

ORIGINAL RESEARCH



Novel BCMA-OR-CD38 tandem-dual chimeric antigen receptor T cells robustly control multiple myeloma

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ABSTRACT

BCMA-targeting chimeric antigen receptor (CAR)-T cell therapy has shown remarkable clinical efficacy against multiple myeloma, yet antigen escape and tumor relapse still occur after the use of these therapies. Designing CAR-T therapies that targets multiple antigens simultaneously seems a feasible way to avoid antigen escape, and it has been extensively studied elsewhere. Here, we report novel BCMA-OR-CD38 Tan CAR T cells that can trigger robust cytotoxicity against target cells expressing either BCMA or CD38. We demonstrate that, in *in vitro* studies, these BCMA-OR-CD38 Tan CAR T cells exhibit similar CAR expression, superior cytotoxicity and antigen-stimulated T cell proliferation as compared to single-targeted CAR T cells or CD38-OR-BCMA Tan CAR T cells. Importantly, these BCMA-OR-CD38 Tan CAR-T cells can achieve complete tumor clearance in myeloma-bearing mice with no relapse observed through the course of these experiments. Finally, this BCMA-OR-CD38 Tan CAR was fully compatible with existing clinical grade T cell manufacturing procedures and can be implemented using current clinical protocols. Taken together, our results present an effective solution to the challenge of antigen escape in BCMA CAR T-cell therapies.

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Introduction

Multiple myeloma (MM) is the second most common hematological malignancy worldwide, and resulted in 113,474 deaths globally in 2019.¹ As a cancer of plasma cells, MM is characterized by the infiltration of malignant plasma cells in the bone marrow and is associated with high levels of monoclonal (M) protein in the blood and/or serum.² In recent years, the introduction of drugs targeting MM to its microenvironments, such as the proteasome inhibitor bortezomib and the immunomodulatory drugs (IMiDs) thalidomide and lenalidomide, have been used in initial, consolidation, maintenance, and salvage therapies, and have markedly improved patient outcomes and survival rates.³ However, MM remains largely incurable, due to the development of resistant and refractory disease.⁴

Chimeric antigen receptor T cell (CAR-T) therapy has emerged as a novel immunotherapy that has tremendous potential for long-term disease control in some hematologic malignancies,^{5–7} which has encouraged the development of CAR-T cells for treating MM, especially in the context of relapsed/refractory disease.⁸ Among the several CAR T-cell therapies, major efforts have been devoted to CAR T-cells targeting B-cell maturation antigen (BCMA), and more than 20 early-phase clinical studies are currently performed.^{9,10} Notably, the bb2121 anti-BCMA CAR-T Cell (Idecabtagene Vicleucel, ide-cel) from Bluebird Bio was approved by the Food and Drug Administration (FDA) as the first BCMA CAR T cell for the treatment of adult patients with relapsed or refractory MM on Mar 26th,

2021.¹¹ However, several clinical trials have reported patients' relapse involving tumor cell BCMA antigen loss or expression down-regulation below the threshold required for CAR T cell activity.^{12–15} Furthermore, a recent report demonstrated that biallelic loss of BCMA is one of the resistance mechanisms to Idecabtagene Vicleucel therapy.¹⁶ T cells expressing a single-chain bispecific CAR, also known as "Tan CAR", was previously proposed in a proof-of-concept approach to increase the specificity of effector cells and to offset antigen escape.¹⁷ Subsequently, multiple studies have demonstrated that Tan CARs reduced the likelihood of antigen escape and potentially improved the efficacy of CAR-T cell therapies.^{18–20} The design of Tan CARs is based on an "OR"-gated signal computation, and when either of two different antigens is present on a target cell, the T-cells will be activated, thus, tumor cells have to lose both antigens simultaneously to escape T cell surveillance.^{17,21} To construct tandem dual CARs for the treatment of MM, it is necessary to choose another antigen that is widely expressed in MM cells. In addition to BCMA, CD38 has recently been identified as a feasible antigen target for MM, since myeloma cells show strong and uniform expression of CD38 at different disease stages.^{22,23} Moreover, anti-CD38 CAR-T-cell therapy is currently being tested in the clinic trials (NCT03464916, NCT03754764). In a previous study, we identified an anti-CD38 scFv from an scFv phage display library, and the therapeutic effects of CD38 CAR-T cells on myeloma cells has been demonstrated with *in vivo* and *in vitro*

experiments (*a manuscript during revision*). We thus reason that simultaneous targeting of BCMA and CD38 would take advantage of the therapeutic efficacy of BCMA targeting while preventing tumor escape due to BCMA loss.

Here, we describe two novel Tan CARs which were composed of an antigen recognition domain (an anti-CD38 and an anti-BCMA scFv in tandem), CD8 hinge and transmembrane domain, CD28 costimulatory domain, and the signaling portion of CD3 ζ . Our results found that Tan-CAR T cells could trigger robust T cell-mediated cytotoxicity and cytokine production when either BCMA or CD38 was present on a target cell. Moreover, these Tan-CAR-T cells had significantly higher cytotoxicity and proliferation than single-targeted CAR T cells when encountering BCMA and CD38 antigens simultaneously. Notably, BCMA-OR-CD38 Tan CAR-T cells exhibited more potent antitumor and proliferative activity than CD38-OR-BCMA Tan CAR-T cells. Next, we assessed the efficacy of BCMA-OR-CD38 Tan CAR-T cells against multiple myeloma cells *in vivo*. The results showed that BCMA-OR-CD38 Tan CAR-T cells had robust therapeutic effects on an NPG mouse model bearing myeloma, and could achieve complete tumor clearance four days after the second dose of Tan-CAR T cells. Thus, our results suggest that this tandem-dual antigen targeting strategy represents an effective anti-neoplastic therapy and may ultimately provide an effective safeguard against antigen escape of BCMA in adoptive T-cell therapies for MM.

