

Utilizing Stim-R™ Technology to Reduce Irradiated Feeder Cell Use in the Tumor Infiltrating Lymphocyte Culture Process

Riley Allen, Jason Duecker, Jessica Ojeda, Benjamin Harris, Mikayla Min, Samantha Chan, Christina Ta, Emily Fu-Sum, Purnima Sundar, Yogin Patel, Aileen Li, Shobha Potluri, Suman Vodhala, Gary Lee, Alexander Cheung

Lyell Immunopharma, Inc., South San Francisco, CA and Seattle, WA

For more information, please contact: rallen@lyell.com

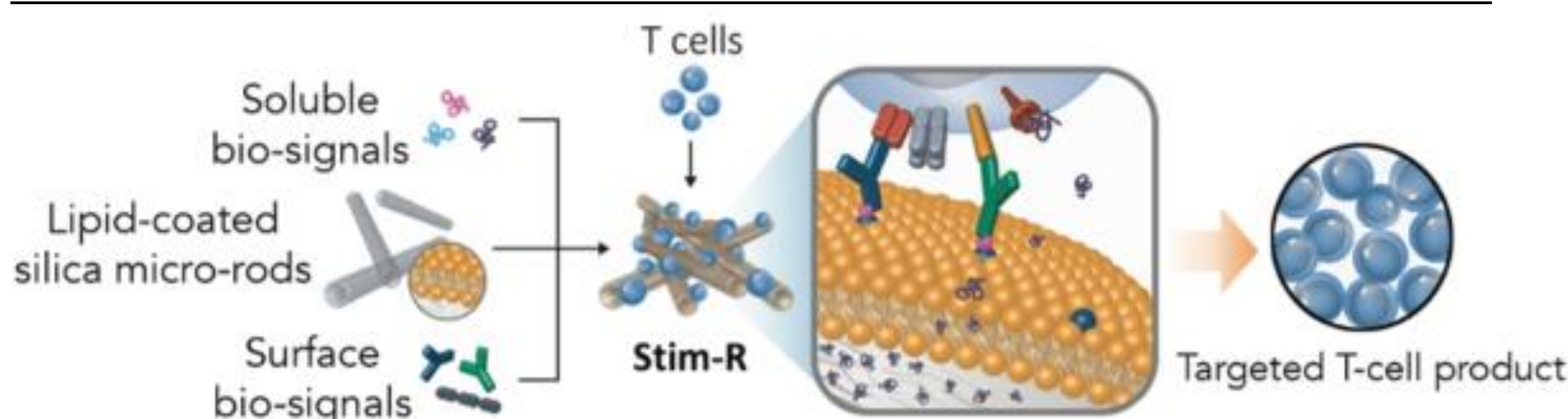
Abstract #440

Background

- Tumor-infiltrating lymphocyte (TIL) therapy can mediate effective responses in solid tumors and is currently FDA approved for metastatic melanoma in patients previously treated with other therapies.
- Conventional TIL manufacturing processes require the use of “feeder cells”, irradiated allogeneic peripheral blood mononuclear cells (PBMCs).
- Sourcing feeder cells from qualified donors may incur logistical and financial obstacles¹.
- Feeder cells introduce donor-to-donor variability¹.
- Stim-R™ is a customizable and degradable biomimetic emulating physiologic, cell-like presentation of signal molecules (Fig. 1) and may be used as a replacement for feeder cells during TIL manufacturing.

