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RECEIVED 31 July 2024 ACCEPTED 20 September 2024 PUBLISHED 18 October 2024

CITATION

King LA, de Jong M, Veth M, Lutje Hulsik D, Yousefi P, Iglesias-Guimarais V, van Helden PM, de Gruijl TD and van der Vliet HJ (2024) V δ 2 T-cell engagers bivalent for V δ 2-TCR binding provide anti-tumor immunity and support robust V γ 9V δ 2 T-cell expansion. *Front. Oncol.* 14:1474007. doi: 10.3389/fonc.2024.1474007

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V δ 2 T-cell engagers bivalent for V δ 2-TCR binding provide antitumor immunity and support robust V γ 9V δ 2 T-cell expansion

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Background: V γ 9V δ 2 T-cells are antitumor immune effector cells that can detect metabolic dysregulation in cancer cells through phosphoantigen-induced conformational changes in the butyrophilin (BTN) 2A1/3A1 complex. In order to clinically exploit the anticancer properties of V γ 9V δ 2 T-cells, various approaches have been studied including phosphoantigen stimulation, agonistic BTN3A-specific antibodies, adoptive transfer of expanded V γ 9V δ 2 T-cells, and more recently bispecific antibodies. While V γ 9V δ 2 T-cells constitute a sizeable population, typically making up ~1-10% of the total T cell population, lower numbers have been observed with increasing age and in the context of disease.

Methods: We evaluated whether bivalent single domain antibodies (VHHs) that link V δ 2-TCR specific VHHs with different affinities could support V γ 9V δ 2 T-cell expansion and could be incorporated in a bispecific engager format when additionally linked to a tumor antigen specific VHH.

Results: Bivalent VHHs that link a high and low affinity V δ 2-TCR specific VHH can support V γ 9V δ 2 T-cell expansion. The majority of V γ 9V δ 2 T-cells that expanded following exposure to these bivalent VHHs had an effector or central memory phenotype and expressed relatively low levels of PD-1. Bispecific engagers that incorporated the bivalent V δ 2-TCR specific VHH as well as a tumor antigen specific VHH triggered antitumor effector functions and supported expansion of V γ 9V δ 2 T-cells *in vitro* and in an *in vivo* model in NOG-hIL-15 mice.

Conclusion: By enhancing the number of $V\gamma 9V\delta 2$ T-cells available to exert antitumor effector functions, these novel V $\delta 2$ -bivalent bispecific T cell engagers may promote the overall efficacy of bispecific V $\gamma 9V\delta 2$ T-cell engagement, particularly in patients with relatively low levels of V $\gamma 9V\delta 2$ T-cells.

KEYWORDS

 $V\gamma9V\delta2$ T-cells, bispecific T-cell engager, single domain antibody, expansion, immunotherapy, cancer

Introduction

V γ 9V δ 2 T-cells constitute a homogeneous unconventional T-cell population that orchestrates both innate and adaptive immunity (1). V γ 9V δ 2 T-cells recognize target cells in an HLA-independent manner through phosphoantigen (pAg)-induced conformational changes in the butyrophilin (BTN) 2A1/3A1 complex (2, 3). pAg are metabolites that accumulate intracellularly due to dysregulation of the mevalonate pathway during cellular stress, caused by such processes as infection or malignant transformation, or upon exposure to aminobisphosphonates (N-BP) (4). Following activation, V γ 9V δ 2 T-cells can cross-present antigens (5–7), produce proinflammatory cytokines and chemokines, and induce cytotoxicity in a wide range of malignancies via membrane bound TRAIL and Fas ligand or through granzyme B/perforin release (8).

Although V γ 9V δ 2 T-cells make up ~1-10% of CD3⁺ T-cells in the circulation, their numbers can be reduced with increasing age and in the context of disease (9-14). Notably, the presence of $\gamma\delta$ T-cells within tumor-infiltrating immune cells (15) and the relative abundance of specifically tumor-infiltrating $V\gamma 9V\delta 2$ T-cells correlate to improved patient outcome (16, 17). Several therapeutic approaches designed to exploit the antitumor properties of Vy9V82 T-cells were explored. N-BPs such as pamidronate and zoledronate, as well as synthetic pAg analogues like bromohydrin pyrophosphate (BrHPP), were tested, either alone or in combination with IL-2, to activate $V\gamma 9V\delta 2$ T-cells in vivo or ex vivo followed by adoptive cell therapy (ACT) in various malignancies. While these strategies were overall well-tolerated and safe, and resulted in increased frequencies of circulating Vγ9Vδ2 T-cells across different types of cancer patients, only a minority of patients participating in these clinical trials exhibited significant signs of anti-tumor efficacy (8, 18, 19).

A novel approach based on V γ 9V δ 2 T-cell stimulation using an agonistic BTN3A-specific monoclonal antibody alone or in combination with IL-2 is currently being evaluated in a phase 2 clinical trial in advanced-stage cancer patients (20, 21). In addition, multiple efforts are ongoing to incorporate tumor-targeting moieties in Vy9V82 T-cell based cancer therapies, including the use of chimeric antigen receptors (CAR) and bispecific antibodies. Multiple studies, both preclinical and clinical, have reported promising antitumor activity upon administration of CARmodified or monoclonal antibody conjugated expanded (V δ 1 or V δ 2) $\gamma\delta$ T cells (22–27). Adoptive cell therapy, and in particular allogeneic CAR T-cell therapy, poses several challenges, not only related to the laborious nature of the manufacturing process with associated high costs, but also to limitations in transduction efficacy, product success rate, and the requirement of lymphodepleting preconditioning chemotherapy regimens to enhance their persistence (28, 29). Bispecific T-cell engagers (bsTCEs), which consist of linked antibody binding domains directed against a tumor associated antigen and a T-cell specific receptor, potentially provide a relatively cost-effective off-the-shelf approach to redirect endogenous T-cells to tumors (30). We and others have developed bispecific engagers to direct Vy9V82 T-cells to tumors and demonstrated that these have the potential to induce robust lysis of various malignancies in a tumor associated antigen restricted fashion (31–37). Importantly, early signs of potential anti-tumor efficacy were also noted during phase 1 dose escalation in patients treated with bispecific V γ 9V δ 2 T-cell engagers with a high-affinity V δ 2-TCR specific VHH (38, 39). One can envision that the antitumor efficacy of these bispecific V γ 9V δ 2 T-cell engagers can be further promoted by enhancing the number of V γ 9V δ 2 T-cells available to exert antitumor effector functions *in vivo*. To this end, we explored whether a V δ 2-TCR binding arm that consisted of two V δ 2-TCR specific VHHs could combine antitumor activity with expansion of V γ 9V δ 2 T-cells.

Here, we report that bispecific $\nabla\gamma9V\delta2$ T-cell engagers that combine a tumor antigen-specific single domain antibody (VHH) with a bivalent VHH, that links a high and low affinity V $\delta2$ -TCR specific VHH, trigger antitumor effector functions and support expansion of $\nabla\gamma9V\delta2$ T-cells *in vitro* and *in vivo*. By enhancing the number of $\nabla\gamma9V\delta2$ T-cells available to exert antitumor effector functions in the tumor microenvironment, these novel V $\delta2$ bivalent bsTCEs may promote the overall efficacy of bispecific $\nabla\gamma9V\delta2$ T-cell engagement, particularly in patients with relatively low levels of $\nabla\gamma9V\delta2$ T-cells.

Materials and methods

Tumor cell lines

SW480 tumor cells (EGFR⁺) were obtained from American Type Culture Collection (ATCC, CCL-228) and maintained in Dulbecco's Modified Eagle Medium (DMEM, 1965-039, Gibco) supplemented with 10% (v/v) fetal calf serum (FCS, 04-007-1A, Biological Industries), 0.05 mM β-mercaptoethanol (β-ME, 200-646-6, Merck), and 100 IU/ml sodium penicillin, 100 µg/ml streptomycin sulphate, and 2.0 mM L-glutamine (PSG, 10378-016, Life technologies). 22Rv1 tumor cells (PSMA⁺) were obtained from the European Collection of Authenticated Cell Cultures (ECACC, 5092802) and MM.1s tumor cells (kind gift from R. Groen, Amsterdam UMC, Vrije Universiteit, Amsterdam, the Netherlands) were transfected with CD1d as described previously (40) (referred to as MM.1s.CD1d). The 22Rv1 and MM.1s tumor cell lines were maintained in Roswell Park Memorial Institute 1640 (RPMI-1640, 22400089, Gibco) medium supplemented with 10% FCS, β-ME, and PSG. Tumor cell lines were tested regularly for Mycoplasma using PCR.

