Poster #457



Engineered tumor-infiltrating lymphocytes (TIL) with membrane-bound IL15 (mbIL15) and LIGHT (TNFSF14) generate significant antitumor efficacy in fibrotic tumor models Benjamin Primack*, Balazs Koscso*, Violet Young, Meghan Langley, Theresa Ross, Ngoc Ly, Carmela Passaro, Zheng Ao, Sean Smith, Nirzari Shah, Dexue Sun, Dan Jun Li, Dhruv Sethi, Jan ter Meulen, Michelle Ols, Jeremy Tchaicha

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Introduction

- Non-engineered TIL cell therapy, requiring concomitant administration of high-dose IL2, has demonstrated efficacy in melanoma, but with substantial toxicity^{1,2}
- To develop IL2-independent TIL to treat immunologically hot tumors, cytoTIL15[™] cells are engineered to express regulatable membrane-bound interleukin 15 (mblL15) using an FDA-approved small-molecule drug, acetazolamide (ACZ), which binds to the drugresponsive-domain (DRD) carbonic anhydrase 2 (CA2; Figure 1), driving pharmacologically regulatable mbIL15-mediated expansion and persistence without the need for exogenous IL2 administration (OBX-115: NCT05470283, NCT06060613)
- To address immunologically cold tumors, another proinflammatory cytokine, LIGHT, was added to the cytoTIL15 backbone. LIGHT, a tumor necrosis factor family member (TNFSF14), interacts with lymphotoxin beta receptor (LT β R) and herpes virus entry mediator (HVEM) found on various tumor microenvironment (TME) cell types, such as cancer-associated fibroblasts (CAF; **Figure 2**)
- Preclinically, intratumoral LIGHT expression via viral or cell therapy leads to tertiary lymphoid structure (TLS) formation³ and vascular normalization,⁴ both associated with better outcomes⁵
- Previously, we showed co-regulated expression of mbIL15 and LIGHT increased tumor cell killing by autologous TIL and efficacy in the B16-F10 syngeneic model⁶
- Herein, efficacy of mbIL15- and LIGHT-engineered TIL was tested in fibroblast-rich patient-derived human xenograft (PDX) and syngeneic tumor models



Figure 1. The cytoDRiVE platform uses a drug-responsive domain (DRD) to enable an FDA-approved small-molecule ligand to regulate expression of a protein fused to the DRD. In the absence of the cognate DRD ligand (drug OFF-state), the fusion protein is degraded by the proteasome. In the presence of the ligand (drug ON-state), the DRD is stabilized and the protein is expressed. In the case of cvtoTIL15 and cvtoTIL15-LIGHT cells, a DRD derived from carbonic anhydrase 2 (CA2) is used to regulate both mbIL15 and LIGHT via delivery of acetazolamide (ACZ).



48-hour co-culture (B). Statistical tests: one-way ANOVA, cytoTIL15+ACZ v cytoTIL15-LIGHT+ACZ,



Figure 4. A CRC lung metastasis was excised from a patient and ~100 mg tumor fragment was implanted subcutaneously within 24 hours of surgery (Passage 0 [P0]). Two NSG female mice were implanted with similarly sized fragments that were allowed to grow for ~150 days. Serial passaging was performed until there was enough viable tumor tissue available to support a large-scale implantation for an efficacy study. Animals (n=8 per treatment group) were randomized on Day 32 following the large-scale PDX tumor implant, and TIL (cytoTIL15 or cytoTIL15-LIGHT cells) were infused IV the following day. Both treatment groups were dosed ACZ daily via oral gavage at 200 mg/kg from day of infusion (Day 0) until termination of the study.



Figure 5. Expression levels of immune cell (CD45+), tumor (EpCAM+), and CAF (aSMA+) surface markers on P0 tumor tissue after growing in an NSG mouse for 150 days. PDX tumor was excised, homogenized, and analyzed by flow cytometry with anti-human CD45, anti-human EpCAM, and antihuman αSMA (A). CRC PDX-bearing NSG mice were adoptively transferred with 25×10^6 engineered autologous TIL exactly 33 days post-tumor implant (**B–C**). Tumor measurements were collected twice a week via digital calipers (B) starting on the day before ACT and continuing for the duration of the study. Whole blood samples were collected weekly (n=4 per treatment group) via submandibular bleeding to quantitate TIL (C). Statistical tests: Mann-Whitney, cytoTIL15-LIGHT v cytoTIL15, *p value<0.05; last two timepoints, cytoTIL15-LIGHT (n=8) v cytoTIL15 (n=7).



Figure 6. Splenocytes from transgenic mice expressing the gp100-specific pmel TCR were retrovirally transduced to express constitutive mbIL15 and LIGHT, or mbIL15 alone. After transduction, these CD8+ splenocytes were then expanded in the presence of murine IL2 for 6 days. The mouse colorectal cancer cell line, MC38, was engineered to express the gp100 melanoma antigen to serve as a tumor-specific target for these adoptively transferred, engineered CD8+ pmel T cells (A). Foldincrease in CD8+ pmel cells was monitored throughout the expansion; transduction efficiency and mblL15 expression levels were determined by flow cytometry on day of ACT (B). Cytotoxicity quantification of CRC cytoTIL15 cells versus cytoTIL15-LIGHT cells against autologous tumor/CAF or tumor-only spheroids over a 24-hour co-culture (C).