Methods

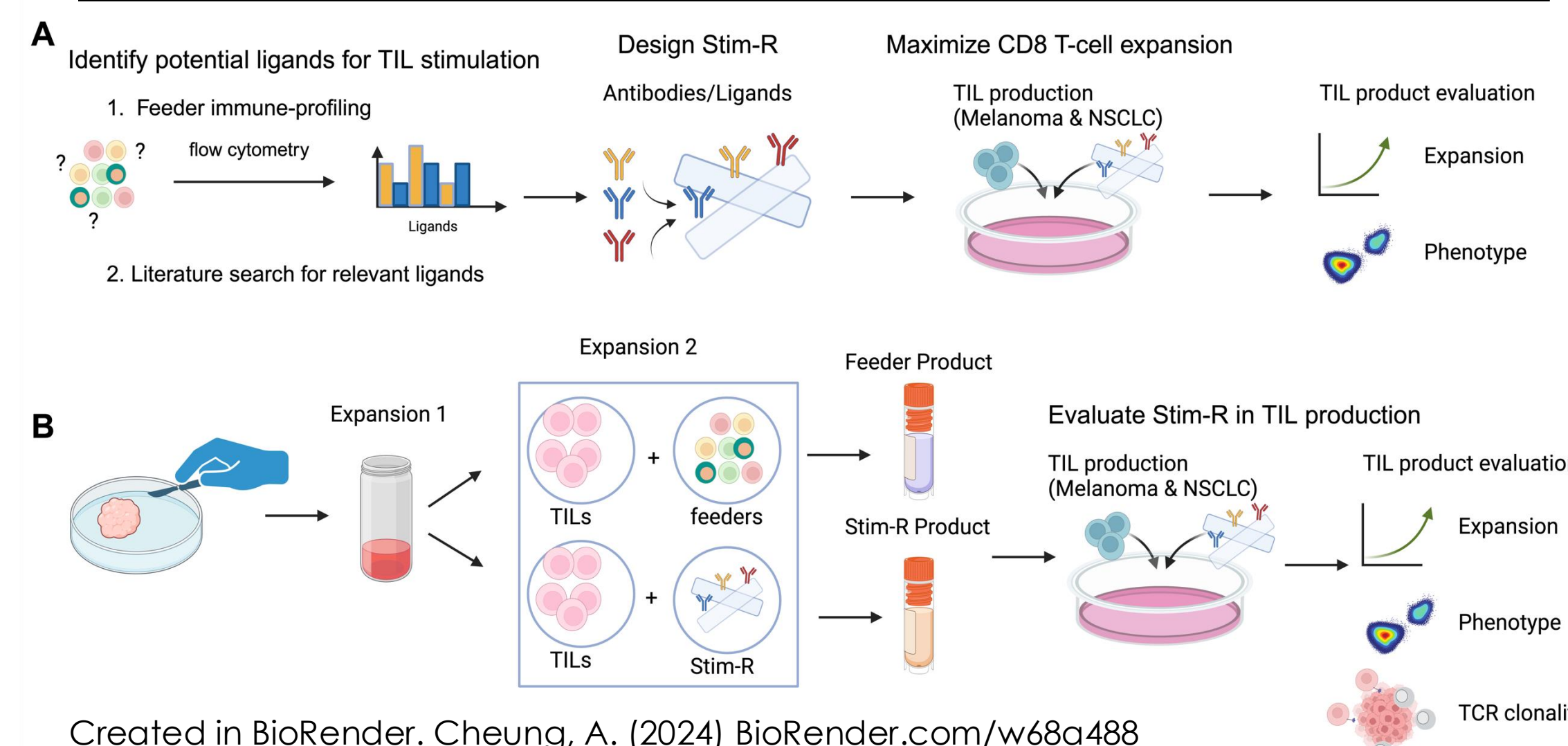
Figure 1: The Stim-R technology is a programmable cell-signaling platform for feeder cell replacement



The Stim-R technology comprises biodegradable lipid-coated silica micro-rods that can present multiple signals in precise densities and stoichiometries² in a way that mimics physiologic presentation.

- Feeder cells were analyzed using flow cytometry to assess the expression of potential T-cell stimulating ligands^{3,4,5}. Based on these data, Stim-R formulations were designed to present these ligands in various combinations and stoichiometries (Fig. 2A).
- Control TILs were produced using Lyell's Epi-R™ P2⁶ protocol at research scale. Stim-R was utilized as a replacement for feeder cells in “expansion 2” (Fig. 2B). A matrix of assays was used to compare the performance of Stim-R to feeder cells:
- TIL expansion and phenotype measured by automated cell counter & spectral cytometer.
- T-Cell Receptor (TCR) sequencing to assess polyclonality and retention of putative tumor-reactive cells (PTRCs) in products. (PTRCs are cells that have TCRs identified from high-frequency T-cell clones in the starting tumor material)⁷.