PBMC isolation and V γ 9V δ 2 T-cell cultures

Healthy donor- and cancer patient peripheral blood mononuclear cells (PBMC) were isolated from whole blood by density gradient centrifugation using LymphoprepTM (AXI-1114547, Fresenius). Healthy donor blood samples were obtained under written informed consent from Sanquin (Amsterdam, the Netherlands). Blood samples from cancer patients were obtained after approval by the institutional review board (medical ethical committee Amsterdam UMC, location VUmc) and written informed consent was obtained from all the participants from the

Amsterdam UMC (location VUmc, Amsterdam, the Netherlands). Expanded Vy9V82 T-cells were generated from healthy donor PBMC-derived Vγ9Vδ2 T-cells as described before (33). In short, $V\delta 2^+$ T cells were isolated from healthy donor PBMC using magnetic bead sorting using FITC-conjugated V82 antibody (Supplementary Table S1) in combination with anti-mouse IgG microbeads (130-048-401, Miltenyi Biotec). Purified Vγ9Vδ2 Tcells were stimulated weekly with irradiated feeder mix consisting of healthy donor PBMC (1 x 10⁶ cells/ml), JY cells (1 x 10⁵ cells/mL, 94022533, ECACC), IL-7 (10 U/mL, 207-IL-025, R&D Systems), IL-15 (10 ng/mL, 34-8159-85, eBioscience), and PHA (50 ng/mL, R30852801, Thermo Fisher Scientific). Purity of Vγ9Vδ2 T-cells used in experiments was ≥95%. All functional experiments were performed in RPMI medium supplemented with 10% FCS, β-ME, and PSG. Cell cultures were kept at 37°C in a humidified atmosphere containing 5% CO₂.

Design, production and purification of V δ 2-TCR-specific constructs

Human Vδ2-TCR specific llama-derived VHHs were generated and screened as previously described (41). In short, two Lama glamas were immunized with human Vy9V82 T-cells and V82-TCR-specific VHHs were selected using phage display followed by confirmation of specific binding to $V\delta 2^+$ T cells by flow cytometry. Bivalent constructs were generated by linking two monovalent V δ 2-TCR-specific VHHs with low (VHH-5C7, Kd ~350nM), intermediate (VHH-5D3, Kd ~21nM) or high (VHH-6H4, Kd ~0.4nM) affinity in various combinations using a Gly4Ser amino acid based linker [(G4S)2; referred to as 10 amino acid (AA) linker], resulting in the following bivalent constructs: 6H4-(G4S)2-5C7, 5C7-(G4S)2-6H4, 6H4-(G4S)2-6H4, 5D3-(G4S)2-6H4, 6H4-(G4S)₂-5D3, 5D3-(G4S)₂-5D3, 5D3-(G4S)₂-5C7, 5C7-(G4S)₂-5D3, 5C7-(G4S)₂-5C7. The bivalent construct 6H4-5C7 was generated with multiple linker lengths; G4S, (G4S)₂ or (G4S)₄ (referred to as 5, 10 or 20 AA linker). Purified protein was produced by ImmunoPrecise Antibodies (IPA, Utrecht, The Netherlands) using DNA transfected HEK293E cells and rmp Protein A affinity chromatography followed by preparative size exclusion. Proteins used were >95% pure and monomeric.

Three different tumor associated antigen (TAA) specific VHHs [i.e. directed against EGFR (VHH-7D12 (42)], PSMA [VHH-JVZ-007 (43)] or CD1d [VHH-1D12 (44)] were linked with a G4S linker to the N-terminus of 6H4-5C7 (termed $V\delta 2^{hi-lo}$): EGFR-G4S-6H4-(G4S)₂-5C7, PSMA-G4S-6H4-(G4S)₂-5C7 and CD1d-G4S-6H4-(G4S)₂-5C7 (referred to as EGFR-V $\delta 2^{hi-lo}$ bsVHH, PSMA-V $\delta 2^{hi-lo}$ bsVHH and CD1d-V $\delta 2^{hi-lo}$ bsVHH). Next, Fc domain and antialbumin VHH containing molecules were created to allow for *in vivo* plasma half-life extension. For the Fc domain, knobs-intoholes technology was used for heterodimerization (HC1, knob mutation T366W and HC2, hole mutations T366S, L368A, Y407V), LFLE mutations were included to silence the Fc domain except for the neonatal Fc receptor (FcRn) (L234F and L235E silencing, EU numbering), a Cys220 deletion was incorporated to avoid an unpaired cysteine and a modified hinge was used (HC1: 5C7-G4S-6H4-hinge-Fc paired with HC2: EGFR-hinge-Fc, PSMAhinge-Fc or CD1d-hinge-Fc; referred to as EGFR-V $\delta 2^{\rm hi-lo}$ bsVHH-Fc, PSMA-Vδ2^{hi-lo} bsVHH-Fc and CD1d-Vδ2^{hi-lo} bsVHH-Fc. Expression of these tumor targeting VHHs was performed via pcDNA3.1(+) (100µg, Genscript) transfection in Expi293F cells (A14527, Gibco) using the ExpiFectamine 293 Transfection Kit (A14524, Gibco). Proteins were purified using rmp Protein A Sepharose Fast Flow resin (17-5138-03, GE healthcare) (EGFR-Vo2^{hi-lo} bsVHH, PSMA-Vo2^{hi-lo} bsVHH and CD1d-Vo2^{hi-lo} bsVHH) or CaptureSelect C-tagXL Affinity Matrix (2943072050, Thermo Scientific) (EGFR-V\delta2^{hi-lo} bsVHH-Fc, PSMA-Vδ2^{hi-lo} bsVHH-Fc or CD1d-Vδ2^{hi-lo} bsVHH-Fc). Quality control was done using size exclusion ultra performance liquid chromatography. For fusion to an anti-albumin (Alb) VHH (45), the following construct was generated: EGFR-(G4S)2-Alb-(G4S)2-6H4-G4S-5C7 (referred to as EGFR-Vδ2^{hi-lo} bsVHH-albumin; illustrated in Supplementary Figure S4A). Purified protein for the construct was produced by IPA as described above. Proteins used were >95% pure and >94% heterodimer shown by native mass spectrometry analysis. Lastly, bispecific engagers containing a monovalent Vδ2-TCR-specific VHH linked with a TAA-specific VHH, i.e. EGFR-G4S-5C8, PSMA-G4S-5C8 and CD1d-G4S-5C8 (referred to as EGFR-Vδ2 bsVHH, PSMA-Vδ2 bsVHH and CD1d-Vδ2 bsVHH), were generated as described before (31, 32) and are illustrated in Supplementary Figure S3A.

Flow cytometry

Cells were resuspended in PBS (1073508600, Fresenius Kabi) supplemented with 0.5% bovine serum albumin (M090001/03, Fisher Scientific) and 20 μ g/ml NaN3 (247-852-1, Merck) and incubated with fluorochrome-labeled monoclonal antibodies and viability dyes (Supplementary Table S1) for 30 min at 4°C. Unbound fluorochrome-labeled antibodies were washed away. Cells were analyzed using the FACS LSRFortessa XL-20 (BD Biosciences) and data analysis was performed using FlowJo v10.8.1 (BD Biosciences).

