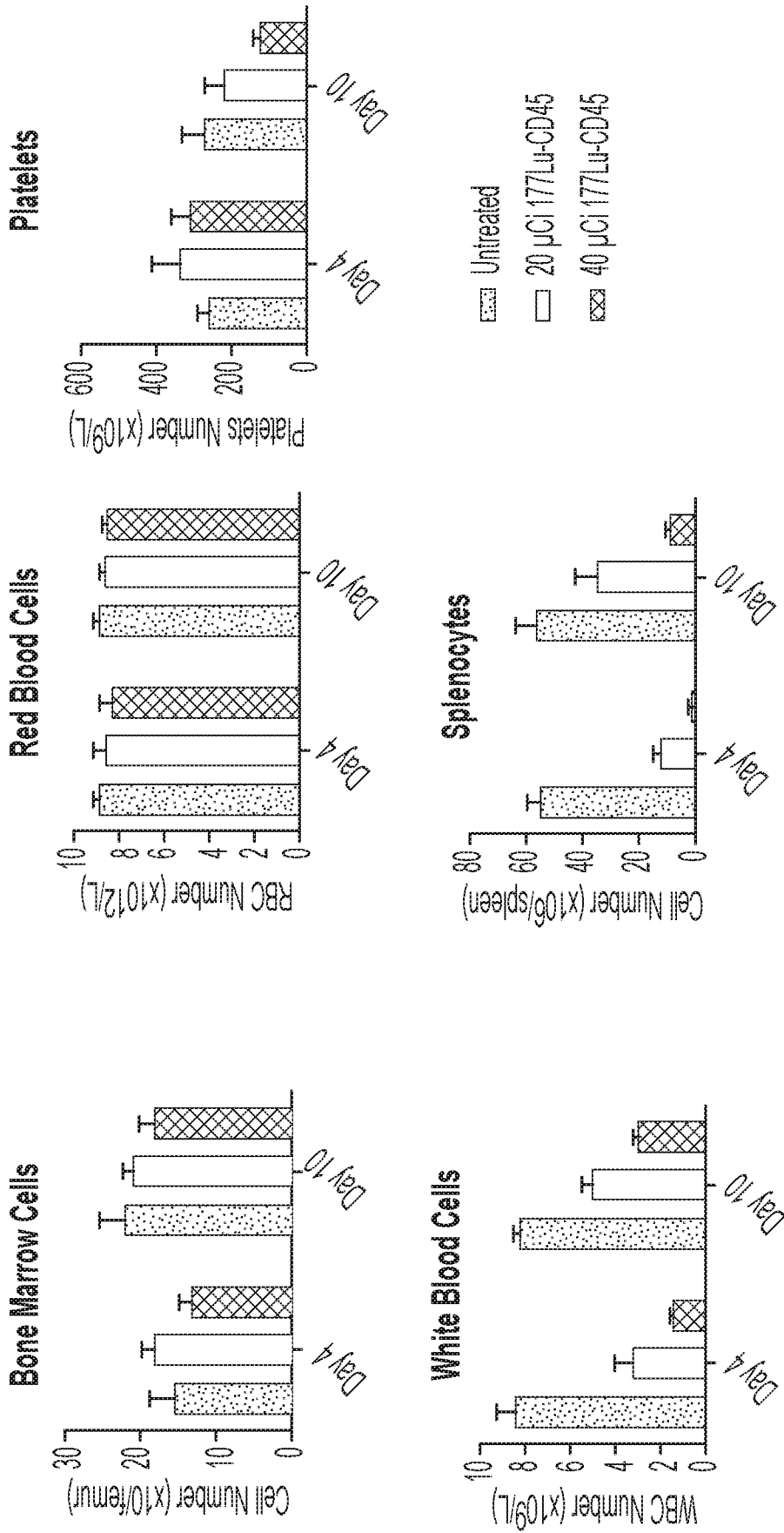


FIG. 21A