Figure 2: Stim-R for TIL formulation design and screening workflow to assess product attributes^{8,9}



Created in BioRender. Cheung, A. (2024) BioRender.com/w68a488

- (A) Ligands that are crucial for T-cell activation were identified and assessed in different combination on Stim-R for their ability to expand CD8⁺ T cells in “expansion 2”. Ref. (Fig. 3 & 4)
- (B) An optimized formulation of Stim-R was further benchmarked against the standard Epi-R P2 protocol in terms of CD8 T-cell stemness, polyclonality, and the retention of PTRCs. Ref. (Fig. 5, 6 & 7)

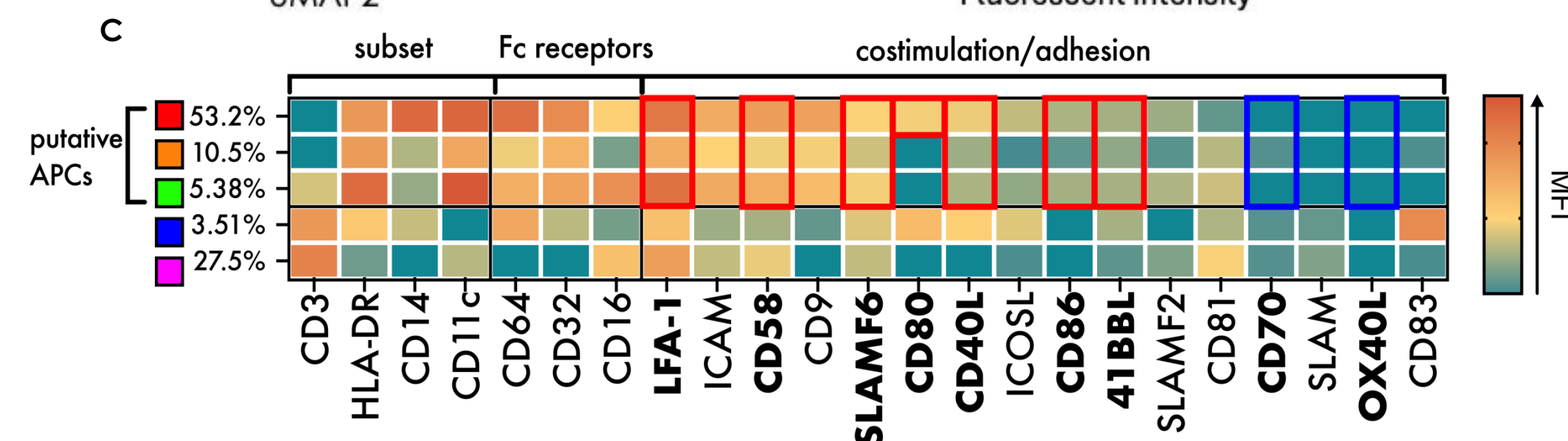
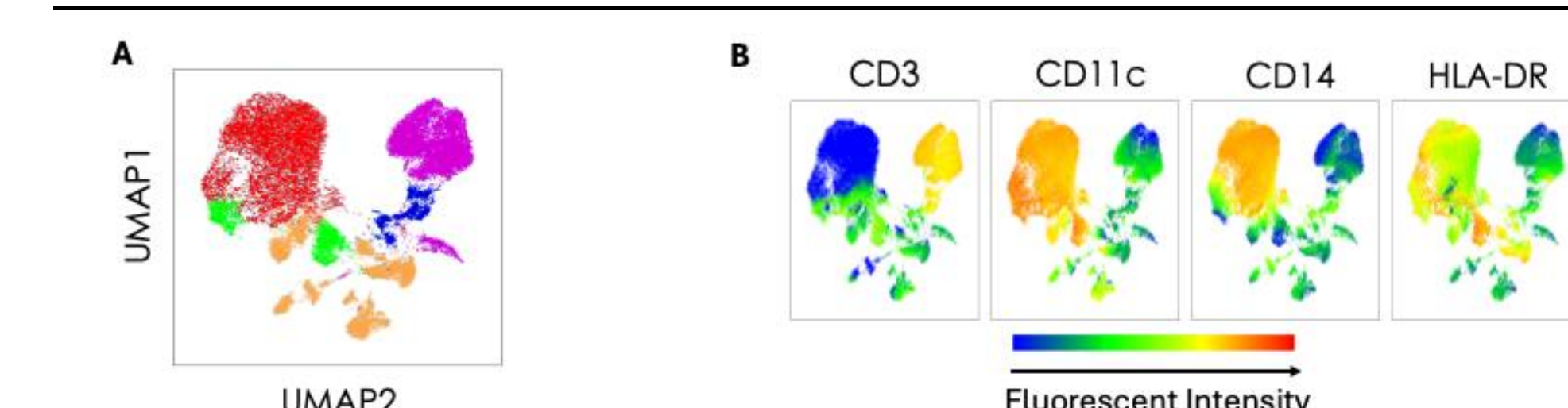
Results