Results

Construction and generation of Tan-CAR-T cells

The antigen-binding domain of BCMA-OR-CD38 Tan-CAR or CD38-OR-BCMA Tan-CAR was composed of an anti-BCMA scFv (VL-linker-VH) and an anti-CD38 scFv (VH-linker-VL) in tandem connected via a long, flexible linker peptide (Gly₄Ser)₄ (Figure 1(b)). The difference between these two Tan CARs is the order of the two scFv domains (Figure 1(c)). The corresponding single-targeted CARs contained only the respective scFvs for antigen binding (Figure 1(a)). The retroviral vector titers of Tan-CAR were high (BCMA-OR-CD38 CAR: 3.59×10^8 CFU/mL; CD38-OR-BCMA CAR: 2.82×10^8 CFU/mL) though this was lower than the CD19 CAR (as retroviral vector standard), BCMA-CAR and CD38-CAR groups (Figure 1(e)). The transduction efficiency of BCMA-OR-CD38 Tan-CAR⁺ T was 85.7%, CD38-OR-BCMA Tan-CAR⁺ T was 74.0%, BCMA-CAR⁺ T was 78.2% and CD38-CAR⁺ T was 87.3% (Figure 1(g)). The Tan CARs were expressed with a similar efficiency as the BCMA CAR or CD38 CAR, implying that the targeting of this dual-antigen CAR modification did not affect the transduction efficiency of this CAR. Notably, the retroviral vector titers and T cell transduction efficiency of BCMA-OR-CD38 CAR were higher than that of CD38-OR-BCMA Tan-CAR. Transgenic copy counts, or vector copy numbers (VCNs), are one of the indicators for evaluating the safety of CAR T cell products. The FDA recommends that the integration copy number should be < 5 copies per genome.²⁴ In our results, the copy number of the CAR molecule (BCMA-CAR, CD38-CAR, or Tan-CARs) in the corresponding CAR-T cells were all less than 5 copies/ CAR-T cell (Figure 1(f)).

Detection of effective Tan CAR-T targeting to cells expressing various antigens

To evaluate specific cytotoxicity of Tan-CAR T cells on myeloma cells, the BCMA-OR-CD38 Tan CAR-T, the CD38-OR-BCMA Tan CAR-T, BCMA CAR-T, CD38 CAR-T and Pan-T cells were co-incubated respectively with BCMA-K562 or CD38-Raji cells for 8 hours. Annexin-V staining was used to determine the apoptosis of tumor cells. The Tan CAR-T cells showed significant cytotoxicity toward BCMA-K562 and CD38-Raji cells, while single-targeted CAR-T cells only respond to BCMA⁺ or CD38⁺ target cells, respectively (Figure 2(a,c)). Notably, the cytotoxicity of BCMA-OR-CD38 Tan CAR-T associated with BCMA-K562 and CD38-Raji cells was significantly higher than CD38-OR-BCMA Tan CAR-T cells (Figure 2(a)). The effective targeting of Tan CAR-T cells toward CD38 and BCMA antigens simultaneously was investigated based on two distinct strategies, the apoptosis or survival rate analysis of target cells after them co-cultured with CAR-T cells for 8 hours. The target cells consisted of BCMA-K562 and CD38-Raji cells mixed in a ratio of 1:1. The results showed that Tan CAR-T cells efficiently lysed both BCMA-K562 and CD38-Raji cells, and the cytotoxicity associated with these cells was significantly higher than single-targeted CAR-T cells (Figure 2(a,c)). We also carried out another experiment to further compare the cytolytic efficiency of single-targeted CAR-T cells with Tan CAR-T cells. RPMI-Luc cells expressing both CD38 and BCMA antigens were utilized for co-culturing with different CAR-T cells, after 8 hours of incubation, the BCMA-OR-CD38 Tan-CAR-T cells showed more significant cytotoxicity than single-targeted CAR-T cells and CD38-OR-BCMA Tan-CAR-T cells (Figure 2(b)). The un-modified K562 cells (CD38⁻BCMA⁻) were negative control cells and there was no significant cytotoxicity in this background (Figure 2(a)). The Pan-T cells (un-transduced T cells) were used as control effective T cells, and their responses to target cells were limited (Figure 2(a-c)).

The release of proinflammatory cytokines by CAR T cells was then measured using the Human CD8/NK Panel CBA Kit. After 8 hours of incubation of effector and target cells together, the cell culture supernatant was harvested and the levels of cytokines IL-10, IL-6, TNF- α , IFN- γ and Granzyme B were measured by flow cytometry. The levels of TNF- α , IFN- γ and Granzyme B released by Tan CAR T cells significantly increased after target cell stimulation to levels that was comparable to that generated by BCMA-CAR or CD38 CAR T cells in response to their respective cognate antigens (Figure 2(d)). The K562 cells that did not express CD38 or BCMA antigen were used as negative control cells. The results showed that Tan CAR T cells released limited proinflammatory cytokines after co-incubated with K562 cells (Figure 2(d)).

Tan CAR-T-cell proliferation in response to CD38⁺ and BCMA⁺ tumor cells

To investigate if tumor cells expressing specific antigens could stimulate CAR-T cell proliferation, we performed a CFSE proliferation assay. CAR-T and Pan-T cells were labeled with CFSE and co-cultured with BCMA K562+ CD38 Raji (mixed