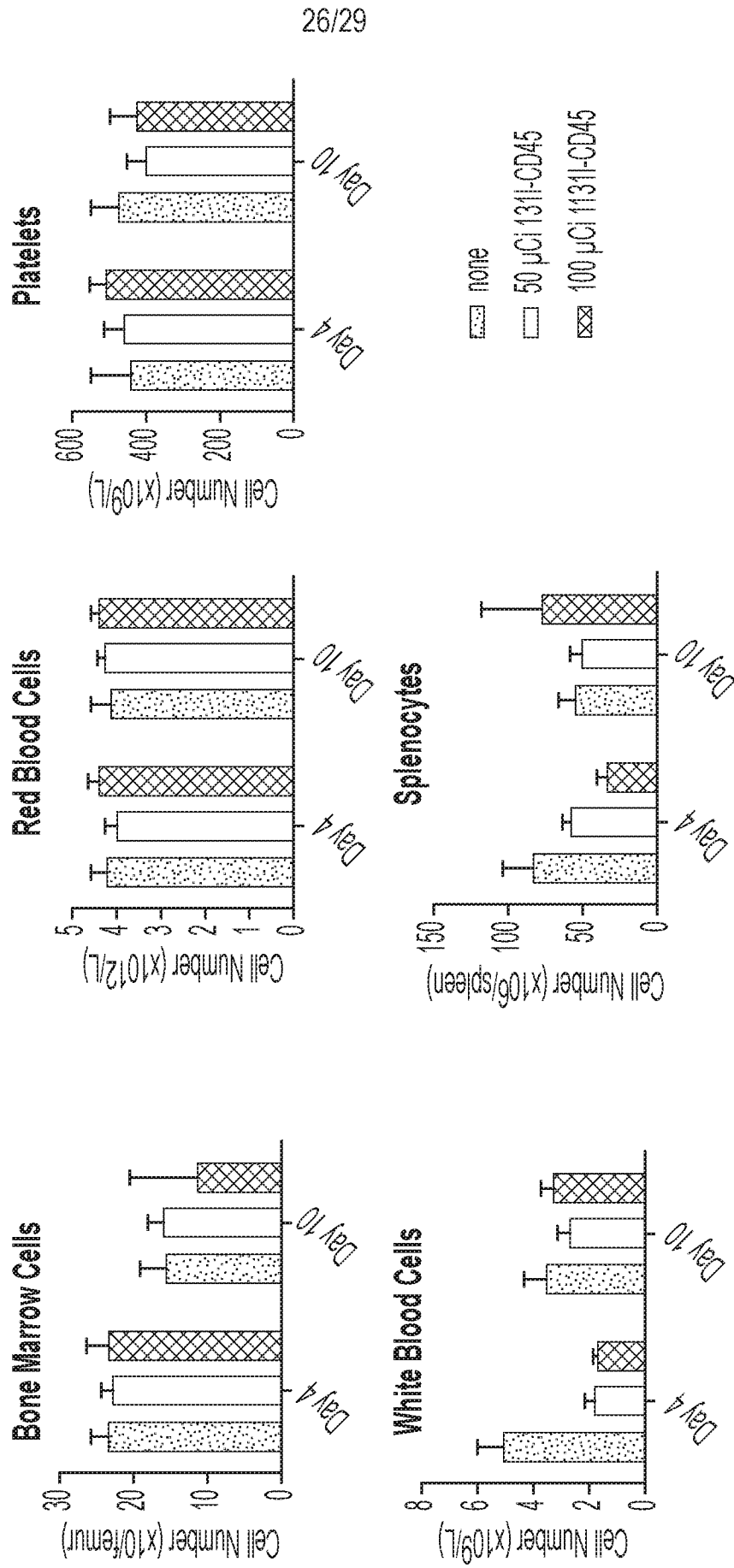


FIG. 21B

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