Key Findings:

Stim-R (EXT221) is able to replace feeder cells in “expansion 2” of Epi-R P2 and produce TIL products with:

- Comparable expansion of CD8⁺ T cells.
- Similar stemlike characteristics measured by the percentage of CD62L⁺CD27⁺ and CD39⁺CD69⁻ cells in the CD8⁺ T-cell population.
- Comparable polyclonality, measured by the Simpson Clonality Index.
- Proportional putative tumor reactive cells (PTRCs).

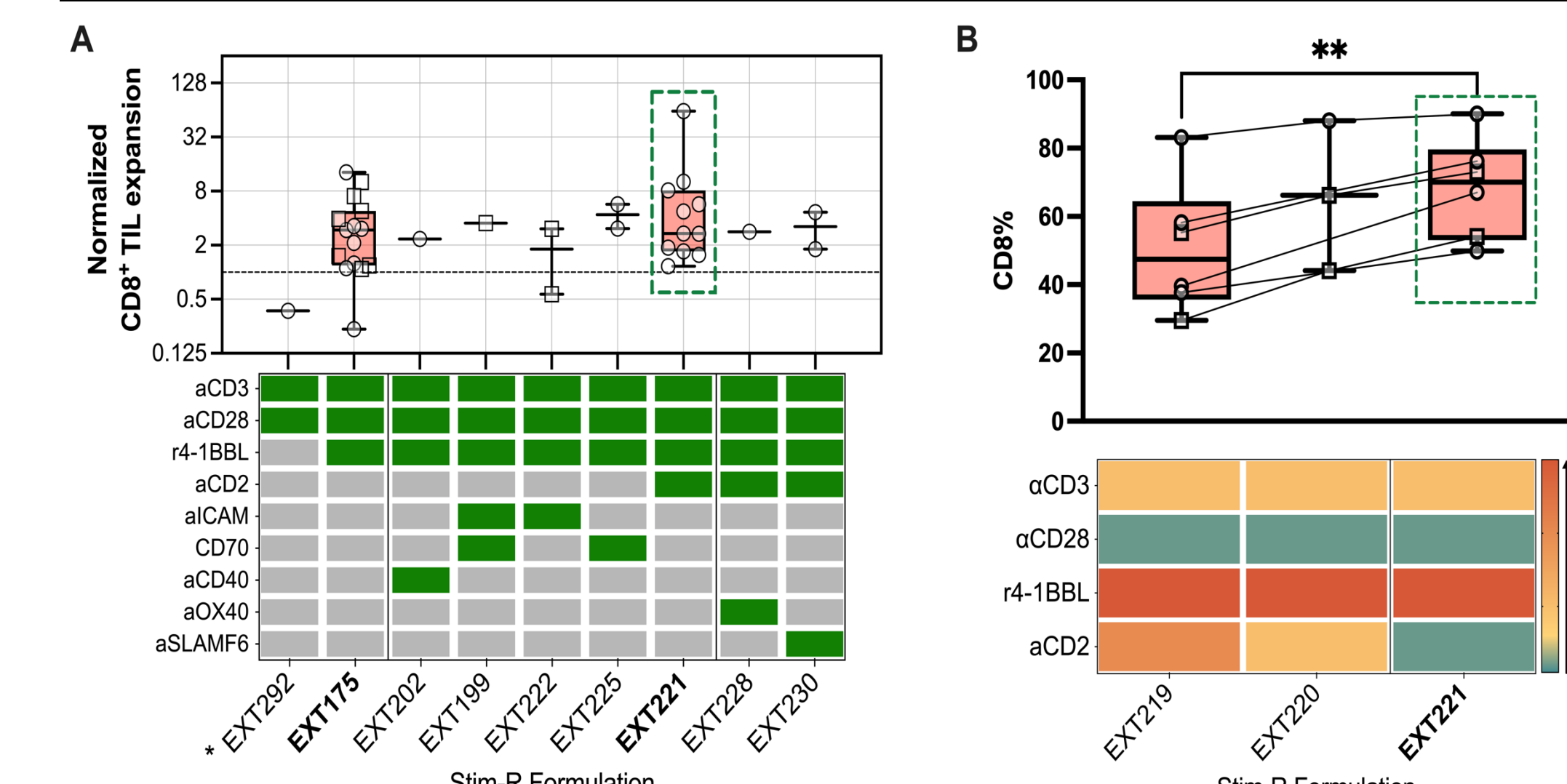
Figure 3: Putative Antigen Presenting Cells (APCs) in bulk feeder cell population express various costimulatory and adhesion molecules potentially important for T-cell activation



