Metabolic Reprogramming Enhances Expansion and Potency of CAR-T cells

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Next steps: Further exploring the effects of O_2 levels and pressure on transduced T cells in vitro and future in vivo experiments

On-going in vitro investigations. More studies are underway to understand the effects of oxygen and pressure on transduction efficiency, expansion, phenotype and potency on T cells grown in the AVATAR.





CD19-CAR T cells grown in 5_5 AVATAR conditions kill better at low E:T ratios than the other conditions. These T cells performed well in an acute (24hr) killing assay and demonstrated good serial killing at 72hrs.

- Introducing pressure in the T cell manufacturing process enhances the overall vield.
- Phenotype and function of CD19-CAR T cells grown in the AVATAR have no significant effect on phenotype and cytotoxic function *in vitro* or *in vivo*.
- Experiments are currently being conducted to further validate these discoveries. The next step is to determine the correlation between in vitro and in vivo efficacy. Additionally, we are investigating solid tumor models using a similar approach.



Phenotypic analysis of circulating CD3/4/8 T cells: Population Distribution and Memory Subsets

CD4 and CD8 distribution. Blood samples from each timepoint were analyzed for T cell populations. The figures on the left show the distribution of CD4+ and CD8+ T cell within the total CD3+ gate. The CD4 population increased overtime, while the CD8+ population decreased. The drop in the number of CD8+ T cells present correlated with the increasing tumor burden.

- Vehicle NI UTD
- NI CAR 2.5e6
- o NI CAR 1.25e6
- CAR 2.5e6 15_2
- △ CAR 1.25e6 15 2 CAR 2.5e6 15 5
- CAR 1.25e6 15_5

Effector memory (EM) and central memory (CM) distribution. Both EM and CM populations were also assessed with in the CD3/4/8 subsets. EM and CM were defined as CD3/4/8+CD45RA-CCR7- and CD45RA-CCR7+ respectively. Overall, the cells remained within the EM and CM populations and did not terminally differentiate or appear exhausted via LAG3+ expression. LAG3+ expression was barely detectable in any of the major subsets.

Conclusions

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