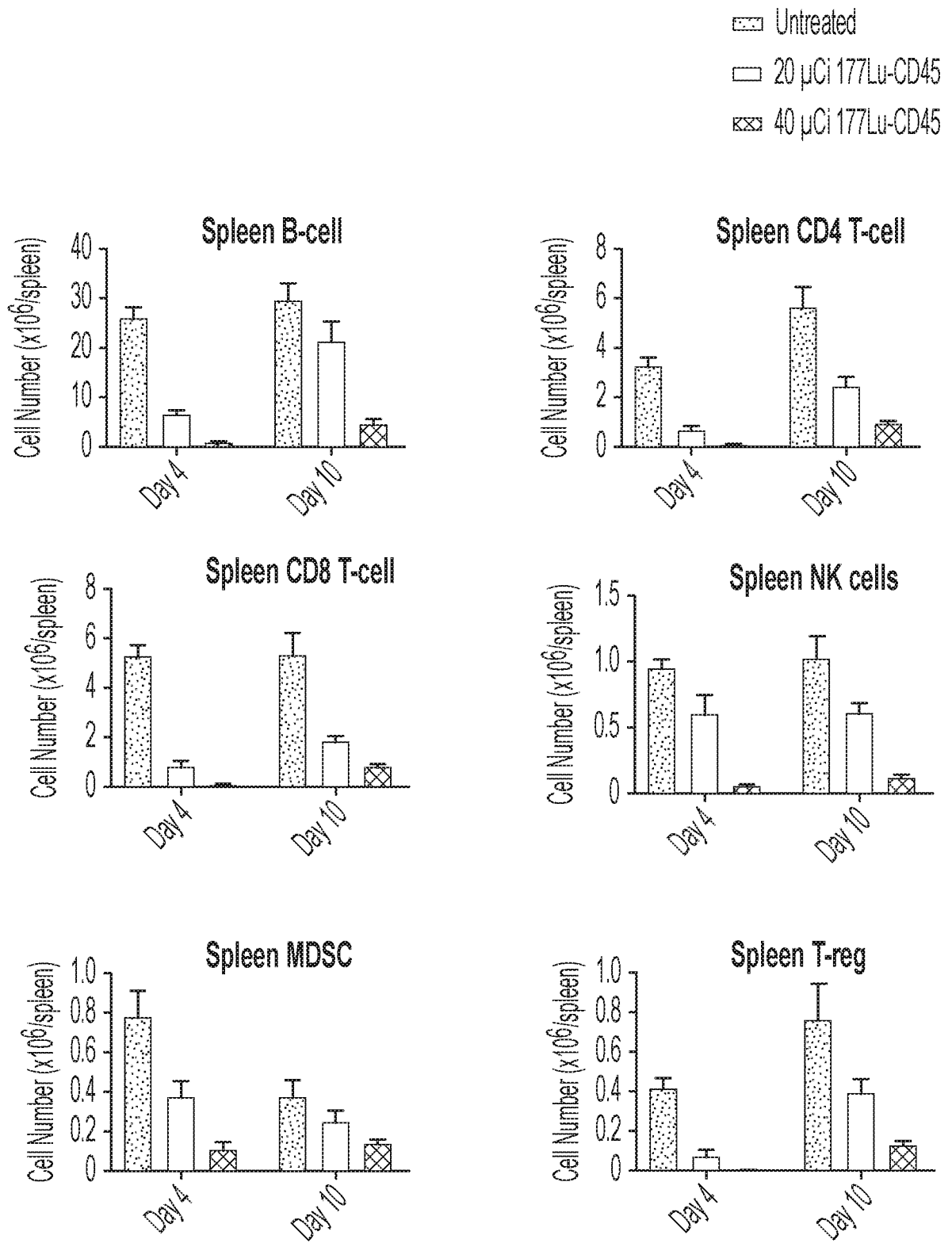