Target cell binding by V δ 2-TCR specific constructs

Binding of the various bivalent and bispecific VHH constructs to V δ 2, EGFR, PSMA or CD1d was evaluated using flow cytometry or ELISA, as indicated. For flow cytometry analysis, V γ 9V δ 2 T-cells, SW480, 22Rv1 or MM.1s.CD1d tumor cells were incubated with a concentration range of the constructs for 45 min at 4°C, followed by extensive washing (5x) and incubation for 30 min at 4°C with FITClabeled rabbit-anti-llama polyclonal Ab (Supplementary Table S1). For the ELISA, wells of a clear bottom F96-well Maxisorp plate (439454, Nunc) were coated overnight with human Gamma9Delta2 TCR (0.5 µg/mL, produced by IPA). Wells were blocked with 2% BSA (A2153, Sigma) for 60 min, and test samples were incubated for 120 min at room temperature. The rabbit anti-camelid VHH cocktail (60-min incubation, 19L002038, GenScript) and swine anti-rabbit immunoglobulins HRP (60-min incubation, 41289300, Dako) were used for detection and 3,3',5,5'-Tetramethylbenzidine (TMB, Life Technologies, #SB02) was used as a substrate for color development. The absorbance (450 nm) was measured using a plate reader (Molecular Devices, iD5) and the data analysis was done using SoftMax Pro software v 7.1.

Assessment of V γ 9V δ 2 T-cell frequency, expansion and phenotype using healthy donor and cancer patient PBMC

Vγ9Vδ2 T-cell percentages in healthy donor and cancer patientderived PBMC (patient characteristics provided in Supplementary Table S2) were assessed using flow cytometry using fluorescentlylabelled antibodies against CD3, Vy9, V82 (Supplementary Table S1). The capacity of the various bivalent and bispecific VHH constructs to induce Vy9V82 T-cell expansion was assessed by incubating healthy donor or cancer patient-derived PBMC (patient characteristics provided in Supplementary Table S3) with indicated concentrations of the bivalent or bispecific constructs, 10 µM pamidronate disodium salt hydrate (referred to as pamidronate, P2371-10MG, Sigma) (positive control) or medium (negative control). Twenty-four hrs later, 100 IU/ml IL-2 (Proleukin, Clinigen) was added to all conditions (including medium, termed IL-2 control). Fold expansion and enrichment of $V\gamma 9V\delta 2$ T-cells (compared to baseline) were assessed using flow cytometry at the end of the 8-day culture period using fluorescently-labelled antibodies against CD3, V γ 9, V δ 2 and 123counting eBeadsTM (01-1234-42, Thermofisher). Depending on the experiment, 7-AAD (A9400-1MG, Sigma), fixable viability dves eFluorTM 506 or eFluorTM 780 (Supplementary Table S1) were used to identify viable cells. The phenotype of expanded Vy9V82 T-cells was determined using fluorescently-labelled antibodies against CD27, CD45RA, CD25, CD69, HLA-DR, DNAM-1, NKG2D, NKG2A, PD-1, CTLA-4 and TIGIT (see Supplementary Table S1 for clones and fluorophores).

Assessment of V γ 9V δ 2 T-cell degranulation and target cell lysis

V γ 9V δ 2 T-cell degranulation and subsequent tumor cell lysis were assessed by flow cytometry. Expanded healthy donor-derived V γ 9V δ 2 T-cells were incubated with SW480, 22Rv1 or MM.1s.CD1d tumor cells (1:1 E:T ratio) for 24hrs with a concentration range of the TAA-V δ 2^{hi-lo} bsVHH with or without Fc domain or anti-albumin-VHH, or medium control. Degranulation and activation of V γ 9V δ 2 T-cells was assessed using fluorescently-labelled antibodies against CD3, V γ 9, V δ 2, CD25 and CD107a (Supplementary Table S1) and tumor cell lysis was determined using 7-AAD and 123counting eBeads^{TM.} To assess degranulation and tumor cell lysis of 8-day expanded V γ 9V δ 2 T- cells in healthy donor derived PBMC, $\gamma\delta$ T-cells were isolated from the PBMC cultures using the untouched human TCR γ/δ^+ T-cell isolation kit (130-092-892, Miltenyi Biotec) and used as described above in the presence or absence of the respective TAA-V δ 2^{hi-lo} bsVHH or TAA-V δ 2^{hi-lo} bsVHH-Fc engagers. Supernatants were collected and stored at -20°C until further analysis for IL-2, TNF and IFN- γ secretion using the human Th1/Th2/Th17 cytometric bead array (CBA) kit (560484, BD). CBA data were analyzed with FCAP Array software v3.0 (BD).

In vivo mouse study

Sub-lethally irradiated (1.75 Gy on day 0) NOD.Cg-Prkdcscid Il2rgtm1Sug/JicTac (NOG)-hIL-15 mice (8.5 weeks old, #13683-F Taconic Biosciences) were kept under pathogen-free conditions (Laboratory Animal Center of the Netherlands Cancer Institute, Amsterdam, The Netherlands) and used for evaluation of in vivo expansion of Vγ9Vδ2 T-cells in PBMC. This in vivo experiment was approved by the Animal Welfare Committee of the Netherlands Cancer Institute and was performed in accordance with national guidelines. Mice were randomized into 2 groups (n=4/group) and intravenously (i.v.) inoculated with 10x10⁶ healthy human donorderived PBMC (n=1 healthy donor with 0.89% Vy9V82 T-cells of total CD3 $^{\rm +}$ T-cells) on day 0. PBS or the CD1d-V $\delta 2^{\rm hi-lo}$ bsVHH (0.5 mg/kg) were intraperitoneally (i.p.) administered on days 0 and 4. Blood (day 1 and 8) and lungs, spleens and livers (day 8) were collected and prepared for flow cytometry analysis. Erythrocytes were lysed using lysis buffer (0.16 M NH₄Cl; 76050139.1000, Boom and 0.011 M KHCO3; 104854, Merck). Lungs were dissociated while stirring in a flask for 45 min at 37°C using Iscove's Modified Dulbecco's Medium (IMDM) supplemented with 5% FCS, PSG, 0.1% DNAse I (10104159001, Roche) and 0.14% collagenase A (10103586001, Roche). Spleens and livers were dissociated using a 100 µM cell strainer (352360, Falcon) and a syringe plunger, and dissociated cells were washed with PBS. Blood and tissue-derived single cell suspensions were used for flow cytometry analysis using a combination of 7-AAD, fluorescently-labelled antibodies against CD45 (mouse), CD45 (human), CD3, Vγ9, Vδ2, CD27, CD45RA and 123 counting eBeads TM for V γ 9V δ 2 T-cell analysis (Supplementary Table S1).

Statistical analysis

Analysis were performed using Prism v.9.1.0 (GraphPad Software). Data were analyzed using unpaired or paired *t* test, one-way analysis of variance (ANOVA) with Tukey's comparisons test, or two-way ANOVA with Tukey's multiple comparisons test, as appropriate. Binding curves and dose–response curves with EC_{50} values were calculated using nonlinear regression analysis. To calculate the correlation between V γ 9V δ 2 T-cell frequency and donor age, Spearman's rank correlation analysis was used. *P*=< 0.05 was considered significant and showed with asterisks: *P*=< 0.05: *, *P*=< 0.01: **, *P*=< 0.001: ***.

Results

$V\gamma 9V\delta 2$ T-cell frequencies in peripheral blood of healthy adult donors and adult cancer patients

Given that the V γ 9V δ 2 T-cell frequency in peripheral blood has been reported to decline with age and can be reduced due to underlying malignancies (9-14), we assessed the frequency of Vγ9Vδ2 T-cells in peripheral blood of healthy donors [n=121; 68 females, 53 males, mean age (range): 41 (18-77)] and cancer patients with various solid tumors (n=91; see Supplementary Table S2 for patient characteristics). As shown in Supplementary Figure S1A, the $V\gamma 9V\delta 2$ T-cell frequency varied substantially between individuals and was significantly higher in healthy donor compared to cancer patient PBMC ($3.5 \pm 0.3\%$ versus $2.2 \pm 0.4\%$ of total T cells; mean \pm SEM, P = 0.01). In healthy female and male donors, the Vy9V82 T-cell frequency negatively correlated with donor age (Supplementary Figure S1B). This negative correlation between Vy9V82 T-cell frequency and age was not (statistically significantly) observed in cancer patients, though the population of adult patients with solid tumors evaluated in our analysis was generally skewed towards older age (66 \pm 0.9 yr versus 40 \pm 1.5 yr in healthy donors; mean ± SEM; Supplementary Figure S1C). Of interest, the frequency of Vγ9Vδ2 T-cells was comparable in cancer patients and healthy donors with age ≥ 39 yr (V γ 9V δ 2 T-cell frequency 2.0 \pm 1.9% versus 2.3 \pm 3.3% of total T cells in healthy donors \geq 39 yr and cancer patients \geq 39 yr respectively; mean \pm SEM; Supplementary Figure S1D), suggesting that the Vγ9Vδ2 Tcell frequencies in peripheral blood of cancer patients are predominantly influenced by age, rather than by the underlying malignancy.

