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Ho Ngai; ... et. al

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IL-21 Selectively Protects CD62L⁺ NKT Cells and Enhances Their Effector Functions for Adoptive Immunotherapy

Ho Ngai,^{*,†,1} Gengwen Tian,^{*,†,1} Amy N. Courtney,* Soodeh B. Ravari,* Linjie Guo,* Bin Liu,* Jingling Jin,* Elise T. Shen,* Erica J. Di Pierro,* and Leonid S. Metelitsa^{*,†,‡}

T cells expressing CD19-specific chimeric Ag receptors (CARs) produce high remission rates in B cell lymphoma, but frequent disease recurrence and challenges in generating sufficient numbers of autologous CAR T cells necessitate the development of alternative therapeutic effectors. Va24-invariant NKTs have intrinsic antitumor properties and are not alloreactive, allowing for off-the-shelf use of CAR-NKTs from healthy donors. We recently reported that CD62L⁺ NKTs persist longer and have more potent antilymphoma activity than CD62L⁻ cells. However, the conditions governing preservation of CD62L⁺ cells during NKT cell expansion remain largely unknown. In this study, we demonstrate that IL-21 preserves this crucial central memorylike NKT subset and enhances its antitumor effector functionality. We found that following antigenic stimulation with α -galactosylceramide, CD62L⁺ NKTs both expressed IL-21R and secreted IL-21, each at significantly higher levels than CD62L⁻ cells. Although IL-21 alone failed to expand stimulated NKTs, combined IL-2/IL-21 treatment produced more NKTs and increased the frequency of CD62L⁺ cells versus IL-2 alone. Gene expression analysis comparing CD62L⁺ and CD62L⁻ cells treated with IL-2 alone or IL-2/IL-21 revealed that the latter condition downregulated the proapoptotic protein BIM selectively in CD62L⁺ NKTs, protecting them from activation-induced cell death. Moreover, IL-2/IL-21–expanded NKTs upregulated granzyme B expression and produced more T_H1 cytokines, leading to enhanced in vitro cytotoxicity of nontransduced and anti-CD19-CARtransduced NKTs against CD1d⁺ and CD19⁺ lymphoma cells, respectively. Further, IL-2/IL-21-expanded CAR-NKTs dramatically increased the survival of lymphoma-bearing NSG mice compared with IL-2-expanded CAR-NKTs. These findings have immediate translational implications for the development of NKT cell-based immunotherapies targeting lymphoma and other malignancies. The Journal of Immunology, 2018, 201: 2141-2153.

cells engineered to express chimeric Ag receptors (CARs) that target the CD19 Ag have substantially improved outcomes in patients with B cell malignancies, particularly acute lymphoblastic leukemia (1). However, CAR T cells are less effective against non-Hodgkin lymphoma, with less than half of patients achieving durable complete responses (2). Moreover, T cells obtained from lymphoma patients, particularly children,

ORCID: 0000-0001-7013-3042 (A.N.C.).

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have been shown to possess reduced proliferative potential and consequently produce limited cell numbers when expanded ex vivo (3). Thus, there is an urgent need to develop alternative strategies for CAR-redirected immunotherapy.

Several allogeneic CAR T cell–based therapeutic approaches have been developed to promote antilymphoma activity and minimize the risk of graft-versus-host disease (GVHD). For example, donor-derived, virus-specific T cells engineered to express anti–CD19 CAR (CD19-CAR) undergo expansion early after hematopoietic stem cell transplantation and produce objective responses without causing GVHD (4). However, with the exception of malignancies caused by EBV, virus-specific T cells are not inherently tumor reactive, and their antitumor potential depends entirely on the specificity of the CAR, which can be circumvented by escape variants. Because of these limitations, other lymphocyte subsets, such as NK, $\gamma\delta$ T, and NKT cells, have been tested. These cells have inherent antitumor properties and can be safely infused into immunosuppressed individuals without causing GVHD (5, 6).

Type 1 NKTs are an evolutionarily conserved subset of innate lymphocytes characterized by expression of invariant TCR α -chain V α 24-J α 18 and reactivity to glycolipids presented by the HLA class-I–like molecule CD1d (7). CD1d is widely expressed by both hematopoietic and nonhematopoietic cell types, including monocytes, primitive hematopoietic stem cells, thymocytes, keratinocytes, hepatocytes, and both normal and malignant B cells (8). It is also found on the surface of tumor-associated macrophages (9), which are associated with poor outcome in lymphoma (10). Unlike HLA molecules, CD1d is monomorphic, with the result that NKTs are not alloreactive; indeed, allo-hematopoietic stem cell transplantation mouse model studies suggest that NKTs suppress GVHD and enhance the graft-versus-leukemia effect

^{*}Texas Children's Cancer Center, Department of Pediatrics, Baylor College of Medicine, Houston, TX 77030; [†]Department of Pathology and Immunology, Baylor College of Medicine, Houston, TX 77030; and [‡]Center for Cell and Gene Therapy, Baylor College of Medicine, Houston, TX 77030

¹H.N. and G.T. contributed equally to this work.

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Address correspondence and reprint requests to Dr. Leonid S. Metelitsa, Department of Pediatrics, Baylor College of Medicine, 1102 Bates Avenue, C.1760.06, Houston, TX 77030. E-mail address: Ismeteli@txch.org

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Abbreviations used in this article: AICD, activation-induced cell death; BCM, Baylor College of Medicine; BIM, Bcl-2–like protein 11; CAR, chimeric Ag receptor; CD19-CAR, anti–CD19 CAR; CTV, CellTrace Violet; αGalCer, α-galactosylceramide; GVHD, graft-versus-host disease, NSG, NOD/SCID/IL-2Rγ^{null}.

(11). Importantly, reconstitution of peripheral blood NKTs has been associated with long-term remission in pediatric leukemia patients receiving haploidentical transplants (12). Thus, NKTs possess natural antitumor effector functions and can be generated from healthy donors for off-the-shelf use.

Although NKTs are found at low frequency in human peripheral blood, we have developed methods for clinical-scale CAR-NKT cell production (NCT03294954) (13, 14). Generation of large numbers of NKTs for clinical applications requires robust ex vivo expansion while simultaneously preserving cell functionality and longevity. Repeated stimulation and extended culture of T cell therapeutic products, for example, have been associated with exhaustion, poor in vivo persistence, and limited objective responses in cancer patients (15). Conversely, a higher frequency of central memory phenotype cells in T cell therapy products has been linked with more potent therapeutic activity in preclinical models and clinical trials (15, 16).

