Efficient preclinical treatment of cortical T cell acute lymphoblastic leukemia with T lymphocytes secreting anti-CD1a T cell engagers

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ABSTRACT

Background The dismal clinical outcome of relapsed/refractory (R/R) T cell acute lymphoblastic leukemia (T-ALL) highlights the need for innovative targeted therapies. Although chimeric antigen receptor (CAR)-engineered T cells have revolutionized the treatment of B cell malignancies, their clinical implementation in T-ALL is in its infancy. CD1a represents a safe target for cortical T-ALL (coT-ALL) patients, and fratricide-resistant CD1a-directed CAR T cells have been preclinically validated as an immunotherapeutic strategy for R/R coT-ALL. Nonetheless, T-ALL relapses are commonly very aggressive and hyperleukocytic, posing a challenge to recover sufficient non-leukemic effector T cells from leukapheresis in R/R T-ALL patients.

Methods We carried out a comprehensive study using robust in vitro and in vivo assays comparing the efficacy of engineered T cells either expressing a second-generation CD1a-CAR or secreting CD1a x CD3 T cell-engaging Antibodies (CD1a-STAb).

Results We show that CD1a-T cell engagers bind to cell surface expressed CD1a and CD3 and induce specific T cell activation. Recruitment of bystander T cells endows CD1a-STAb with an enhanced in vitro cytotoxicity than CD1a-CAR T cells at lower effector:target ratios. CD1a-STAb T cells are as effective as CD1a-CAR T cells in cutting-edge in vivo T-ALL patient-derived xenograft models.

Conclusions Our data suggest that CD1a-STAb T cells could be an alternative to CD1a-CAR T cells in coT-ALL patients with aggressive and hyperleukocytic relapses with limited numbers of non-leukemic effector T cells.

INTRODUCTION

T cell acute lymphoblastic leukemia (T-ALL) is a hematological malignancy resulting from the transformation and accumulation of T lineage precursor cells.1 T-ALL is phenotypically and genetically very heterogeneous, with frequent genetic mutations in transcription factors and signaling pathways involved in hematopoietic homeostasis and T cell development.2,3 T-ALL accounts for around 10%-15% and 20%-25% of all acute leukemias diagnosed in children and adults, respectively, with a median age at presentation of 9 years.4 Although intensive chemotherapy regimens developed over the last two decades have allowed improved clinical management and survival rates, the 5-year event-free and overall survival rates are still low, especially in adult patients. More importantly, relapse/refractory (R/R) T-ALL remains a challenge with a particularly dismal outcome and lack of approved potentially curative options beyond hematopoietic stem cell transplantation and conventional chemotherapy, thus highlighting the need for novel targeted therapies.5-8

Immunotherapeutic strategies based on the redirection of the immune effector cells to
efficiently recognize and eliminate tumor cells has revolutionized cancer treatment. In recent years, adoptive cell immunotherapies based on T cells bearing chimeric antigen receptors (CAR T) or systemic administration of bispecific T cell-engaging (TCE) antibodies have shown outstanding response rates in B cell malignancies, mainly B-ALL. However, T cell redirecting strategies for T cell malignancies raise additional challenges such as fratricide of effector T cells and potential life-threatening T cell aplasia due to shared antigen expression between effector T cells and T cell blasts, reinforcing the need of both complex genome editing approaches of uncertain safety/efficacy and novel target antigens differentially expressed between normal T cells and T cell blasts. In this sense, we have previously identified CD1a as a surface antigen with barely expression across human cells/tissues but highly and consistently expressed in blasts from patients suffering from cortical T-ALL (coT-ALL), a major subgroup of T-ALL, thus representing a therapeutic target for R/R coT-ALL patients while preventing effector T cell fratricide and T cell aplasia. We consequently generated and characterized CD1a-directed CAR T cells for the treatment of coT-ALL with robust and specific cytotoxicity against CD1a+ T-ALL samples both in vitro and in vivo model.

An emerging strategy which combines advantages of antibody-based and T cell-based therapies, termed Secreting T cell-redirecting Antibodies (STAb)-T immunotherapy, involves the use of engineered T cells secreting small-sized bispecific anti-TAA (tumor-associated antigen) x anti-CD5 antibodies, such as diabodies or tandem scFvs. In contrast to CAR T cell therapies, T cell recruitment is not restricted to engineered T cells when using STAb T strategies. The polyclonal recruitment by secreted TCEs of both engineered and unmodified bystander T cells present at the tumor site might boost the antitumor T cell response. In fact, several groups have shown promising therapeutic effects of STAb T cells in CD19+ B cell malignancies.

Here, we report for the first time the generation of STAb T cells secreting an anti-CD1a x anti-CD5 TCE (CD1a-STAb T cells) and demonstrate their efficacy in several in vitro and in vivo cutting-edge models of coT-ALL. Our results indicate that STAb T therapy controls tumor progression similar to CD1a-CAR T therapy and exhibits slightly higher persistence of T cells in vivo. Our study suggests that CD1a-STAb T cells represent an alternative to CD1a-CAR T cells in coT-ALL, especially in R/R patients with leukapheresis products showing limited numbers of non-tumoral effector T cells.