A bivalent V δ 2-TCR specific VHH that combines high and low affinity V δ 2 binding domains supports V γ 9V δ 2 T-cell expansion

As TCR cross-linking, which is not achieved with monovalent TCR binding, initiates signaling typically resulting in activation, differentiation, cytokine production and proliferation, we evaluated whether linkage of two V82-TCR specific VHHs could promote Vγ9Vδ2 T-cell expansion. Three Vδ2-TCR specific VHHs were selected based on affinity (i.e. VHH-5C7, Kd ~350 nM; VHH-5D3, Kd ~21 nM; and VHH-6H4, Kd ~0.4 nM) (ref (41) and not shown), and differentially combined and linked using a 10 AA linker into nine different bivalent VHHs. Binding of the various bivalent Vδ2-VHHs to Vγ9Vδ2 T-cells was assessed using flow cytometry or ELISA and showed that the binding EC50 was comparable for compounds that contained a high affinity VHH linked to either a low or intermediate affinity VHH or two intermediate-binding VHHs ($EC_{50} = 0.05-0.43$ nM) (Figure 1A). For the high and low affinity VHH combination, N-terminal positioning of the high affinity VHH-6H4 resulted in a lower EC₅₀ than when this VHH was positioned C-terminally. Compounds consisting of an intermediate and a low affinity VHH bound with higher EC₅₀ values of 17-22 nM. The binding EC₅₀ of the linked low affinity VHHs could not be reliably assessed using flow cytometry and was therefore determined using ELISA, resulting in a value of 211 nM (Figure 1A right panel). To assess whether the bivalent V δ 2-VHHs could support Vy9V82 T-cell expansion, healthy donor derived PBMC were cultured for 8 days in the presence or absence of 1 nM (optimal concentration across the compounds as determined by titration, data not shown) of the bivalent Vδ2-VHHs or 10 μM pamidronate. After 24 hrs, 100 IU/ml of recombinant human (rh) IL-2 was added to the cultures (including the medium control, termed IL-2 control). Figure 1B illustrates the design of the experiment and Figure 1C shows a representative gating strategy for the assessment of the $V\gamma 9V\delta 2$ T-cell frequency. As shown in Figure 1D, maintained or increased Vy9V82 T-cell numbers were only observed in the conditions exposed to bivalent VHHs containing either 2 linked low affinity VHHs or the low affinity VHH linked to the high affinity VHH. When enrichment of Vγ9Vδ2 T-cells was assessed, a similar pattern was observed. Enrichment and expansion were most consistent and pronounced when PBMC were cultured with the bivalent high-low affinity 6H4-5C7 (i.e. N-terminally positioned 6H4) VHH (enrichment 35.3 ± 49.4%; median \pm IQR; P = 0.03, and expansion 5.6 \pm 12.9 fold; median \pm IQR; P = 0.07, paired t tests). Most of the other bivalent Vδ2-VHHs did not support enrichment and expansion and in some cases even resulted in a reduction of $V\gamma 9V\delta 2$ T-cell numbers, possibly due to fratricide. As expected, enrichment and expansion were also observed when PBMC were cultured in the presence of pamidronate. In the absence of IL-2, neither pamidronate nor bivalent Vδ2-VHHs triggered Vγ9Vδ2 T-cell expansion (Supplementary Figure S2). To assess the impact of different linker lengths, bivalent VHHs with the high affinity V82-TCR VHH-6H4 and the low affinity V δ 2-TCR VHH-5C7 were generated with either a G4S, (G4S)₂ or (G4S)₄ linker (referred to as 5, 10 or 20 AA linker). As shown in Figures 1E, F, these bivalent (6H4-5C7) V δ 2-VHHs showed a similar binding profile (i.e. similar EC₅₀ and maximum binding) and similar enrichment and expansion of V γ 9V δ 2 T-cells in the 8 day PBMC cultures. Enrichment and expansion of $V\gamma 9V\delta 2$ T-cells was already observed using 0.1 nM of the bivalent (6H4-5C7) Vδ2-VHHs and not further enhanced with higher concentrations. The bivalent V δ 2-VHH with the high affinity VHH-6H4 (N-terminal) a 10 AA linker and the low affinity VHH-5C7 was selected for further experiments and termed bivalent $V\delta 2^{hi-lo}$ VHH.

Bispecific engagers that incorporate a bivalent V δ 2-TCR-specific VHH combine V γ 9V δ 2 T-cell expansion and tumor associated antigen specific effector functions

To explore whether a bispecific engager that includes the bivalent $V\delta 2^{hi-lo}$ VHH as well as a VHH specific for a tumor associated antigen (TAA) could combine TAA directed V γ 9V δ 2 T-cell effector functions and V γ 9V δ 2 T-cell expansion, six bispecific engagers were generated (see Figure 2A for design). The engagers



using flow cytometry (n=3) or ELISA (n=2). (B) Overview of experimental design to assess expansion. (C) Representative gating strategy to asses enriched and expanded $V\gamma9V\delta2$ T-cells within PBMC. (D) Enrichment (left panel) and fold expansion (right panel) of $V\gamma9V\delta2$ T-cells during an 8 day culture of healthy donor PBMC in the presence or absence of 1 nM bivalent $V\delta2$ -VHHs or 10 µM pamidronate (n=3-6). (E) Binding of the 6H4-5C7 bivalent VHH with either a 5, 10 or 20 AA linker to $V\gamma9V\delta2$ T-cells (n=3). (F) Enrichment (left panel) and fold expansion (right panel) of $V\gamma9V\delta2$ T-cells during an 8 day culture of healthy donor PBMC in the presence or absence of 0.001, 0.1, 1 or 100 nM of the 6H4-5C7 bivalent VHH with either 5, 10 or 20 AA linker or 10 µM pamidronate. Data in (A) (left panel) and (D–F) assessed using flow cytometry. Data in A (right panel) assessed using ELISA. Data represent mean and SEM (A, E) or individual data-points are indicated using open circles and box and whisker plots indicate the median, 25th to 75th percentiles and minimum to maximum (D, F). Two-way ANOVA with Tukey's multiple comparisons test (F) were used for statistical analysis; P=<0.05: *.

included the bivalent $V\delta 2^{hi-lo}$ VHH either directly linked to a VHH directed against EGFR [clone 7D12 (42)], PSMA [clone JVZ-007 (43)] or CD1d [clone 1D12 (44)] at the N-terminus or (indirectly) linked to these VHHs using Fc domains that heterodimerize using knobs-into-holes technology (see Materials and Methods). The latter formats would allow for half-life extension of the engagers when applied *in vivo*.

All six constructs bound with similar EC_{50} values of 0.2-0.5 nM to V γ 9V δ 2 T-cells (Figure 2B) and supported significant V γ 9V δ 2 T-cell enrichment (median 20.6-58.6% of total T cells, versus 2.9% in IL-2 control) and expansion (median 12.2-32.4 fold expansion, versus 1.4 fold in IL-2 control) when added at a concentration of 1 nM (TAA-V δ 2^{hi-lo} bsVHH) or 100 nM (TAA-V δ 2^{hi-lo} bsVHH-Fc) to healthy donor derived PBMC for 8 days (optimal concentrations determined by titration, not shown), illustrating that the ability to induce V γ 9V δ 2 T-cell expansion was retained in these molecular formats (Figure 2C). Enrichment and expansion of V γ 9V δ 2 T-cells was not observed when a monovalent V δ 2-VHH was linked to the

EGFR or PSMA-specific VHHs, underscoring the importance of including the bivalent $V\delta 2^{hi-lo}$ VHH when expansion is desired (Supplementary Figures S3A, B). Of note, enrichment and expansion of V γ 9V δ 2 T-cells was observed when PBMC were cultured with CD1d specific engagers containing either the monovalent V δ 2-VHH or the bivalent V δ 2^{hi-lo} VHH, which was likely related to CD1d expressing cells in PBMC (e.g. monocytes, B cells, circulating dendritic cells) that allowed V δ 2-TCR crosslinking to support V γ 9V δ 2 T-cell expansion (32).