Despite significant progress delineating NKT cell development and maintenance in mice (17), the homeostatic requirements for human NKTs remain largely unexplored, particularly in the context of therapeutic applications. Unlike peripheral blood T cells, freshly isolated NKTs do not have a clear hierarchy of naivecentral-effector differentiation and instead display an effector memory-like phenotype (18). However, we recently reported that NKT cell ex vivo expansion in response to stimulation with a specific Ag, *α*-galactosylceramide (*α*GalCer), resulted in accumulation of CD62L⁺ central memory-like cells, which were found to be progressively lost upon subsequent TCR stimulation (14). Importantly, when NKTs were transduced with a CD19-CAR, only the CD62L⁺ subset induced long-term, disease-free survival in lymphoma-bearing NOD/SCID/IL-2R γ^{null} (NSG) mice (14), indicating that maintenance of the central memory-like phenotype is critical for generating effective, long-acting NKT cell-based therapies.

Our initial analysis of immune-related genes differentially expressed in CD62L⁺ versus CD62L⁻ NKT subsets revealed significantly higher levels of *IL-7R* and *IL-21R* mRNA expression in the former (14), suggesting that IL-7 and/or IL-21 may support central memory-like differentiation and superior therapeutic activity of NKTs. IL-7 and IL-21 play important roles at different stages of B, T, and NKT cell development and function (19, 20). Although IL-7R deficiency impacts development of T and NKT cells similarly (21), a recent report showed that IL-21/ IL-21R-induced Stat3 signaling is selectively required for NKT development in humans (22). However, the role of IL-21 in supporting differentiation and maintenance of central memory-like NKTs has not been examined. In this study, we evaluated the effects of IL-21 on the number, phenotype, and functional properties of ex vivo-expanded NKTs and CAR-NKTs. Our results demonstrate that IL-21 selectively protects CD62L⁺ NKTs from activation-induced cell death (AICD) by downregulating expression of proapoptotic protein Bcl-2-like protein 11 (BIM). Moreover, IL-21 supports expansion of highly cytotoxic, T_H1-polarized CAR-NKTs that promote long-term survival of lymphomabearing mice.

Materials and Methods

Cell lines

K562, Daudi, Raji, Ramos, and 293T cells were purchased from the American Type Culture Collection (Manassas, VA). The B-8-2 clone of the K562 cell line was derived previously in our laboratory (14). All lines except 293T were cultured in RPMI 1640 (HyClone, Logan, UT); 293T cells were maintained in IMDM (HyClone). Both types of medium were supplemented with 10% FBS (Life Technologies, Carlsbad, CA) and 2 mM GlutaMAX-I (Life Technologies, Waltham, MA).

NKT cell isolation, transduction, expansion, and sorting

Peripheral blood of healthy donors was purchased from Gulf Coast Regional Blood Center. PBMCs were isolated from buffy coats by density gradient centrifugation. NKTs were purified using anti-iNKT microbeads (Miltenyi Biotec), and the negative PBMC fraction was irradiated (40 Gy) and aliquoted. NKTs were stimulated with an aliquot of autologous PBMCs pulsed with aGalCer (100 ng/ml; Kyowa Hakko Kirin). The culture was supplemented every other day with rIL-2 (200 U/ml; National Cancer Institute, Frederick, MD), rIL-7 (10 ng/ml; PeproTech), and/or rIL-21 (10 ng/ml; PeproTech), where indicated, in complete RPMI (RPMI 1640, 10% heat-inactivated FBS, 2 mM GlutaMAX-1; HyClone). NKTs were expanded for 10 to 12 d and then restimulated with B-8-2 cells (irradiated with 100 Gy). Twenty four-well, nontissue culture plates were coated with retronectin (Takara Bio) overnight. On day 2 after restimulation, the retronectin-coated plates were washed, inoculated with 1 ml of retroviral supernatant containing CD19-CAR, and spun for 60 min at 4600 g. Viral supernatant was then removed, and stimulated NKTs were added to the wells in complete media and 200 U/ml rIL-2, with or without 10 ng/ml rIL-21. Cells were removed from the plate after 48 h, washed, resuspended at 106 cells/ml in complete RPMI with IL-2 or IL-2/IL-21 combination, and plated for continued expansion. NKT number was determined by trypan blue (Life Technologies) counting. When indicated, NKTs were labeled with CD62L-PE mAb (GREG-56; BD Biosciences) and anti-PE microbeads (Miltenyi Biotec) followed by magnetic sorting into CD62L⁺ and CD62L⁻ subsets according to the manufacturer's instructions. The phenotype of the sorted cells was determined by FACS.

Retro- and lentiviral constructs and retrovirus production

The CD19-CAR construct was made as previously described (14, 23). The construct consists of a single-chain variable fragment from the CD19specific Ab FMC-63 that is connected via a short spacer derived from the IgG1 hinge region to the transmembrane domain derived from CD8a, followed by the signaling endodomain of 4-1BB fused with the CD3 ζ signaling chain. Retroviral supernatants were produced by transfecting 293T cells with a combination of CD19-CAR-containing plasmid, RDF plasmid encoding the RD114 envelope protein, and PeqPam3 plasmid encoding the Moloney murine leukemia virus gag-pol fusion, as previously described (14, 23). The lentiviral construct encoding BIM variant 9 (24), envelope plasmid pMD2.G, and packaging plasmid & 8.2 were generous gifts from Dr. K. Scott and Dr. Y.-H. Tsang (Baylor College of Medicine [BCM]), and the control lentiviral plasmid encoding nontargeting short hairpin RNA was obtained from Origene. Lentiviral supernatants were generated from 293T cells transfected with relevant lentiviral construct(s), pMD2.G, and δ 8.2.

Proliferation and apoptosis assays

NKTs were labeled with CellTrace Violet (CTV; Thermo Fisher Scientific, Waltham, MA) and stimulated with α GalCer-pulsed B-8-2 cells. Cell proliferation was examined on day 6 by measuring CTV dilution using flow cytometry. Early and late apoptosis were measured on day 3 post–NKT stimulation by staining for annexin V and 7-AAD (BD Biosciences, Franklin Lakes, NJ), respectively, followed by flow cytometry.

Multiplex cytokine quantification assay

CD19-CAR–NKTs were stimulated for 24 h by Daudi lymphoma cells at a 1:1 ratio. Supernatants were collected and analyzed using the MILLIPLEX MAP Human Cytokine/Chemokine Immunoassay panel (Millipore) for Luminex analysis according to the manufacturer's protocol.

