Melanoma reactive TCR-modified T cells generated without activation retain a less differentiated phenotype and mediate a superior in vivo response

Siao-Yi Wang, Tamson V. Moore, Annika V. Dalheim, Gina M. Scurti & Michael I. Nishimura

Adoptive T cell therapy with T cell receptor (TCR)-modified T cells has shown promise in treating metastatic melanoma and other malignancies. However, studies are needed to improve the efficacy and durability of responses of TCR-modified T cells. Standard protocols for generating TCR-modified T cells involve activating T cells through CD3 stimulation to allow for the efficient transfer of tumor-reactive receptors with viral vectors. T cell activation results in terminal differentiation and shortening of telomeres, which are likely suboptimal for therapy. In these studies, we demonstrate efficient T cell transduction with the melanoma-reactive TIL1383I TCR through culturing with interleukin 7 (IL-7) in the absence of CD3 activation. The TIL1383I TCR-modified T cells generated following IL-7 culture were enriched with naïve (T_N) and memory stem cell populations (T_SCM) while maintaining longer telomere lengths. Furthermore, we demonstrated melanoma-reactivity of TIL1383I TCR-modified cells generated following IL-7 culture using in vitro assays and a superior response in an in vivo melanoma model. These results suggest that utilizing IL-7 to generate TCR-modified T cells in the absence of activation is a feasible strategy to improve adoptive T cell therapies for melanoma and other malignancies.

The introduction of immunotherapies has dramatically improved the treatment of metastatic melanoma over the past two decades. High-dose interleukin-2 (IL-2), immune checkpoint inhibitors, and oncolytic viral therapy have all demonstrated clinical benefit in patients with metastatic melanoma. Despite these successes, a significant number of patients fail to respond and the majority of responding patients relapse. Adoptive T cell therapy (ACT) has become an exciting form of treatment for malignant disease. Tumor-infiltrating lymphocytes (TIL) have been shown to induce significant clinical responses in melanoma patients. However, the process of TIL preparation is laborious, time-consuming, and dependent on the presence of preexisting anti-tumor reactivity. In addition, not every patient has an accessible tumor to develop TIL cultures. Gene-modified T cells were developed to overcome the limitations of TIL. By transducing cells with tumor-reactive receptors, therapeutic potential is effectively transferred to patient peripheral lymphocytes. We previously described clinical and biological responses in a clinical trial of gene-modified T cells expressing a T cell receptor (TCR) reactive to the melanoma antigen, tyrosinase (TIL1383I). While this and other trials using melanoma-reactive TCRs show promise, further studies are needed to improve the efficacy and durability of responses of TCR-modified T cells for melanoma.

Studies that characterized T cell subpopulations suggest that stimulation of naïve cells (T_N) induces differentiation into T memory stem cells (T_SCM), followed by central memory T cells (T_CDM), and subsequently shorter-lived effector memory T cells (T_EM) and effector T cells (T_EF) in a linear manner. Both preclinical and clinical data suggest that less-differentiated populations have superior therapeutic potential. In melanoma and mesothelioma mouse models, T_SCM have been shown to be more effective than other subpopulations in mediating tumor regression. Clinical trials of chimeric antigen receptor (CAR) T cells for neuroblastoma and
B cell malignancies both demonstrate that frequencies of TSCM and TCM correlate with expansion, persistence, and response. Most recently, a study demonstrated that a small stem-like population in TIL product was associated with complete response in patients with metastatic melanoma while terminally differentiated TIL were associated poor persistence. Besides T cell phenotype, proliferation potential and the telomere length of cell products have also been correlated with therapeutic efficacy. A trial utilizing TIL in metastatic melanoma patients demonstrated a correlation between clinical response and longer telomere lengths in infused products. Despite these data, current methods of generating both TCR-modified and CAR T cells involve activation through CD3 stimulation to allow for efficient viral gene transfer. This stimulation results in differentiation, shortening of telomeres, and replicative senescence, which are likely suboptimal for therapy.

Interleukin 7 (IL-7) is a T cell homeostatic regulator that induces G0 to G1 progression without inducing cell division or affecting T cell phenotype. Culturing with IL-7 has also been shown to allow for transduction of T cells with lentiviral vectors in the absence of activation. In these studies, we show that utilizing IL-7 to generate TIL1383I TCR-modified T cells in the absence of activation results in a cellular product with less-differentiated subpopulations, longer telomere lengths, and antigen-reactivity in vitro and in vivo.

Results
Generating TIL1383I TCR-modified T cells in the absence of activation. As T cell activation likely yields a suboptimal therapeutic product, we aimed to generate melanoma-reactive TCR-modified T cells following culture with IL-7 rather than CD3 stimulation. Prior studies have suggested that progression from G0 to G1 phase of the cell cycle correlates with lentiviral transduction susceptibility. We therefore performed cell cycle analyses using acridine orange on T cells from 5 individual donors following no treatment, IL-7 culture, or CD3 activation. A representative cell staining is depicted in Fig. 1a. Analyses show that IL-7 cultured T cells...
had significantly more cells in the G1 phase of the cell cycle compared to untreated cells (means of 21% vs. 2%, p = 0.0079; Fig. 1b). CD3 activated T cells had a mean of 28% in G1, which was comparable to IL-7 cultured cells. In contrast, there was no significant difference between IL-7 cultured and untreated T cells in S phase. CD3 activated T cells had significantly more cells in S phase compared to IL-7 cultured cells (means of 10% vs. 3%, p = 0.0079; Fig. 1b). Altogether, these data suggest that culturing with IL-7 induces G0 to G1 progression, potentially rendering T cells susceptible to lentiviral transduction without commitment to S phase and cell division.