Feeder cells pooled from three healthy donors were profiled using flow cytometry.

- (A) & (B) Putative APCs, the hypothetical stimulatory population in the feeder cells, were identified based on high HLA-DR and low CD3 expression.
- (C) Heat map of the expression of various differentiation, Fc receptors, and costimulatory/adhesion molecules on the surface of the APC population.

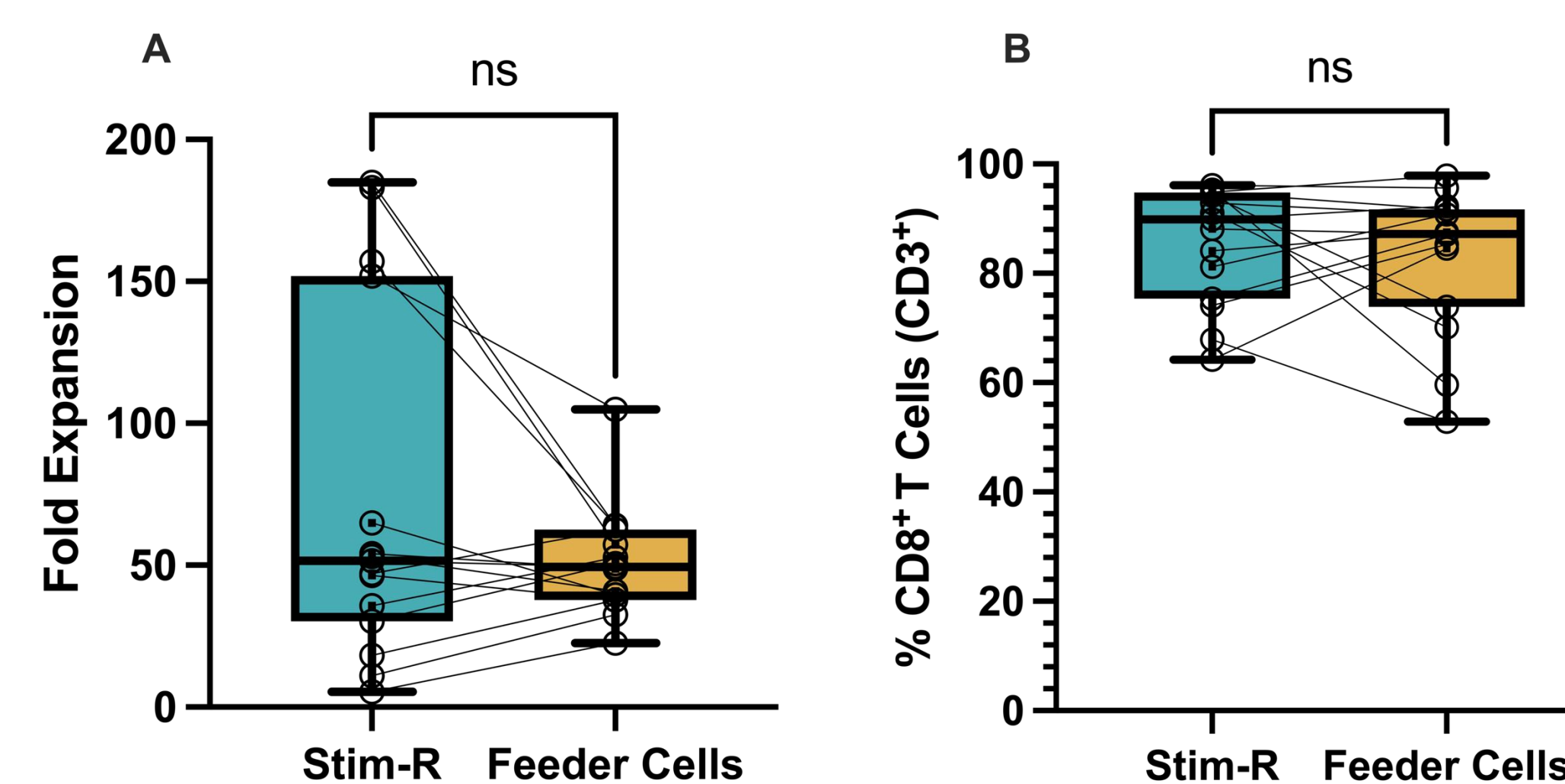
Figure 4: Screening based on CD8 T-cell expansion identified an optimal combination of ligands



