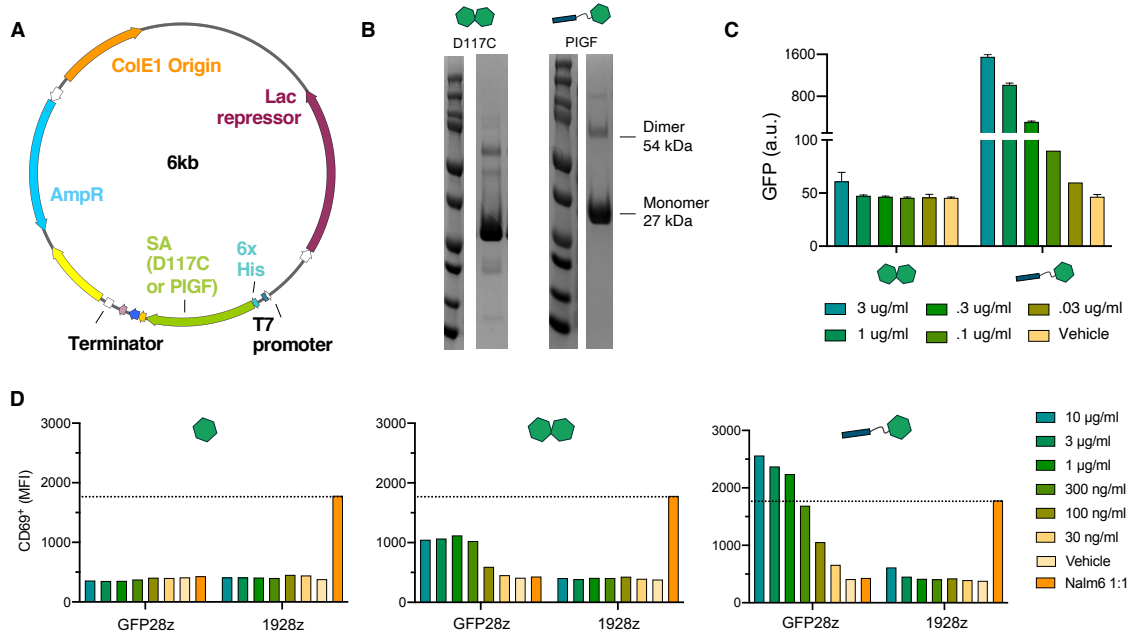
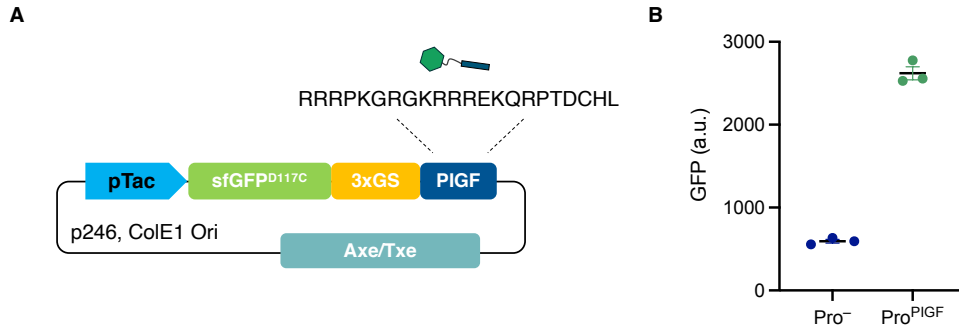