We then used our lentiviral vector containing the melanoma-reactive TIL1383I TCR and a non-signaling truncated CD34 transduction marker to transduce T cells, generating TIL1383I TCR-modified T cells (Fig. 2a). As previously described, CD34 expression is used as a marker for gene transfer and additionally allows for the selection of transduced cells using magnetic-based technologies. We transduced T cells from 5 individual donors using our lentiviral vector either after no treatment, IL-7 culture, or CD3 activation. Flow cytometry analysis demonstrated a low level of CD34 expression in untreated CD3+ cells. In contrast, CD3+ cells demonstrated substantial CD34 expression when transduced after culturing with IL-7 (range of 21.7–55.3%). As expected, activated CD3+ cells demonstrated considerable CD34 expression (range of 27.1–62.3%; Fig. 2b). These findings suggest that culturing with IL-7 allows for efficient gene-transfer of TIL1383I TCR to T cells in the absence of CD3 activation.

Characterizing the phenotypes of TIL1383I TCR-modified T cells generated in the absence of activation. We next evaluated the phenotypes of TIL1383I TCR-modified T cells generated in the absence of activation to assess for potential therapeutic benefits. As studies have demonstrated that CD4+ cells play an important role in the anti-tumor response of gene-modified T cells through both direct cytotoxicity and maintaining persistence, we evaluated the distribution of CD4+ and CD8+ subsets in CD3+ cells transduced either following IL-7 culture or CD3 activation. Although there was heterogeneity between donors, flow cytometry analysis on 5 donors demonstrated that CD3+CD34+ cells transduced after culturing with IL-7 maintained a relatively balanced CD4+/CD8+ ratio (CD4+ cells ranging from 36.3 to 70.3% and CD8+ cells ranging from 30.6 to 48.0%). Interestingly, in the donors we tested, CD3+CD34+ cells transduced following CD3 activation were predominantly CD8+ in our experiments (CD4+ cells ranging from 5.6 to 23.3% and CD8+ cells ranging from 66.4 to 84.2%; Fig. 3a).

In addition, we assessed the subset composition of TIL1383I TCR-modified T cells generated in the absence of activation as data suggest superior efficacy in less-differentiated populations. The differentiation markers, CD45RA, CD62L, CCR7, and CD95 were evaluated in CD3+CD34+ cells transduced either following IL-7 culture or CD3 activation. Even with heterogeneity between donors, CD3+CD34+ cells transduced following IL-7 culture maintained a significantly higher percentage of CD45RA+CD62L− cells (T_N, T_SCM) when compared
to cells transduced following CD3 activation (means of 58.3% vs. 19.7%, \( p = 0.0079 \); Fig. 3b). Furthermore, a significant proportion of the CD45RA+CD62L-CD4+CD34+ cells transduced following CD3 activation was also CCR7+CD95+ (TSCM) (range of 56.9–89.5%). In contrast, CD3+CD34+ cells transduced following CD3 activation demonstrated a significantly higher percentage of CD45RA-CD62L+ (TEM) cells when compared to cells generated following IL-7 culture (means of 40.5% vs. 15.1%, \( p = 0.0317 \); Fig. 3b). These findings were consistent when evaluating either CD8+ or CD4+ subsets individually (Fig. 3c). Notably, there was a lower absolute cell number in the CD4+ subset of CD3+CD34+ cells transduced following CD3 activation. A representation of the phenotypic composition of CD3+CD34+ cells transduced following IL-7 treatment or CD3 activation is depicted in Fig. 3d. Taken together, these results suggest that generating TIL1383I TCR-modified T cells in the absence

Figure 3. Phenotypic characterization of TIL1383I TCR-modified T cells. (a) Flow cytometry analysis depicting CD4+ and CD8+ populations of CD3+CD34+ cells transduced either following IL-7 culture or CD3 activation. (b) Percentage of T(N/TSCM) (CD62L+CD45RA+) populations and the TEM (CD62L-CD45RA-) population in CD3+CD34+ cells transduced either following IL-7 culture or CD3 activation. The graph depicts each data point from 5 donors with lines connecting data points from an individual donor. Statistics were calculated by two-tailed Wilcoxon rank sum test. *\( P = 0.0317 \); **\( P = 0.0079 \). (c) Flow cytometry analysis depicting the T(N) (CD62L+CD45RA+CCR7+CD95-), TSCM (CD62L+CD45RA+CCR7+CD95+), TEM (CD62L-CD45RA-), and T(TEFF) (CD62L+CD45RA+) populations of CD3+CD34+CD8+ and CD3+CD34+CD4+ cells transduced either following IL-7 culture or CD3 activation. (d) Chart depicting the T cell subpopulation composition of CD3+CD34+CD8+ and CD3+CD34+CD4+ cells transduced either following IL-7 culture or CD3 activation. Graphs in (a), (c), and (d) are representative of 5 experiments using cells from 5 different donors.
of activation leads to a product with a balanced CD4+/CD8+ ratio and a less-differentiated phenotype, which may be advantageous for therapy.