After binding of these TAA-V $\delta 2^{hi-lo}$ bsVHH and TAA-V $\delta 2^{hi-lo}$ bsVHH-Fc molecules to the TAA expressing tumor cells was confirmed (Figure 2D), the antitumor effector functions of these molecules were evaluated in co-cultures of V $\gamma 9V\delta 2$ T-cells and tumor cells expressing the respective TAAs (i.e. SW480 (EGFR⁺), 22Rv1 (PSMA⁺) or MM.1s.CD1d (CD1d⁺)). A representative gating strategy is shown in Supplementary Figure S5A. As shown in Figures 2E, F, all constructs were able to induce V $\gamma 9V\delta 2$ T-cell degranulation and subsequent tumor cell lysis with EC_{50s} in the

low-picomolar range. Addition of the Fc domain resulted in a 3-9 fold reduced potency for tumor lysis, possibly due to steric hindrance or an increased intermembrane distance between the $V\gamma 9V\delta 2$ T-cell and TAA⁺ tumor target cell which may interfere with efficient immune synapse formation (46).

Incorporation of an anti-albumin binding domain can also provide a means to extend *in vivo* half-life (45). To explore this, we added an anti-albumin VHH to the EGFR-V $\delta 2^{hi-lo}$ bsVHH (see

Supplementary Figure S4A for design) and confirmed binding to both V γ 9V δ 2 T-cells and EGFR⁺ tumor cells (Supplementary Figure S4B). The EGFR-V δ 2^{hi-lo} bsVHH-albumin retained the ability to support V γ 9V δ 2 T-cell enrichment and expansion in healthy donor PBMC (Supplementary Figure S4C) and also triggered V γ 9V δ 2 T-cell degranulation and lysis of SW480 tumor cells (Supplementary Figure S4D). The potency (i.e. EC₅₀ for degranulation and tumor lysis) of the EGFR-V δ 2^{hi-lo} bsVHH-



FIGURE 2

TAA-V $\delta 2^{hi-lo}$ bsVHH and TAA-V $\delta 2^{hi-lo}$ bsVHH-Fc support V $\gamma 9V\delta 2$ T-cell expansion and trigger V $\gamma 9V\delta 2$ T-cell degranulation and tumor cell tysis. (A) Illustration of the TAA-V $\delta 2^{hi-lo}$ bsVHH with and without Fc domain. (B) Binding of the TAA-V $\delta 2^{hi-lo}$ bsVHH with and without Fc to V $\gamma 9V\delta 2$ T-cells. Data represent mean and SEM (n=3). (C) Enrichment (left panel) and fold expansion (right panel) of V $\gamma 9V\delta 2$ T-cells during an 8 day culture of healthy donor PBMC in the presence or absence of 1 nM TAA-V $\delta 2^{hi-lo}$ bsVHH, 100nM TAA-V $\delta 2^{hi-lo}$ bsVHH-Fc or 10 µM pamidronate. Individual data-points are indicated using open circles and box and whisker plots indicate the median, 25th to 75th percentiles and minimum to maximum (n=15-30). (D) Binding of the TAA-V $\delta 2^{hi-lo}$ bsVHH and TAA-V $\delta 2^{hi-lo}$ bsVHH-Fc to SW480 (EGFR⁺), 22Rv1 (PSMA⁺) or MM.1s.CD1d (CD1d⁺) tumor cells. Data represent mean and SEM (n=3). (E) V $\gamma 9V\delta 2$ T-cell CD107a expression after 24 hr co-cultures of V $\gamma 9V\delta 2$ T-cells and SW480, 22Rv1 or MM.1s.CD1d tumor cells (1:1 E:T ratio) \pm concentration range of the TAA-V $\delta 2^{hi-lo}$ bsVHH and TAA-V $\delta 2^{hi$ albumin construct was ~10 fold lower compared to the EGFR-V $\delta 2^{hi-lo}$ bsVHH-Fc perhaps as a result of impaired immune synapse formation when albumin was bound to the engager in this specific orientation.

Vγ9Vδ2 T-cells expanded using TAA-Vδ2^{hilo} bsVHH and bsVHH-Fc are activated and perform cytotoxic effector functions when exposed to TAA expressing tumor cells

The functional properties of the expanded V γ 9V δ 2 T-cells were next explored. For this purpose, expanded Vy9V82 T-cells were enriched from the PBMC cultures using negative MACS isolation and co-cultured with either EGFR⁺ SW480 colorectal cancer cells, PSMA⁺ 22Rv1 prostate cancer cells, or CD1d⁺ MM.1s multiple myeloma cells for an additional 24 hr; the TAA-Vδ2^{hi-lo} bsVHH/ bsVHH-Fc engagers were re-added to the respective conditions. Figure 3A illustrates the experimental design and Supplementary Figures S5B, C show the Vy9V82 T-cell frequency post expansion and post enrichment and a representative dot-plot thereof. Compared to Vy9V82 T-cells cultured with IL-2 only (IL-2 control), V γ 9V δ 2 T-cells expanded using pamidronate, the V δ 2^{hi-} ^{lo} bivalent VHH, or the V $\delta 2^{hi-lo}$ bsVHH/bsVHH-Fc typically expressed higher levels of the activation marker CD25 upon coculture with tumor cells (Figure 3B). In the presence of the TAA- $V\delta 2^{hi-lo}$ bsVHH and bsVHH-Fc molecules, the expanded V $\gamma 9V\delta 2$ T-cells expressed significantly higher levels of the degranulation marker CD107a and, in line, triggered more robust tumor lysis (Figures 3C, D; plots from a representative donor are shown in Supplementary Figures S5D, E). Supernatants obtained from these 24 hr co-cultures demonstrated that the TAA-V $\delta 2^{hi\text{-lo}}$ bsVHH molecules promoted the production of higher levels of various proinflammatory cytokines, i.e. IL-2, TNF and IFN-γ (Figure 3E). Overall, the increase in proinflammatory mediators and cytolytic effects were most pronounced when expanded $V\gamma 9V\delta 2$ T-cells were cultured with TAA-Vδ2^{hi-lo} bsVHH. Taken together, Vγ9Vδ2 Tcells expanded from PBMC with the TAA-V82^{hi-lo} bsVHH and bsVHH-Fc exhibit an activated effector phenotype and mediate tumor lysis when exposed to TAA expressing tumor cells.

$V\gamma 9V\delta 2$ T-cells can be expanded from cancer patient PBMC using TAA-V $\delta 2^{hi\text{-lo}}$ bsVHH and bsVHH-Fc

We next evaluated whether the TAA-V $\delta 2^{hi-lo}$ bsVHH and bsVHH-Fc molecules could also support expansion of V γ 9V $\delta 2$ Tcells from cancer patient PBMC. For this purpose, PBMC were obtained from patients with gastric cancer, esophageal cancer or melanoma. The majority of these patients had advanced-stage disease and was treatment naïve (see Supplementary Table S3 for patient characteristics). As shown in Figure 4A, pamidronate, the bivalent V $\delta 2^{hi-lo}$ VHH as well as the TAA-V $\delta 2^{hi-lo}$ bsVHH and TAA-V $\delta 2^{hi-lo}$ bsVHH-Fc constructs supported the enrichment and expansion of V γ 9V $\delta 2$ T-cells in 8 day PBMC cultures. Enrichment (55.9 \pm 33.2%; median \pm IQR) and expansion (69.8 \pm 79.5 fold; median \pm IQR) was most pronounced with the CD1d-V82^hi-lo bsVHH and less striking when the Fc domain was connected to this bsVHH. With the EGFR and PSMA specific V82^hi-lo engagers, expansion was similar in the presence and absence of the Fc domain.