METHODS

Cell lines and culture conditions

HEK293T (CRL-3216), MOLT4 (CRL-1582, ACC 362), NALM6 (CRL-9273), and K562 (CCL-243) cell lines were purchased either from the American Type Culture Collection (Rockville, Maryland, USA) or the DSMZ (Braunschweig, Germany). Target cells expressing the firefly luciferase gene were either produced in house (NALM6Luc, K562Luc) or a gift from Jan Cools Laboratory (MOLT4Luc). HEK293T cells stably expressing extracellular CD1a (HEK293TCD1a) were generated in house by transduction with pCCL lentiviral vectors encoding CD1a cDNA and cell sorting with anti-CD1a antibodies. HEK293T (WT and CD1a) were cultured in Dulbecco’s modified Eagle’s medium (DMEM) (Lonza, Walkersville, Maryland, USA) supplemented with 2 mM L-glutamine (Life Technologies, Paisley, UK), 10% (vol/vol) heat inactivated fetal bovine serum (FBS) and antibiotics (100 units/mL penicillin, 100 µg/mL streptomycin) (both from Sigma-Aldrich, St. Louis, Missouri, USA), referred to as DMEM complete medium. MOLT4, NALM6, and K562 cells were cultured in RPMI-1640 (Lonza) supplemented with 2 mM L-glutamine, 10% heat-inactivated FBS and antibiotics, referred to as RPMI complete medium (RCM). All the cell lines were grown at 37°C in 5% CO2 and routinely screened for mycoplasma contamination by PCR using the Mycoplasma Gel Detection Kit (Biotools, Madrid, Spain).

Vector construction

The pCDNA3.1-CD1a-scFv expression vector encoding the human kappa (κ) light chain signal peptide L1, followed by the NA1/34.HLK clone-derived CD1a scFv (VH-VL) and a C-terminal polyHis tag was synthesized by GenArt AG (ThermoFisher Scientific, Regensburg, Germany). To generate the CD1a-TCE-encoding lentiviral transfer vector, a synthetic gene encoding the L1-CD1a-scFv flanked by Mlu and Afl was synthesized by GenArt AG and cloned into the vector pCLL-EFIα-LiTE-T2A-EGFP (unpublished), obtaining the plasmid pCCL-EFla-CD1a-TCE-T2A-EGFP. The lentiviral transfer vector pCCL-EFla-CD1a-CAR-T2A-EGFP encoding the CD1a-CAR was previously described. pCCL lentiviral vectors encoding CD1a cDNA were obtained by blunt-XhoI and BamHI subcloning from CD1A_OHu13436C_pcDNA3.1 (+) (GenScript) into blunt-BspEI and BamHI pCCL.

Cell transfection, T cell binding and activation assays

HEK293TWT cells were transfected with the appropriate expression vectors using Lipofectamine 3000 Reagent (ThermoFisher Scientific, Waltham, Massachusetts, USA) according to the manufacturer’s protocol. After 48 hours, transiently transfected HEK293TWT cells were analyzed by flow cytometry and conditioned media were collected and stored at −20°C for western blotting, TCE binding assays and T cell activation studies. For CD1a-TCE binding assay, conditioned media from transiently transfected HEK293TWT cells were incubated with CD1a-negative and CD1a-positive cells and analyzed with an APC-conjugated anti-His mAb (Miltenyi Biotec, Auburn, CA, USA) by flow cytometry. For T cell activation assays, CD1a-negative and CD1a-positive cells were cocultured with freshly isolated T cells at a 1:1 effector:target (E:T) ratio in the presence of conditioned media from transiently transfected HEK293TWT cells. After 24 hours, T cell activation was
analyzed by flow cytometry using PE-conjugated anti-
CD69 mAb.

**Western blotting**

Samples were separated under reducing conditions on
10%–20% Tris-glycine gels (Life Technologies, Carlsbad, California, USA), transferred onto PVDF membranes (Merck Millipore, Tullagreen, Carrigtwohill, Ireland) and probed with 200ng/mL anti-His mAb (Qiagen, Hilden Germany), followed by incubation with 1.6µg/mL horseradish peroxidase (HRP)-conjugated goat anti-
mouse IgG, Fc specific (Sigma-Aldrich). Visualization of
protein bands was performed with Pierce ECL Western Blotting substrate (Rockford, IL, USA) and ChemiDoc MP Imaging System machine (Bio-Rad Laboratories, Hercules, California, USA).

**Lentivirus production and titration**
CD1a, CD1a-CAR- or CD1a-TCE-encoding viral particles pseudotyped with vesicular stomatitis virus G glycoprotein were generated in HEK293T cells by using standard poly-
ethylenimine transfection protocols and concentrated by
ultracentrifugation, as previously described.

Viral titers were consistently in the range of 1×10⁴ transducing units/mL. Functional titers of CD1a-CAR- and CD1a-
TCE-encoding lentiviruses were determined by limiting dilution in HEK293T cells and analyzed using green fluo-
rescent protein (EGFP) expression by flow cytometry.