Assessing the proliferative potential of TIL1383i TCR-modified T cells generated in the absence of activation. Studies have shown an established correlation between telomere length and the proliferative potential of peripheral blood lymphocytes (PBLs)\(^{20,26}\). We therefore compared the telomere lengths of TIL1383I TCR-modified T cells generated either following IL-7 culture or CD3 activation. We performed fluorescence in situ hybridization (FISH) using a FITC labeled peptide nucleic acid (PNA) probe recognizing the six nucleotide telomere sequence on 5 donors. Although there was variability between donors, we found that CD3+CD34+ cells transduced following IL-7 culture demonstrated higher fluorescence when compared to cells transduced following CD3 activation, suggesting an increased presence of telomeres (Fig. 4a). Using the melanoma cell line, 624 MEL, as a control, we calculated the relative telomere length of the CD3+CD34+ cells transduced following each condition. Analysis of the 5 donors demonstrated that the relative telomere length of CD3+CD34+ cells was

![Figure 4](image_url)

Figure 4. Telomere length analysis of TIL1383I TCR-modified T cells. (a) Flow cytometry analysis of depicting fluorescence of cells transduced either following IL-7 culture or CD3 activation after in situ hybridization with a FITC-labelled PNA telomere probe. (b) Mean relative telomere length calculated using 624 MEL cells as the control population. The graph depicts each data point from 5 experiments using 5 different donors with lines connecting data points from an individual experiment. Statistics were calculated by two-tailed Wilcoxon rank sum test. ** P = 0.0079.
significantly higher when transduced following IL-7 treatment compared to CD3 activation (p = 0.0079; Fig. 4b). These findings suggest that TIL1383I TCR-modified T cells generated following IL-7 culture have a higher proliferative potential than cells generated following CD3 activation, which may lead to a more effective therapy.

Evaluating antigen-specific reactivity of TIL1383I TCR-modified T cells generated in the absence of activation. To assess the tyrosinase-specific reactivity of TIL1383I TCR-modified T cells generated in the absence of activation, we performed intracellular cytokine assays following peptide stimulation on 3 donors. CD3+CD34+ cells transduced following either IL-7 culture or CD3 activation were co-cultured with either unloaded T2 cells, T2 cells loaded with an irrelevant GP100 peptide, or T2 cells loaded with the tyrosinase peptide. Following overnight incubation, intracellular staining of CD3+CD34+ cells transduced following IL-7 treatment demonstrated a substantial percentage of cells over background producing both IFN-α and TNF-β when stimulated with tyrosinase-loaded T2 cells (Fig. 5a). As expected, CD3+CD34+ cells transduced following CD3 activation had a high frequency of cells producing both IFN-α and TNF-β when stimulated with tyrosinase-loaded T2 cells. There was expected background cytokine production by cells incubated with media alone, unloaded T2 cells, and T2 cells loaded with an irrelevant GP100 peptide. We then assessed the melanoma reactivity of TIL1383I TCR-modified T cells generated without activation by quantifying cytokine release following stimulation with a melanoma cell line. CD3+CD34+ cells transduced following either IL-7 culture or CD3 activation from 3 donors were co-cultured with either 624 MEL cells or unloaded T2 cells for 24 h. ELISAs performed on supernatants demonstrated significant presence of IFN-α and TNF-β above background in both cells transduced following IL-7 culture and following CD3 activation (Fig. 5b). These results demonstrate that TIL1383I TCR-modified T cells generated in the absence of activation are reactive against the tyrosinase peptide and a melanoma cell line.
Evaluating the in vivo anti-melanoma activity of TIL1383I TCR-modified T cells generated in the absence of activation.

To assess the in vivo activity of TIL1383I TCR-modified cells, we established tumors through subcutaneous injection of 5 × 10^6 624 MEL into NOD/SCID/γ−/− (NSG) mice (n = 12 per treatment group). Upon development of palpable tumors (Day 7 post injection), 5 × 10^6 CD3+CD34+ cells transduced either following IL-7 culture or CD3 activation were adoptively transferred intravenously into tumor-bearing mice. IL-2 was administered twice a day for 5 days following adoptive transfer to enhance T cell responses as previously described and as done in clinical trials. Tumor-bearing mice receiving no treatment were used as controls. Mice treated with CD3+CD34+ cells transduced following IL-7 culture demonstrated a significant delay in tumor progression when compared to untreated mice (p = 0.0025; Fig. 6). Furthermore, mice treated with CD3+CD34+ cells transduced following IL-7 culture had lower tumor volumes when compared to those treated with activated CD3+CD34+ cells (p = 0.0480; Fig. 6). In contrast, mice treated with activated CD3+CD34+ cells did not have a statistically significant difference when compared to untreated mice. IL-2 treatment alone in mice did not have a significant effect on tumor growth (data not shown). Altogether, these data suggest that TIL1383I TCR-modified T cells generated following IL-7 culture have superior anti-melanoma activity in vivo on a per cell basis compared to cells generated following activation.