***p value<0.005. Previously shown at AACR 2024.6





day before ACT and continuing for the duration of the study (C), and overall survival was tracked based on tumor volume threshold (D). Statistical tests: Mann-Whitney, mblL15-LIGHT v mblL15, *p value<0.05, **p value<0.01; mblL15 v Vector control, [#]p value<0.05, ^{##}p value<0.01, ^{###}p value<0.005 **(C)**; log-rank (Mantel-Cox), mblL15-LIGHT v mblL15, *p value<0.05 (D).

Figure 8. mblL15- and LIGHT-expressing pmel T cells create a pro-inflammatory TME in MC38-gp100 tumors



Figure 8. Fibrotic MC38-gp100 tumor-bearing mice were adoptively transferred with 5×10^6 engineered CD8+ pmel cells 26 days post-tumor implant. Tumors were collected on Day 7 post-ACT and homogenized using GentleMACS with Mouse Tumor Dissociation Kit (Miltenvi Biotec) and TME phenotype was characterized using flow cytometry. Transduced pmel cells were identified as single live CD45+ CD3+ CD8+ Thy1.1+ Thy1.2+ cells; M1 phenotype of total TAM population were identified as single live CD45+ CD3- CD11b+ F4/80+ iNOS+ cells (A). Tumor IFNy and MCP-1 levels were determined using MSD (Mesocale Discovery) on supernatants collected during tumor homogenization (B). Each bar represents the mean ± SEM from 5 individual tumors. Statistical tests: ordinary one-way ANOVA, all comparisons, *p value<0.05, **p value<0.01, ***p value<0.005.

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Results

Day 7 in Tumor Supernatant





Figure 9. The mouse CRC cell line MC38 engineered to express gp100 was implanted subcutaneously in C57BL/6J mice. After 26 days, implants developed into large (~300 mm³ average) fibrotic tumors and mouse CD8+ pmel cells engineered with either mblL15 alone or mblL15-LIGHT were adoptively transferred into the tumor-bearing mice. Tumors were harvested at Day 20 (A) and Day 26 (B) post-ACT. After overnight incubation in 10% formalin, tumors were bathed in 30% sucrose solution for 3–5 days before being frozen in OCT cups. Fluorescent staining was performed on 10–12µm thick cryosections using anti-B220 (B cells) and anti-CD3 (T cells) antibodies. *Images in (B)* previously shown at AACR 2024.6



Figure 10. The mouse CRC cell line MC38 engineered to express gp100 was implanted subcutaneously in C57BL/6J mice. After 26 days, implants developed into large (~300 mm³ average) fibrotic tumors and 5×10^{6} CD8+ pmel cells engineered with either mblL15 alone or mblL15-LIGHT were adoptively transferred into the tumor-bearing mice. Tumors were collected on Day 26 post-ACT and homogenized using GentleMACS to characterize the TME using flow cytometry. CD69 expressio of transduced pmel cells identified as single live CD45+ CD3+ CD8+ Thy1.1+ Thy1.2+ cells (A). Pie charts represent the percentage of host CD8+ cells expressing 0, 1, 2, or 3 of the exhaustion markers (TIM3, LAG3, PD1, TIGIT) analyzed (B). Host CD8+ cells identified as single live CD45+ CD3+ CD8+ Thy1.1- Thy1.2+ cells. Statistical tests: Unpaired t test, mblL15-LIGHT v mblL15, *p value<0.05 (A); Chi-square test, mblL15-LIGHT v mblL15, p-value = 0.194 (B).

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Figure 11. Tumor-draining (TDLN) and non-draining lymph nodes (NDLN) from opposite flank were collected on Day 20 post-ACT from mice described in Figure 10. To detect epitope spread, TDLN and NDLN were first homogenized by pushing through a 70-µm cell strainer using a syringe plunger. Singlepeptides relevant to MC38 tumor (Cpne1, Irgq, Reps1, Rp18 P15e, Zbtb40, Adpgk) or gp100 for 24 hours followed by the detection of IFNy-producing cells. Data representative of the average of two

- Adding LIGHT to mbIL15 in an autologous TIL product yields
- Pmel cells expressing both mbIL15 and LIGHT exhibited robust, durable tumor growth inhibition and provided a survival advantage in vivo over cells expressing mblL15
- Syngeneic mbIL15 and LIGHT co-expressing pmel cells induce B-cell aggregates and TLS formation in MC38 tumors in vivo
- LIGHT induces spreading of the T-cell response to MC38specific epitopes in tumor-draining lymph nodes, corroborating its ability to promote priming of new endogenous T-cell immune responses
- These preclinical results suggest that engineering TIL with regulatable mbIL15 and LIGHT has the potential to enhance their efficacy against challenging-to-treat tumors with a fibrotic TME, providing an opportunity to address unmet medical need

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	Abbrev	/ia	ations
ACT, adoptive cell transfer; ACZ, acetazolamide; CAF, cancer-associated fibroblast; CRC, colorectal cancer; DRD, drug- responsive domain; HEV, high endothelial venule; IV, intravenous; LIGHT, homologous to Lymphotoxin, exhibits Inducible expression and competes with HSV <u>G</u> lycoprotein D to <u>H</u> erpesvirus entry mediator, a receptor expressed on <u>T</u> lymphocytes; mbIL15, membrane-bound IL15; NDLN, non-draining lymph node; NT, no transfer; P0, passage 0; PDX, patient-derived xenograft; TCR, T-Cell Receptor; TDLN, tumor-draining lymph node; TIL, tumor-infiltrating lymphocytes; TLS, tertiary lymphoid structure; TME, tumor microenvironment; TNFSF14, tumor necrosis factor (ligand) superfamily member 14.			
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	Disclosures		
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