Flow cytometry

NKT cell phenotype was assessed using mAbs for CD3 (UCHT1), V α 24-J α 18 (6B11), CD4 (RPA-T4), granzyme B (GB11), CD62L (DREG-56; BD Biosciences, San Jose, CA), V β 11 (C21; Beckman Coulter, Brea, CA), and IL-21R (17A12; BioLegend, San Diego, CA, and BD Biosciences). CD19-CAR expression by transduced NKTs was detected using antidid mAb (clone 136.20.1) (25), a gift from Dr. B. Jena (MD Anderson Cancer Center, Houston, TX). Intracellular staining was performed using a fixation/permeabilization solution kit (BD Biosciences) with mAbs for Bcl2 (N46-467; BD Biosciences) and BIM (Y36; Abcam, Cambridge, MA) followed by staining with a secondary goat anti-rabbit IgG-AF488 mAb (Abcam). Phosflow staining was performed using Cytofix Buffer (BD Biosciences) and Perm Buffer III (BD Biosciences) with a mAb for Stat3 (pY705; Clone 4; BD Biosciences). Detection of Stat3 phosphorylation was performed after 15 min of treatment with IL-21. Fluorochrome- and isotype-matching Abs suggested by BD Biosciences or R&D Systems were

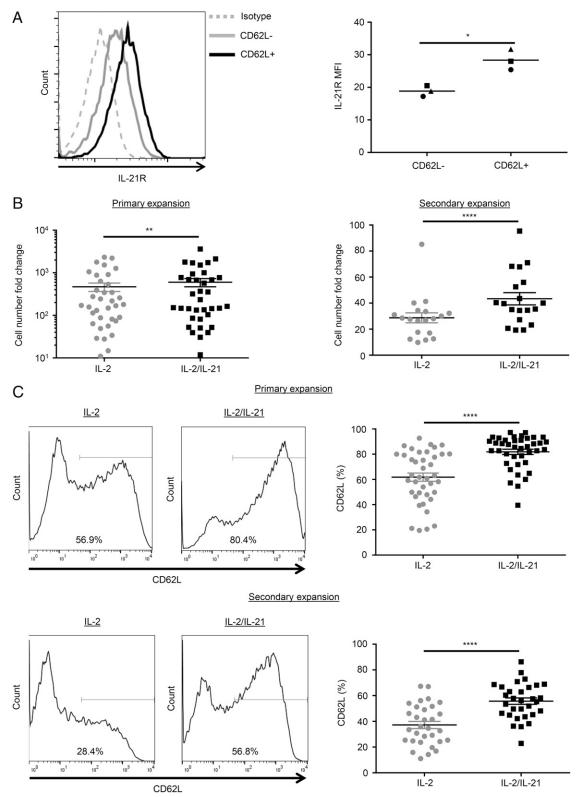


FIGURE 1. Combined treatment with IL-2 and IL-21 produces more NKTs and increases the frequency of CD62L⁺ NKTs compared with IL-2 alone. (**A**) IL-21R expression of ex vivo–expanded primary NKTs was examined using flow cytometry at day 4 after antigenic stimulation with irradiated, α GalCer-pulsed B-8-2 APCs. NKTs were gated into CD62L⁺ and CD62L⁻ populations, and their respective IL-21R expression was measured. A representative histogram from one of three donors (left) and mean of mean fluorescence intensity (MFI) for all donors (n = 3, right) are shown. Each symbol denotes an individual donor. *p < 0.05, paired Student *t* test. (**B**) Following primary or secondary stimulation as described in (A), NKTs were cultured with the indicated cytokines for 12 or 10 d, respectively. NKT cell number was determined using the trypan blue exclusion assay. Cell count fold change compared with day 0 is shown as the mean \pm SEM for all donors following primary expansion (n = 36, left) and secondary expansion (n = 19, right). (**C**) NKTs were expanded as described in (B). CD62L expression of NKTs was examined using flow cytometry on days 12 and 10 after primary and secondary stimulation, respectively. Representative histograms (left) or the mean \pm SEM of percentage of CD62L⁺ NKTs for all donors (right) following primary expansion (n = 40, top panel) and secondary expansion (n = 31, bottom panel) are shown. **p < 0.001, ****p < 0.0001, Mann–Whitney U test.

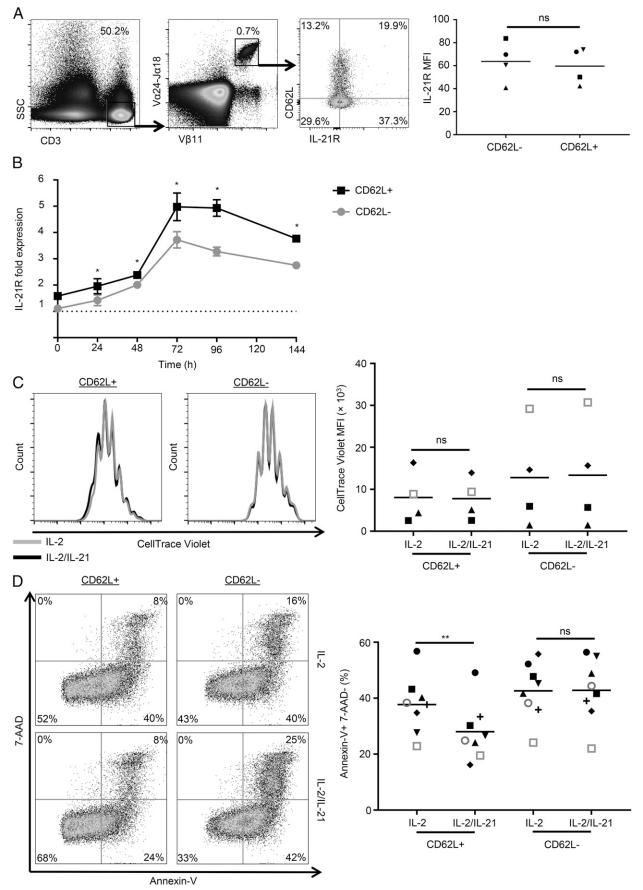


FIGURE 2. $CD62L^+$ NKTs upregulate IL-21R following antigenic stimulation and are selectively protected by IL-21. (**A**) IL-21R expression of $CD62L^+$ and $CD62L^-$ NKTs was analyzed within populations of freshly isolated PBMCs using flow cytometry. 7-AAD⁺ dead cells were excluded, and the remainder were gated for $CD3^+/V\beta11^+/V\alpha24$ -J $\alpha18^+$ NKTs. Representative flow cytometry plots are from one of four healthy *(Figure legend continues)*

used as negative controls. Analysis was performed on an LSR II five-laser flow cytometer (BD Biosciences) using BD FACSDiva software version 6.0 and FlowJo 10.1 (Tree Star, Ashland, OR).

In vitro cytotoxicity assay

Cytotoxicity of parental and CD19-CAR–NKTs against Ramos and Raji cells, respectively, was evaluated using a 4-h luciferase assay as previously described (13).

Gene expression analysis

Total RNA was collected using the Direct-zol RNA MiniPrep Kit (Zymo Research, Irvine, CA). Gene expression analysis was performed using the Immunology Panel version 2 (NanoString, Seattle, WA) with the nCounter Analysis System by the BCM Genomic and RNA Profiling Core. Data were analyzed using nSolver 3.0 software (NanoString). Differences in gene expression levels between CD62L⁺ and CD62L⁻ subsets in the two culture conditions were evaluated using the paired, moderated *t* statistic of the Linear Models for Microarray Data (Limma) analysis package (26).