Discussion
Adoptive T cell therapy remains a promising form of treatment for metastatic melanoma. As highlighted above, T cell activation used to generate tumor-reactive T cells results in a phenotype that is likely suboptimal for therapy. In addition, TIL, TCR-modified T cells, and CAR T cells all typically undergo in vitro expansion to obtain a sufficient number of cells for clinical response. Continued stimulation during expansion leads to further T cell differentiation, shortening of telomeres, and activation induced cell death (AICD). As cytokines have been shown to render T cells susceptible to lentiviral transduction in the absence of CD3 stimulation, there have been efforts to generate melanoma-reactive T cells in the absence of activation. One group has previously demonstrated successful transfer of a TCR reactive to the melanoma antigen, MART-1, without prior stimulation of CD3. However, TCR-modified or CAR T cells generated in the absence of activation have yet to be evaluated in vivo. This is likely due to the low absolute yield and the inability to select for transduced cells. In these studies, we demonstrate the efficient transfer of TIL1383I TCR and a CD34 marker allowing for magnetic-based sorting.
without CD3 activation. In addition, we evaluate TCR-modified T cells generated in the absence of activation using an in vivo system for the first time.

IL-2, IL-7, and IL-15 have been demonstrated to enhance lentiviral transduction of T cells without CD3 stimulation. However, IL-2 has been known to result in terminal differentiation and enhance AICD while both IL-2 and IL-15 induce T cell proliferation. In contrast, IL-7 is a T cell homeostatic regulator that promotes T cell survival without stimulating cell division. Cell cycle analysis in our studies confirmed that IL-7 culture transitions lymphocytes from G0 to G1 without commitment to S phase and therefore proliferation (Fig. 1). Thus, we utilized IL-7 in our studies to facilitate gene transfer in attempt to preserve maximal immune-competence.

Using our lentiviral construct, we were able to efficiently generate TIL1383I TCR-modified T cells following IL-7 culture (Fig. 2b). In the donors we tested, cells transduced following IL-7 culture maintained a relatively balanced CD4+/CD8+ ratio compared to the low CD4+/CD8+ ratio we observed in activated cells. Prior studies from another group found a similarly high proportion of CD8+ cells when transducing stimulated T cells. Infusion products used in different CD19 CAR T cell trials also frequently skewed toward an either predominantly CD4+ or CD8+ population. Although the causes are not completely clear, differences in transduction and in vitro expansion protocols, including methods of T cell stimulation, cytokines, feeder cells, viral vectors, and length of expansion, all may account for inconsistent CD4+/CD8+ ratios. Patient sex, age, ethnicities, or prior exposures may also play a role. As variability in the infusion product likely accounts for the unpredictable clinical responses to adoptive cell therapy, there have been efforts in generating a more consistent product. Studies have shown that an infusion product formulated with defined CD4+ and CD8+ subsets in a 1:1 ratio enhances in vivo anti-tumor activity. This has led to a CAR T trial in which CD4+ and specific CD8+ subsets were purified and transduced separately before formulated in a 1:1 ratio. Although the trial demonstrated significant clinical responses, the manufacturing process remains laborious. Generating TCR-modified T cells following IL-7 culture appears to be a straightforward strategy to achieve a similar CD4+/CD8+ composition without the potential consequences of T cell activation.

As evidence suggests that less-differentiated T cells have higher therapeutic potential, there are ongoing efforts to preserve less-differentiated phenotypes of TCR-modified or CAR T cells. The group that formulated an infusion product with a defined CD4+/CD8+ ratio used a purified CD8+ TCM starting population for transduction. In addition, the use of IL-7 and IL-15 during in vitro expansion has been used to maintain less-differentiated T cell populations. However, both of these strategies continue to rely on CD3 stimulation for efficient gene transfer. Even when cell products were generated from a less-differentiated CD8+ TCM population, the patients received a product that consisted of CD8+ cells that were mostly TCM, which is consistent with our studies that demonstrate significant T cell differentiation following activation (Fig. 3b–d). In addition, a recent study evaluating TCR-modified T cells specific to the cancer-testis antigen, NY-ESO-1, following expansion with IL-7 and IL-15 demonstrated a limited enrichment of less-differentiated cells and no absolute benefit. By transducing T cells in the complete absence of activation, we obtain TIL1383I TCR-modified T cells with preserved TN and TCM populations and retained telomere lengths.

