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Do Not Post

Decoy10 or Decoy10 + IL-2 Induce

Activation Markers in CD8 T Cells

CD69 Expression

Treatment

CD226 Expression

in CD8+ T Cells

Treatment

CICON 2025 Abstract #203 (Friday Sept. 12 Session B 12:00 pm - 3:00 pm)

Background

Activation of immune receptors, such as Toll-like (TLR), NOD-like (NLR) and Stimulator of Interferon Genes (STING) is required for efficient innate and adaptive immunity. Gram-negative bacteria (G-NB) contain multiple TLR, NOD and STING agonists. Potential utility of G-NB for cancer immunotherapy is supported by observations of tumor regression in the setting of infection and Coley's Toxins. Coley reported that intravenous (i.v.) administration was likely most effective but produced toxicity. The discovery of TLRs and their agonists, particularly the potent TLR4 agonist lipopolysaccharide (LPS)-endotoxin, comprising ~75% of the outer membrane of G-NB, suggests that it may be both a critical active ingredient and responsible for dose-limiting toxicity of i.v. G-NB. LPS analogues are also clinically validated vaccine adjuvants

We have produced killed, intact bacteria products from nonpathogenic G-NB with ~95% reduction of LPS activity. One product, Decoy10, contained TLR2,4,8,9, NOD2 and STING agonist activity and exhibited reduced i.v. toxicity in mice and rabbits relative to unprocessed cells. Decoy10 and closely related Decoy20 produced single agent activity or combination-mediated durable regressions, with immunological memory, in mice with syngeneic or human colorectal, hepatocellular, pancreatic carcinomas or non-Hodgkin's lymphoma. Regressions were observed in combination with chemotherapy, a non-steroidal anti-inflammatory drug, anti-PD-1, or rituximab, were associated with induction of 18-26 plasma cytokines/chemokines, activation of innate and adaptive immune pathways in tumors, and were dependent on NK, CD4+ and CD8+ T

Preliminary clinical results demonstrating rapid clearance of Decoy20 with transient induction of >50 plasma cytokines and chemokines (2-4) are supportive of a "pulse-prime" mechanism, whereby Decoy bacteria produce transient, but broad innate and adaptive immune activation (NCT05651022).

Results Summary

We have analyzed immune activation by Decoy bacteria in vitro using human peripheral blood mononuclear cells (PBMCs). Decoy10 induced immature to mature M1 macrophage maturation, monocyte-derived dendritic cell maturation, Th1 CD4⁺ T cell polarization and inhibited Treg production. Decoy10 induced activation markers in Natural Killer (NK), Natural Killer T (NKT), NKT-Like, CD4⁺ T, CD8⁺ T and CD4⁺CD8⁺ double positive T cells, and stimulated production of gamma-delta ($\gamma\delta$) T cells. Some results were dependent on or enhanced by Decoy10 combination with IL-2. Finally, Decoy10 enhanced PBMC-mediated tumor cell killing. The results demonstrate that Decoy bacteria stimulate maturation, polarization or activation of cellular mediators of innate and adaptive anti-tumor immune responses, inhibit an immune-suppressive mechanism, and enhance immune cell killing of tumor cells.

Methods

Decoy10 was prepared as described (1) with final assay concentrations representing per mL values. Experiments were carried out at HD Biosciences (San Diego, CA) in triplicate or duplicate when PBMCs from three donors were tested (all 96-well format) using complete (GlutaMAX) RPMI medium (supplemented with L-glutamine, sodium pyruvate, non-essential amino acids, Hepes, penicillin/streptomycin, and β-2 ME) with fetal bovine serum (human serum for the macrophage assay). Frozen human PBMCs were obtained from StemExpress (Placerville, CA) or AllCells (Alameda, CA), or isolated from whole human blood purchased from the San Diego Blood Bank using LymphoPrep/SepMate from StemCell Technologies. Flow cytometry studies were carried out with an Attune Flow Cytometer from ThermoFisher.