**T cell transduction and culture conditions**
Peripheral blood mononuclear cells (PBMC) were
isolated from volunteer healthy donors’ peripheral blood (PB) or buffy coats by density-gradient centrifuga-
tion using Lymphoprep (Axis-Shield, Oslo, Norway) or
Ficoll Paque Plus (GE Healthcare, Little Chalfont, UK).
PBMC were plate-coated activated with 1µg/mL anti-CD3
(OKT3) and 1µg/mL anti-CD28 (CD28.2) mAbs (BD
Biosciences) for 2 days and transduced (MOI of 10) with
CD1a-CAR- or CD1a-TCE-encoding lentiviruses in the
presence of 10ng/mL interleukin (IL)-7 and 10ng/mL
IL-15 (Miltenyi Biotec, Bergisch Gladbach, Germany).
As negative controls, non-transduced or GFP-transduced
T cells were used (NT). T cells were expanded in RCM
supplemented with IL-7 and IL-15 (10ng/mL) (Miltenyi
Biotec).

**Cytotoxicity assays**
For cytotoxicity assays, target cells (cell lines and primary
T-ALL blasts, 1x10⁵ cells/well in a 96-well plate for cell
lines and 2x10⁶ for primary blasts) were labeled with
µM cell proliferation dye eFluor 670 (ThermoFisher
Scientific) following manufacturer’s instructions and
cultured with NT, CD1a-CAR, or CD1a-STAB T cells
at different E:T ratios for the indicated time periods.
Effector cell-mediated cytotoxicity was assessed by flow
cytometry analyzing the residual alive, non-apoptotic
(7-aminoactinomycin D; AnnexinV) eFluor 670-positive
target cells in each condition. For primary T-ALL blasts,
absolute counts of target cells were also determined by
using Trucount absolute counting tubes (BD Biosci-
ences).

Additional wells with only target cells were always plated as controls. For bystander cytotoxicity assays, CD1a-CAR or CD1a-STAB T cells were co-
cultured with or without non-transduced activated T cells
(NT) and luciferase-expressing target cells (K562Luc or
MOLT4Luc) at the indicated E:T ratios. As controls, NT
tcells were cultured with target cells. After 48 hours, supernatants were collected and stored at −20°C for cytokine
secretion analysis, and 20µg/mL D-luciferin (Promega)
was added before bioluminescence quantification using a
Victor luminometer (PerkinElmer, Waltham, Massa-
chemus, USA). Percent-specific cytotoxicity was calcu-
lated using the formula: 100 – [(bioluminescence of each
sample×100) / mean bioluminescence of NT-target cells].
Specific lysis was established as 100% of cell viability, and
100% lysis was established by adding 5% Triton X-100
to target cells. For cytotoxic studies using transwell
non-contacting system, 5×10⁴ K562Luc or MOLT4Luc cells
and 1×10⁵ NT cells were plated on bottom wells, and
CD1a-CAR, CD1a-STAB or NT T cells were added at the
indicated ratios into 0.4µm-pore polycarbonate insert
wells (Corning, Kennebunk, Maine, USA). Biolumines-
cence was quantified after 48 hours. For real-time cyto-
toxicity assays, the xCELLigence RTCA DP system (Acea
Biosciences, San Diego, California, USA) was used. At day
0 1x10⁵ HEK293TWT or HEK293TCD1a cells were plated in
an E-Plate 16 (Acea Biosciences) and cultured at 37°C
and 5% CO₂. After 24 hours, NT, CD1a-CAR or CD1a-
STAB T cells were added at different E:T ratios and cell
index values were measured every 15 min for 80 hours
using RTCA Software 2.0 (Acea Biosciences). Specific
lysis was established as 100% of cell viability of target cells,
and 100% lysis was established by adding 10-fold diluted
(in RCM) Cytosilysis Reagent (Acea Biosciences) instead of
effector cells.

**Cytokine secretion analysis**
IFN-γ, TNFα, and IL-2 secretion was analyzed by ELISA
(Diacline, Besancón Cedex, France; BD Biosciences),
following manufacturer’s instructions.

**Flow cytometry**
Antibodies used for flow cytometry analysis are detailed in
online supplemental table S1. Cell surface expression of
CD1a-CAR and cell surface-bound CD1a-TCE were
detected using a biotin-SP goat anti-mouse IgG, F(ab’)₂
(Jackson ImmunoResearch, West Grove, Pennsylvania,
USA) and PE-conjugated streptavidin (ThermoFisher
Scientific). Cell acquisition was performed in a BD FACS-
Canto II flow cytometer using BD FACSDiva software (BD
Biosciences). Analysis was performed using FlowJo V10
software (Tree Star, Ashland, Oregon, USA).