In vivo experiments

NSG mice were obtained from The Jackson Laboratory and maintained at the BCM animal care facility. Mice were injected i.v. with 2×10^5 luciferase-transduced Daudi lymphoma cells to initiate tumor growth. On day 3, mice were injected i.v. with 4×10^6 to 8×10^6 CD19-CAR–NKTs followed by i.p. injection of IL-2 (1000 U/mouse) only or a combination of IL-2 (1000 U/mouse) and IL-21 (50 ng/mouse) every other day for 2 wk. Tumor growth was assessed once per week by bioluminescent imaging (Small Animal Imaging Facility, Texas Children's Hospital).

Statistics

We used the Shapiro–Wilk test to assess normality of continuous variables. Normality was rejected when the *p* value was <0.05. For nonnormally distributed data, we used the Mann–Whitney *U* test to evaluate differences in continuous variables between two groups. To evaluate differences in continuous variables, we used two-sided, paired Student *t* tests to compare two groups, one-way ANOVA with posttest Bonferroni correction to compare more than two groups, and two-way ANOVA with the Sidak post hoc test to compare in a two-by-two setting. Survival was analyzed using the Kaplan–Meier method with the log-rank (Mantel–Cox) test to compare two groups. Statistics were computed using GraphPad Prism 7 (GraphPad Software, San Diego, CA). Differences were considered significant when the *p* value was <0.05.

Results

Combined IL-2/IL-21 treatment promotes NKT cell expansion and increases CD62L⁺ frequency

We recently demonstrated that ex vivo–expanded CD62L⁺ NKTs express significantly higher levels of *IL-21R* and *IL-7R* mRNA compared with CD62L⁻ NKTs (14). We confirmed elevated cellsurface expression of both IL-7R (14) and IL-21R in CD62L⁺ versus CD62L⁻ NKTs (Fig. 1A), suggesting that IL-7 and/or IL-21 signaling may preferentially support maintenance of the CD62L⁺ NKT subset. To test this hypothesis, we isolated and expanded human peripheral blood NKTs by stimulating them with α GalCer-pulsed CD1d⁺ B-8-2 APCs (14) and supplementing the medium with various combinations of IL-2, IL-21, and IL-7. In contrast to IL-2, IL-7 or IL-21 alone induced little to no NKT cell expansion. However, combination of IL-21 and/or IL-7 with IL-2 significantly increased NKT numbers compared with the IL-2 only condition (Fig. 1B, Supplemental Fig. 1A). Although both IL-2/ IL-7 and IL-2/IL-21 cytokine combinations boosted NKT numbers, only IL-2/IL-21 consistently increased the frequency of CD62L⁺ NKTs compared with IL-2 alone (Fig. 1C, Supplemental Fig. 1B), leading us to shift focus solely to IL-21. We also observed accumulation of CD4⁺ NKTs in cells cultured with IL-2/ IL-21 versus IL-2 alone (Supplemental Fig. 1C), but CD62L expression was elevated in both CD4⁺ and CD4⁻ cells grown with the cytokine combination (Supplemental Fig. 1D). This suggests that the increased frequency of CD62L⁺ cells in the IL-2/IL-21 growth condition occurs independently of the preferential expansion of CD4⁺ NKTs. Overall, these results demonstrate that both IL-2/IL-7 and IL-2/IL-21 combinations increase the number of NKTs generated in response to antigenic stimulation compared with IL-2 alone, but only the latter increases prevalence of the CD62L⁺ subset.

CD62L⁺ NKTs preferentially upregulate IL-21R following antigenic stimulation and are selectively protected from AICD by IL-21

To determine whether the differential IL-21R expression observed in CD62L⁺ versus CD62L⁻ subsets is present in primary, unmanipulated NKTs or induced by antigenic stimulation, we performed a multiparameter flow cytometry analysis on freshly isolated peripheral blood NKTs. We found that IL-21R was expressed at comparable levels in CD62L⁺ and CD62L⁻ subsets of freshly isolated NKTs (p = 0.71) (Fig. 2A) and that NKTs upregulated IL-21R expression within 24 h of α GalCer stimulation, reaching maximum expression by 72 h (Fig. 2B). CD62L⁺ NKTs achieved significantly higher levels of IL-21R expression and sustained peak-level expression longer than CD62L⁻ NKTs (p < 0.05). In line with this, CD62L⁺ NKTs demonstrated higher Stat3 phosphorylation levels than CD62L⁻ NKTs in response to a range of IL-21 concentrations (Supplemental Fig. 1E). These data indicate that antigenic stimulation preferentially sensitizes CD62L⁺ NKTs to IL-21.

To determine how IL-21 increases the frequency of CD62L⁺ NKTs, we sorted NKTs into CD62L⁺ and CD62L⁻ subsets, stimulated them with α GalCer, and measured their proliferation and apoptosis rates after culture with IL-2 or IL-2/IL-21. Cotreatment with IL-2 and IL-21 did not affect NKT cell proliferation in either subset 6 d after stimulation (Fig. 2C). By contrast, in the CD62L⁺ subset, IL-2/IL-21 treatment reduced the frequency of early apoptotic cells (annexin V⁺/7-AAD⁻) at 3 d poststimulation compared with IL-2 treatment alone (Fig. 2D). These findings show that IL-21 preserves CD62L⁺ NKTs by selectively reducing the occurrence of AICD in this subset.

IL-2/IL-21 downregulates the proapoptotic factor BIM in CD62L⁺ NKTs

To understand how IL-21 reduces AICD in $CD62L^+$ NKTs, we evaluated the mRNA expression levels of immune-related genes in stimulated $CD62L^+$ and $CD62L^-$ NKTs grown with IL-2/IL-21 or

donors. Mean of mean fluorescence intensity (MFI) for all donors is shown (n = 4). Each symbol represents an individual donor. Paired Student t test. (**B**) NKTs were stimulated with irradiated, α GalCer-pulsed B-8-2 cells and cultured in medium supplemented with IL-2. IL-21R expression was analyzed in CD62L⁺ and CD62L⁻ NKTs using flow cytometry at the indicated time intervals following stimulation. Fold MFI increase was calculated by dividing the MFI value at each time point by the MFI value of the isotype control. Shown are the mean ± SEM from three donors. *p < 0.05, paired Student t test. (**C**) CTV-labeled NKTs were magnetically sorted into CD62L⁺ and CD62L⁻ subsets and stimulated with α GalCer-pulsed B-8-2 cells cultured in medium with IL-2 or IL-2/IL-21. Cell proliferation was assessed on day 6 after stimulation. Shown are representative results from one of four donors (left) and mean of CTV MFI for all donors (n = 4 donors, right). Each symbol represents an individual donor. Two-way ANOVA with Sidak post hoc test. (**D**) NKTs were sorted and stimulated as described in (C). Annexin V and 7-AAD staining was performed on day 3 after stimulation. Shown are representative results from one of eight donors tested (left) and mean of percentage of annexin V⁺/7-AAD⁻ NKTs for all donors (n = 8 donors, right). Each symbol represents an individual donor. **p < 0.01, two-way ANOVA with Sidak post hoc test. no, not significant.