SUPPLEMENTARY MATERIALS

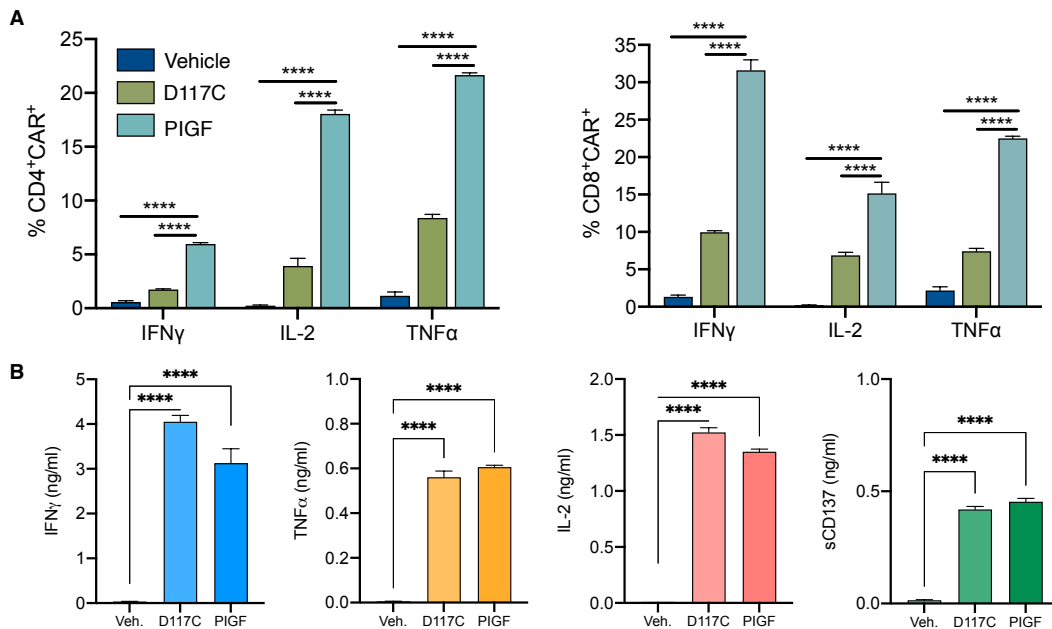


Supplementary Figure 1. Recombinantly produced, His-tag purified PIGF binds to collagen coated plates and strongly activates GFP-CAR⁺ Jurkats. (A) Plasmid map of protein expression vector transformed into eNiCo21(DE3) *E. coli* cells for protein purification of D117C and PIGF sfGFP variants. (B) Elutions of His-tag purified sfGFP variants, sizes show monomeric and dimeric molecules. (C) D117C and PIGF variants were plated in half log dilutions in PBS on collagen coated plates, incubated for 30 m at 37°C and washed 2X with PBS to dislodge any unbound protein. Fluorescence intensity was read at 488 nm on a standard Tecan plate reader. (D) CD69 expression on GFP CAR⁺ or CD19 CAR⁺ Jurkat cells. Jurkats were plated on collagen coated plates and assessed for CD69 expression by flow cytometry following 16-hour incubation with purified GFP antigens or co-culture with CD19⁺ target cells (Nalm6). Error bars represent s.d. of biological replicates.

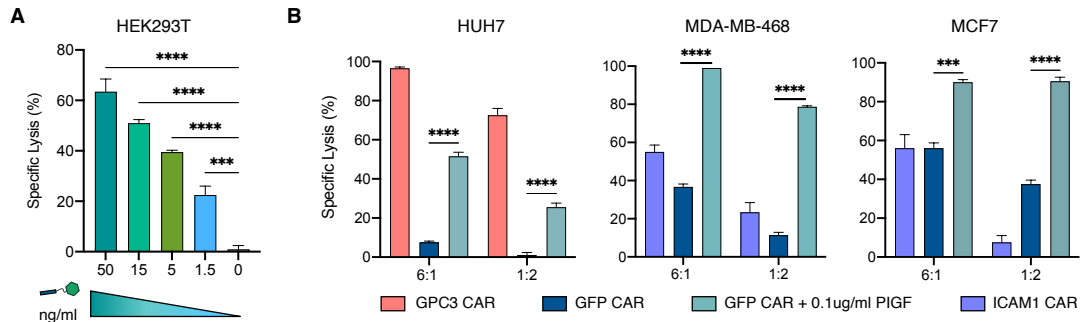


Supplementary Figure 2. Therapeutic *E. coli* Nissle 1917 (EcN) produces sfGFP-PIGF SA.

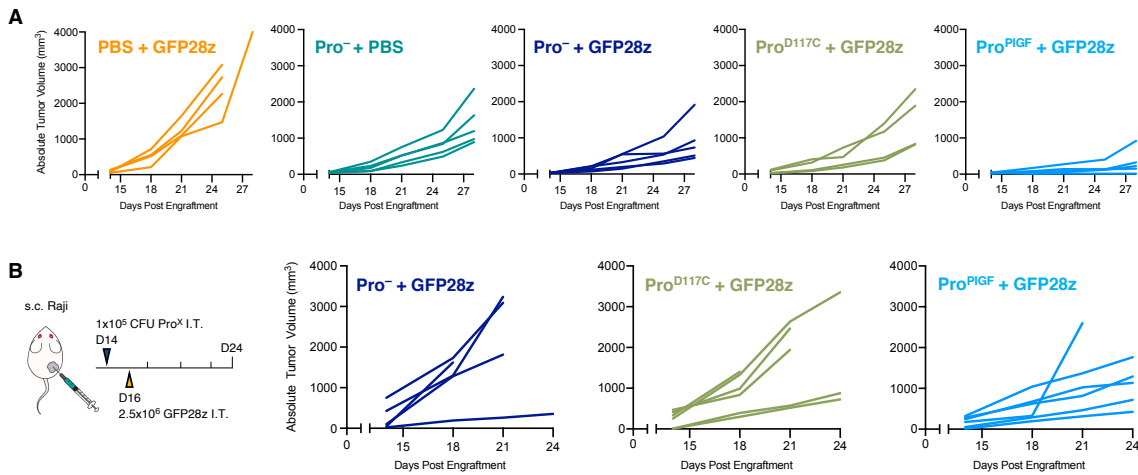
(A) PIGF SA variant is expressed from an Axe/Txe stabilized, high copy number plasmid under a constitutive *tac* promoter. (B) PIGF production by EcN (Pro^{PIGF}) as measured by GFP-fluorescence on a Tecan plate-reader and shown relative to an empty EcN control (Pro⁻). Error bars represent s.d. of biological replicates. a.u. arbitrary units.