CD16-CD14+ monocytes were isolated from PBMCs with the StemCell 19359 EasySep Kit. M1 macrophage maturation was assessed with immature M1 macrophages produced by incubating 5x10⁴/well monocytes for 4 days with 50 ng/mL GM-CSF. After centrifugation-based washing the cells were incubated with Decoy10 or a positive control (50 ng/mL IFN-y + IL-6, 20 ng/mL LPS) for 3 days followed by analysis of surface CD80 and intracellular iNOS.

Dendritic cell maturation was assessed with immature DCs by incubating 1.25x10⁵/well CD16-CD14+ monocytes for 2 days with 500 ng/mL GM-CSF + IL-4. After centrifugation-based washing, the immature DCs were incubated for 3 days in fresh medium containing GM-CSF + IL-4 and maturation cytokines (2,000 ng/mL PGE2, 10 ng/mL IL-6, IL-1β, TNF-α; positive control), or GM-CSF + IL-4 and Decoy10 followed by analysis of surface CD80, CD83 and CD86. Greater than 90% of double positive cells were also triple positive (not shown).

CD4 Th1 polarization was assessed after 6-day PBMC incubation (2.5x10⁵/well) without or with 1 μg/mL anti-CD28, 1 μg/mL IL-4, 20 U/mL IL-2 and 20 ng/mL IL-12p70 (positive control), or with Decoy10 (± 20 U/mL IL-2 + 1 µg/mL anti-CD28), followed by analysis of surface CD4 and intracellular IFN-y. The effect of Decoy10 on CD4 Treg polarization was assessed by incubating PBMCs as above with 1 µg/mL anti-CD28, 100 U/mL IL-2 and 5 ng/mL TGF-β ±Decoy10 followed by analysis of CD4, CD25, and

FoxP3 expression. Activation of NK (CD56+, CD4-, CD8-), NKT (CD56+, CD4+, CD8-), NKT-Like (CD56+, CD4-, CD8+), CD4+ T (CD56-, CD4+, CD8-), CD8+ T (CD56-, CD4-, CD8+) and CD4+/CD8+ double positive T (CD56-, CD4+, CD8+) cells was assessed by incubating PBMCs (2.5x10⁵/well) for two days with 100 U/mL IL-2 (positive control for NK cells and all T cells), or with 2.5 µg/mL anti-CD28 in wells precoated overnight with 100 µL of 5 µg/mL OKT3 anti-CD3 antibody (additional positive control for all T cells) ± 1x10⁷ Decoy10/mL, or with Decoy10 alone. Cell activation was assessed by analysis of surface CD69, CD226, and intracellular IFN-y, Perforin, and Granzyme B. Immune activation studies were carried out with three independent PBMC donors (Red, Green, Blue) in duplicate. Results for one donor were not included for NKT-Like cells due to significant discrepancies between duplicates.

Induction of CD3+CD4-CD8- y9δ2 T cells was carried out by incubating PBMCs at 3x10⁵ per well for 6 days with 1-Hydroxy-2methyl-2-butenyl 4-pyrophosphate (HMBPP, Sigma #95098) or Decoy10 followed by staining for CD3, CD4, CD8, V γ 9, and V δ 2. Killing of human MDA-MB-231 breast carcinoma cells (ATCC) by Decoy10 + PBMCs was assessed with IncuCyteTM NucLightTM Red Lentivirus-transfected tumor cells (reagents from Essen Biosciences, Ann Arbor, Michigan). Tumor cells were plated at 2x10⁴/well and incubated overnight in complete RPMI medium. PBMCs (1x10⁴ per well) and/or Decoy10 (3x10⁶ per mL) were added, and the incubation was continued for 7 days followed by image analysis using the IncuCyteTM system (Sartorius).

Statistical significance for flow cytometry experiments was analyzed with GraphPad Prism software using a standard, unpaired t-test with comparison to the appropriate no addition, vehicle, or single agent control, with results indicated by standard p≤0.05-0.0001 (*, **, ***, ****) notation.