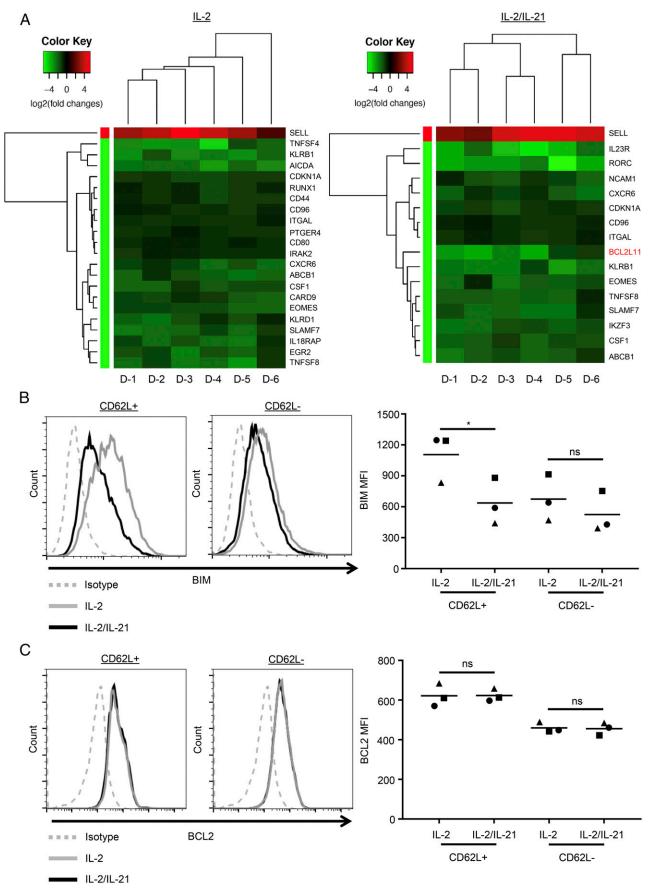


FIGURE 3. Alleviation of AICD by IL-2/IL-21 in $CD62L^+$ NKTs is associated with selective downregulation of BIM. (**A**) NKTs were magnetically sorted into $CD62L^+$ (left) and $CD62L^-$ (right) subsets; stimulated with plate-coated agonistic anti-CD3, anti-CD28, and anti-4-1BB mAbs; and cultured with IL-2 or IL-2/IL-21. mRNA was isolated after 24 h, and gene expression analysis was performed using the Human Immunology Panel version 2 and the nCounter Analysis System (NanoString). Heat maps show log_2 -fold gene expression changes (false discovery rate values (*Figure legend continues*)

IL-2 only for 24 h. We observed that culturing with IL-2/IL-21 versus IL-2 resulted in significant downregulation of multiple genes in both NKT subsets.

Using stringent NanoString gene expression analysis parameters (false discovery rate <0.01), we found that *BCL2L11* was downregulated in the CD62L⁺ versus CD62L⁻ subset, specifically in the IL-2/IL-21 and not the IL-2 only condition (Fig. 3A, Supplemental Fig. 2A). BCL2L11 encodes the proapoptotic protein BIM, a member of the BCL2 family that acts in conjunction with other BCL2 proteins to initiate the intrinsic apoptotic pathway (27). Intracellular flow cytometry confirmed that BIM expression was markedly downregulated in CD62L⁺ but not CD62L⁻ NKTs cultured with IL-2/IL-21 versus those cultured with IL-2 alone (Fig. 3B). Because intrinsic apoptosis initiation depends on the ratio of pro- to antiapoptotic BCL2 proteins (28), we measured expression of BCL2, the primary antiapoptotic regulator, in NKTs under the same experimental conditions as above. Fig. 3C shows that BCL2 expression levels were not affected by IL-21. To determine whether BIM downregulation is required for IL-21-mediated reduction of AICD in the CD62L⁺ subset, we transduced CD62L⁺ NKTs with lentiviral vectors expressing either BIM splice variant 9 (24) or a control construct (Supplemental Fig. 2B). As expected, NKTs transduced with the control construct showed decreased levels of BIM expression in the IL-2/IL-21 condition versus IL-2 alone. Overexpression of BIM variant 9 in IL-2/IL-21-cultured NKTs restored BIM expression to a level similar to that observed in control NKTs stimulated with IL-2 alone. Although control CD62L⁺ NKTs demonstrated significantly lower rates of apoptosis in the IL-2/ IL-21 versus IL-2 condition, overexpression of BIM completely abrogated the protective effect of IL-21 (Supplemental Fig. 2C). In contrast, neither IL-21 treatment nor BIM overexpression affected the high rate of apoptosis observed in CD62L⁻ NKTs (Supplemental Fig. 2D). Thus, IL-21 selectively inhibits the expression of the proapoptotic factor BIM in CD62L⁺ NKTs, without affecting expression of antiapoptotic BCL2, to mediate protection of the CD62L⁺ subset from AICD.

IL-21 enhances TCR- and CAR-mediated NKT cell cytotoxicity by upregulating granzyme B expression

To examine the effect of IL-21 treatment on TCR-dependent NKT cell cytotoxicity, we cocultured NKTs with CD1d⁺ Ramos lymphoma target cells pulsed with aGalCer. Fig. 4A shows that addition of IL-21 to the coculture significantly enhanced the CD1d-restricted cytotoxicity of NKTs against lymphoma cells at all tested E:T ratios. Next, we transduced NKTs with a previously described CD19-specific CAR containing a 4-1BB costimulatory endodomain (14) and evaluated their cytotoxicity against CD19⁺ CD1d⁻ Raji lymphoma cells. CAR-NKTs killed Raji cells more effectively at E:T ratios <10:1 when they were expanded with IL-2/IL-21 versus IL-2 alone (Fig. 4B). As short-term in vitro cytotoxicity of human NKTs is primarily mediated by cytotoxic granule exocytosis (29), we performed intracellular staining for two major components of this pathway: perforin and granzyme B. Perforin was highly expressed in both subsets independent of cotreatment with IL-21 (data not shown). When cultured with IL-2 alone, CD62L⁻ cells expressed a higher level of granzyme B than

CD62L⁺ cells (p < 0.01, Fig. 4C), and IL-2/IL-21 treatment significantly increased granzyme B expression to similarly high levels in both CD62L⁺ and CD62L⁻ NKTs. In line with this, IL-21 preferentially increased the cytotoxicity of CD62L⁺ NKTs, which are typically less cytotoxic than CD62L⁻ cells when cultured with IL-2 alone (Fig. 4D). Therefore, IL-21-mediated upregulation of granzyme B expression in NKTs maximizes their cytotoxic potential.