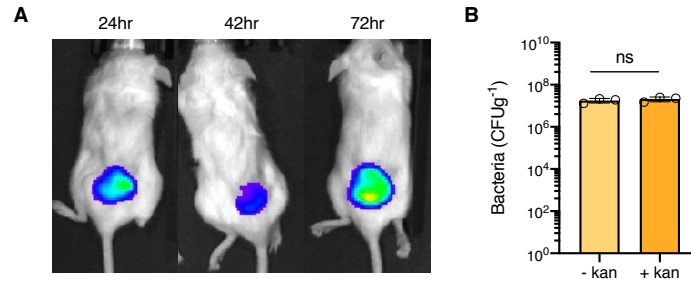
Supplementary Figure 3. GFP-directed CAR-T cells activate in response to soluble and collagen-bound sfGFP. (A) Quantification of flow cytometric analysis of intracellular staining for pro-inflammatory cytokines in response to 0.1 ng/ml soluble D117C, or collagen bound PIGF. **(B)** Quantification of cytokine production as measured in cell culture supernatants from GFP28z exposed to a PBS vehicle, D117C, or collagen-bound PIGF for 24hr. Error bars represent s.d. of biological replicates, **** $p < 0.0001$ 2way (A) or 1way (B) ANOVA, Holm-Sidak multiple comparison correction.



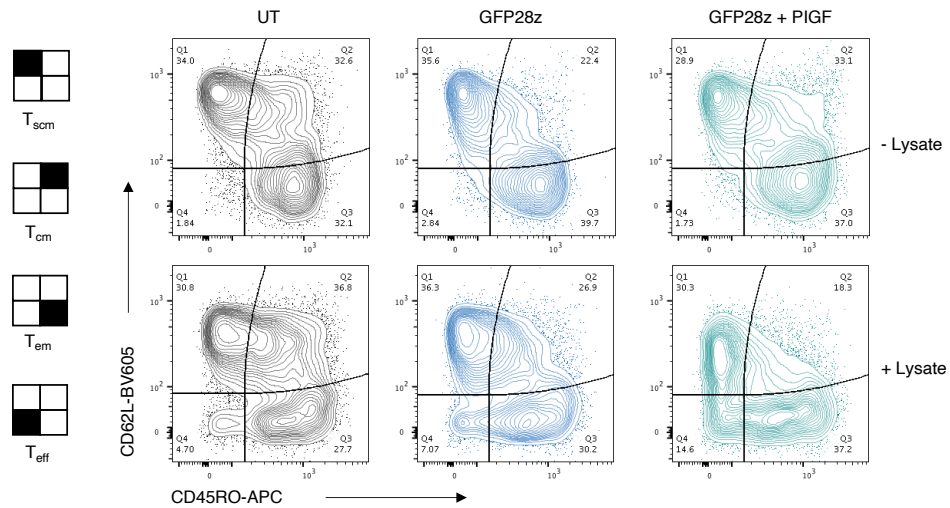
Supplementary Figure 4. GFP-directed CAR-T cells mediate killing of target cells in response to collagen-bound sfGFP. (A) Overnight killing assays against *fluc*⁺ HEK293T at a fixed effector to target (E:T) ratio of 1:3 and half log dilutions of collagen-bound PIGF from 50-1.5 ng/ml. **(B)** Overnight killing assays against *fluc*⁺ HUH7, MDA-MB-468, or MCF7 target cells at defined effector to target (E:T). CAR-T cells were co-cultured with target cells on collagen coated plates +/- 0.1 ug/ml sfGFP-PIGF for 20 hours before lysis and addition of luciferin. Luminescence (RLU) was detected with a Tecan plate reader within 20 minutes of lysis. Specific lysis (%) was determined by normalizing RLU to co-cultures with untransduced T cells. Error bars represent s.d. of biological replicates, *** p<0.001, **** p<0.0001 1way (A) or 2way ANOVA (B), Holm-Sidak multiple comparison correction.