TAA-V $\delta 2^{hi-lo}$ bsVHH and bsVHH-Fc expanded V $\gamma 9V\delta 2$ T-cells display a central/ effector memory phenotype in healthy donors and retain a diverse memory dominated phenotype in cancer patients

The phenotype of Vy9V82 T-cells expanded from healthy donor and cancer patient PBMC using pamidronate, the $V\delta 2^{hi-lo}$ bivalent VHH, the various TAA-V $\delta 2^{\rm hi-lo}$ bsVHH and TAA-V $\delta 2^{\rm hi-lo}$ bsVHH-Fc molecules and the IL-2 control was characterized (representative gating strategy shown in Supplementary Figure S6). At baseline, most Vγ9Vδ2 T-cells in healthy donor PBMC had a central memory (CD27⁺CD45RA⁻, 47.1 \pm 4.5%; mean \pm SEM) phenotype with naïve (CD27⁺CD45RA⁺, 23.6 ± 4.2%; mean ± SEM), effector memory (CD27⁻ CD45RA⁻, 17.1 ± 4.6%; mean ± SEM) and terminally differentiated (CD27⁻CD45RA⁺, 12.1 ± 1.9%; mean \pm SEM) V γ 9V δ 2 T-cell populations representing smaller fractions (Figure 4B). After an 8-day culture of PBMC with 100 IU/ml IL-2, pamidronate or the various TAA-Vδ2^{hi-lo} bsVHH and bsVHH-Fc engagers, expanded Vy9V82 T-cells exhibited a shift towards an effector memory phenotype with a concomitant reduction in the frequency of naïve and terminally differentiated subsets while the fraction of central memory cells overall remained similar in size. Of interest, $V\gamma 9V\delta 2$ T-cells expanded with the TAA-V $\delta 2^{hi-lo}$ bsVHH-Fc molecules retained larger proportions of $V\gamma 9V\delta 2$ T-cells with a terminally differentiated phenotype (EGFR-V $\delta 2^{hi-lo}$ bsVHH; P = 0.0007, PSMA-V $\delta 2^{hi-lo}$ bsVHH; P =0.02, CD1d-V $\delta 2^{hi-lo}$ bsVHH; P = 0.01, unpaired t tests) and naïve phenotype (EGFR-V $\delta 2^{hi-lo}$ bsVHH; P = 0.008, PSMA-V $\delta 2^{hi-lo}$ bsVHH; P = 0.0002, CD1d-V $\delta 2^{hi-lo}$ bsVHH; P = 0.002, unpaired t tests). When compared to $V\gamma 9V\delta 2$ T-cells in healthy donor PBMC, the proportion of central memory $V\gamma 9V\delta 2$ T-cells was increased in cancer patient PBMC at baseline (72.2 \pm 6.8% versus 47.1 \pm 4.4% in healthy donors; mean \pm SEM, P = 0.001; unpaired t test). This was accompanied by a lower proportion of $V\gamma 9V\delta 2$ T-cells with a naïve phenotype (9.7 \pm 2.6% versus 23.6 \pm 4.2% in healthy donors; mean \pm SEM, *P* = 0.02; unpaired *t* test), which may suggest their priming and involvement in the antitumor immune response (Figure 4C). The V γ 9V δ 2 T-cells expanded from cancer patient PBMC using the $TAA\text{-}V\delta2^{hi\text{-}lo}$ bsVHH and bsVHH-Fc were in general skewed to an effector memory phenotype. However, while the proportion of naïve and terminally differentiated effector memory cells was strongly reduced upon expansion in $V\gamma 9V\delta 2$ T-cells expanded from healthy donor PBMC, these were maintained in Vy9V82 Tcells expanded from cancer patient PBMC (Figure 4C).

Healthy donor V γ 9V δ 2 T-cells expanded by the TAA-V δ 2^{hi-lo} bsVHH and bsVHH-Fc constructs as well as pamidronate were highly activated as shown by strong increases in CD25 and HLA-

DR expression (Figure 4D). During culture, expression of DNAM-1, an activating receptor expressed by most $V\gamma9V\delta2$ T-cells, remained highly expressed and was even expressed by a larger proportion in several cases. Of interest, NKG2D, an activating receptor also expressed by most $V\gamma9V\delta2$ T-cells, remained highly expressed when $V\gamma9V\delta2$ T-cells were expanded with pamidronate or the TAA-V $\delta2^{hi-lo}$ bsVHH, but was expressed by a lower proportion of $V\gamma9V\delta2$ T-cells when these were expanded with the TAA-V $\delta 2^{hi-lo}$ bsVHH-Fc engagers, though this did not reach statistical significance. The upregulation of activation markers on expanded V γ 9V $\delta 2$ T-cells was accompanied by an increase in the proportion of V γ 9V $\delta 2$ T-cells expressing co-inhibitory receptors including NKG2A, CTLA-4, and TIGIT. Of note, the proportion of V γ 9V $\delta 2$ T-cells expressing the inhibitory immune checkpoint receptor PD-1 was not altered. Overall, the increased expression of multiple inhibitory receptors on expanded V γ 9V $\delta 2$ T-cells likely



FIGURE 3

 $V\gamma9V\delta2$ T-cells expanded using TAA- $V\delta2^{hi-lo}$ bsVHH and TAA- $V\delta2^{hi-lo}$ bsVHH-Fc are activated and induce tumor lysis when exposed to TAAexpressing tumor cells. **(A)** Schematic overview of method used for PBMC culture, subsequent assessment of $V\gamma9V\delta2$ T-cell expansion, as well as co-culture of enriched $V\gamma9V\delta2$ T-cells and tumor cells. **(B–E)** $V\gamma9V\delta2$ T-cell CD25 expression **(A**; n=4-12), CD107a expression **(B**; n=4-12), specific tumor cell lysis **(C**; n=4-12) and levels of IL-2, TNF and IFN- γ (pg/ml, **D**; n=4-5) in 24hr co-cultures of SW480 (EGFR⁺), 22Rv1 (PSMA⁺) or MM.1s.CD1d (CD1d⁺) tumor cells with $V\gamma9V\delta2$ T-cells enriched (purity > 60%) from 8 day cultures of healthy donor PBMC in the presence or absence of 1 nM TAA- $V\delta2^{hi-lo}$ bsVHH, 100 nM TAA- $V\delta2^{hi-lo}$ bsVHH-Fc or 10 μ M pamidronate (1:1 E:T ratio). Data generated through flow cytometry **(B–D)** or CBA **(E)**. Individual data-points are indicated using open circles and box and whisker plots indicate the median, 25th to 75th percentiles and minimum to maximum. One-way ANOVA with Tukey's multiple comparisons test was used for statistical analysis; P=< 0.001: ***, P=< 0.001: ***.



FIGURE 4

Bivalent V82^{hi-lo} VHH, TAA-V82^{hi-lo} bsVHH and TAA-V82^{hi-lo} bsVHH-Fc support V γ 9V82 T-cell expansion in cancer patient PBMC and expanded V γ 9V82 T-cells display a memory dominated phenotype. (**A**) Enrichment (left panel) and fold expansion (right panel) of V γ 9V82 T-cells during an 8 day culture of cancer patient PBMC in the presence or absence of 1nM bivalent V82^{hi-lo} VHH, 1 nM TAA-V82^{hi-lo} bsVHH, 100nM TAA-V82^{hi-lo} bsVHH-Fc or 10 μ M pamidronate (n=5-10). (**B**, **C**) Proportion (%) of central memory (CD27⁺ CD45RA⁻), effector memory (CD27⁻ CD45RA⁺), terminally differentiated (CD27⁻ CD45RA⁺) or naïve (CD27⁺ CD45RA⁺) cells within total V γ 9V82 T-cell population before (baseline) and after an 8 day culture in the presence or absence of 1 nM TAA-V82^{hi-lo} bsVHH.Fc or 10 μ M pamidronate (n=5-16) using healthy donor PBMC (**B**) or cancer patient PBMC (**C**). (**D**, **E**) Expression of CD25, HLA-DR, DNAM-1, NKG2D, NKG2A, PD-1, CTLA-4 and TIGIT on V γ 9V82 T-cells expanded for 8 days in the presence or absence of 1 nM TAA-V82^{hi-lo} bsVHH, 100nM TAA-V82^{hi-lo} bsVHH.Fc or 10 μ M pamidronate (data reflect % positive of total V γ 9V82 T-cell fraction; n=5-17) using healthy donor PBMC (**D**) or cancer patient PBMC (**E**). Data generated using flow cytometry. Individual datapoints are indicated using open circles and whisker plots indicate the median, 25th to 75th percentiles and minimum to maximum (**A**, **D**, **E**). One-way ANOVA with Tukey's multiple comparisons test was used for statistical analysis and asterisks are shown compared to IL-2 control; P=< 0.05: *, P=< 0.001: ***, P=< 0.001: ****.