**In vivo T-ALL xenograft models**
Seven to twelve-week-old NOD.Cg-Prkdcscid Il2rgtm1Wjl/SzJ
mice (NSG; The Jackson Laboratory, USA) were bred and
housed under pathogen-free conditions. Mice were


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infused intravenously with 3×10^6 MOLT4^Luc cells and 3 days later received 5×10^6 NT, CD1a-CAR, or CD1a-STAb T cells. Tumor growth was evaluated twice a week by bioluminescence imaging as previously described. Tumor burden and T cell persistence was evaluated by flow cytometry in PB, and bone marrow (BM) after sacrifice at week 3. For T-ALL patient-derived xenograft (PDx) models, seven-to-twelve-week-old NSG mice were sublethally irradiated (2 Gy) and intravenously transplanted with 1×10^6 CD1a^+ T-ALL PDx blasts. Two weeks later, mice were intravenously infused with 3–4×10^6 NT, CAR-CD1a, or STAb-CD1a T cells. Tumor burden and effector T cell persistence was followed up every 2 weeks by bleeding and by BM analysis at different time points, and subsequent flow cytometry analysis. In vivo studies were carried out at the Barcelona Biomedical Research Park (PRBB) in accordance with the guidelines of the Animal Experimentation Ethics Committee. All procedures were performed in compliance with the institutional animal care committee of the PRBB (DAAM9624).

**Statistical analysis**

Results of experiments are expressed as mean or mean±SE of the mean (SEM). Statistical tests indicated in figure legends were performed using Prism V6 (GraphPad Software, La Jolla, California, USA). Significance was considered only when p values were less than 0.05 (*p<0.05; **p<0.01; ***p<0.001, ****p<0.0001).

**RESULTS**

The CD1a-TCE binds to cell surface expressed CD1a and CD3 and induces specific T cell activation

To generate a small-sized Fc-free CD1a-directed TCE, the NA1/34.HLK scFv and the OKT3 scFv were fused in tandem via a G4S peptide linker and cloned under the control of the EF1α promoter in a T2A-based bicistronic lentiviral vector (pCCL-EF1α-CD1a-TCE-T2A-EGFP, figure 1A,B). The vector encoding an anti-CD1a second-generation (4-1BB-based) CAR (pCCL-EF1α-CD1a-CAR-T2A-EGFP, figure 1C,D) has been described previously. The CD1a-TCE was efficiently secreted by transduced HEK293TWT cells with the expected molecular weight of 58 kDa (online supplemental figure S1A). Binding assays using CD1a/CD3 K562 cells, CD1a/CD3 MOLT4 cells, and CD1a/CD3 primary PB lymphocytes (online supplemental figure S2) demonstrated the bispecificity of the secreted CD1a-TCE (online supplemental figure S1B). To study the biological activity of the secreted CD1a-TCE on T cell activation, primary T cells were co-cultured with K562 or MOLT4 cells in the presence of cell-free conditioned medium (CM) derived from non-transfected (NT) or transiently transfected (CD1a-CAR or CD1a-TCE) HEK293TWT cells. High expression of CD69 was detected only when T cells were co-cultured with CD1a-positive cells in the presence of CD1a-TCE CM. T cell activation was not detected when CM from NT or CD1a-CAR-transfected HEK293TWT cells were used (online supplemental figure S1C).

**Generation of human primary CD1a-STAb T cells**

We next transduced primary T cells with CD1a-CAR- or CD1a-TCE-encoding lentiviruses. Transduction efficiencies were determined by flow cytometry after 4–8 days according to the percentage of GFP^CD3^ cells (figure 1E). CD1a-CAR surface expression as well as CD1a-TCE decoration in CD1a-STAb T cells were successfully detected using a polyclonal anti-F(ab')_2 antibody (figure 1F) recognizing the scFv domains of both CD1a-CAR and CD1a-TCE constructs. Transduction efficiencies were 40%–50% and 20%–30% for CD1a-CAR and CD1a-TCE-transduced T cells, respectively. While in CD1a-CAR-transduced T cells the expression of EGFP and F(ab')_2 mainly identifies a single transduced population (figure 1F, upper panel), in CD1a-TCE-transduced T cells several subpopulations are distinguished: transduced/undecorated T cells (GFP^F(ab')_2^-), non-transduced/undecorated T cells (GFP^F(ab')_2^+), and transduced/non-decorated T cells (GFP^F(ab')_2^-) (figure 1F, lower panel). Transduced CD1a-CAR T cells and CD1a-STAb T cells exhibited a similar proportion of CD4^+ and CD8^+ cells (figure 1G). The relative distribution of naïve, central memory, effector memory, and effector T cell subsets was similar in NT, CD1a-CAR^+/- and CD1a-STAb^+/- T cells, with the most prevalent subset being effector memory T cells (figure 1H).

