# Single cell RNA sequencing reveals functionally validated signatures of cytotoxicity in anti-FRα CoStimulatory Antigen Receptor (CoStAR™) activated CD4+ T cells. Instilbio

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### **INTRODUCTION**

#### • There are several limiting factors for the development of cell therapy modalities in solid tumors, including tumour cell clonal heterogeneity, high expression of inhibitory receptors and poor endogenous costimulation (1-5).

- Tumor-infiltrating lymphocytes (TILs) contain a hugely diverse T-cell receptor (TCR) repertoire (6), thereby offering the broadest diversity of antitumor reactivity.
- TIL therapy has a proven track record of effective clinical efficacy (7-9).
- We have developed the CoStimulatory Antigen Receptor (CoStAR) platform to overcome the

## RESULTS



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- suppressive TME.
- The CoStAR described here consists of a FRαspecific single chain antibody fragment (scFv) fused to the signaling domains from CD28 and CD40.
- We have previously shown that engagement of CoStAR amplifies TCR signalling leading to enhanced T cell activity (10,11).
- Here, we use coculture assays and subsequent scRNA-seq analysis with functional in vitro validation techniques to explore the relative contribution of CoStAR to TCR activity in an effort to further understand CoStAR function.

### **METHODS**

- T cells from three healthy donors were engineered with an HLA-A\*02/CEA (CEA:691-699 IMIGVLVGV) specific TCR and/or FRα-specific CoStAR and subsequently enriched for expression.
- T cells were cocultured with HLA-A\*02+/CEA+ NCI-H508 tumour cells overexpressing human FRa (H508-FRa) before processing through 10X Genomics 5' GEX scRNA-seq workflow.
- Bioinformatic analysis was used to compare TCR and/or CoStAR+ populations, as well as CD4+ and CD8+ populations.
- Functional validation of gene signatures was performed by isolating CD4+ and CD8+ cells and coculturing with H508-FRa, before analysis of

#### $Log_2$ fold change cutoff, 0.25; p-value cutoff, 0.05

#### Figure 2: Differential expression analysis reveals CoStAR activity leads to different activation and differentiation states in CD4+ and CD8+ T cells.

- TCR+CoStAR signaling resulted in increased expression of genes associated with activation (e.g. TNFRSF4, TNFRSF9) and a cytotoxic phenotype (e.g. GZMA, GZMB, GNLY) in CD4+ cells compared to TCR alone signaling.
- TCR+CoStAR signaling resulted in increased expression of genes associated with a less differentiated phenotype (e.g. CCR7, SELL) and increased cytotoxicity and decreased expression of genes associated with exhaustion (e.g. HAVCR2, TIGIT, PDCD1) in CD8+ cells compared to TCR alone signaling.

TCR+CoStAR-Td vs TCR-Td CD4+ T cells CD8+ T cells KEGG\_DNA\_REPLICATION GOBP\_CHROMOSOME\_SEGREGATION padj GOBP\_CELL\_KILLING · 0.03 GOBP\_RESPONSE\_TO\_INTERFERON\_GAMMA KEGG\_CYTOKINE\_CYTOKINE\_RECEPTOR\_INTERACTION · 0.02 KEGG\_CHEMOKINE\_SIGNALING\_PATHWAY GOMF\_CYTOKINE\_ACTIVITY · 0.01 GOMF\_CHEMOKINE\_ACTIVITY GOBP\_POSITIVE\_REGULATION\_OF\_CYTOKINE\_PRODUCTION · GOBP\_CYTOKINE\_MEDIATED\_SIGNALING\_PATHWAY -GOBP\_LEUKOCYTE\_MIGRATION -Normalized Enrichment Score

Figure 3: Pathway analysis shows enrichment of pathways associated with cytokine signaling and cell killing in CD4+ T cells and with migration and cell cycle in CD8+ T cells.

- Pathway analysis indicated enhancement of cytotoxic phenotype and cell killing in CD4+ T cells upon CoStAR-mediated costimulation.
- In CD8+ T cells, TCR+CoStAR signaling resulted positive in enrichment of cell cycle related pathways and negative enrichment of leukocyte migration in line with a less differentiated phenotype.