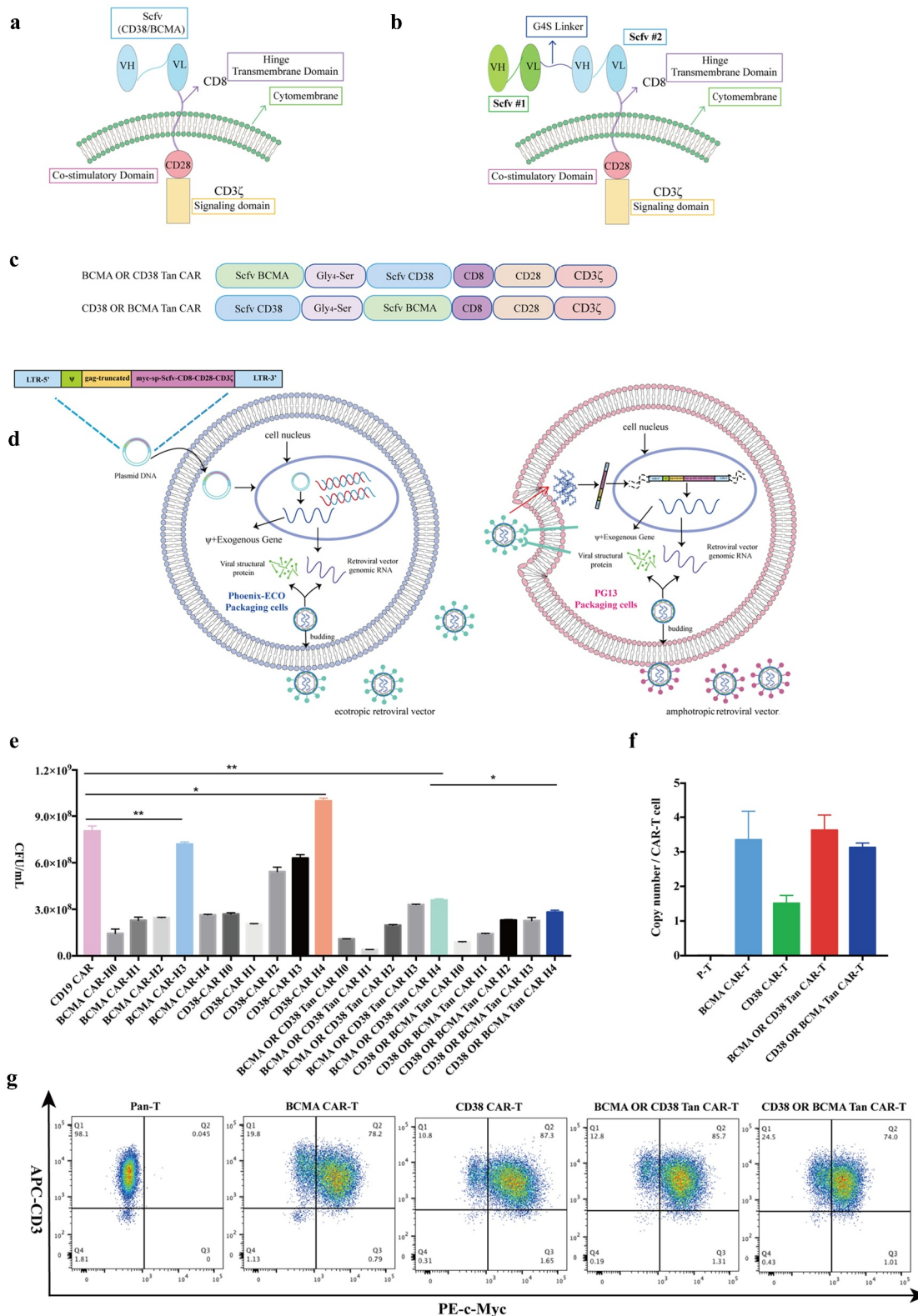


Figure 1. Construction and generation of retroviral vectors. (a) Schematic of a second-generation, single-targeted CAR (BCMA-CAR or CD38-CAR). (b) Schematic of a bi-specific, OR-gate tandem-CAR. (c) Schematic of two Tan CARs containing variations in the order of scFv domains. CD8 is the hinge and transmembrane domain, CD28 is the costimulatory domain. (d) Schematic of the retroviral vector packaging process. First, the Phoenix-ECO cells produced ecotropic retroviral vectors by transient transfection and then integrated them into the packaging cell line PG13 to produce amphotropic retroviral vectors that can transduce human primary T cells. (e) The titer detection of amphotropic retroviral vectors by Q-PCR. The amphotropic retroviral vector supernatants from CD38-CAR, BCMA-CAR, or Tan-CARs were harvested every 24 h for a total of 5 days, named H0, H1, H2, H3, and H4. (f) Detection of the copy number of CAR molecules. Q-PCR was performed using primers specific for this transgene, and genomic DNA extracted from the isolated and activated human T cells 48 h after transducing was used as templates. (g) Analysis of transduction efficiency. The retroviral vectors BCMA-CAR, CD38-CAR, or Tan-CARs transduced the isolated and activated human T cells, and the transduction efficiency was evaluated by detection of *c-Myc* protein expression using flow cytometry. The data shown is the mean \pm standard deviation (SD). The results were analyzed with a student's *t* test. Error bars represent the SD, * $p < .05$, ** $p < .01$.

