

# Adapting T cells for the Tumor Microenvironment (TME) During Manufacturing for Improved Anti-Tumor Potency

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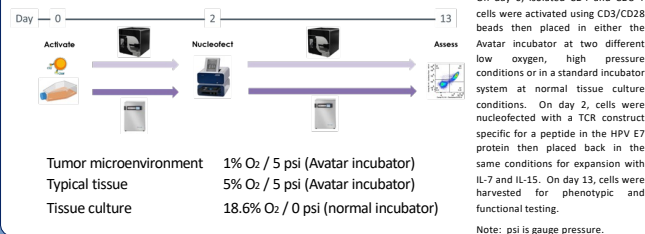
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## Introduction

The microenvironment of solid tumors is very challenging for anti-tumor T cells. Low oxygen, low pH, inhibitory signals (e.g. cell surface receptors, metabolites, lipids), and high intratumoral pressure drive infiltrating T cells toward inhibition, exhaustion and death. Improved T cell fitness in this environment is critical to efficacious and durable T cell therapies. The goal of this work was to improve the fitness of therapeutic T cells in the tumor microenvironment by changing the conditions under which a T cell therapy is manufactured. To this end, we utilized a low-oxygen, high-pressure environment created by the Avatar incubator system and AmplifyBio's small-scale Non-viral Gene Editing (NVGE™) platform to generate TCR-T cells with a metabolic program that improves their fitness and potency in the harsh TME.

## Methods

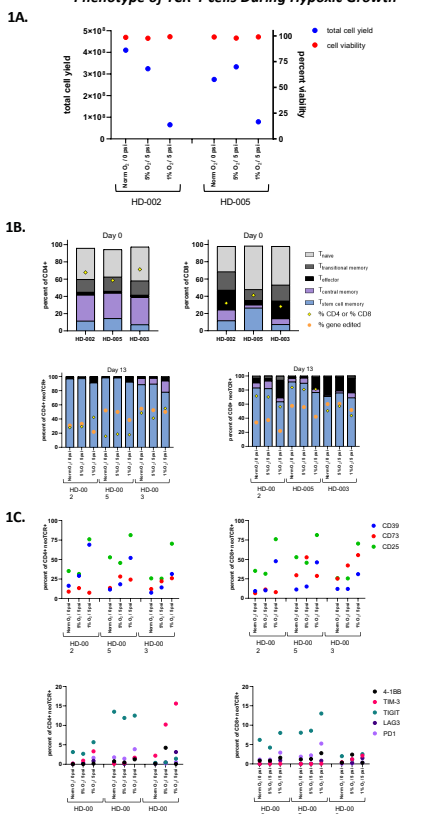


## Conclusions

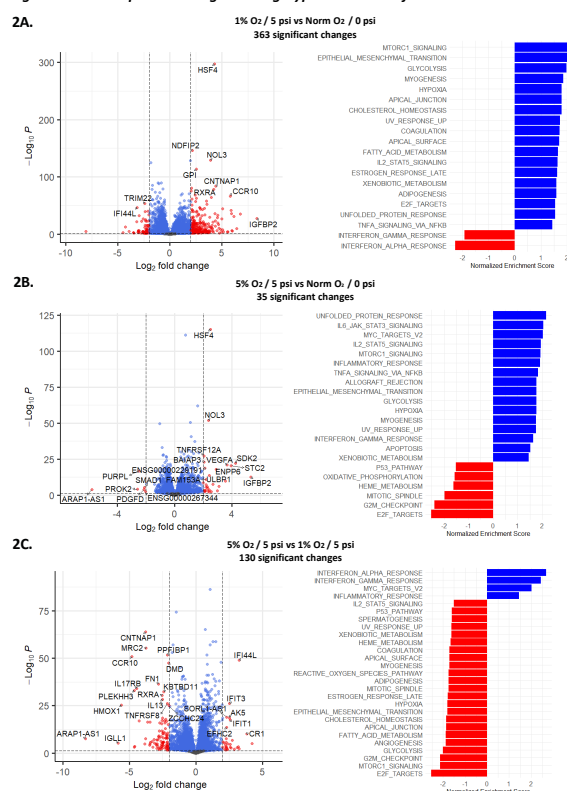
- TCR-T cells generated using AmplifyBio's small-scale NVGE™ platform in the Avatar at 5% O<sub>2</sub> / 5 psi (similar to a typical tissue environment) show growth, gene editing rates, memory subsets, and activation/exhaustion phenotypes similar to TCR-T cells expanded under standard tissue culture (normoxic) conditions.
- The TCR-T cells grown in a tissue environment exhibit changes in their transcriptome indicative of adaptation to lower oxygen levels, a shift toward more glycolytic metabolism while conserving their overall spare respiratory capacity, and consistently improved anti-tumor cell cytotoxicity at low effector-to-target ratios in a hypoxic environment relative to cells expanded under normoxic conditions.
- Growing TCR-T cells under extreme hypoxia (1% O<sub>2</sub> / 5 psi), as found in a tumor, was deleterious to all measures of phenotype and function.
- These results suggest there is a benefit in anti-tumor efficacy to manufacture TCR-T therapies under more physiological tissue oxygen and pressure conditions.