$$\text{Normalized CD8}^+ \text{ TIL expansion} = \frac{\text{fold expansion} \times \text{CD8}^+ \text{ Stim-R product}}{\text{fold expansion} \times \text{CD8}^+ \text{ Epi-R P2 feeder product}}$$

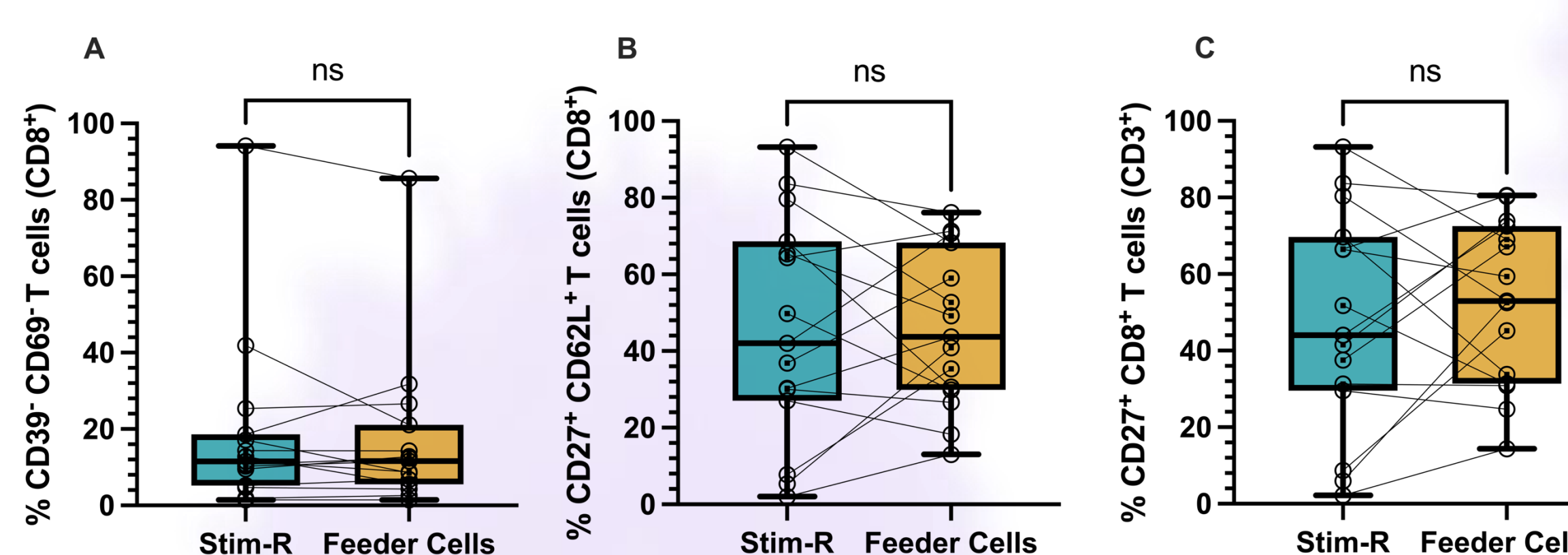
- (A) Preliminary experiments identified r4-1BB signaling as critical to TIL expansion (EXT175). Subsequently, the addition of aCD2 signaling allowed for consistent normalized CD8⁺ TIL expansion over a diverse set of tumors (EXT221). The * designates that EXT292 was tried in n=3 additional tumors but expansion was not high enough to obtain a CD8⁺ percentage from the TIL product.
- (B) Titration of aCD2 on Stim-R was performed to optimize aCD2 intensity. EXT221 was identified as the optimal formulation for feeder replacement based on higher CD8 T cell percentage in the expanded product.