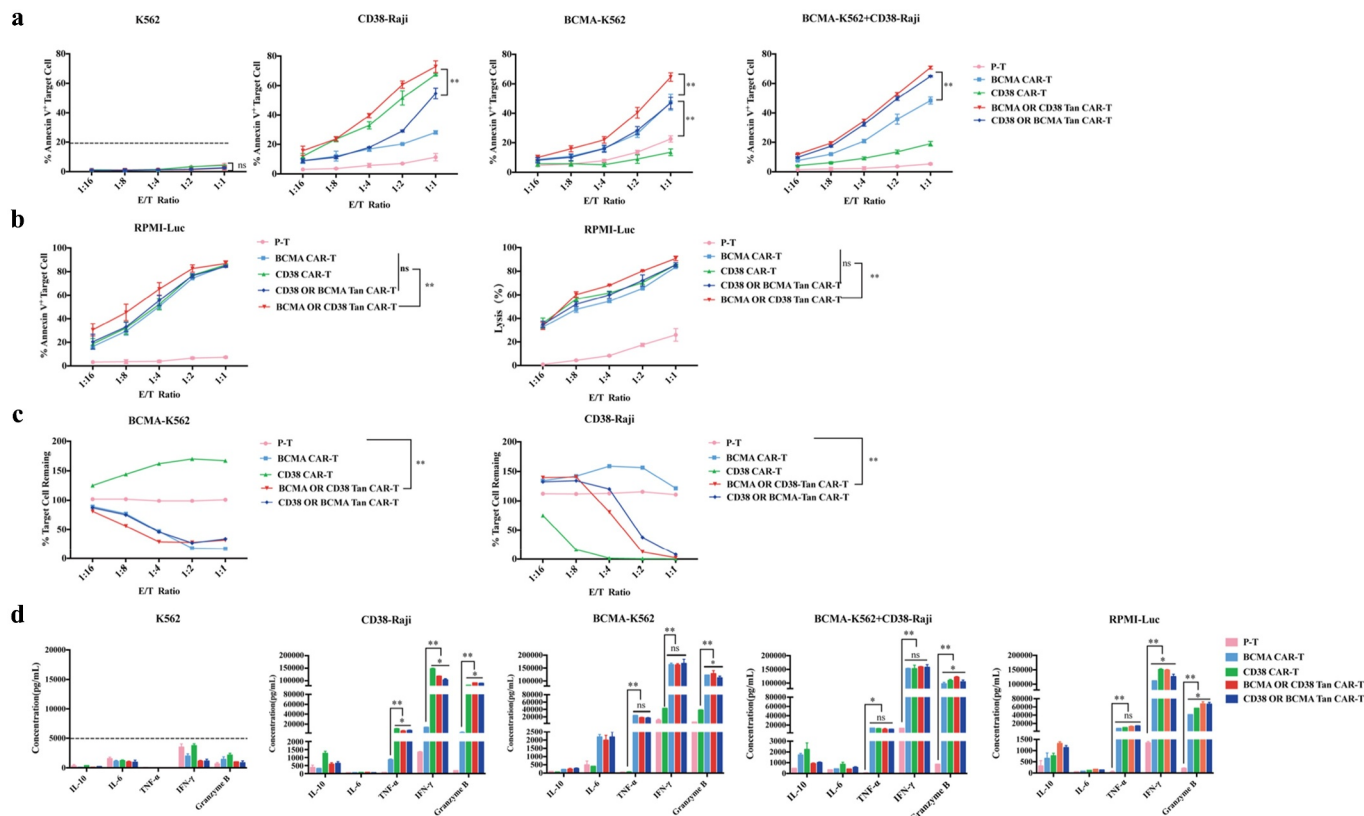


Figure 2. The specific cytotoxicity of Tan CAR-T cells after antigen stimulation. (a) The cytotoxicity of single-targeted CAR-T and Tan CAR-T cells after 8 hours of co-cubation with BCMA-K562, CD38-Raji, or BCMA-K562+ CD38-Raji cells (mixed in 1:1) at different effect-to-target ratios. Apoptosis was determined using an Annexin-V apoptosis Kit. (b) The cytotoxicity of single-targeted CAR-T and Tan CAR-T cells after 8 hours of co-cubation with RPMI-Luc was detected with two different methods, Annexin-V apoptosis (left panel) and luciferase assay (right panel). (c) The specific cytotoxicity of Tan CAR-T cells for BCMA-K562 and CD38-Raji cells. BCMA-K562+ CD38-Raji cells (mixed in 1:1) were co-cultured with single-targeted CAR-T and Tan CAR-T cells for 48 hours, and the survival of BCMA-K562 or CD38-Raji cells was determined. (d) Cytokine release from Tan-CAR T cells in response to target cells as indicated, with the cytokines evaluated including IL-6, IL-10, TNF- α , IFN- γ , and Granzyme B. Data shown is the mean \pm standard deviation (SD). The results were analyzed by two-way ANOVA, and error bars represent the SD, * $p < .05$, ** $p < .01$.

in 1:1) or PRMI-Luc cells, and the T-cell proliferation was evaluated via detection of decreasing CFSE fluorescence intensity using flow cytometry. The results showed that Tan-CAR T cells, particularly BCMA-OR-CD38 Tan-CAR T cells, proliferated significantly more than single-targeted CAR-T cells and CD38-OR-BCMA Tan-CAR T cells in response to activation by target cells at a 1:1 effect-to-target ratio (Figure 3(a)). Furthermore, we also plotted the T cell proliferation and viability curves without target cell stimulation. These showed that

Tan-CAR T cells proliferated to an extent that was similar to single-targeted CAR-T cells (Figure 3(b)).

BCMA-OR-CD38 Tan CAR-T cells are effective in the treatment of tumors *in vivo*

As illustrated above, BCMA-OR-CD38 Tan CAR-T cells exhibited more potent antitumor and proliferative activity than CD38-OR-BCMA Tan CAR-T cells. Thus, we next assessed

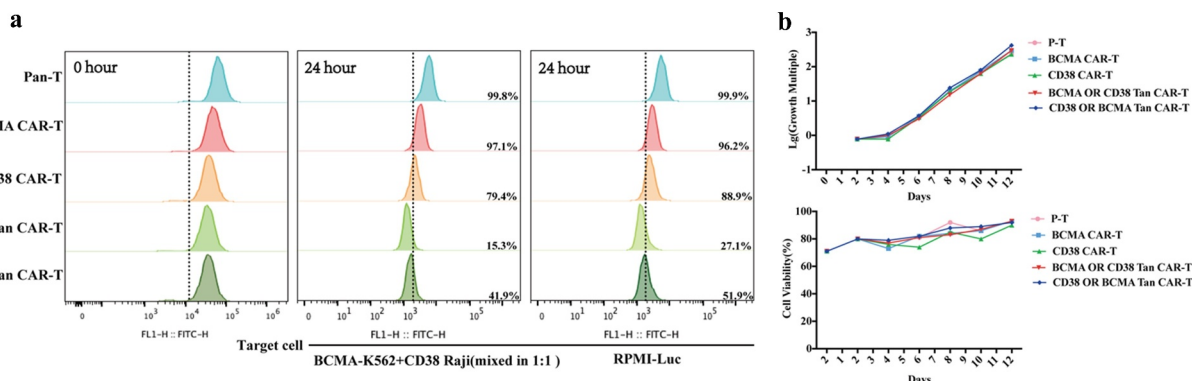


Figure 3. T cell proliferation detection. (a) CAR T cells labeled with CFSE were co-cultured with BCMA K562+ CD38 Raji (mixed in 1:1) or PRMI-Luc cells for 24 hours at a 1:1 effect-to-target ratio, and the CFSE fluorescence intensity was determined by flow cytometry. (b) Pan-T and CAR-T cells were counted every 2 days using an automated cell counter and cell growth and viability curves were plotted here.

the efficacy of BCMA-OR-CD38 Tan CAR-T cells against multiple myeloma cells *in vivo* in an intravenous RPMI-Luc tumor xenograft model. After tumor injection, tumor-bearing mice with similar tumor sizes were randomly divided into 5 groups ($n = 8$ per group), and Pan-T, BCMA CAR-T, CD38 CAR-T, or BCMA-OR-CD38 Tan CAR-T cells were provided by intravenous injection on days 12 and 19 (Figure 4(a)). Bioluminescence imaging revealed that either single-targeted CAR-T cells or BCMA-OR-CD38 Tan CAR-T cells yielded complete tumor clearance by day 23 (Figure 4(b)). Correspondingly, we also detected tumor cell residues in the bone marrow (BM), liver, spleen, and blood of mice by flow cytometry. The result showed that there were almost no tumor cells observable in the mice receiving BCMA-OR-CD38 Tan CAR-T or single-targeted CAR-T treatment (Figure 4(c)). Notably, two weeks after the second CAR-T injection, we measured the levels of human IFN- γ in the blood serum using ELSA assay, which were significantly higher for BCMA-OR-CD38 Tan CAR-T cell-treated mice than for single-targeted CAR-T cell-treated mice, indicative of greater IFN- γ production by the Tan CAR-T cells (Figure 4(e)). Next, we evaluated the expansion of CAR-T cells by detecting the percentage of CD3⁺ T cells in the peripheral blood. The results indicated that BCMA-OR-CD38 Tan CAR-T cells had experienced more extensive expansion in mice than either the Pan-T or CD38 CAR-T cells did (Figure 4(d)). Finally, the safety of this Tan CAR-T cell therapy was evaluated, and the results showed that mice in the BCMA-OR-CD38 Tan CAR-T cells-treated groups had significantly longer median survival times than those from the Untreated or Pan-T groups (Figure 4(f)). However, there was no significant difference between BCMA-OR-CD38 Tan CAR-T and single-targeted CAR-T cells in terms of tumor clearance or median survival times.