Discussion

Induction of maturation, polarization or activation of multiple innate and adaptive immune cell types, and enhancement of PBMC-mediated tumor cell killing by Decoy10 is consistent with our pre-clinical in vivo data demonstrating that tumor eradication by Decoy10 combination therapy is dependent on NK, CD4 T and CD8 T cells, and is associated with innate and adaptive immune pathway activation (1).

The data are also consistent with our preliminary Phase 1 clinical trial results, where we have observed broad plasma cytokine and chemokine-associated immune activation (2-4).

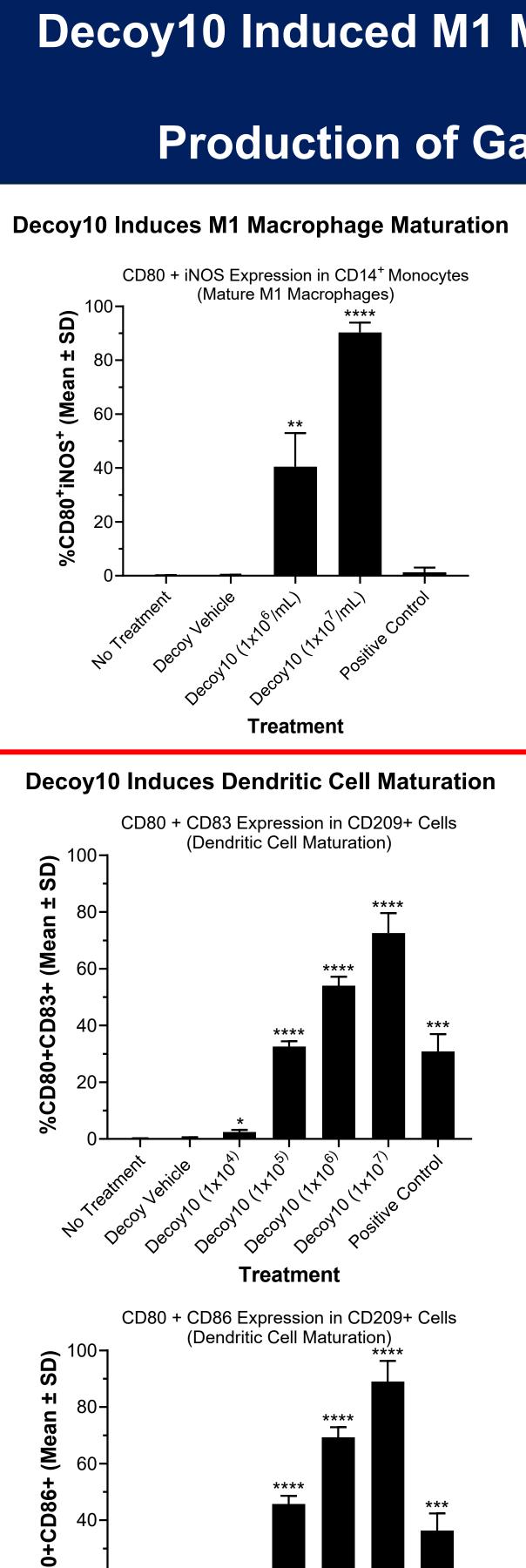
The current data also suggest that Decoy bacteria may produce additive or synergistic effects with IL-2 in some settings.