Figure 5: Utilizing Stim-R in “expansion 2” of Epi-R P2 maintains equivalent CD8⁺ T-cell expansion



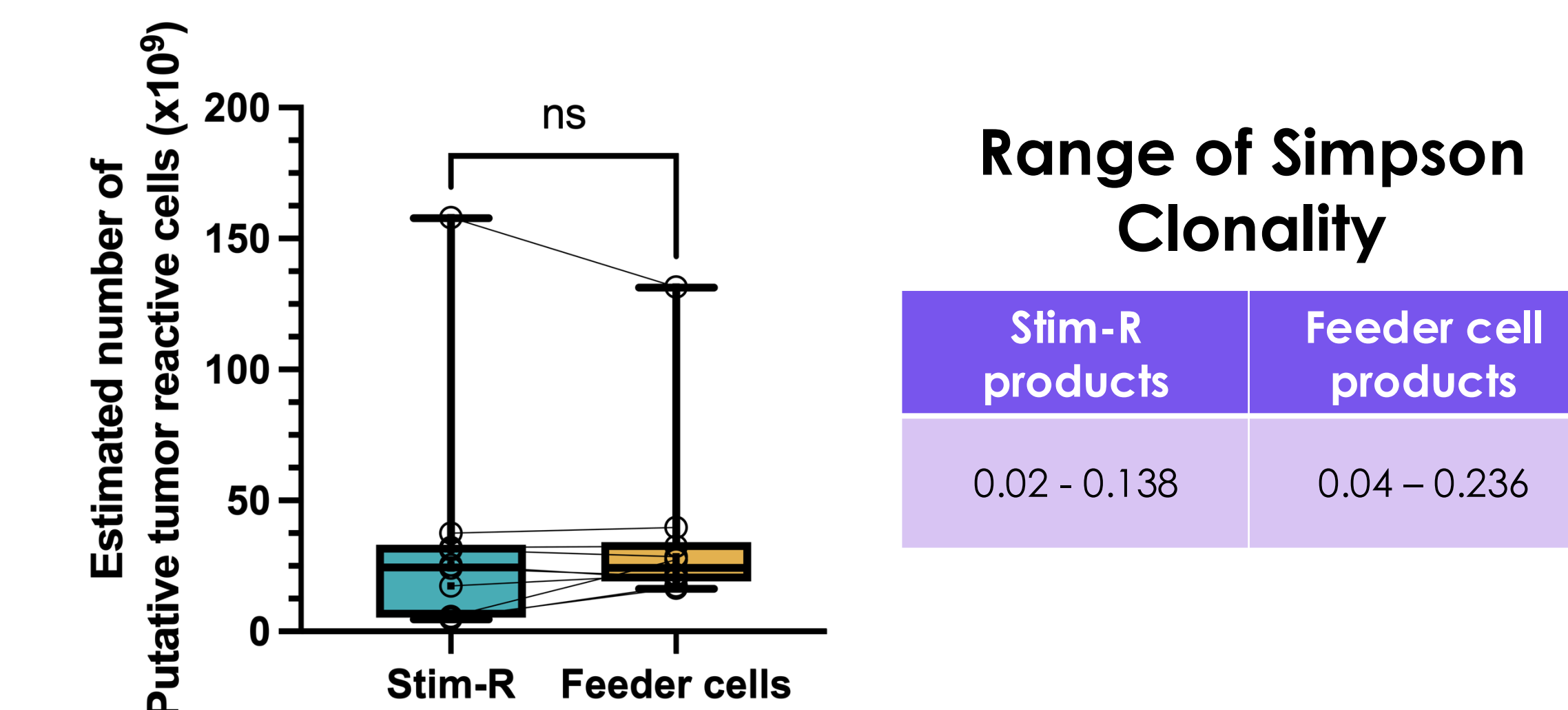
- (A) & (B) Comparison of fold expansion and CD8⁺ percentage of TIL products stimulated with either Stim-R or Feeder cells in “expansion 2” of TIL production. Statistical significance was determined via paired t-test, n=14.