We confirmed the melanoma-specific reactivity of TIL1383I TCR-modified T cells generated following IL-7 culture using both in vitro and in vivo systems. Intracellular cytokine assays demonstrated cytokine production when TIL1383I TCR-modified cells were incubated with tyrosinase-loaded cells for infusion, but this comes at the expense of telomere shortening and reduced proliferative potential. In vitro expansion commonly used to increase the percentage of transduced cells and obtain the required number of cells for infusion, and this comes at the expense of telomere shortening and reduced proliferative potential. It would be difficult to obtain comparable absolute numbers of gene-modified cells typical for infusion without in vitro expansion. However, TIL1383I TCR-modified T cells generated in the absence of activation have desirable qualities for therapy, so the number of cells required to see a clinical response may not be as high. Our in vivo experiments used a number of cells that was 50% of what we previously used when evaluating TCR-modified T cells in vivo to assess for tumor reactivity at a lower than standard dose. Indeed, our in vivo experiments demonstrate significant melanoma reactivity at a lower than standard cell number, justifying a lower starting dose for future clinical trials. Furthermore, generating a potent therapy without in vitro expansion would shorten production time and permit more timely treatment.

In conclusion, we demonstrate efficient generation of melanoma-reactive TIL1383I TCR-modified T cells in the absence of activation by utilizing IL-7 treatment and a lentiviral vector. TIL1383I TCR-modified T cells generated following IL-7 culture have a balanced CD4+/CD8+ ratio, less-differentiated phenotype, and increased proliferative potential. These TIL1383I TCR-modified T cells generated in the absence of activation have anti-melanoma reactivity both in vitro and in vivo. Studies investigating the in vivo persistence of TIL1383I TCR-modified T cells generated without activation and the phenotype of the persisting cells are currently underway. A growing body of data suggests that less-differentiated phenotypes and longer telomeres correlate with clinical response. Thus, generating TIL1383I TCR-modified T cells in the complete absence of activation is a feasible approach.
strategy to improve therapeutic potential. These findings may be applied to other tumor reactive TCRs and CARs and improve adoptive T cell therapy for melanoma and other malignancies.

Materials and methods

Cells lines. T2, a human TAP-deficient HLA-A2⁺ cell line, and HEK 293 T were obtained from ATCC (Rockford, MD). 624 MEL, a melanoma cell line was previously described (Bethesda, MD)40. GPRTG, a 293 T-based packaging cell line for production of lentiviral vector particles, was obtained from the National Gene Vector Biorepository (Indiana University, Indianapolis, IN)42. T2 cells were maintained in RPMI (Thermo Fisher, Walther, MA) with 10% heat-inactivated fetal bovine serum (HI-FBS; Tissue Culture Biologics, Long Beach, CA). All other cell lines were maintained in DMEM (Thermo Fisher) with 10% HI-FBS. 1 ng/mL doxycycline (Clontech Laboratories, Palo Alto, CA) was added to DMEM for GPRTG cells. Cells were maintained at 37 °C in a humidified 5% CO₂ incubator.

T cells. Peripheral blood lymphocytes (PBLs) from independent normal human donors were isolated from purchased apheresis products (Key Biologics, Memphis, TN) using Ficoll-Hypaque (Sigma-Aldrich, St. Louis, MO) density gradients40,41. T cells were activated by stimulating PBLs from an individual donor with 50 ng/mL anti-CD3 mAb (Miltenyi Biotec, Bergisch Gladbach, Germany), 300 IU/mL recombinant human IL-2 (rhIL-2; Novartis Pharmaceuticals, East Hanover, NJ), and 100 ng/mL recombinant human IL-15 (rhIL-15; Biological Resources Branch, National Cancer Institute, Bethesda, MD) for three days prior to transduction43. Alternatively, PBLs from the same donor were cultured with 20 ng/mL recombinant human IL-7 (rhIL-7; Biological Resources Branch, National Cancer Institute) for seven days prior to transduction.

Cell cycle analysis. Cell cycle analysis was performed using acridine orange as previously described44. 10⁶ cells were resuspended in 0.1 ml of PBS. 0.5 ml of Buffer I (20 mM Citrate–Phosphate, pH 3.0, 0.1 mM EDTA, 0.2 M Sucrose, 0.1% Triton X-100) was added to the suspension and gently agitated. 2 mg/ml acridine orange (Sigma-Aldrich) was diluted 1:100 in Buffer II (10 mM Citrate–Phosphate, pH 3.8, 0.1 M NaCl). 0.5 ml of Buffer II containing acridine orange was subsequently added to the suspension and gently agitated. Cell luminescence was measured using the LSR Canto flow cytometer (BD Biosciences, San Jose, CA) and data was analyzed with FlowJo software V10.1 (BD Biosciences, https://www.flowjo.com). The fraction of cells in G0, G1, S, and G2/M phases of the cell cycle was determined by DNA and RNA contents.

Lentivirus production. A lentiviral vector was constructed to include the melanoma antigen (tyrosinase)-reactive TIL1383I TCR, and a truncated CD34 molecule (CD34t) as a transgene expression marker as previously described40. The pLVX-E1a-N1 (Clontech, Mountain View, CA) vector was modified to include the TCR alpha chain, P2A self-scleaving linker, TCR beta chain, T2A linker, and CD34t. This modified vector was used to generate a stable GPRTG lentiviral producer cell line as follows. Viral supernatants were produced by transfecting 293 T cells with the pLVX-TIL1383I TCR plasmid using Lenti-X single shots following the manufacturer's directions (Takara, Mountain View CA). GPRTG cells were spinoculated (1000 X G for 2 h at 32 °C) with viral supernatants containing 8 µg/ml of polybrene (Sigma-Aldrich), cultured for an additional 72 h, and resuspended in DMEM with doxycycline. TIL1383I TCR lentiviral supernatants for T cell transductions were produced as follows. Transduced GPRTG cells were seeded in doxycycline-free DMEM in 10 cm tissue culture dishes. DMEM was replaced with Freestyle 293 expression media (Gibco, Gaithersburg, MD) 24 h later. TIL1383I TCR lentiviral supernatant was then collected following a 48 h incubation.