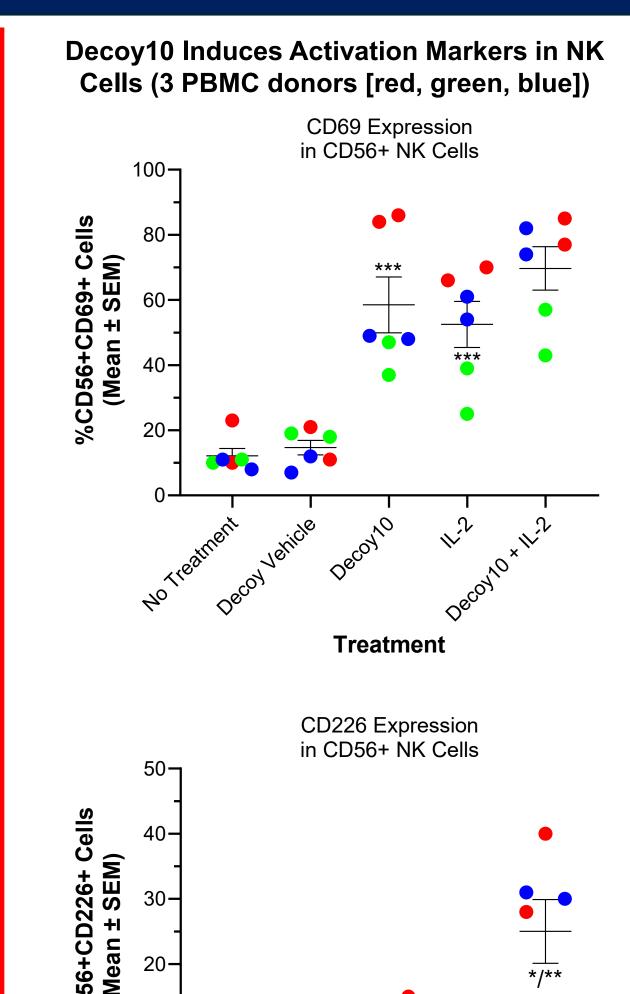
References

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Acknowledgement: We thank Dr. Ramesh Halder and Dr. Mengling Liu of HD Biosciences, a WuXi AppTec company, for carrying out the experiments reported in this poster.

Decoy10 Induced M1 Macrophage Maturation, DC Maturation, CD4⁺ Th1 Polarization, Treg Inhibition, Activation Markers in NK, NKT, CD4⁺ and CD8⁺ T Cells, Production of Gamma-Delta T Cells and Enhanced PBMC-Mediated Killing of Tumor Cells