10.3389/fonc.2024.1474007

reflects their increased activation state (47, 48). Indeed, NKG2A was previously reported to identify a population of $V\gamma 9V\delta 2$ T-cells with greater cytotoxic potential (11). In the conditions where cancer patient PBMC were cultured with pamidronate or the $V\delta 2^{hi-lo}$ bsVHH containing constructs, Vγ9Vδ2 T-cells expressed increased levels of the activation markers CD25 and HLA-DR (Figure 4E). At baseline, expression of DNAM-1 and NKG2D was more variable on cancer patient Vy9V82 T-cells and expression did not significantly change upon expansion with the TAA-V82^{hi-lo} bsVHH and bsVHH-Fc molecules. Of interest, while PD-1 was typically expressed by a low proportion of Vγ9Vδ2 T-cells in healthy donor PBMC at baseline, more variable expression was noted on $V\gamma 9V\delta 2$ T-cells in cancer patient PBMC with a further increase in the proportion of PD-1 expressing Vγ9Vδ2 T-cells noted in several patients after expansion with either pamidronate or the Vδ2^{hi-lo} bsVHH constructs. NKG2A expression by Vγ9Vδ2 T-cells was variable at both baseline and upon expansion. CTLA-4 and TIGIT were expressed by an approximately similar proportion of Vγ9Vδ2 T-cells in cancer patient and healthy donor PBMC and were in both cases upregulated upon expansion, though the proportion of Vy9V82 T-cells expressing CTLA-4 appeared to be more consistently increased upon activation and expansion in cancer patient PBMC.

CD1d-V δ 2^{hi-lo} bsVHH construct induces expansion of V γ 9V δ 2 T-cells *in vivo*

While all TAA-V $\delta 2^{hi-lo}$ bsVHHs supported expansion of Vγ9Vδ2 T-cells, results were consistently robust with the CD1d- $V\delta2^{hi\text{-}lo}$ bsVHH and this engager was therefore selected to assess whether $V\gamma 9V\delta 2$ T-cell expansion could also be induced in vivo. As mice lack the phosphoantigen responsive $\gamma\delta$ T-cell population that humans have, an immunodeficient NOG-hIL-15 mouse model was selected as this allowed for inoculation of human PBMC and could simultaneously provide a relatively low level of IL-15 cytokine support. Mice (n=4/group) were i.v. inoculated with human PBMC (n=1 healthy donor with 0.89% Vγ9Vδ2 T-cell of total CD3⁺ T-cells) on day 0 followed by i.p. administration of either PBS or CD1d-V $\delta 2^{hi-lo}$ bsVHH on days 0 and 4 (Figure 5A). On day 8, animals were sacrificed and the proportion of $V\gamma 9V\delta 2$ T-cells was assessed in peripheral blood, spleen, liver and lungs (gating strategy illustrated in Supplementary Figure S7). As shown in Figures 5B, C, the proportion of Vγ9Vδ2 T-cells was significantly higher in mice treated with the CD1d-V82hi-lo bsVHH compared to the PBS control. Administration of the CD1d-V82^{hi-lo} bsVHH resulted in an impressive 11525 (± 7692; median ± IQR, P = 0.004) fold expansion of the V γ 9V δ 2 T-cells in peripheral blood. CD1d-V δ 2^{hi-} $^{\rm lo}$ bsVHH expanded Vy9V 2 T-cells isolated from spleen, liver and lungs all expressed a central or effector memory phenotype, whereas 4.8% (± 1.0; mean ± SEM) of the V γ 9V δ 2 T-cells in peripheral blood retained a naïve phenotype (Figure 5D). In summary, these results indicate that $V\delta 2^{hi-lo}$ bsVHH based engagers can also trigger expansion of Vγ9Vδ2 T-cells in a NOG-hIL-15 mouse in vivo model.

Discussion

Vy9V82 T-cells are a relatively homogeneous population of antitumor immune effector cells capable of inducing cytotoxicity in a wide range of malignancies independently of HLA, which makes them highly promising for cancer immunotherapy (8). Early clinical trials that attempted to exploit the therapeutic potential of V γ 9V δ 2 Tcells were based on activating and expanding Vy9V82 T-cells either ex vivo, followed by ACT, or directly in vivo using N-BP or BrHPP alone or in combination with IL-2. These approaches were shown to be safe and tolerable (49-62), although clinical results lacked consistency (49, 51, 52, 59, 62, 63), which may have been related to the absence of a specific tumor targeting moiety, blunting of $V\gamma 9V\delta 2$ T-cell responsiveness with repeated administration of N-BP/BrHPP and the donor-to-donor variability in the size of the V γ 9V δ 2 T-cell population (49, 51, 54, 56). Recently, strategies that drive tumor selective activation of Vy9V82 T-cells, e.g. using CARs or bispecific antibodies, have emerged as promising novel therapeutic approaches (8). While production of CAR (V γ 9V δ 2) T-cells is considered laborious and costly, and requires preconditioning lymphodepleting chemotherapy regimens before administration to patients, bispecific antibodies could provide a more straightforward off-the-shelf approach. We and others have recently demonstrated the potential of bispecific Vy9V82 T-cell engagers to elicit potent tumor cell lysis (31-37, 64, 65). As these engagers rely on the endogenous V γ 9V δ 2 Tcell population, which, as this study confirms, varies between individuals and with age (9-14), and do not consistently induce Vγ9Vδ2 T-cell expansion, we evaluated whether a bispecific antibody that is bivalent, rather than monovalent, for V δ 2-TCR binding, could combine tumor specific activation with consistent expansion of $V\gamma 9V\delta 2$ T-cells. Our data show that a bivalent VHH that links a high and low affinity Vδ2-TCR specific VHH can support Vγ9Vδ2 Tcell expansion in vitro, ex vivo and in vivo and that this ability is maintained when incorporated in bispecific formats, allowing lysis of TAA expressing tumor cells, also in combination with an Fc domain or anti-albumin binding unit for plasma half-life extension.

From an available panel of V δ 2-TCR specific VHHs, three were selected based on low (VHH-5C7), intermediate (VHH-5D3) and high (VHH-6H4) affinity (41) and linked in different combinations and orientations. Only bivalent VHHs that combined a high and low affinity V δ 2-TCR specific VHH supported consistent and potent V γ 9V δ 2 T-cell enrichment and expansion, particularly with N-terminal positioning of the high affinity V δ 2-VHH and independent of the explored linker lengths. As V γ 9V δ 2 T-cells are potent effector cells, one can envision that the bivalent V δ 2-VHHs containing combinations of high and intermediate affinity V δ 2-VHHs resulted in clustering of activated V γ 9V δ 2 T-cells and limited subsequent expansion through fratricide or activationinduced cell death due to too strong TCR triggering.