**STAb-CD1a T cells induce a more potent and rapid cytotoxic responses than CAR-CD1a T cells**

To test the ability of CD1a-CAR and CD1a-STAb T cells to kill CD1a^+ T-ALL cells, several cytotoxicity assays were conducted. First, we studied the killing capacity of non-transduced (NT), CD1a-CAR, and CD1a-STAb transduced T cells at different E:T ratios after 24 hour co-culture with CD1a^+ NALM6 or CD1a^+ MOLT4 cells (online supplemental figure S2). CD1a-STAb T cells were able to significantly eliminate CD1a^+ cells even at a 1:16 E:T ratio and induce ~90% cytotoxicity at a 1:1 E:T ratio. In contrast, CD1a-CAR T cells only exhibited significant cytotoxicity at high E:T ratios (figure 1I). Similar results were obtained using primary T-ALL samples in 24 hour assays, where co-culture with CD1a-STAb T cells induced a slight increase in target cell death compared with CD1a-CAR T cells (figure 1J, online supplemental figure S2). In short-time co-culture systems with CD1a^+ or CD1a^+ target cells at a 1:4 E:T ratio, CD1a-STAb T cells killed, in clear contrast to CD1a-CAR T cells, a significant proportion of leukemic cells after 2 hours (35%) and 4 hours (70%) (figure 1K). Next, using an impedance-based real-time cytotoxicity assay, CD1a-STAb T cells mediated a rapid reduction of CD1a^+ target cell viability (online supplemental figure S3A), whereas CD1a-CAR T cells showed a significantly lower cytotoxic effect that required higher E:T ratios (figure 1L). Target cells cultured alone (online supplemental figures S3B–S3D).
Figure 1  continued on next page
supplemental figure S3B) revealed similar viability kinetics to the coculture of NT cells with CD1α+ cells as well as the coculture of NT, CD1α- CAR or CD1α-STAB T cells with CD1α- cells (online supplemental figure S3C).

After 24 hour coculture at a 1:1 E:T ratio, TNFα was significantly increased only in cocultures of CD1α+ target cells with CD1α- STAB T cells. However, IL-2 secretion was significantly higher in cocultures of CD1α+ target cells with CD1α- CAR T cells (online supplemental figure S3C). IFNγ levels were similar in cocultures of CD1α- CAR or CD1α-STAB T cells with MOLT4 and primary T-ALL cells (online supplemental figure S3D), revealing different proinflammatory cytokine profiles depending on whether the CD1α-specific interaction was triggered by CAR or STAB T cells.

**CD1α-STAB T cells eliminate T-ALL cells through the granular exocytosis pathway**

To determine the effector mechanisms involved in CD1α- CAR and CD1α-STAB killing of T-ALL cells, we used the Ca2+-chelating agent EGTA to inhibit granular exocytosis and/or a blocking anti-Fas mAb (figure 1M). In control (untreated) conditions, CD1α-STAB T cells induce higher cytotoxicity than CD1α-CAR T cells on MOLT4 cells at the different time points tested (2, 4, and 24 hours). The cytotoxic action of both CD1α-CAR and CD1α-STAB T cells was completely ablated on EGTA treatment, indicating their dependence on the granular exocytosis pathway to induce target cell death (figure 1N). However, Fas blockage did not diminish lysis levels, showing no influence in the cytotoxic process (figure 1N). These data suggest that CD1α-directed CAR and STAB T cells follow a different kinetic profile but share the same cytolytic effector mechanisms.

**Recruitment of bystander T cells provides STAB-CD1α T cells with greater in vitro tumor cell killing efficiency than CD1α-CAR T cells**

To study the ability of CD1α-STAB T cells to recruit non-engineered bystander T cells, direct contacting co-culture systems were performed (figure 2A). Keeping a constant number of 5×10⁴ CD1α+ MOLT4Luc cells, decreasing numbers of activated effector T cells (AT: NT, CD1α-CAR or CD1α-STAB T cells) were added to the culture, resulting in different AT:Target ratios (from 1:50000 to 2:1). Increasing numbers of NT bystander T cells were added to maintain a constant 2:1 E(AT+bystander):T ratio (figure 2C). The bystander recruitment ability of CD1α-STAB T cells was demonstrated by their enhanced specific cytotoxicity achieved at an E:T ratio as low as 1:50 after 48 hour co-culture with MOLT4Luc cells. In contrast, CD1α-CAR T cell-mediated cytotoxicity against CD1α+ cells only reached that shown by CD1α-STAB T cells at the highest E:T ratio (2:1), with significantly reduced cytotoxicity across lower E:T ratios.

Interestingly, the levels of IFNγ secretion by CD1α-CAR T cells were higher than in CD1α-STAB T cells at the highest E:T condition (figure 2C), but these levels rapidly decreased at lower E:T ratios, indicating that the bystander effect mediated from CD1α-STAB T cells offers equal or superior cytotoxicity capacity than CD1α-CAR T cells without an enhanced cytokine release. No cytotoxic, bystander effect or IFNγ secretion were detected after 48 hours direct co-culture of activated effector cells with CD1α+ (K562Luc) cells (online supplemental figure S3E).