Lentiviral transduction of T cells. PBLs from an individual donor that were either cultured with IL-7 or activated with anti-CD3, IL-2, and IL-15 were resuspended in TIL1383I TCR lentiviral supernatant containing 8 µg/ml of polybrene and plated in 24 well plates. Plates were spinoculated at 2000 X G at 32 °C for 2 h. TIL1383I TCR-transduced cells were harvested and resuspended in RPMI supplemented with either IL-7 or IL-2 and IL-15.

T cell phenotypic analysis. Evaluation of T cell surface markers was performed via immunofluorescence staining and analyzed via flow cytometry. Monoclonal antibodies (mAbs) used to characterize transduced T cells include: anti-CD3-APC Cy7, anti-CD4-APC, anti-CD8-PerCP/Cy5.5, anti-CD34-PE, anti-CCR7-BV785, anti-CD45RA-PE Cy7 (Biolegend, San Diego, CA), and anti-CD62L-BV650, anti-CD95-FITC (BD Biosciences). Flow cytometry was performed using the LSR Fortessa flow cytometer (BD Biosciences) and data was analyzed with FlowJo software V10.1.

Telomere analysis. The flow-FISH method using the telomere PNA Kit/FITC for Flow Cytometry Kit (Dako, Glostrup, Denmark) was used to analyze telomere length per manufacturer's protocol. 624 MEL cells were used as the control. Fluorescence was measured using an LSR Canto flow cytometer (BD Biosciences) and data was analyzed with FlowJo software V10.1.

Cytokine production and release assays. Antigen reactivity of transduced cells was measured through cytokine production and cytokine release as previously described40. Tyrosinase-specific reactivity was assessed through an intracellular cytokine assay as follows. Gp100:209–217 (gp100) and tyrosinase:368–376 (tyrosinase) peptides were obtained from Synthetic Biomolecules (San Diego, CA). T2 cells were pulsed with 10 µg/ml of peptide for 2 h prior to co-culture. 10⁵ TIL1383I TCR-modified cells were co-cultured with the targets at a 1:1
ratio in a 96 well U-bottom tissue culture plate in 200 µl media. Following overnight incubation, Brefeldin-A and Monensin (Biolegend) were added to the culture. After five hours, cells were stained for surface markers for 20 min at room temperature. Cells were fixed, permeabilized, and stained for intracellular IFN-α and TNF-β.

Data was collected using an LSR Fortessa flow cytometer and analyzed with FlowJo software V10.1. Alternatively, melanoma reactivity was assessed through a cytokine release assay as follows. 105 TIL1383I TCR-modified cells were co-cultured in triplicate with 624 MEL cells at a 1:1 ratio in a 96 well U-bottom tissue culture plate in 200 µl media. Supernatants were harvested following incubation for 24 h. Cytokine release was quantified by sandwich ELISA using monoclonal antibodies to IFN-α or TNF-β (Biolegend).

Melanoma xenograft model. 8–12-week old NOD/SCID/γ−/− (NSG) male and female mice (Jackson Laboratory, Bar Harbor, ME) were treated under an approved IACUC protocol and complied with ARRIVE guidelines. Mice were injected subcutaneously with 5 × 106 624 MEL cells into flanks, forming palpable tumors by day 7. At 7 days post-challenge, 5 × 106 TIL1383I TCR-modified cells were injected intravenously in 100 µl of PBS. Intraperitoneal injections of IL-2 (60,000 IU) were administered twice a day for 5 days as previously described27. Tumor bearing mice either treated with IL-2 alone or untreated were used as controls. Tumor size in 3 perpendicular dimensions was measured twice a week using calipers. Two-tailed Wilcoxon rank sum test was used to determine significant difference between treatment groups.

Ethics approval. All applicable international, national, and/or institutional guidelines for the care and use of animals were followed. All animal studies were conducted under approved Loyola University Institutional Animal Care and Use Committee (IACUC) protocols and complied with ARRIVE guidelines. All recombinant DNA and lentiviral transductions work was done under approved Loyola Institutional Biosafety Committee (IBC) protocols. This article does not contain any studies with human participants performed by any of the authors. All human materials used were from de-identified apheresis products purchased from commercial sources.

Data availability
The data that support the findings of this study and materials are available from the corresponding author, SW, upon reasonable request.