FIG. 22

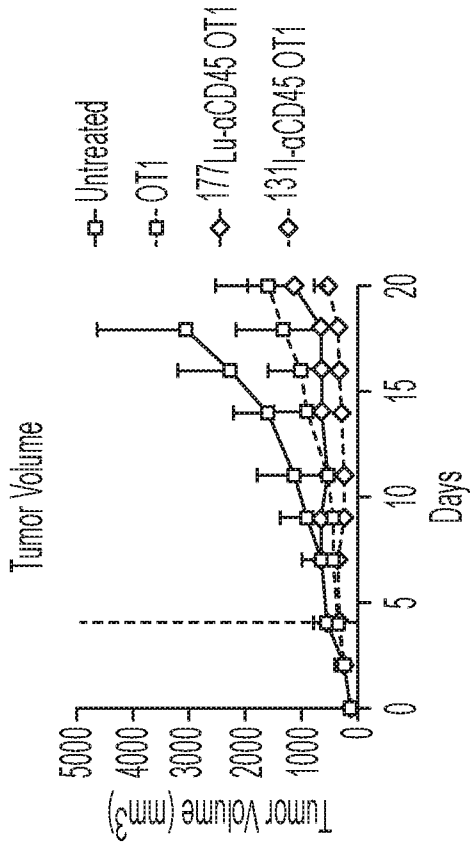


FIG. 23A

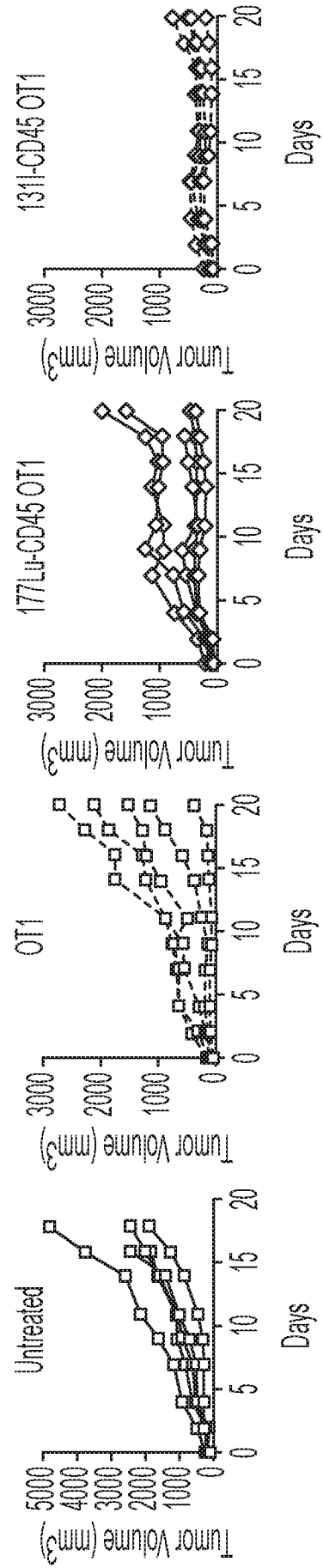


FIG. 23B

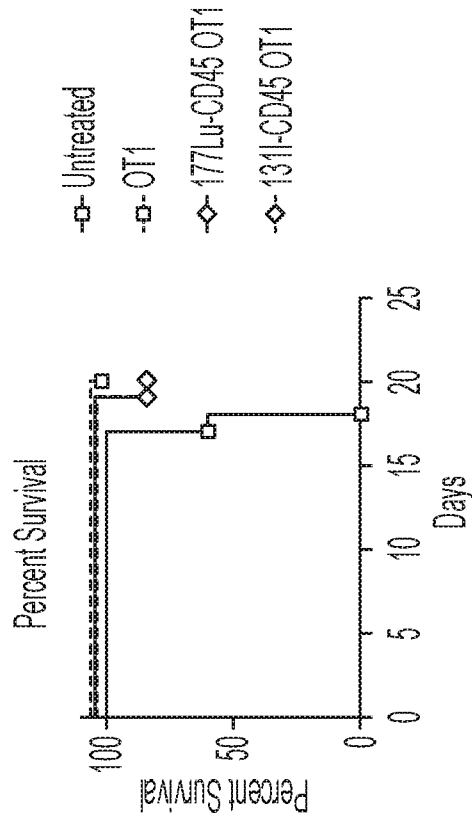


FIG. 23C