#### Figure 5: Cytotoxicity of CD4+ and CD8+ T cells expressing TCR and/or CoStAR using xCELLigence.

- The unexpected observation of enhancement of cytotoxicity signatures in CD4+ T cells with CoStARcostimulation, led us to explore CoStAR-mediated cytotoxicity in vitro.
- TCR and/or CoStAR+ CD4+ and CD8+ T cells were negatively enriched and added to NCI-H508-FRa tumour cells at the indicated effector to target ratios.
- xCELLigence was used to track killing and measurements made at 24, 48 and 72h.
- At a 10:1 E:T we observed efficient killing of target cells by CD4+ TCR+CoStAR-Td T cells vs TCR+ alone at 48 and 72h (P<0.05).



- CD8+ T cells engineered with TCR and/ or CoStAR
- Area under the curve analysis was performed for CD4+ and CD8+ cells across each E:T ratio over 100h.
- We observed significantly increased cytotoxicity with CD4+ cells expressing CEA TCR + CoStAR vs TCR alone (P<0.05) or CoStAR alone (0.0005).

cytokine and chemokine by bead arrays and multiplexed analyte analysis (Mesoscale Discovery), flow cytometric staining and xCELLigence cytotoxicity assays.



### **RESULTS**

- A custom gene reference was created and could successfully quantify scRNA-seq reads of anti-FRa-CoStAR, as well as the recombinant TCR introduced into the T cells.
- A cell is CoStAR+ if at least 1 mRNA copy maps to the CoStAR scFv nucleotide sequence included in the custom reference.
- This was compared with flow cytometric analysis of TCR and CoStAR, with GEX and flow cytometry



### CONCLUSION

- Anti-FRα CoStAR provide functional benefit to CD4+ and CD8+ T cells in overlapping and distinct ways.
- CoStAR enhancement of TCR signalling enhanced signatures of activation and cytotoxicity in CD4+ cells, and enhancement of genes associated with less differentiation and cytotoxicity in CD8+ T cells.
- Functional validation of these gene changes using flow cytometry and multiplexed immunoassay demonstrated strong concordance between gene and protein expression, including IP-10, GM-CSF and GZMB.
- Furthermore, certain proteins including CD134 could be observed to be upregulated following CoStAR engagement alone, suggesting CoStAR activity can have a broader positive impact on T cell function.
- The cytotoxicity profile seen in CD4+ T cells was functionally validated in isolated CD4+ T cells, demonstrating that CoStAR can imbue CD4+ T cells with cytotoxic activity.
- These results support the clinical exploration of anti-FRa CoStAR in FRa+ tumor indications.

### REFERENCES



### showing good concordance.



**Figure 1: Measurements of CoStAR transduction** efficiency are concordant between scRNA-seq and flow cytometry.

• We used flow cytometry and multiplexed immunoassay to measure several cell surface markers and soluble effector molecules produced by negatively enriched CD4+ and CD8+ T cells under conditions of stimulation of TCR and/or CoStAR following coculture with NCI-H508-FRa tumour cells.

Figure 4: Protein analysis shows good concordance of multiple genes identified

through single cell RNA sequencing analysis

- We were able to functionally validate the upregulation of CD134 (TNFRSF4) and CD137 (TNFRSF9) in CD4+ and CD8+ T cells following CoStAR stimulation alone. In fact, in CD4+ T cells both markers were upregulated by CoStAR engagement to a higher degree than following TCR alone stimulation.
- Many other proteins had good concordance with our gene expression data, including the cytokines GM-CSF (CSF2), IL13, and IFNy; and the chemokines IP10 (CXCL10) and MIP1a (CCL3).
- Our analysis also revealed CoStAR-enhanced upregulation of FasL (FASLG), GZMA and GZMB in CD4+ and CD8+ T cells.

1. Marofi F, et al. Stem Cell Res Ther. 2021;12:81. 2. Albelda SM. Cancer Immunol Res. 2020;8:2. 3. Schnell A et al. Cell Research 2020;30:285 4. Truxova I, et al. J Immunother Cancer. 2018;6:139. 5. Bandola-Simon J, Roche PA. Mol Immunol. 2019;113:31-37. 6. Spindler MJ, et al. Nat Biotechnol. 2020;38:609-619. 7. Rohaan MW, et al. J Immunother Cancer. 2018;6:102. 8. Betof Warner A, et al. Clin. Cancer. Res. 2023;29:1835. 9. Pillai M, et al. Am. J. Cancer. Res. 2022;12:3967 10. Sukumaran S, et al. JITC 2021; 2021-SITC 2021.198 11. Sykorova M, et al. JITC2022; SITC-2022.0282 All authors are current or former employees of Instil Bio and may ahve stock and/or stock options

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