Figure 6: Utilizing Stim-R in “expansion 2” of Epi-R P2 maintains comparable stemness of expanded product



- (A), (B) & (C) Comparison of stemness phenotype of TIL products stimulated with either Stim-R or Feeder cells in “expansion 2” of TIL production. Statistical significance was determined via a paired t-test, n=14.

Figure 7: Using Stim-R in place of feeder cells in the “expansion 2” step of Epi-R P2 results in a polyclonal product that retains a similar number of estimated putative tumor reactive cells



Estimated number of PTRCs in TIL products that were activated with either Stim-R or feeder cells in “expansion 2.” A paired t-test was run; no statistical difference was observed (n=10). Estimated number of PTRCs was calculated by extrapolating the number of PTRCs detected to a clinical scale production. Both methods of stimulation resulted in a polyclonal product; the range of Simpson Clonality across all samples in both groups is shown.

Conclusions

- A Stim-R formulation engaging CD2, CD3, CD28 and 4-1BB, designated “EXT221”, was identified based on its ability to mediate robust CD8⁺ TIL expansion in “expansion 2” of the Epi-R P2 protocol.
- Stemlike qualities, polyclonality, and number of PTRCs were similar in the TIL products expanded with Stim-R compared to control TILs.
- Our results suggest that Stim-R is a viable replacement for feeder cells in “expansion 2” of the Epi-R P2 manufacturing protocol, and has the potential to overcome limitations that may arise in feeder cell-mediated TIL expansion.

Acknowledgments

We would like to thank the members of Lyell's Flow Cytometry core (Ken Xiong, Sahithi Cheemalamari) as well as Claire Toney, Melissa Bedard, Audrey Garces, Stefan Siebert, Omar Ali, and Lara Zhao.

References

- Ye Q et al. *J Translational Medicine*. 2011; (9): 131
- Li A et al. SITC Poster 2022. Abstract 252.
- Forget MA, et al. *J Immunother*. 2014;37(9):448-460.
- Ma V, et al. *Sci Adv*. 8(8):4485-4501.
- Yigit B, et al. *Cancer Immunol Res*. 2019;7(9):1485-1496.
- Patel Y, et al. SITC Poster 2023. Abstract 279.
- Olivera G, et al. *Nature*. 2021;596(7870):119-125.
- Krishna S, et al. *Science*. 2020 370, 1328-1334.
- Van den Berg JH, et al. *J Immunother*. 2020;848-859.