IL-21 skews NKT cell cytokine secretion toward a T_H 1-like profile

We previously demonstrated that the CD62L⁺ subset is the main source of cytokines produced by NKTs (14). In this study, we evaluated the impact of IL-2/IL-21 on CD19-CAR-NKT cytokine production following coculture with CD19⁺ Daudi cells. Luminex analysis of coculture supernatants collected after 24 h revealed that IL-2/IL-21 treatment increased GM-CSF and IFN-y production in NKTs derived from three different donors (Fig. 5A). Cocultured NKTs from two of three donors showed increased IL-4 production, whereas NKTs from one donor showed decreased IL-4 production. Despite this interindividual variability, IL-2/ IL-21 treatment consistently increased the IFN-y-to-IL-4 ratio, a measure of $T_{\rm H}1$ versus $T_{\rm H}2$ polarization compared with IL-2 alone. Following CAR stimulation, NKTs produced high levels of another $T_H 1$ cytokine, TNF- α , which increased significantly in the presence of IL-21 compared with IL-2 alone in two of three donors (Supplemental Fig. 3). IL-2/IL-21 treatment also enhanced IL-10 production in all three donors, although absolute concentrations of this cytokine were manyfold lower than those of the T_H1 cytokines. IL-17A was minimally detectable (two donors) or undetectable (one donor) regardless of culture with IL-21. Previous work has shown that murine NKTs produce endogenous IL-21 following in vitro stimulation via CD3 and CD28 (30). Consistent with this observation, we found that antigenic stimulation of human NKTs with aGalCer induced a modest level of IL-21 production that peaked within 24 h and became undetectable by 72 h (Fig. 5B). Furthermore, the CD62L⁺ subset was overwhelmingly the primary source of IL-21 production in three donors tested (Fig. 5C). In all, these findings indicate that IL-21 polarizes NKT cytokine production toward a T_H1-like profile and that transient production of endogenous IL-21 by Ag-stimulated NKTs may further support NKT cell functionality through autocrine signaling.

IL-2/IL-21-expanded CAR-NKTs have superior antitumor activity in vivo

Next, we examined the effect of IL-21 on the in vivo therapeutic potential of CAR-redirected NKTs in a murine model of lymphoma. We i.v. injected NSG mice with luciferase-transduced CD19⁺ Daudi lymphoma cells followed 4 d later by adoptive transfer of CD19-CAR–NKTs expanded with IL-2 or IL-2/IL-21. To evaluate the potential benefit of in vivo IL-21 supplementation, mice from the two groups were further subdivided to receive i.p. injections of either IL-2 or IL-2/IL-21. IL-2–expanded CAR-NKTs had marked therapeutic activity and doubled mouse survival compared with the untreated control group; however, there were few long-term survivors with undetectable disease as

<0.01) in CD62L⁺ versus CD62L⁻ subsets. Data were generated from the NKTs of six donors (24 paired samples). (**B** and **C**) NKTs were stimulated with plate-coated agonistic anti-CD3, anti-CD28, and anti-4-1BB mAbs and cultured with IL-2 or IL-2/IL-21 for 24 h. BIM (B) and BCL2 (C) expression in CD62L^{+/-} gated subsets was measured using intracellular staining and flow cytometry. Shown are representative results from one of three donors (left) and mean of mean fluorescence intensity (MFI) for all donors (n = 3 donors, right). Each symbol represents an individual donor. *p < 0.05, two-way ANOVA with Sidak post hoc test. ns, not significant.

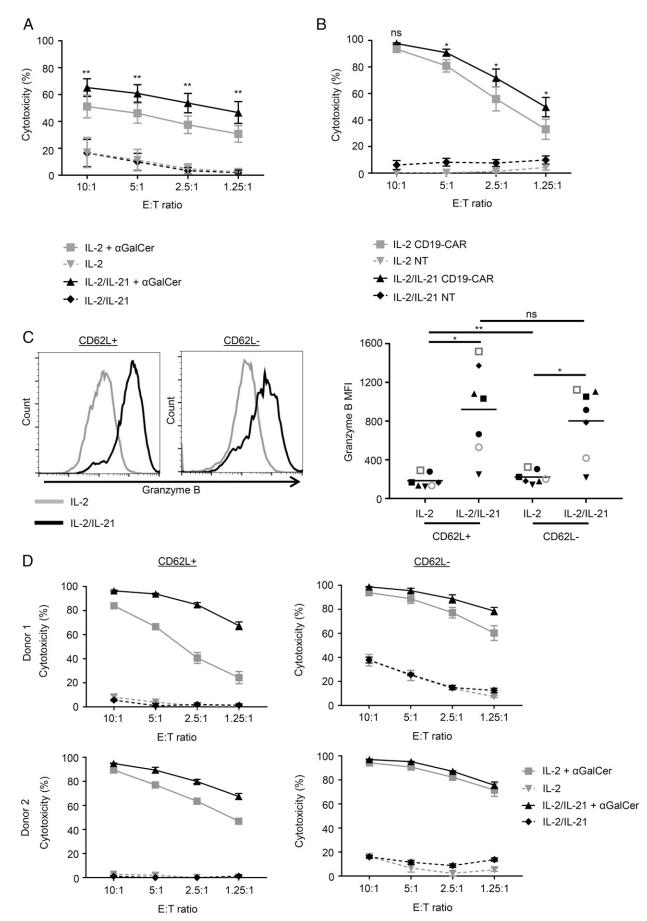


FIGURE 4. IL-21 enhances TCR- and CAR-mediated NKT cell cytotoxicity via upregulation of granzyme B. (A) Luciferase-transduced CD1d⁺ Ramos cells were cocultured with IL-2- or IL-2/IL-21–expanded NKTs. Cytotoxicity was calculated after 4 h by measuring (*Figure legend continues*)

measured at 13 wk after tumor cell injection (p < 0.0001, Fig. 6). IL-2/IL-21-expanded CAR-NKTs instead mediated striking longterm, disease-free survival in treated mice, significantly improving their survival probability compared with those treated with IL-2expanded cells (p < 0.01). In vivo supplementation with IL-2/ IL-21 versus IL-2 did not significantly improve the survival probability of mice receiving either IL-2- or IL-2/IL-21-expanded CAR-NKTs (p = 0.65 and p = 0.32, respectively). Similar results were observed in an independent repeat of this experiment (Supplemental Fig. 4). Notably, CAR-NKT-treated mice displayed no clinical or postmortem pathological signs of xeno-GVHD or other toxic side effects (data not shown). Overall, our results indicate that addition of IL-21 to the growth culture of NKTs strongly enhances their therapeutic potential and should be considered for inclusion in CAR-NKT cell manufacturing processes for lymphoma immunotherapy.