Materials and methods

Plasmid construction and production of retroviral vectors

The tandem-CARs retroviral vector consisted of the following components in-frame from the 5' end to the 3' end: a signal peptide sequence derived from the murine Ig-H (immunoglobulin heavy chain), a human c-Myc tag, an anti-BCMA-scFv derived from the C11D5.3 monoclonal antibody (mAb),²⁵ an anti-CD38 scFv identified from an scFv phage display library in our previous study (*a manuscript during revision*), the hinge and transmembrane regions of the CD8 molecule, the costimulatory domain of CD28, and the CD3 ζ signaling domain. The codon optimization and sequence synthesis for this was carried out at General Biosystems (China). The anti-CD38 and anti-BCMA scFv fragments were assembled by overlap extension PCR and cloned to an MFG retroviral backbone, which is referred to as the BCMA-OR-CD38 Tan-CAR or CD38-OR-BCMA Tan-CAR retroviral vector. The difference between these two Tan CARs is the order of the two scFv domains. We also prepared CD38-CAR and BCMA-CAR retroviral vectors, which had a single CD38 or BCMA scFv domain. The process for the production of Tan CARs and single-targeted CARs retroviral vectors was consistent with the manufacture of clinical grade anti-CD19-CAR retroviral vectors as previously

described.^{26,27} Retroviral vectors producer cell lines were established using two different packaging cell lines. The plasmids encoding these CAR genes were introduced into the human-derived ecotropic packaging cell line Phoenix-eco, and the retroviral vector supernatant transiently produced by these transfected cells was harvested and integrated stably into the genomic DNA of the murine-derived amphotropic packaging cell line PG13 (Figure 1(d)). The retroviral vector supernatant produced by PG13 cells was harvested every 24 hours for a total of five days. The collected samples were called H0, H1, H2, H3, and H4, and were utilized for the transduction of human primary T cells.

Cell-line generation and maintenance

K562 cells were obtained from the Cell Resource Center, Peking Union Medical College (Beijing, China). Raji cells are a CD38⁺ human B lymphocyte cell line that was obtained from the American Type Culture Collection (ATCC, United States). BCMA⁺ GFP⁺-K562 cells were a generous gift from Dr. Wu (China Agricultural University) received in 2018. GFP-Luciferase RPMI8226 cells (RPMI-Luc) were a CD38⁺ and BCMA⁺ myeloma cell line that were obtained from the ATCC. The PG13 gibbon ape leukemia virus packaging cell line and the human ecotropic packaging cell line Phoenix-ECO were obtained from ATCC. K562, Raji, and RPMI cells were maintained in RPMI-1640 medium (Gibco, United States) supplemented with 10% fetal bovine serum (FBS) (Gibco, United States), with 1% penicillin-streptomycin (P/S) solution (Gibco, United States). PG13 and Phoenix ECO cells were cultured in DMEM (Gibco, United States) supplemented with 10% FBS and 1% P/S. All cells were cultured in an incubator (Thermo Fisher, United States) at 37°C and 5% CO₂. All cell lines were tested negative for the mycoplasma contamination and cell-surface markers for these cell lines were validated by flow cytometry.

The generation of CAR-T cells

Peripheral blood mononuclear cells (PBMCs) were isolated from healthy donors by gradient centrifugation using LymphoprepTM (STEMCELL Technologies, Canada). The use of human PBMCs was approved by the Ethics Committee of Beijing University of Chinese Medicine, and all donors gave informed consent. After stimulating these PBMCs with 100 ng/mL of the anti-CD3 monoclonal antibody OKT3 (Sino Biological, China) and 100 U/mL IL-2 (Sino Biological, China) for 48 hours, these T cells were transduced using Tan CARs and single-targeted CARs retroviral vectors. The transduction efficiency was determined using flow cytometry. All T cells were expanded in AIM V medium (Gibco, United States) supplemented with 10% FBS, 1% P/S and IL-2 (100 U/mL), which was renewed every 2 days or as necessary.

Quantitative real-time PCR (Q-PCR)

The titers of retroviral vectors produced by PG13 were determined using Q-PCR. The RNA from retroviral vectors (H0, H1, H2, H3, and H4) was extracted using a QIAamp Viral RNA