Received: 12 February 2021; Accepted: 16 June 2021
Published online: 25 June 2021

References
1. Atkins, M. B. et al. High-dose recombinant interleukin 2 therapy for patients with metastatic melanoma: analysis of 270 patients treated between 1985 and 1993. J. Clin. Oncol. 17, 2105–2116 (1999).
2. Schadenhof, D. et al. Pooled analysis of long-term survival data from phase ii and phase iii trials of ipilimumab in unresectable or metastatic melanoma. J. Clin. Oncol. 33, 1889–1894 (2015).
3. Larkin, J. et al. Combined nivolumab and ipilimumab or monotherapy in untreated melanoma. N. Engl. J. Med. 373, 23–34 (2015).
4. Ribas, A. et al. Pembrolizumab versus investigator-choice chemotherapy for ipilimumab-refractory melanoma (KEYNOTE-001): a randomised, controlled, phase 2 trial. Lancet Oncol. 16, 908–918 (2015).
5. Andtbacka, R. H. I. et al. Final analyses of OPTiM: a randomized phase III trial of talimogene laherparepvec versus granulocyte-macrophage colony-stimulating factor in unresectable stage III-IV melanoma. J. Immunother. Cancer 7, 145 (2019).
6. Rosenberg, S. A., Packard, B. S., Aebersold, P. M. et al. Use of tumor-infiltrating lymphocytes and interleukin-2 in the immunotherapy of patients with metastatic melanoma. A preliminary report. N. Engl. J. Med. 319, 1676–1680 (1988).
7. Dudley, M. E. et al. Adoptive cell therapy for patients with metastatic melanoma: evaluation of intensive myeloablative chemoradiotherapy preparative regimens. J. Clin. Oncol. 26, 5233–5239 (2008).
8. Rosenberg, S. A. et al. Durable complete responses in heavily pretreated patients with metastatic melanoma using T-cell transfer immunotherapy. Clin. Cancer Res. 17, 4350–4357 (2011).
9. Duong, C. P. et al. Cancer immunotherapy utilizing gene-modified T cells: from the bench to the clinic. Mol. Immunol. 67, 46–57 (2015).
10. Moore, T. et al. Clinical and immunologic evaluation of three metastatic melanoma patients treated with autologous melanoma-reactive TCR-transduced T cells. Cancer Immunol. Immunother. 67, 311–325 (2018).
11. Kaech, S. M. & Cui, W. Transcriptional control of effector and memory CD8+ T cell differentiation. Nat. Rev. Immunol. 12, 749–761 (2012).
12. Gattinoni, L., Klebanoff, C. A. & Restifo, N. P. Paths to stemness: building the ultimate antitumour T cell. Nat. Rev. Cancer 12, 671–684 (2012).
13. Gattinoni, L. et al. A human memory T cell subset with stem cell-like properties. Nat. Med. 17, 1290–1297 (2011).
14. Klebanoff, C. A. et al. Determinants of successful CD8+ T-cell adoptive immunotherapy for large established tumors in mice. Clin. Cancer Res. 17, 5343–5352 (2011).
15. Louis, C. U. et al. Antitumor activity and long-term fate of chimeric antigen receptor-positive T cells in patients with neuroblastoma. Blood 118, 6050–6056 (2011).
16. Xu, Y., Zhang, M., Ramos, C. A. et al. Closely related T-memory stem cells correlate with in vivo expansion of CAR.CD19-T cells and are preserved by IL-7 and IL-15. Blood 123, 3750–3759 (2014).
17. Krishna, S. et al. Stem-like CD8 T cells mediate response of adoptive cell immunotherapy against human cancer. Science 370, 1328–1334 (2020).
18. Zhou, J. et al. Telomere length of transferred lymphocytes correlates with in vivo persistence and tumor regression in melanoma patients receiving cell transfer therapy. J. Immunol. 175, 7046–7052 (2005).
19. Feinak, A. D., June, C. H. & Levine, B. L. Engineered T cells: the promise and challenges of cancer immunotherapy. Nat. Rev. Cancer 16, 566–581 (2016).
20. Hodes, R. J., Hathcock, K. S. & Weng, N. P. Telomerases in T and B cells. Nat. Rev. Immunol. 2, 699–706 (2002).
21. Boise, L. H. et al. Growth factors can enhance lymphocyte survival without committing the cell to undergo cell division. Proc. Natl. Acad. Sci. USA 92, 5491–5495 (1995).
22. Cavaliere, S. et al. Human T lymphocytes transduced by lentiviral vectors in the absence of TCR activation maintain an intact immune competence. Blood 102, 497–503 (2003).

23. Circosta, P. et al. T cell receptor (TCR) gene transfer with lentiviral vectors allows efficient redirection of tumor specificity in naive and memory T cells without prior stimulation of endogenous TCR. Hum. Gene. Ther. 20, 1576–1588 (2009).

24. Norell, H. et al. CD34-based enrichment of genetically engineered human T cells for clinical use results in dramatically enhanced tumor targeting. Cancer Immunol. Immunother. 59, 851–862 (2010).

25. Adusumilli, P. S., Cherkassky, L., Villena-Vargas, J. et al. Regional delivery of mesothelin-targeted CAR T cell therapy generates potent and long-lasting CD4-dependent tumor immunity. Sci. Transl. Med. 6, 261 (2014).