Discussion

αGalCer-specific NKTs are an attractive cellular platform for CARredirected immunotherapy as an alternative or partner approach to polyclonal T cells. We previously demonstrated that human NKTs can be engineered to express tumor-specific CARs and subsequently expanded ex vivo on a clinical scale (13). We also discovered that the CD62L⁺ subset is required for NKT and CAR-NKT cell numeric expansion, in vivo persistence, and therapeutic activity (14). However, the requirements for maintaining, functionally preserving, or further enhancing the therapeutic potential of CD62L⁺ NKTs remain largely unknown, limiting the development of NKT cellbased adoptive immunotherapy. In this study, we demonstrate that following antigenic stimulation, CD62L⁺ NKTs preferentially upregulate IL-21R and are selectively protected from AICD by IL-21 via downregulation of BIM expression. Whereas IL-21 alone was unable to support the proliferation of aGalCer-stimulated NKTs, the combination of IL-2 and IL-21 increased the yield of total expanded NKTs or CD19-CAR-NKTs, with a significant increase in frequency of CD62L⁺ NKTs. Importantly, IL-21 increased granzyme B expression in NKTs, making NKT and CAR-NKT cell preparations significantly more cytotoxic against lymphoma cells in a CD1d- and CD19-dependent manner, respectively. We also observed that IL-21 increased the T_H1-like polarization of the NKT cell cytokine profile secreted in response to specific stimulation, further boosting antitumor effector functionality. Finally, IL-2/ IL-21-expanded CAR-NKTs significantly increased the rate of tumor-free long-term survival of lymphoma-bearing NSG mice.

Although IL-21R expression has been reported previously in both human and murine NKTs (22, 30), in this study we demonstrate for the first time to our knowledge that IL-21R is expressed at a higher level in the CD62L⁺ subset of human ex vivo–expanded NKTs compared with the CD62L⁻ subset. This observation has crucial implications for the development of NKT cell–based cancer immunotherapy because the CD62L⁺ subset promotes in vivo persistence and antitumor activity of adoptively transferred NKTs or CAR-NKTs (14). Interestingly, we found uniform IL-21R expression levels across both CD62L⁺ and CD62L⁻ subsets of freshly isolated peripheral blood NKTs, but CD62L⁺ NKTs preferentially upregulated IL-21R following antigenic stimulation. One possible explanation for this phenomenon is that PLZF, a transcriptional master regulator of NKT cell differentiation (31-33), represses gene expression of both CD62L and IL-21R. TCR stimulation could interfere with the PLZFmediated repression, resulting in a coordinated upregulation of CD62L and IL-21R in NKTs. However, repeated TCR stimulation has been shown to induce terminal differentiation in T and NKT cells (34), and, consistent with this observation, we found that IL-21R and CD62L expression were downregulated in NKTs after repeated aGalCer stimulation. Importantly, addition of IL-21 to the NKT culture medium helped maintain the CD62L⁺ subset, even after repeated stimulation, allowing the expansion of central memory-like cells with high therapeutic potential.

We found that IL-21 supported the accumulation of CD62L⁺ NKTs without affecting their proliferation rate by inhibiting AICD in response to aGalCer/IL-2 stimulation. This mode of action in human NKTs differs from that reported by Coquet et al. (30) in a murine system in which IL-21 promoted proliferation of IL-2-stimulated thymic or aGalCer-stimulated splenic NKTs. These differences could be species-specific and/or the result of different anatomical sites from which NKT cells originate. Consistent with our observations, IL-21 has been found to promote CD62L expression and central memory-like differentiation in both human and mouse naive CD8 T cells and to counteract terminal differentiation/exhaustion of these cells following antigenic stimulation with IL-2 or IL-15 (35-37). Further, priming of naive, tumor-specific T cells with IL-21 has been shown to potently enhance their antitumor activity following adoptive transfer into mice (37), and cells produced using this technique have shown promising results in cancer immunotherapy clinical trials (38, 39). However, IL-21 has also been reported to promote terminal differentiation of B cells and Ag-experienced T cells (40-43). These findings suggest that IL-21 acts on lymphocyte subsets in a context-dependent manner that can be influenced by varying states of cell differentiation or activation, as highlighted in recent reviews (20, 44).

Mechanistically, we present the original finding that IL-21 inhibits AICD selectively in CD62L⁺ NKTs by downregulating proapoptotic BIM expression without affecting antiapoptotic BCL2, ultimately shifting the balance of BCL2 family proteins in favor of cell survival (45, 46). Interestingly, IL-21 can induce BIM-dependent apoptosis in LPS-activated B cells (47) but also promotes survival of vaccinia virus-specific CD8 T cells by upregulating BCL2 and BCL-XL expression (48). These disparate observations reinforce the idea that IL-21 signaling has cell typespecific effects. Although our results indicate that IL-21 maintains CD62L⁺ NKTs by preventing AICD, we cannot exclude the possibility that IL-21 also inhibits NKT terminal differentiation

luminescence intensity with a plate reader. Ramos cells not pulsed with α GalCer were used as a control. Shown are data pooled from two independent experiments with three donors each (mean \pm SEM [n = 6]). **p < 0.01, paired Student t test. (**B**) CD19⁺ Raji cells were cocultured with CD19-CAR–NKTs, and CD19-CAR–dependent cytotoxicity was measured as described in (A). Nontransduced (NT) NKTs were used as a control. Shown are data pooled from two independent experiments with three donors each (mean \pm SEM [n = 6]). *p < 0.05, paired Student t test. (**C**) Granzyme B expression in CD62L⁺ and CD62L⁻ subsets supplemented with IL-2 or IL-2/IL-21 was examined at days 10–12 after primary stimulation. Shown are representative results from one of seven donors (left) and mean of mean fluorescence intensity (MFI) for all donors (n = 7, right). Each symbol represents an individual donor. *p < 0.05, **p < 0.01, one-way ANOVA with Bonferroni correction. (**D**) NKTs were magnetically sorted into CD62L⁺ and CD62L⁻ subsets, stimulated with IL-2 or IL-2/IL-21 – expanded NKTs. Cytotoxicity was calculated after 4 h by measuring luminescence intensity with a plate reader. NT NKTs were used as a control. Shown are the results of the two donors tested. ns, not significant.