Supplementary Figure 5. Individual growth trajectories of human tumors treated with the ProCAR system. (A) Nalm6 tumors were established in NSG mice and treated as in Fig. 3A. Individual tumor trajectories are shown. (B) Raji lymphoma cells (5×10^5) were implanted subcutaneously into the hind flank of NSG mice. When tumor volumes reached $\sim 100 \text{ mm}^3$, mice were intratumorally (I.T.) injected with 1×10^5 CFU of Pro⁻, Pro^{D117C}, or Pro^{PIGF} strains. 2.5×10^6 GFP28z ProCAR-T cells were then I.T. delivered 48-hours post bacterial injection. Tumor growth was monitored by caliper measurements every 3-4 days, individual tumor trajectories are shown.

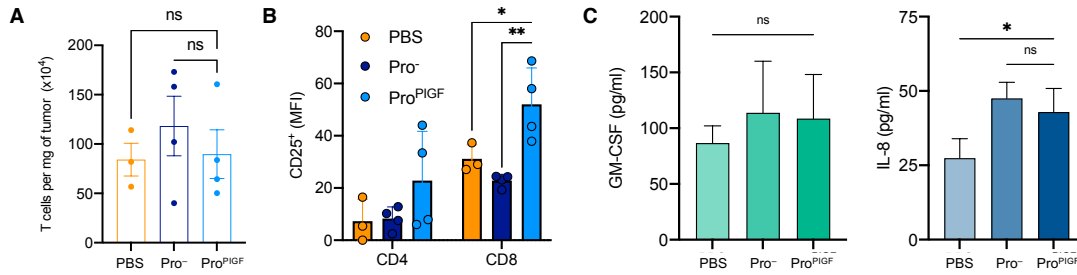


Supplementary Figure 6. Probiotic EcN remains localized to tumors in immunocompromised NSG mice. (A) IVIS images showing bioluminescent Pro⁻ populations over time following intratumoral injection of Raji tumors subcutaneously established as in Fig. S5B. **(B)** At day 14 post treatment, Pro^{PIGF}-treated tumors were homogenized and plated on LB agar plates containing the appropriate antibiotics (+/- kanamycin) for bacteria colony quantification. Error bars represent s.d. of biological replicates, student's t test; ns, not significant



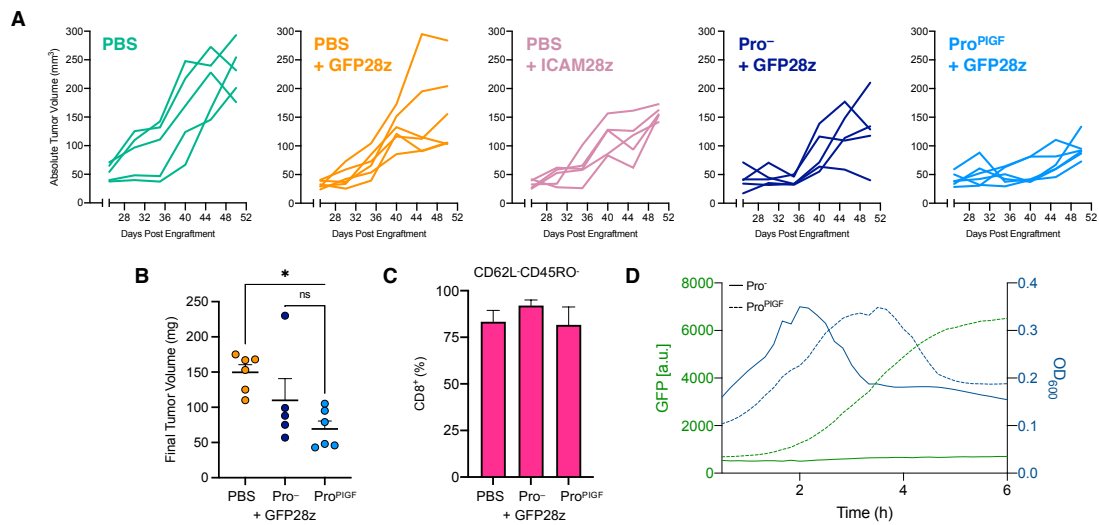
Supplementary Figure 7. *E. Coli* Nissle (EcN) lysate drives T cell effector phenotype.