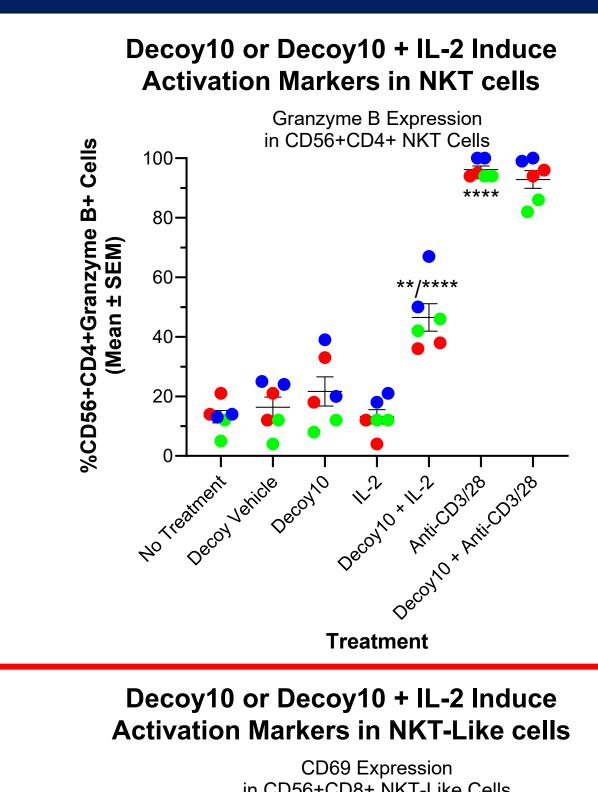


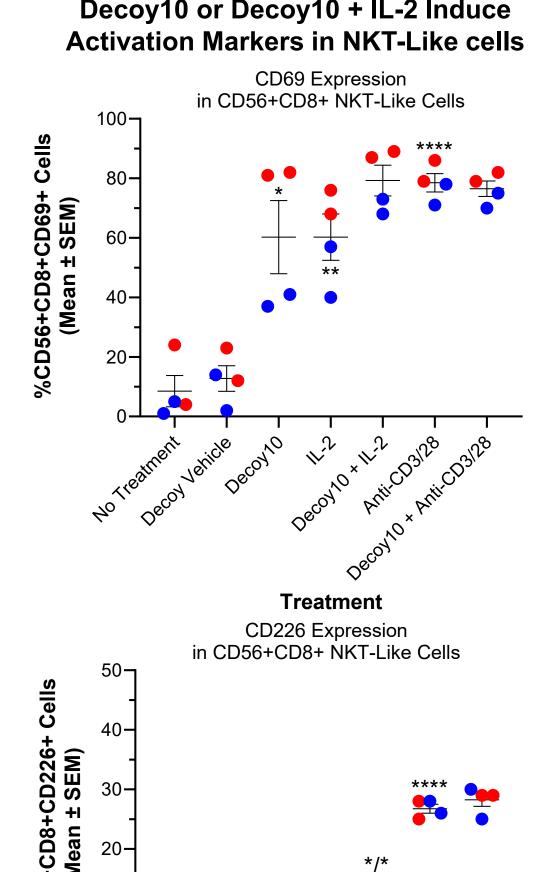


Treatment

Perforin Expression

in CD56+ NK Cells



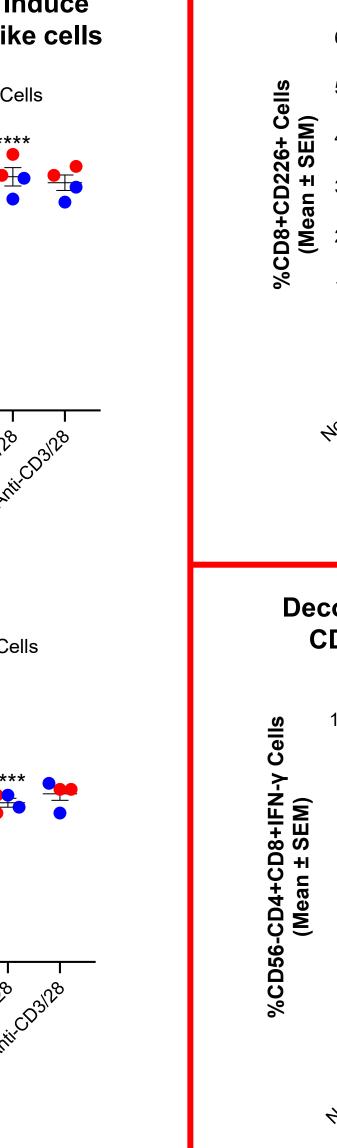