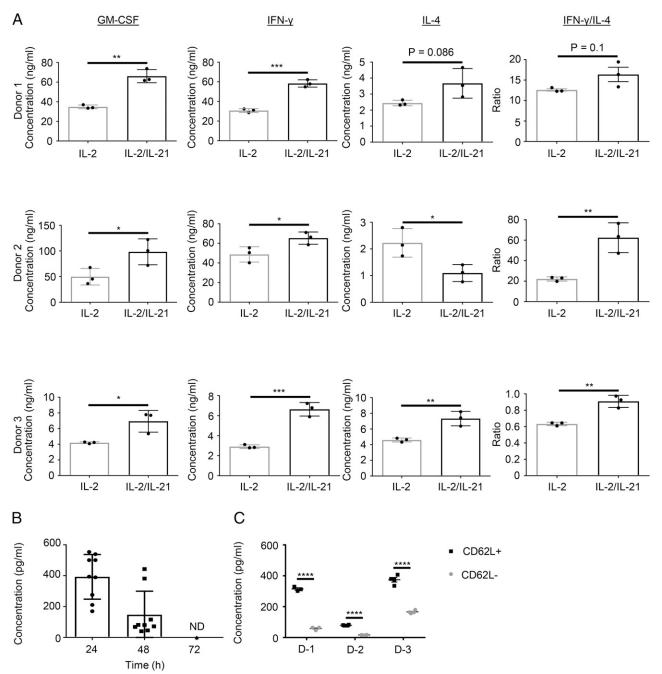


FIGURE 5. IL-21 skews NKT cell cytokine secretion toward a T_{H} 1-like profile. (**A**) IL-2– or IL-2/IL-21–expanded CD19-CAR–NKTs were stimulated with CD19⁺ Daudi lymphoma cells, and supernatants were collected at 24 h. Concentrations of GM-CSF, IFN- γ , and IL-4 in supernatants were measured using a Luminex assay. Data are from two independent experiments with three donors in total, and the IFN- γ -to–IL-4 ratio was calculated. Mean \pm SD of three donors are shown. *p < 0.05, **p < 0.01, ***p < 0.001, unpaired Student *t* test. (**B**) NKTs were stimulated in vitro with irradiated, α GalCer-pulsed B-8-2 cells and IL-2. Culture supernatants were collected 24, 48, and 72 h after stimulation, and IL-21 levels were assessed by ELISA. Shown is the mean IL-21 concentration (picograms per milliliter) \pm SD of three donors. (**C**) NKTs were magnetically sorted into CD62L⁺ and CD62L⁻ subsets and stimulated with α GalCer-pulsed B-8-2 cells. Culture supernatants were collected 24 h after stimulation, and IL-21 levels were assessed by ELISA. Shown are mean IL-21 concentrations \pm SD for each of the three donors. ****p < 0.0001, unpaired Student *t* test.

given that this occurs in Ag-stimulated naive T cells through repression of *Eomes* (37). However, IL-21–mediated suppression of T cell differentiation in that instance was associated with granzyme B downregulation and decreased cytolytic function, whereas we found the opposite effect in both cases for NKTs. This suggests that IL-21 could benefit the antitumor therapeutic potential of NKTs more than T cells.

With respect to cytokine production, IL-21 skewed NKTs toward a T_H 1-like profile, as evidenced by increased production of IFN- γ , GM-CSF, and TNF- α and an increased IFN- γ -to-IL-4 ratio. NKTs are known to produce large amounts of numerous cytokines, which individually may exert opposing effects on tumor immunity. Ultimately, the balance of these cytokines may determine the outcome of NKT cell activation in vivo. In mice, IL-21 has been shown instead to promote a T_H2 -like cytokine secretion profile in NKT thymocytes and to increase IL-4 and IFN- γ production in liver NKTs (30). In a recent phase IIA clinical trial of rIL-21 in melanoma patients, IFN- γ and TNF- α levels produced by peripheral blood NKTs decreased over the course of 5 d, whereas IL-4 levels increased (49). These seemingly contradictory

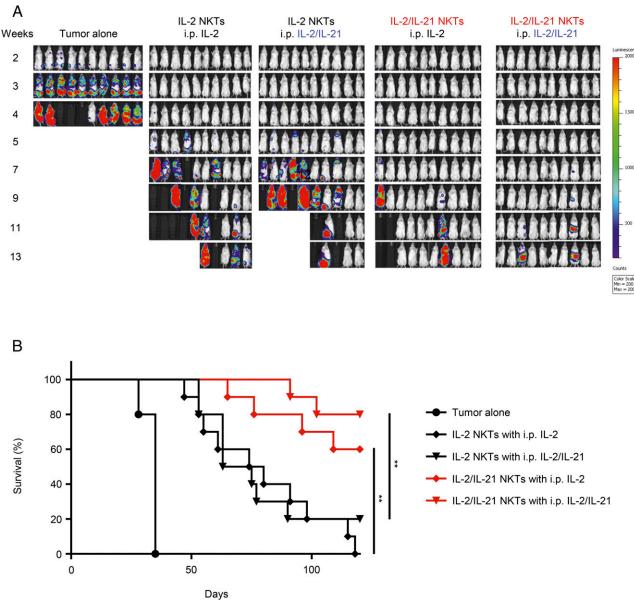


FIGURE 6. IL-2/IL-21–expanded NKTs have superior in vivo antitumor activity. (**A**) NKTs expanded with IL-2 (IL-2 NKTs) or the combination of IL-2/IL-21 (IL-2/IL-21 NKTs) were transduced with a CD19-CAR. Mice were separated into five groups (n = 10 mice per group) before receiving an i.v. injection of 2×10^5 luciferase-transduced Daudi cells on day 0. On day 3, mice received an i.v. injection of the indicated NKT preparations (10^7 cells per mouse) and an i.p. injection of IL-2 (i.p. IL-2) or IL-2/IL-21 (i.p. IL-2/IL-21). Tumor growth was monitored using bioluminescence imaging once per week. (**B**) A survival plot was generated using the Kaplan–Meier method. Shown are data from one of two independent experiments with similar results. Differences in survival probability were compared using the log-rank test. **p < 0.01.

observations may be explained by the fact that NKT IFN- γ production is impaired in cancer patients, particularly those with advanced disease, but can be restored after ex vivo stimulation and expansion (50–52). Therefore, culture conditions play an important role in modulating the therapeutic potential of NKTs, and our results indicate that IL-21–induced T_H1-like cytokine polarization enhances NKT antitumor effector function.

Importantly, we observed a striking enhancement of IL-2/IL-21– expanded CD19-CAR–NKT therapeutic efficacy compared with IL-2–expanded CD19-CAR–NKTs in an aggressive model of B cell lymphoma in NSG mice. Indeed, supplementing the culture with IL-21 promoted the same level of CAR-NKT antilymphoma activity as we previously observed after sorting the cells into a 100% pure CD62L⁺ population (14). By contrast, i.p. administration of IL-21 following CAR-NKT cell transfer had little to no effect on treatment outcome. There are limitations to the use of a xenogenic model system in immune-deficient mice; specifically, we cannot evaluate the impact of CAR-NKT cell and/or IL-21 therapy on the host immune system.

As both NKTs and IL-21 can potently activate innate and adaptive tumor immune responses and have been proven safe in clinical trials (52, 53), there is sound scientific basis for combining NKT cell therapy with IL-21 administration in future clinical studies. Overall, our results support the inclusion of IL-21 in NKT cell ex vivo expansion protocols for lymphoma adoptive immunotherapy.

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Disclosures

G.T., A.N.C., and L.S.M are coinventors on pending patent applications that relate to the use of NKTs in cancer immunotherapy and have been licensed by BCM to Cell Medica, Ltd., for commercial development. Cell Medica, Ltd., provided research support for this project (to L.S.M.) via a sponsored research agreement with BCM. The other authors have no financial conflicts of interest.

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