When the bivalent $V\delta 2^{hi-lo}$ VHHs were linked to VHHs directed against a variety of TAA specific VHHs (i.e. EGFR, PSMA or CD1d, molecules that can be over-expressed by tumor cells (31, 32, 66)) alone or with an Fc or anti-albumin binding domain for half-life extension, enrichment and expansion were maintained at levels similar to those obtained with the N-BP pamidronate. In contrast to N-BP expanded



(A) Timeline of the *in vivo* study. (B) Enrichment (left panel) and fold expansion (right panel) of human PBMC-derived Vy9Vo2 1-cells *in vivo* 8 days after PBMC inoculation and two doses of i.p. injection with PBS or CD1d-Vδ2^{hi-lo} bsVHH. (C) Enrichment of human PBMC-derived Vy9Vδ2 T-cells in spleen, liver and lungs 8 days after PBMC inoculation and two doses of i.p. injection with PBS or CD1d-Vδ2^{hi-lo} bsVHH. (D) Phenotype of expanded human PBMC-derived Vy9Vδ2 T-cells *in vivo* 8 for spleen, liver and lungs 8 days after PBMC inoculation and two doses of i.p. injection with PBS or CD1d-Vδ2^{hi-lo} bsVHH. (D) Phenotype of expanded human PBMC-derived Vy9Vδ2 T-cells *in vivo* from blood, spleen, liver and lungs 8 days after PBMC inoculation and two doses of i.p. injection with PBS or CD1d-Vδ2^{hi-lo} bsVHH. (D) Phenotype of expanded human PBMC-derived Vy9Vδ2 T-cells *in vivo* from blood, spleen, liver and lungs 8 days after PBMC inoculation and two doses of i.p. injection with PBS or CD1d-Vδ2^{hi-lo} bsVHH. Data generated using flow cytometry. Individual data-points are indicated using open circles and box and whisker plots indicate the median, 25th to 75th percentiles and minimum to maximum. Unpaired *t* tests were used for statistical analysis; P=< 0.01: ***.

Vγ9Vδ2 T-cells, TAA-Vδ2^{hi-lo} VHH and bsVHH-Fc expanded Vy9V82 T-cells had the additional ability to specifically engage and lyse TAA expressing tumor cells. Importantly, the TAA-V $\delta 2^{hi-lo}$ bsVHH and bsVHH-Fc also supported Vy9V82 T-cell expansion using PBMC from patients with gastric cancer, esophageal cancer and melanoma. At baseline, cancer patient Vy9V82 T-cells contained a higher proportion of central memory and a lower proportion of naïve Vy9V82 T-cells compared to healthy individuals, which may reflect a natural interaction of $V\gamma 9V\delta 2$ T-cells with tumor cells and differentiation induction through pAg/BTNs. PD-1 was typically expressed by a low proportion of Vγ9Vδ2 T-cells in healthy individuals but was more variable among cancer patients. Upon expansion by TAA-V $\delta 2^{hi-lo}$ bsVHH and bsVHH-Fc, activation markers CD25 and HLA-DR were more frequently expressed as were inhibitory receptors CTLA-4 and TIGIT. In healthy donor PBMC, this was accompanied by an increase in the proportion of NKG2A expressing Vγ9Vδ2 T-cells, while the fraction of Vγ9Vδ2 Tcells expressing PD-1 did not change. In cancer patient PBMC, expression of PD-1 and NKG2A did not statistically significantly

change on V γ 9V δ 2 T-cells upon expansion. The proportion of V γ 9V δ 2 T-cells expressing activating receptors NKG2D and DNAM-1 remained relatively high, and this in combination with an increase in the expression of activation markers and several inhibitory receptors likely reflects an overall cytotoxic phenotype of the expanded V γ 9V δ 2 T-cells. Indeed, NKG2A⁺ V γ 9V δ 2 T-cells were found to be highly cytolytic (11). In line, we demonstrated that V γ 9V δ 2 T-cells expanded with the TAA-V δ 2^{hi-lo} bsVHH and bsVHH-Fc molecules retained the potential to degranulate, produce proinflammatory cytokines and lyse tumor cells in a TAA specific fashion. Combining immune checkpoint inhibitors with TAA-V δ 2^{hi-lo} bsVHH or bsVHH-Fc molecules could be of potential interest, as this strategy may further enhance the antitumor response of the engaged V γ 9V δ 2 T-cells.

The addition of a half-life extension domain to the TAA-V $\delta 2^{hi-lo}$ bsVHH did typically impact the V $\gamma 9V\delta 2$ T-cell expansion, phenotypic alterations and/or effector functions. This may be due to steric hindrance or an increased intermembrane distance between the V $\gamma 9V\delta 2$ T-cell and TAA⁺ tumor target cell interfering with efficient immune synapse formation (46) as a result of either the Fc

domain or a bound albumin molecule, translating into higher EC_{50s} for degranulation and tumor lysis compared to the (non-Fc, non-anti-albumin VHH bound) TAA-V $\delta 2^{hi-lo}$ bsVHH.

To explore the ability of the bivalent $V\delta 2^{hi-lo}$ VHH based approach to similarly support expansion of V γ 9V δ 2 T-cells *in vivo*, one of the TAA-V $\delta 2^{hi-lo}$ bsVHH was also administered to NOG-hIL-15 mice inoculated with human PBMC. Robust expansion of V γ 9V δ 2 T-cells with a predominant central and effector memory phenotype was observed in peripheral blood as well as in spleen, liver and lungs. Notably, although a small fraction of the TAA-V $\delta 2^{hi-lo}$ bsVHH expanded V γ 9V δ 2 T-cells in peripheral blood retained a naïve phenotype, no naïve V γ 9V δ 2 Tcells were observed in the spleen, liver or lungs likely indicating that only V γ 9V δ 2 T-cells with a central memory and effector memory phenotype infiltrated these organs after expansion.

Overall, our results demonstrate that bispecific V δ 2 T-cell engagers that combine a tumor (TAA) specific VHH and a V δ 2^{hi-lo} bivalent VHH can uniquely trigger both TAA specific lysis and support expansion of V γ 9V δ 2 T-cells. As V γ 9V δ 2 T-cell frequencies are variable and may be impacted by the generally advanced age of patients with cancer and by cancer related therapies (67), the here reported approach may promote the antitumor activity of bispecific V γ 9V δ 2 T-cell engagers by increasing the number of functional effector V γ 9V δ 2 T-cells available for tumor lysis and warrants further exploration in the context of both solid and hematologic malignancies.

Data availability statement

The original contributions presented in the study are included in the article/Supplementary Material, further inquiries can be directed to the corresponding author.

Ethics statement

The studies involving humans were approved by medical ethical committee Amsterdam UMC, location VUmc. The studies were conducted in accordance with the local legislation and institutional requirements. The participants provided their written informed consent to participate in this study. The animal study was approved by Animal Welfare Committee of the Netherlands Cancer Institute. The study was conducted in accordance with the local legislation and institutional requirements.

Author contributions

LK: Writing – original draft, Conceptualization, Data curation, Formal analysis, Investigation, Methodology, Project administration, Resources, Visualization, Writing – review & editing. MJ: Writing – original draft, Conceptualization, Data curation, Formal analysis, Investigation, Methodology, Project administration, Resources, Visualization, Writing – review & editing. MV: Writing – review & editing, Investigation. DH: Writing – review & editing. PY: Writing – review & editing. VI-G: Writing – review & editing, Methodology. PH: Writing – review & editing, Conceptualization, Methodology, Supervision. TG: Writing – review & editing, Conceptualization, Methodology, Resources, Supervision. HV: Writing – review & editing, Conceptualization, Investigation, Methodology, Resources, Supervision, Writing – original draft.

Funding

The author(s) declare financial support was received for the research, authorship, and/or publication of this article. This research was funded by Lava Therapeutics NV, Utrecht, The Netherlands.

Acknowledgments

We thank the MCCA Intervention Unit of the Netherlands Cancer Institute for performing the *in vivo* study.

Conflict of interest

NV. TG and HV own LAVA Therapeutics NV shares. DH, PY, VI-G, PH and HV are/were employed by LAVA Therapeutics NV. TG is scientific advisor to LAVA Therapeutics NV.

The authors declare that this study received funding from Lava Therapeutics NV. LK, MJ and MV were funded by LAVA Therapeutics NV.

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Supplementary material

The Supplementary Material for this article can be found online at: https://www.frontiersin.org/articles/10.3389/fonc.2024.1474007/ full#supplementary-material

10.3389/fonc.2024.1474007

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