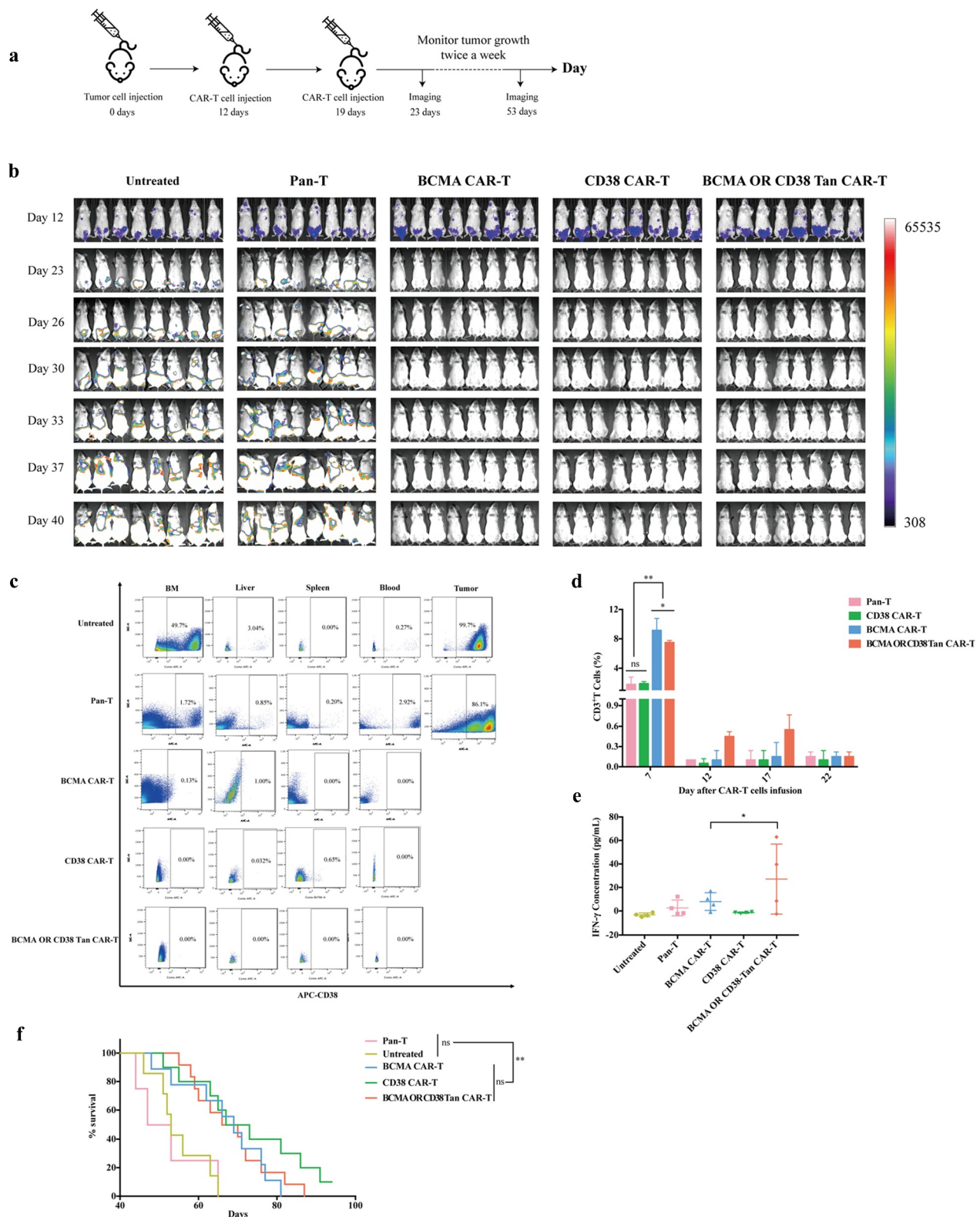


Figure 4. BCMA-OR-CD38 Tan CAR-T cells showed significant anti-tumor activity *in vivo*. (a) Outline of *in vivo* xenograft tumor model treatment schedule: NPG mice were i.v. injected with 2×10^6 RPMI-Luc cells on day 0. After that, tumor-bearing mice were randomly divided into 5 groups ($n = 8$ per group) and received Pan-T, BCMA CAR-T, CD38 CAR-T, or BCMA-OR-CD38 Tan CAR-T cells at an i.v. dose of 1×10^8 /kg on days 12 and 19. Tumor growth was followed by MIIIS living imaging every 3 days beginning on day 4 after the second CAR-T injection. (b) The mouse tumor burden at specific time points is indicated by bioluminescence radiance ($n = 8$ mice per group), and the results were pooled from two independent experiments, with the CAR-T cells being prepared with PBMCs obtained from two health donors. (c) Mouse BM, liver, spleen, blood and tumor cells were stained with anti-human CD38 APC to evaluate whether there were still residual RPMI-Luc cells in mice after CAR-T cells infusion. (d) The percentage of CD3⁺ T cells in peripheral blood was used to evaluate the expansion of CAR T cells. Blood was collected by tail vein and stained with the anti-human CD3 BV786. (e) Two weeks after the second CAR-T injection, the levels of IFN- γ in serum from each group were measured using ELISA assay. (f) Survival analysis of mice treated with Pan-T, BCMA CAR-T, CD38 CAR-T, or BCMA-OR-CD38 Tan CAR-T cells. Kaplan-Meier survival curves were tested using the Mantel-Cox log rank test. The data is shown as the mean \pm standard deviation (SD). The results were analyzed by two-way ANOVA, error bars represent the SD, and * $p < .05$, ** $p < .01$.

Mini Kit (QIAGEN, Germany) and reverse transcribed using the QuantiNova Reverse Transcription Kit (QIAGEN, Germany), all of which followed the protocol described by the manufacturer. A five-point standard curve that consisted of 10^4 to 10^8 copies/ μ L of CAR plasmid DNA was prepared, and the copy number of these retroviral vectors were calculated by the absolute quantitative method. A 95 bp fragment containing portions of the MFG retroviral backbone sequence was amplified using a forward primer (5'-GACACCAGACTAAGAACCTAGAAC-3') and a reverse primer (5'-AGCTGCGATGCCGTCTACTTTGAG-3').

The copy number of the CAR gene was quantified by Q-PCR according to the protocol described previously.²⁸ The genomic DNAs from PBMCs 48 h post transduction were extracted using the QIAamp DNA Mini and Blood Mini Kit (Qiagen, Germany). A five-point standard curve was generated for analyses, consisting of 10^4 to 10^8 copies/ μ L of CAR plasmid DNA spiked into 100 ng of non-transduced T cells genomic DNA to control for background signals. The amplification of *GAPDH* was used as an internal control for the normalization of DNA quantities. The CAR transgene forward primer used was: 5'-ATCGCTCACAACCAGTCG-3'; while the reverse primer used was: 5'-GGTCAGGGAAGTTTACAAGG-3'. Q-PCR was performed with 100 ng of genomic DNA in each reaction using a QuantStudio™ 6 Flex Real-Time PCR System (Life technologies, United States).

Flow cytometry

For the determination of the transduction efficiency of T cells and retroviral vector titers, one million CAR-T cells or PG13 cells were harvested and stained with human c-Myc PE-conjugated antibody (R&D System, United States) for 30–60 mins at 4°C. The uncombined antibody solution was washed away using phosphate-buffered saline (PBS) (Gibco, United States). Fluorescent signals were detected using CytoFLEX (Beckman, United States) and analyzed in CytoFLEX analysis software.

Cytotoxicity assays

BCMA-K562, CD38-Raji, BCMA-K562+ CD38-Raji (mixed in 1:1) or RPMI-Luc target cells were seeded in a 96-well plate at 8×10^4 cells/well and co-incubated with Pan-T (un-transduced T cells), CD38 CAR-T, BCMA CAR-T, BCMA-OR-CD38 Tan CAR-T, or CD38-OR-BCMA Tan CAR-T cells at different effect-to-target ratios (1:16, 1:8, 1:4, 1:2, and 1:1). The effector cell seeding density was based on CAR⁺ cell counts. After 8 hours of incubation, these cells were stained with CD3-BV421 (BD, United States) for 60 mins. The cells were then washed with PBS and stained with Annexin V-Alexa Fluor 647 (Bio Friend, China) for 30 mins at room temperature. These cells were immediately analyzed by CytoFLEX flow cytometry.

The survival rates of BCMA-K562 or CD38-Raji cells were analyzed after BCMA-K562+ CD38-Raji cells (mixed in 1:1) were co-cultured with Pan-T and CAR-T cells at different effect-to-target ratios. After 48 hours of incubation, cells were harvested and stained with CD3-BV421 or CD38-APC (Invitrogen, United States) for 60 mins. Fluorescent signals

were detected using a CytoFLEX (Beckman, United States) and analyzed using FlowJo software (FlowJo LLC).

For RPMI-Luc cells, a luciferase assay was carried using the ONE-Glo™ EX Luciferase Assay System (Promega, United States) after co-culturing with CAR-T cells for 8 hours. The cells were reacted with substrate solution for 3 mins, and the relative luminescence units (RLU) were determined using a SpectraMax i3x Multi-Mode Microplate Reader (Molecular Devices, United States). The lysis rate of tumor cells was calculated according to the following formula:

$$\text{LysisRate(\%)} = 1 - \frac{(\text{RLU of experiment well} - \text{RLU of complete medium})}{(\text{RLU of maximum release well} - \text{RLU of complete medium})} \times 100\%$$

Cytokine release assays

Target cells were seeded in 96-well plates at 8×10^4 cells/well and co-cultured with Pan-T or CAR-T cells at a ratio of 1:1 for 8 hours. The effector cell seeding density was based on the CAR⁺ cells count. The cell culture supernatant samples were harvested, and the levels of various cytokines were evaluated using the LEGENDplex™ Multi-Analyte Flow Assay Kit (Human CD8/NK Panel) (BioLegend, United States). Capture beads consisting of IL-10, IL-6, TNF- α , IFN- γ , and Granzyme B were mixed and co-incubated with supernatant samples for 2 hours at room temperature on a shaker. After washing, the detection antibody cocktail and streptavidin-phycoerythrin (SA-PE) were added sequentially. The concentration of cytokines was determined using a standard curve generated in the same assay. Fluorescence signals were detected using a BD LSRFortessa™ Cell Analyzer (BD, United States) and flow data was analyzed using LEGENDplex™ online analysis software.

