In Vivo Pooled Screening Platform for the Discovery of **Optimized Chimeric Antigen Receptor (CAR) Design in T cells**

Joshua Mace¹, Matthew Forsberg², Nina La Vonne Denne^{3,4}, Siqi Zhao¹, Khloe Gordon Wei¹, Jai Raman¹, Narendra Maheshri¹, Christian Capitini², Krishanu Saha^{3,4}, Shawdee Eshghi¹, Taeyoon Kyung¹ ¹ Ginkgo Bioworks, Boston, MA, ² University of Wisconsin School of Medicine and Public Health, Madison, WI, ⁴ Wisconsin Institute for Discovery, University of Wisconsin-Madison, WI, ⁴ Wisconsin Institute for Discovery, University of Wisconsin-Madison, WI, ⁴ Wisconsin Institute for Discovery, University of Wisconsin-Madison, WI, ⁴ Wisconsin Institute for Discovery, University of Wisconsin-Madison, WI, ⁴ Wisconsin Institute for Discovery, University of Wisconsin-Madison, WI, ⁴ Wisconsin Institute for Discovery, University of Wisconsin-Madison, WI, ⁴ Wisconsin Institute for Discovery, University of Wisconsin-Madison, WI, ⁴ Wisconsin Institute for Discovery, University of Wisconsin-Madison, WI, ⁴ Wisconsin Institute for Discovery, University of Wisconsin-Madison, WI, ⁴ Wisconsin Institute for Discovery, University of Wisconsin-Madison, WI, ⁴ Wisconsin Institute for Discovery, University of Wisconsin-Madison, WI, ⁴ Wisconsin Institute for Discovery, University of Wisconsin-Madison, WI, ⁴ Wisconsin Institute for Discovery, University of Wisconsin-Madison, WI, ⁴ Wisconsin Institute for Discovery, University of Wisconsin-Madison, WI, ⁴ Wisconsin Institute for Discovery, University of Wisconsin-Madison, WI, ⁴ Wisconsin Institute for Discovery, University of Wisconsin-Madison, WI, ⁴ Wisconsin Institute for Discovery, University of Wisconsin-Madison, WI, ⁴ Wisconsin Institute for Discovery, University of W

Background

While Chimeric Antigen Receptor expressing T cells (CAR-T cells) have shown high response rates against hematological cancers in the clinic, they have shown limited efficacy against solid tumors due to lack of persistence and immunosuppression in the tumor microenvironment (TME). To address the TME, recent efforts have focused on discovering novel intracellular signaling domains (ICDs) with enhanced anti-tumor efficacy at scale. However, in vitro screens are inherently limited in mimicking the TME and other physiologically relevant aspects of adoptive cell therapy (ACT).

Here, we developed an in vivo pooled library screen approach alongside our existing in vitro platform to identify potent CAR designs in a CHLA-20 neuroblastoma xenograft model. We cloned a DNA-barcoded 10,000-member CAR ICD library into an anti-GD2 CAR backbone. GD2 CAR library-expressing primary T cells were injected into tumor-bearing mice at >100x library coverage, and spleens and tumors were collected 2 weeks post-ACT. Sequencing genomic DNA quantified the abundance of barcodes linked to each CAR ICD combination. We hypothesized that any remaining CAR-T cells at the tumor site and spleen would represent T cells with long-term persistence elicited by antigen-specific CAR ICD signaling. 103 novel CAR constructs were synthesized based on barcode enrichment from both in vitro and in vivo pooled screens, and we validated their anti-tumor function in a serial tumor rechallenge assay.





Relative frequency of ICD appearance among enriched barcodes from in vitro and in vivo pooled screens