## Results

**Figure 1. Cell Growth Characteristics, Memory Subsets, and Phenotype of TCR-T cells During Hypoxic Growth**



**Figure 2. Transcriptome Changes During Hypoxic Growth of TCR-T cells**

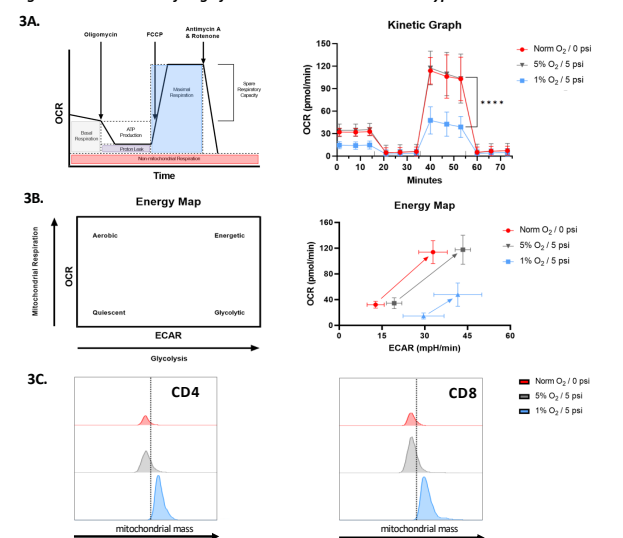


**2A.** Volcano plots of gene expression levels demonstrate the 1% O<sub>2</sub> / 5 psi condition has the largest number of significant changes in mRNA expression relative to the norm O<sub>2</sub> / 0 psi and 5% O<sub>2</sub> / 5 psi conditions including upregulation of known hypoxia-inducible genes like HSF4, IGF1R, VEGFA and NOL3. Normalized Enrichment Scores of pathway gene expression indicate that TCR-T cells expanded in 1% O<sub>2</sub> / 5 psi exhibit increased mRNA expression in pathways related to hypoxia, glycolysis, and cell stress, and decreased expression of the IFN- $\gamma$  pathway.

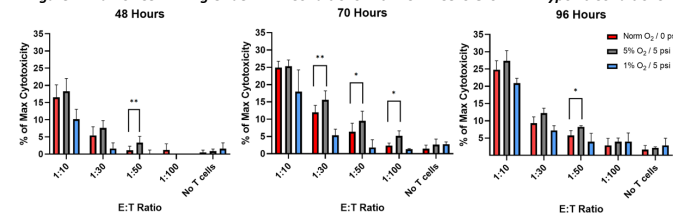
**2B.** The number of transcriptome differences between 5% O<sub>2</sub> / 5 psi and norm O<sub>2</sub> / 0 psi conditions was 35 total genes, suggesting that this condition does not perturb the T cells as much as the full hypoxia condition. Nevertheless, many of the same pathways were upregulated as in the 1% O<sub>2</sub> / 5 psi condition. However, significant upregulated pathways included anti-tumor and inflammatory pathways such as IL-2, IFN- $\gamma$ , and TNF- $\alpha$ , suggesting improved function.

**2C.** Comparison of the TCR-T cells expanded in either 5% O<sub>2</sub> / 5 psi or 1% O<sub>2</sub> / 5 psi confirms the upregulation of inflammatory and IFN- $\gamma$  pathways in the 5% O<sub>2</sub> / 5 psi condition, and down-regulation of glycolysis in the 1% O<sub>2</sub> / 5 psi condition. RNA-seq was used to assess the effect of the different conditions on the transcriptomes of the TCR-T cells (2 donors) at day 14. Differential expression analysis was performed using DESeq2, with a  $|\log_2(\text{foldchange})| > 2$  considered significant. Using Gene Set Enrichment Analysis (GSEA), Hallmark pathways from the Human MSigDB Collection were analyzed for up- or down-regulation at day 14, with an adjusted p value < 0.01 defining significance.

**Figure 3. Metabolic Profiling of TCR-T Cells Generated Under Hypoxic Conditions**



**Figure 4. Tumor Cell Killing Under TME Conditions with TCR-T Cells Grown in Hypoxic Conditions**



TCR-T cell cytotoxicity against CaSki tumor cells tested under TME conditions (1% O<sub>2</sub> / 2 psi) shows that TCR-T cells cultured at 5% O<sub>2</sub> / 5 psi demonstrate consistently higher cytotoxicity at multiple E:T ratios relative to norm O<sub>2</sub> / 0 psi and 1% O<sub>2</sub> / 5 psi conditions. Data shown are the average plus standard deviation of 3 separate donors. Significance assessed by a paired Student's t test at each E:T ratio. Cytotoxicity of the CaSki cells assessed by release of lactate dehydrogenase (LDH) in a plate-based format.