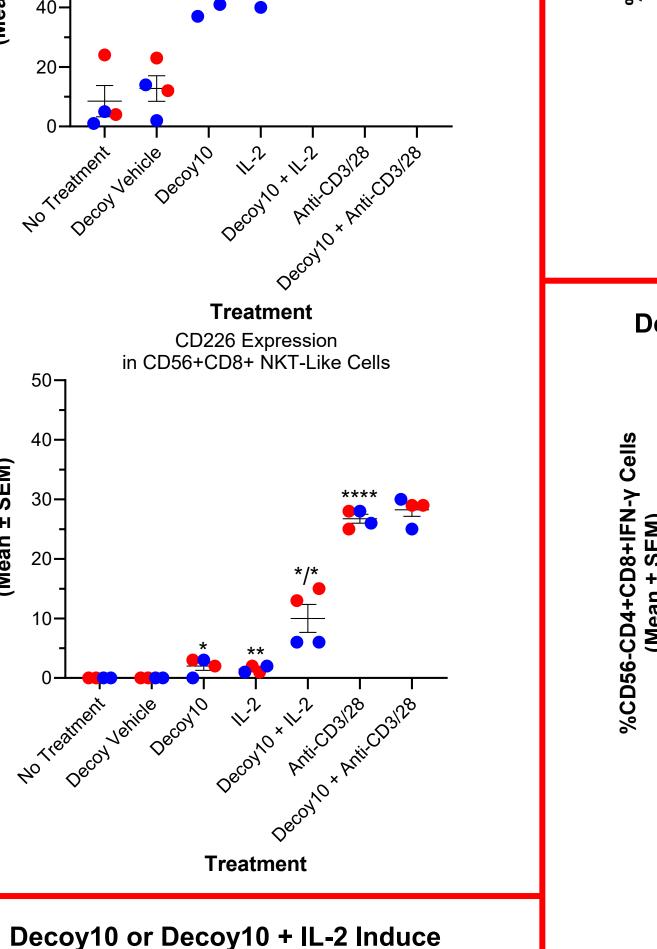
Activation Markers in CD4 T Cells

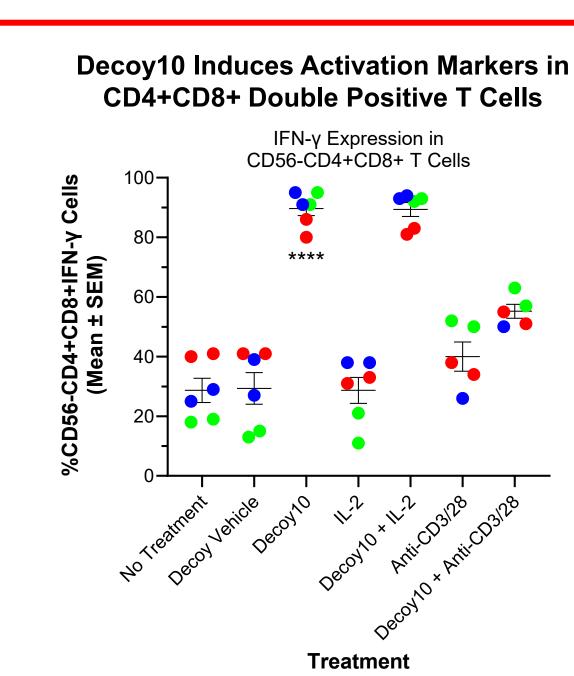
CD4+CD69+ Cell (Mean ± SEM)