To further demonstrate the bystander effect of CD1α-STAB T cells, similar co-cultures were performed in a non-contacting transwell system (figure 2B). Keeping
Figure 2  STAb-CD1a T cells display enhanced tumor cell killing by recruiting bystander T cells. (A, B) Schematic representation of the direct contact (A) and the non-contacting Transwell (B) co-culture systems used to study the ability of secreted CD1a-STAb to induce bystander T cell cytotoxicity. (C) Decreasing numbers of activated effector T (AT) cells (NT, CD1a-CAR or CD1a-STAb) were co-cultured with $5 \times 10^4$ MOLT4Luc target cells and increasing numbers of NT T cells from the same donor (bystander T cells), resulting in the indicated AT:T ratios but maintaining a constant 2:1 effector (AT+bystander):Target ratio. (D) $5 \times 10^4$ MOLT4Luc cells and $1 \times 10^5$ bystander T cells were plated in the bottom well and decreasing numbers (from $1 \times 10^5$ to $1 \times 10^1$) of activated T (AT) cells (NT, CD1a-CAR or CD1a-STAb) in the upper well. After 48 hours, the percentage of specific cytotoxicity was calculated by adding D-luciferin to detect bioluminescence, and IFNγ secretion was determined by ELISA (C, D). (E) MOLT4 cells were co-cultured with NT, CD1a-CAR or CD1a-STAb T cells at the indicated E:T ratios, and the expression of CD3 and CD1a was analyzed by flow cytometry after 4 and 11 days to assess potential leukemia escape. Data represent mean±SEM of at least three independent experiments by triplicates. Significance was calculated by a two-way ANOVA test corrected with a Tukey's multiple comparisons test (*p<0.05; **p<0.01; ***p<0.001, ****p<0.0001). ANOVA, analysis of variance; E:T, effector:target; NT, non-transduced; CAR, chimeric antigen receptor; STAb, secreting T cell-redirecting antibodies.
Figure 3  CD1a-STAb T cells control the progression of coT-ALL cells in vivo. (A) Experimental design of in vivo cytotoxicity in NSG mice intravenously engrafted with MOLT4Luc cells followed by infusion of NT, CD1a-CAR, or CD1a-STAb T cells. (B, C) Bioluminescence images monitoring disease progression (B) and total RADIANCE quantification at the indicated time points (C). (D) Percentage of MOLT4 cells, identified as HLA-ABC+CD45+CD1a+CD3– by flow cytometry, in peripheral blood (PB) and bone marrow (BM) at sacrifice. (E) Percentage of T cells, identified as HLA-ABC+CD45+CD1a–CD3+ by flow cytometry, in PB, BM and spleen at sacrifice. Plots from (C, D, E) show mean±SEM of at least 5 mice per group. Statistical significance was calculated by an one-way ANOVA test corrected with a Tukey’s multiple comparisons test (*p<0.05; **p<0.01; ***p<0.001, ****p<0.0001). ALL, acute lymphoblastic leukemia; ANOVA, analysis of variance; NT, non-transduced; CAR, chimeric antigen receptor; STAb, secreting T cell-redirecing antibodies.

A constant number of 5×10⁴ MOLT4Luc cells and 1×10⁵ non-engineered bystander T cells both plated in the bottom well, decreasing numbers (from 1×10⁵ to 1 × 10³) of AT (NT, CD1a-CAR or CD1a-STAb T cells) were plated in the insert upper well of the transwell system. After 48 hours, cell killing was only detected when CD1a-STAb T cells were present, indicating that secreted CD1a-TCEs effectively redirected non-engineered bystander T cells toward CD1a⁺ target cells in the bottom wells (figure 2D). IFNγ secretion was also dependent on the presence of CD1a-STAb T cells in the transwell system (figure 2D). In contrast, no cytotoxicity or IFNγ secretion were detected in the presence of CD1a-CAR T cells, or when the target cells were CD1a⁻ (online supplemental figure S3F).

In addition, we studied whether cortical T-ALL cells were able to escape from immune control by co-culturing either NT, CD1a-CAR or CD1a-STAb T cells with MOLT4 cells at low E:T ratios. After 4 days coculture, CD1a-STAb T cells completely eliminated leukemic cells at 2:1, 1:1 and 1:2 E:T ratios; even after 11 days, leukemic cell numbers were restrained below 10% of total cells (figure 2E). In contrast, CD1a-CAR T cells were not able to control MOLT4 growth below the 1:1 E:T ratio. No cell surface CD1a down-modulation was detected when MOLT4 cells were co-cultured with CD1a-CAR or CD1a-STAb T cells (online supplemental figure S3G).