Representative flow cytometry contour plots assessing the phenotype of T-cells following stimulation with either media alone or EcN lysate +/- 0.1 ug/ml PIGF for 48hr. CD8⁺ T cell populations were stained for CD45RO and CD62L expression to determine effector T cell differentiation, from stem cell memory (T_{scm}) CD62L⁺CD45RO⁻, central memory (T_{cm}) CD62L⁺CD45RO⁺, effector memory (T_{em}) CD62L⁻CD45RO⁺, and terminal effector (T_{eff}) CD62L⁻CD45RO⁻.

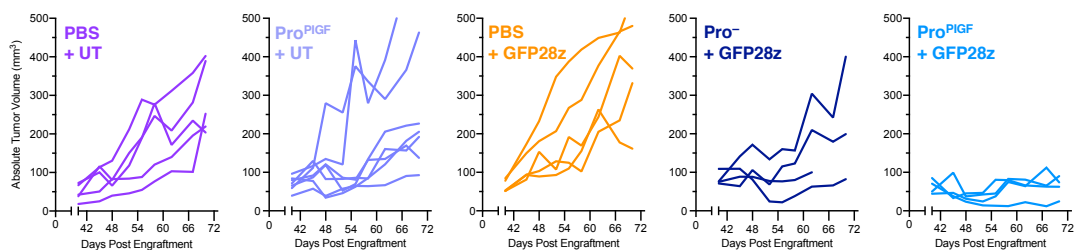


Supplementary Figure 8. Characterizing the effect of EcN strains on GFP28z cells *in vivo*.

Nalm6 cells (5×10^5) were implanted subcutaneously into the hind flank of NSG mice. When tumor volumes reached $\sim 100 \text{mm}^3$, mice were I.T. injected with either PBS or 1×10^5 CFU of Pro^{PIGF} or Pro⁻. On day 2 post Pro^X injection, all groups received an I.T. injection of 2.5×10^6 GFP28z ProCAR-T cells. Tumors were harvested and homogenized on day 4 for analysis. **(A)** Absolute counts of hCD45⁺CD3⁺ cells per mg of tumor. **(B)** Flow cytometric quantification of CD25 surface expression on intratumoral hCD45⁺CD3⁺ CD8⁺ or CD4⁺ cells from each treatment group. **(C)** Quantification of cytokine levels from tumor homogenates. Error bars represent s.d. of biological replicates, * p < 0.05, ** p < 0.01 1way (A) or 2way ANOVA (B), Holm-Sidak multiple comparison correction.



Supplementary Figure 9. Characterization of T cell exhaustion in a triple negative breast cancer (TNBC) model. (A) Subcutaneous MDA-MB-468 tumors were established in NSG mice prior to I.T. injection with PBS, Pro⁻, or Pro^{PIGF} on day 26 post tumor engraftment. On day 28, mice received a single I.T. injection of either PBS, 2.5x10⁶ GFP28z ProCAR-T cells, or 2.5x10⁶ ICAM1-specific CAR-T cells (1CAM28z), as in Fig. 5A. Tumor growth was monitored by caliper measurements every 3-4 days, individual tumor trajectories are shown. (B) On day 55 tumors were taken from mice treated with PBS, Pro⁻, or Pro^{PIGF} strains in combination with GFP28z and weighed *ex vivo*. (C) Frequency of intratumoral hCD45⁺CD3⁺CD8⁺ cells displaying a terminally differentiated effector phenotype (T_{eff}, CD62L⁻CD45RO⁻). (D) Pro^X strains were isolated from day 55 tumor homogenates and grown overnight on the plate reader to measure OD₆₀₀ and GFP fluorescence intensity. Error bars represent s.d. of biological replicates, * p<0.05 1way ANOVA; ns, not significant



Supplementary Figure 10. Individual growth trajectories of MDA-MB-468 subcutaneous TNBC tumors treated with the ProCAR system. Subcutaneous MDA-MB-468 tumors were established in NSG mice prior to I.T. injection with PBS, Pro⁻, or Pro^{PIGF} on day 40 post tumor engraftment. On day 44 Mice received an initial I.T. injection of 2.5×10^6 untransduced (UT), or 2.5×10^6 GFP28z ProCAR-T cells, followed by a second I.T. dose of UT or ProCAR-T cells 14 days later (day 58), as in Fig. 5F. Tumor growth was monitored by caliper measurements every 3-4 days, individual tumor trajectories are shown.