26. Weng, N. P. et al. Tales of tails: regulation of telomere length and telomerase activity during lymphocyte development, differentiation, activation, and aging. Immunol. Rev. 160, 43–54 (1997).

27. Shilyansky, J. et al. Identification of a T-cell receptor from a therapeutic murine T-cell clone. J. Immunother. 20, 247–255 (1997).

28. Grabstein, K. H. et al. Cloning of a T cell growth factor that interacts with the beta chain of the interleukin-2 receptor. Science 264, 965–968 (1994).

29. Waldmann, T. A. The biology of interleukin-2 and interleukin-15: implications for cancer therapy and vaccine design. Nat. Rev. Immuno. 6, 595–601 (2006).

30. Moeller, M. et al. Adoptive transfer of gene-engineered CD4+ helper T cells induces potent primary and secondary tumor rejection. Blood 106, 2995–3003 (2005).

31. Brentjens, R. I. et al. Safety and persistence of adoptively transferred autologous CD19-targeted T cells in patients with relapsed or chemotherapy refractory B-cell leukemias. Blood 118, 4817–4828 (2011).

32. Kochenderfer, J. N. et al. B-cell depletion and remissions of malignancy along with cytokine-associated toxicity in a clinical trial of anti-CD19 chimeric-antigen-receptor-transduced T cells. Blood 119, 2709–2720 (2012).

33. Porter, D. L., Hwang, W. T., Frey, N. V. et al. Chimeric antigen receptor T cells persist and induce sustained remissions in relapsed refractory chronic lymphocytic leukemia. Sci. Transl. Med. 7, 303 (2015).

34. Sommermeyer, D. et al. Chimeric antigen receptor-modified T cells derived from defined CD8+ and CD4+ subsets confer superior antitumor reactivity in vivo. Leukemia 30, 492–500 (2016).

35. Turtle, C. J. et al. CD19 CAR-T cells of defined CD4+CD8+ composition in adult B cell ALL patients. J. Clin. Invest. 126, 2123–2138 (2016).

36. Turtle, C. J. et al. Durable molecular remissions in chronic lymphocytic leukemia treated with CD19-specific chimeric antigen receptor-modified T cells after failure of ibrutinib. J. Clin. Oncol. 35, 3010–3020 (2017).

37. Turtle, C. J., Hamaï, L. A., Berger, C. et al. Immunotherapy of non-Hodgkin’s lymphoma with a defined ratio of CD8+ and CD4+ CD19-specific chimeric antigen receptor-modified T cells. Sci. Transl. Med. 8, 355 (2016).

38. Curti, N. et al. IL-7 and IL-15 instruct the generation of human memory stem T cells from naive precursors. Blood 121, 573–584 (2013).

39. Gong, W. et al. Comparison of IL-2 vs IL-7/IL-15 for the generation of NY-ESO-1-specific T cells. Cancer Immunol. Immunother. 68, 1195–1209 (2019).

40. Spear, T. T. et al. TCR gene-modified T cells efficiently treat established hepatitis C-associated hepatocellular carcinoma tumors. Cancer Immunol. Immunother. 65, 293–304 (2016).

41. Rivoltini, L. et al. Quantitative correlation between HLA class I allele expression and recognition of melanoma cells by antigen-specific cytotoxic T lymphocytes. Cancer Res. 55, 3149–3157 (1995).

42. Throm, R. E. et al. Efficient construction of producer cell lines for a SIN lentiviral vector for SCID-X1 gene therapy by concatemeric array transfection. Blood 113, 5104–5110 (2009).

43. Spear, T. T. et al. Altered peptide ligands impact the diversity of polyfunctional phenotypes in T cell receptor gene-modified T cells. Mol. Ther. 26, 996–1007 (2018).

44. Darzynkiewicz, Z. Differential staining of DNA and RNA in intact cells and isolated cell nuclei with acridine orange. Methods Cell Biol. 33, 285–298 (1990).

Acknowledgements
This research was funded by the National Institutes of Health R21 CA153789 and R01 AI129543.

Author contributions
S.W. and M.N. designed the study; S.W., T.M., A.D., and G.S. performed experiments; S.W., T.M., and M.N. analyzed the data; S.W., T.M. and M.N. wrote the manuscript; all authors reviewed the manuscript.

Competing interests
The authors declare no competing interests.

Additional information
Correspondence and requests for materials should be addressed to S.-Y.W.

Reprints and permissions information
is available at www.nature.com/reprints.

Publisher’s note
Springer Nature remains neutral with regard to jurisdictional claims in published maps and institutional affiliations.

Open Access
This article is licensed under a Creative Commons Attribution 4.0 International License, which permits use, sharing, adaptation, distribution and reproduction in any medium or format, as long as you give appropriate credit to the original author(s) and the source, provide a link to the Creative Commons licence, and indicate if changes were made. The images or other third party material in this article are included in the article's Creative Commons licence, unless indicated otherwise in a credit line to the material. If material is not included in the article's Creative Commons licence and your intended use is not permitted by statutory regulation or exceeds the permitted use, you will need to obtain permission directly from the copyright holder. To view a copy of this licence, visit http://creativecommons.org/licenses/by/4.0/.

© The Author(s) 2021