**CD1a-STAb T cells are as effective as CD1a-CAR T cells in short-term in vivo T-ALL models**

The antitumor effect of CD1a-STAb T cells was evaluated in a coT-ALL xenograft model. 3×10⁶ MOLT4Luc cells were intravenously injected in NSG mice, followed by intravenously administration of 5×10⁶ NT, CD1a-CAR, or CD1a-STAb T cells 3 days later (figure 3A). BLI of the mice was done twice per week at the indicated time points to assess leukemia progression (figure 3A). While all NT-treated mice showed unrestricted leukemia development,
Figure 4 CD1a-STAb and CD1a-CAR T cells are effective in eliminating primary coT-ALL in long-term in vivo models. (A, D) Experimental design of in vivo cytotoxicity in NSG mice receiving intravenous coT-ALL patient-derived xenograft (PDX-A in A and PDX-B in D) cells (1×10⁶) followed by NT, CD1a-CAR or CD1a-STAb T cells 2 weeks later (3×10⁶ in A and 4×10⁶ in D). In (A), disease-free mice were rechallenged with further 1×10⁶ PDX-A cells on week 5. (B, C) Percentage of leukemic (B) and T cells (C) in PB and BM at the indicated time points in the PDX-A model. (E, F) Percentage of leukemic (E) and T cells (F) in PB and BM at the indicated time points in the PDX-B model. Numbers of mice with leukemic graft at endpoint, determined as >1% blasts, are indicated. Leukemic blasts were identified by flow cytometry as HLA-ABC⁺CD45⁺CD1a⁺CD3⁻ and CD34⁺ (PDX-A) or CD38⁺ (PDX-B), and T cells as HLA-ABC⁺CD45⁺CD1a⁻CD3⁺. BM plots from B, C) and E, F) show mean±SEM of at least 5 mice per group. ALL, acute lymphoblastic leukemia; BM, bone marrow; PB, peripheral blood; NT, non-transduced; CAR, chimeric antigen receptor; STab, secreting T cell-directing antibodies.

Both the CD1a-CAR- and CD1a-STAb-treated groups were equally able to control disease progression, as evidenced by BLI (figure 3B,C). Even though bioluminescence analysis showed some tumor burden in 4/6 mice in the CD1a-CAR group and only in 1/5 in the CD1a-STAb group, flow cytometry analysis of PB and BM at sacrifice revealed complete control of the disease in both CD1a-CAR-treated and CD1a-STAb-treated mice (figure 3D). Regarding T cell persistence, we observed similar levels across all treatments in all tissues analyzed,
with a non-statistically significant tendency for higher persistence in the CD1a-STAb group (figure 3E).

**CD1a-STAb and CD1a-CAR T cells are effective in eliminating primary T-ALL in long-term in vivo models**

The effectiveness of CD1a-STAb T cells and CD1a-CAR T cells were also compared side-by-side against primary samples in vitro. Two independent CD1a+ coT-ALL PDX models with different aggressiveness were used. We intravenously injected 1×10⁶ leukemic blasts for both models in NSG mice, and 2 weeks later, after confirming leukemic engraftment in the BM, 3×10⁶ (PDX-A) or 4×10⁶ (PDX-B) effect T cells were intravenously injected (figure 4A,D). In the PDX-A model we tested the efficacy of both T cell-redirecting strategies in a setting with relatively low tumor burden, around 2% blasts in the BM at the time of T cells transfer. Both CD1a-CAR and CD1a-STAb T cells were able to ablate leukemic graft in PB and BM in contrast to NT-treated mice, which showed uncontrolled leukemia progression and had to be sacrificed by week 4. To evaluate the effectiveness of the remaining effector cells in the CAR and STAb groups, mice were re-challenged with 1×10⁶ PDX-A cells to simulate a relapse in week 5 and followed up until week 12. In the CD1a-CAR group 2/6 (33%) and 1/6 (17%) mice showed significant leukemic engraftment (>1% blasts) in PB and BM, respectively, whereas in the CD1a-STAb group only 1/5 (20%) mice did (figure 4B). Of note, leukemia relapses in independent mice correlated with decreased numbers of effector T cells in the PB and BM compartments (figure 4C).

In a second PDX model (PDX-B), with higher tumor burden at the time of T cell transfer (engraftment of ~20% in BM), we assessed the effectiveness of CD1a-STAb T cells in a more aggressive, highly active disease setting. Because of its higher aggressiveness (higher leukemic graft at week 0), mice were injected with 4×10⁶ rather than 3×10⁶ effector T cells. Although CD1a-CAR T cells were slightly more effective than CD1a-STAb T cells in achieving minimal residual disease negative in this aggressive PDX-B, both CD1a-CAR- and CD1a-STAb treatments were similarly effective and able to reduce leukemic burden in the BM to ~0.5% by week 4 and further down (<0.02%) by week 8 (figure 4E). Similar T cell persistence levels were observed in CAR- or STAb-treated mice (figure 4F). Taken together, both strategies show robust anti-leukemic activity in long-term cutting-edge in vivo models, and slight differences in the efficacy might be attributed to differential persistence of CD1a-CAR and CD1a-STAb T cells in independent mice.