T cell proliferation assays

BCMA-K562+ CD38-Raji (mixed in 1:1) or RPMI-Luc target cells were seeded in 96-well plates at 8×10^4 cells/well. These Pan-T and CAR-T cells were stained with Carbo-xyfluorescein Diacetate Succinimidyl Ester (CFSE) (BD, United States) for 30–60 mins at 37°C. After washing with AIM-V medium, these CAR-T cells were incubated with target cells at a ratio of 1:1. The effector cell seeding density was based on the CAR⁺ cells count. After 24 h of incubation, the cells were stained with CD3-APC, the CFSE fluorescence intensity was determined using a CytoFLEX and the data was analyzed using FlowJo software. The Pan-T and CAR-T cell counts were performed using a Countess™ II Automated Cell Counter (Thermo Fisher, United States) every 2 days, and the T cell proliferation and viability curves were plotted using GraphPad Prism 7.0a software.

CAR-T cell activity detection in vivo

All animal experiments were performed with the approval of the Institutional Animal Care and Use Committee of Beijing University of Chinese Medicine. Six- to seven-week-old female NOD.Cg-Prkdc^{scid} Il2rg^{tmVst/Vst} (NPG) mice were obtained

from Vitalstar Biotechnology Co. (Beijing, China). To establish xenograft models, these NPG mice were intravenously injected (i.v.) with a RPMI-Luc tumor cell line (2×10^6 cells/mouse) on day 0. Then these mice were randomly divided into five groups (Untreated, Pan-T, CD38 CAR-T, BCMA CAR-T, BCMA-OR-CD38 Tan CAR-T and CD38-OR-BCMA Tan CAR-T) on day 12 after tumor injection. CAR T cells were injected intravenously into tumor-bearing mice at a dose of 1×10^8 /kg on days 12 and 19. The tumor progression was monitored using the MIIS living image system (Molecular Devices, United States) every 3 days beginning on day 4 after the second CAR-T injection. For *in vivo* imaging, the mice were injected intraperitoneally with 150 mg/kg VivoGlo™ Luciferin (Promega, United States) and imaged under isoflurane anesthesia using the MIIS living imaging system. We also utilized flow cytometry to further evaluate whether there were still residual tumor cells in these mice. In this experiment, blood was collected by removing the eyeball, liver, spleen, and bone marrow, and tumor tissues were isolated after these mice were sacrificed. The liver, spleen and tumor cells were obtained by grinding tissues carefully, while bone marrow cells were aspirated from femurs and tibias, and the lymphocytes were isolated by lysing red blood cells. Next, the cells were stained with anti-human CD38-APC (BD, United States) for 60 mins. After washing with PBS, the cells were stained with 7-AAD (5 μ l/test) (BD, United States) for 10 mins before flow cytometry (BD LSRFortessa™ Cell Analyzer) analysis. At different time points after the second CAR-T cell infusion, the percentage of CD3⁺ T cells in the peripheral blood was evaluated by flow cytometry. The blood was collected from the tail vein and stained with an anti-human CD3-BV786 antibody (BD, United States) and 7-AAD, this detection was carried out using a BD LSRFortessa™ Cell Analyzer. Two weeks after the second CAR-T injection, blood was collected via the tail vein, after at least 2 hours standing, the serums were obtained by centrifuging at $3000 \times g$ for 20 minutes at 4°C. The levels of human IFN- γ in serum from each group were measured using Human IFN-gamma ELISA Kit (Proteintech, United States). Finally, the mice were sacrificed when they lost the ability to eat and exercise spontaneously, and the survival time for each mouse was recorded to plot Kaplan-Meier survival curves.

Statistical analyses

Statistical analyses were performed using GraphPad Prism 7.0a software. For the comparisons of two groups, a two-tailed Student *t* test or nonparametric test was performed. When comparing multiple groups, a one-way ANOVA or a two-way ANOVA test, as appropriate, was applied. The significances of the differences in the Kaplan-Meier survival curves were analyzed using the Mantel-Cox log rank test. A *P* value of < 0.05 was considered statistically significant.

Discussion

Inspired by the success of CD19 CAR T-cells therapy for B-cell leukemia and lymphoma, several CAR T-cell therapies are also being developed for cancer, among which BCMA CAR-T has received much attention.^{29,30} BCMA, also known as tumor

necrosis factor receptor superfamily member 17 (TNFRSF17), is normally expressed solely on a subset of mature B-cells, whereas it is uniformly expressed by malignant plasma cells in many cases of MM.³¹ However, recent clinical trials have reported that individual patients with MM who received BCMA-targeted CAR-T cell therapy suffered relapses due to the loss of the BCMA antigen.^{12–15} To find other target antigens for CAR-T cells and develop new strategies for MM treatment, previously, we identified an anti-CD38 scFv from a scFv phage display library, and found that novel CD38 CAR-T cells had robust therapeutic effects on myeloma cells. Based on this, we further engineered single-chain bispecific (OR-gate) CARs that efficiently targeted not only BCMA, but also CD38, and evaluated the specificity and efficacy of these tan CAR T cells *in vitro* and *in vivo*.

Human CD38 antigen (45 kDa) is a single-chain type II transmembrane glycoprotein, and $> 90\%$ of malignant plasma cells from patients with MM show surface expression of CD38.³² In addition, CD38 is also expressed by cells of the immune system, including T-cells, B-cells, NK-cells, macrophages, and dendritic cells,³³ but the expression levels are lower compared to MM cells.³⁴ Thus, CD38 has been regarded as a feasible target for the treatment of MM, and several studies have indicated the efficacy and safety of CD38 monoclonal antibodies in clinical applications.^{35,36} Remarkably, an anti-CD38 antibody (Daratumumab) has been approved for the treatment of RRMM by the FDA.³⁷ Encouraged by these results, many researchers have explored the feasibility of developing CAR-T cell therapy targeting CD38 molecules. Early preclinical studies had shown that CD38 CAR-T cells were capable of proliferating, producing cytokines and effectively eliminating CD38⁺ myeloma cells.²² Recently, several clinical trials on anti-CD38 CAR-T-cell treatments of MM are in progress (NCT03464916, NCT03754764). Our study found that CD38-specific CAR-T cells showed significant lytic activity against myeloma cells *in vivo* and *in vitro*, and these cells showed no defects in *ex vivo* expansion.

Currently, tandem-dual CARs targeting two distinct antigens has become an effective way that can avoid tumor relapse due to antigen escape, potentially improve the efficacy issues with anti-BCMA CAR-T.³⁸ As illustrated above, CD38 is an attractive target antigen that is combinable with BCMA CAR in dual CAR T-cell targeting strategies to prevent MM relapse caused by loss of BCMA antigen expression. Thus, we designed Tan CARs that target BCMA and CD38, and evaluated the anti-tumor activity of these CAR T-cells against multiple myeloma *in vitro* and *in vivo*. The dual target specificity of Tan CAR was achieved via inserting dual-antigen recognition domains into a single CAR molecule. Besides Tan CAR, there are multiple approaches to achieving bi-specific signal recognition, such as co-expressing two different CARs in one T cell (called Dual CAR)³⁹ or mixing two CAR-T cell lines, each targeting a different antigen (called CAR pool).⁴⁰ Compared to Dual CAR, Tan CAR has a significantly smaller genetic payload ($\sim 40\%$ smaller in DNA length), which leads to more efficient viral vector packaging, higher transduction efficiency, and increased antigen-stimulated proliferation.^{20,41,42} These factors are beneficial for clinical T cell production and genetic modification of CAR-T cells.^{27,43} A CAR pool strategy avoids

the issue of poor transduction efficiency, but requires manufacturing two CAR-T products that significantly increases treatment cost and reduces the probability of successful T-cell production within a short clinical timeframe. More importantly, the two engineered T-cell populations may compete for the limited nutrients and homeostatic cytokines available in circulation.¹⁹ Finally, as mentioned above, CD38 is also expressed on the surface of T cells, which may cause the fratricide of CAR-T cells. Therefore, it is possible that co-administering BCMA and CD38 CAR-T cell populations might result in the disproportionate expansion of CD38 CAR-T cells at the expense of BCMA CAR-T cells, thereby compromising this strategy's ability to safeguard against the loss of BCMA. For this reason, we chose to engineer Tan CARs capable of dual-antigen recognition by attaching two tandem scFv domains to a single CAR molecule. Our data indicates that Tan-CAR T cells were indeed insensitive to the loss of BCMA on target cells, and triggered robust target-cell lysis capabilities in response to either BCMA or CD38 stimulation.

In this study, we designed two Tan CARs that had the antigen recognition domain from BCMA scFv and CD38 scFv in tandem via glycine and serine residues, a long, flexible linker peptide that has been utilized in the clinic since it has lower immunogenicity than other linkers.^{44,45} The difference between these two Tan CARs is the order of the two scFv domains. These Tan CAR T cells were transduced with retroviral vectors harvested from the PG13 retroviral vector producer cell line and the transduction efficiency exceeded 70%. We evaluated the effective targeting of Tan CAR-T on BCMA and CD38 both *in vivo* and *in vitro*. Our results showed that Tan-CAR-T cells could trigger robust T cell-mediated cytotoxicity and cytokine production after being stimulated with either BCMA or CD38. Moreover, these Tan-CAR-T cells had significantly higher cytotoxicity and proliferation than single-targeted CAR-T cells when encountering BCMA and CD38 antigens simultaneously. Notably, BCMA-OR-CD38 Tan CAR-T cells exhibited more potent antitumor and proliferative activity than CD38-OR-BCMA Tan CAR-T cells when they interacted with BCMA-K562, CD38-Raji, BCMA-K562+ CD38-Raji (mixed in 1:1), or RPMI-Luc cells. Thus, the therapeutic effect of BCMA-OR-CD38 Tan CAR T-cells was investigated using an immunodeficient mouse model bearing RPMI-Luc tumor cells. Our data showed that BCMA-OR-CD38 Tan CAR-T cells could achieve complete clearance of myeloma cells with no tumor relapse observed. Moreover, the levels of IFN- γ in serum were significantly higher for BCMA-OR-CD38 Tan CAR-T cell-treated mice than for single-targeted CAR-T cell-treated mice. Unfortunately, due to the lack of corresponding target cells, we were unable to further verify the specificity of BCMA-OR-CD38 Tan-CAR-T cells *in vivo*. Finally, the overall survival rate analysis showed that BCMA-OR-CD38 Tan CAR-T cells had similar safety as single-targeted CAR-T cells, no additional toxicities were observed. However, for a more reliable outcome further research has to include more detailed validation experiments. Since CD38 is also expressed by some key immune cell population, the on-target off-tumor effects of CD38-targeted CAR-T cells including Tan CAR-T in the treatment of MM has become

a problem that cannot be neglected. Researchers have made many attempts to address this issue, such as optimizing the design of the antigen recognition domain of CD38 CARs,⁴⁶ designing doxycycline (DOX) inducible Tet-on CD38-CARs,⁴⁷ knocking out⁴⁸ or block CD38 molecules.⁴⁹ In previous research, we found that shRNA-mediated knockdown of CD38 for CD38 CAR-T cells exhibited similar proliferative capacity with CD38 CAR-T cells, possibly due to the lack of expression of CD38 in CD38 CAR-T cells (*unpublished observation*).

Recently, studies about Tan-CAR-T cells targeting CD38 and BCMA for the treatment of MM have been increasing, and some of them have being test in clinical trials (ChiCTR1800018143, ChiCTR1900026286, NCT03767751). Notably, a clinical trial demonstrated that the bispecific anti-BCMA/anti-CD38 CAR-T cells had an improved efficacy and manageable safety profile for the therapy of relapsed or refractory multiple myeloma (RRMM).⁵⁰ In this study, we reported a novel BCMA-OR-CD38 Tan CAR T cell line that can robustly eliminate MM cells *in vitro* and *in vivo*. This BCMA-OR-CD38 Tan CAR-T may provide an effective and clinically applicable solution to the challenge of antigen escape, which has been observed in several clinical trials of BCMA CAR-T cell therapies. The process of the production of BCMA-OR-CD38 Tan CAR-T cells was consistent with the current clinical grade anti-CD19-CAR T cell manufacturing process, without the extra burden caused by large viral packaging and transduction payloads, which enabled the production of a single T-cell product targeting two clinically validated antigens associated with MM. In the future, after the efficacy and safety of BCMA-OR-CD38 Tan CAR-T cells *in vitro* and *in vivo* were sufficiently assessed, we also wish that the therapeutic effect of BCMA-OR-CD38 Tan CAR-T cells on MM can be tested in clinical trials.

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Disclosure statement

The authors declared they had no conflict of interests.

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Data availability statement

All of the data and materials are available from the corresponding author upon reasonable ask.

Authors' contributions

Jianxun Wang and Yaru Feng conceived and designed this study; Yaru Feng wrote the manuscript; Yaru Feng and Xiuying Liu performed the

most experiments; Xiaorui Li, Yating Zhou, Zhiru Song, Jing Zhang and Bingjie Shi assisted in the experiments. All the authors read and approved the final version of the manuscript.

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