**DISCUSSION**

The development of safer and efficacious immunotherapies for T-ALL remains challenging because the shared expression of target antigens between CAR T cells and T-ALL blasts leads to either CAR T cell fratricide or immunodeficiency, but also because of potential T-ALL blast contamination during the manufacturing process. We report the first CD1a x CD3 TCE immunotherapy strategy for the treatment of CD1a⁺ coT-ALL. We have engineered T cells to express soluble CD1a x CD3 TCEs which successfully bind to cell surface expressed CD1a and CD3, resulting in the specific activation of the T cells. In contrast to membrane-anchored CD1a-CAR-transduced T cells, flow cytometry analysis several subpopulations in the CD1a-TCE-transduced T cell preparation, transduced and decorated T cells (GFp⁺/F(ab'²)²⁺), non-transduced but decorated T cells (GFp⁺/F(ab')²⁺) and transduced but non-decorated T cells (GFp⁺/F(ab')²⁺), thus confirming the secretion of functional CD1a x CD3 TCE and their ability to decorate surrounding bystander T cells. In vitro short-term and long-term co-culture assays revealed that CD1a-STAb T cells induce a more potent and rapid cytotoxic responses than CD1a-CAR T cells. Mechanistically, the CD1a-specific interaction triggered by either CAR or STAb T cells resulted in different proinflammatory cytokine profile whereas both CD1a-CAR and CD1a-STAb T cells use the granular exocytosis pathway as a common cytolytic effector mechanism. Both contacting and non-contacting co-culture systems confirmed the bystander recruitment ability of CD1a-STAb T cells, a major biological feature providing STAb-CD1a T cells with greater in vivo tumor cell killing efficiency than CD1a-CAR T cells. Interestingly, the bystander effect mediated from CD1a-STAb T cells offers equal or superior cytotoxicity capacity than CD1a-CAR T cells without an enhanced IFNγ release, thus reducing potential cytokine release-associated side effects and offering a safer therapeutic profile than CD1a-CAR T cells. Finally, CD1a-STAb T cells are as effective as CD1a-CAR T cells in cutting-edge in vivo T-ALL cell line and PDX models. Although similar T cell persistence levels were observed in CAR- or STAb-treated mice, leukemia relapses correlated with decreased numbers of effector T cells in the PB and BM. Our data suggest that CD1a-STAb T cells could be an alternative to CD1a-CAR T cells for treating coT-ALL patients.

STAb T cells represent a next-generation T cell-redireacting immunotherapy for B-ALL, and coT-ALL, being easily applicable to other cancers for which a suitable immunotherapy target is available. A major advantage for STAb T cells over CAR T cells lies in the fact that an effective treatment with STAb T cells might require lower T cell doses, which could be of particular relevance when an adequate number of mature effector T cells cannot be engineered due to either the lymphopenic status of many multi-treated patients or manufacturing constrains in patients with aggressive and hyperleukocytic relapses. In this regard, a reduction in the therapeutically effective effector T cell dose to be transferred into the patients may increase the number of patients benefiting from STAb T cell therapy, and significantly reduce the manufacturing costs.

Immune and phenotypic escape mechanisms to anti-CD19 immunotherapies have been experimentally and clinically demonstrated in B-ALL, and commonly lead...
to CD19-resistant leukemias with dismal prognosis. 34-40 Our previous work in B-ALL showed that CD19-STAR T cell therapy could prevent CD19 downregulation and subsequent tumor escape more efficiently and at lower E:T ratios than CD19-CAR T cells.30, 41 In contrast, loss of CD1a expression was not detected on cell surface of target cells co-cultured with either CD1a-CAR or CD1a-STAR T cells. These differences may be attributable to the impact that the density and biology of the targeted antigen plays on T cell activation. 42 In addition, it is worth mentioning that CD1a expression was not detected on cell surface of target cells cultured with CAR or CD19-STAR T cells in vitro models despite the apparently more potent and rapid in vitro cytotoxic responses of CD1a-STAR T cells. In summary, CD1a-STAR T therapy could be an alternative to CD1a-CAR T cells in T-ALL, especially in R/R patients with leukapheresis products showing limited numbers of non-tumoral effector T cells.

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Funding
Research in LA-V laboratory is funded by the Spanish Ministry of Science and Innovation (PID2020-117323RB-I00 and PID2021-121711-T), and the Carlos III Health Institute (DTS20-000089), with European Regional Development Fund (FEDER) cofinancing; the Spanish Association Against Cancer (AEC, PROYB190844/UAL) and the CRIS Cancer Foundation (FCRIS-2018-0042 and FCRIS-2021-0090). Research in PM laboratory is supported by CERCA/Generalitat de Catalunya and Fundació Josep Carreras-Obra Social la Caixa for core support; ‘la Caixa’ Foundation under the agreement LCF/PR/H19/0152160011; the European Research Council grant (ERC-POC-957466); the Spanish Ministry of Science and Innovation (PID2019-108160RB-I00) and the ISCIII-RCORCS within the Next Generation EU program (plan de Recuperación, Transformación y Resiliencia). MLT is supported by Spanish Ministry of Science and Innovation (PID2019-105623R-B00) and the Spanish Association Against Cancer (CICFPI80307R0). PP is supported by Carlos III Health Institute (PG1-01934), with FEDER cofinancing and Fundación Ramón Areces. NT was supported by an FPU PhD fellowship from Spain’s Ministerio de Universidades (FPU19/00039), OH was supported by an industrial PhD fellowship from the Comunidad de Madrid (IND2020/BMD-17668). LA-D was supported by a Rio Hortega fellowship from the Carlos III Health Institute (CM20/00004). VMD is supported by the Torres Quevedo subprogram of the State Research Agency of the Ministry of Science, Innovation and Universities (Ref. PTG2020-011056). DSM is partially supported by a Sara Borrell fellowship from Carlos III Health Institute (CD19/00013).

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