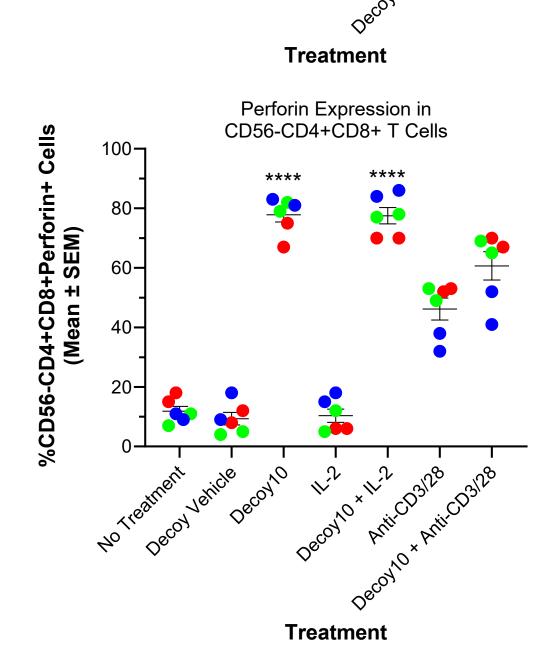
CD69 Expression

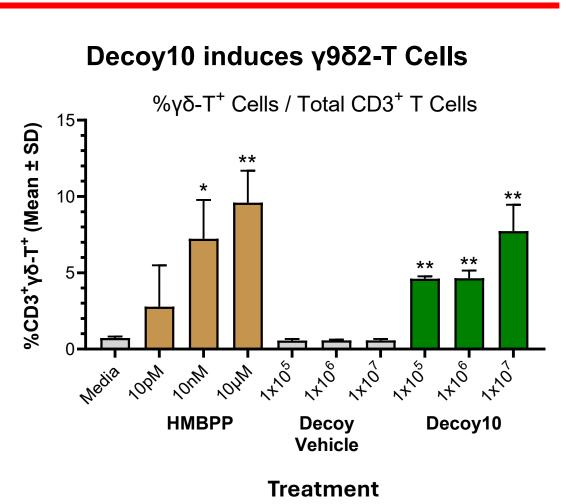
in CD4+ T Cells

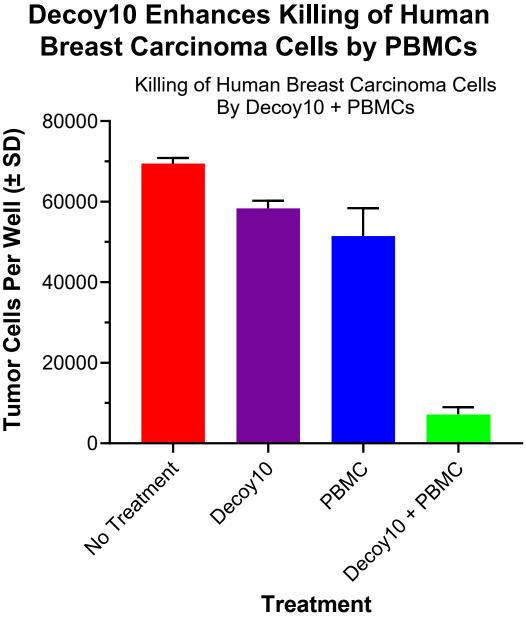


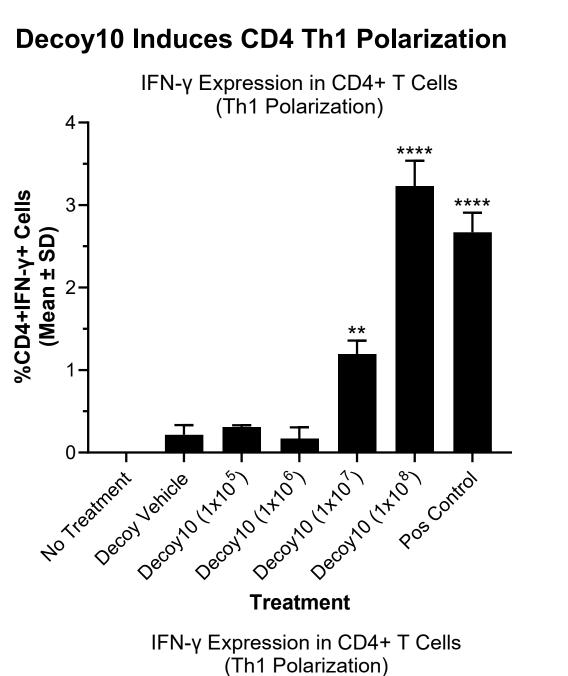




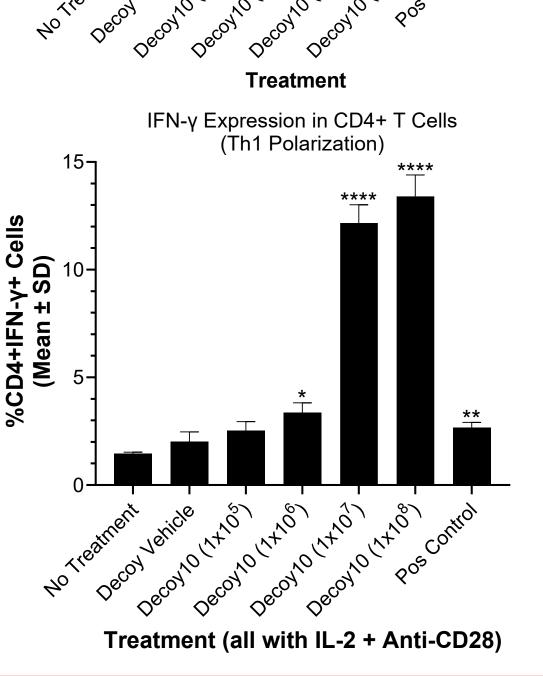


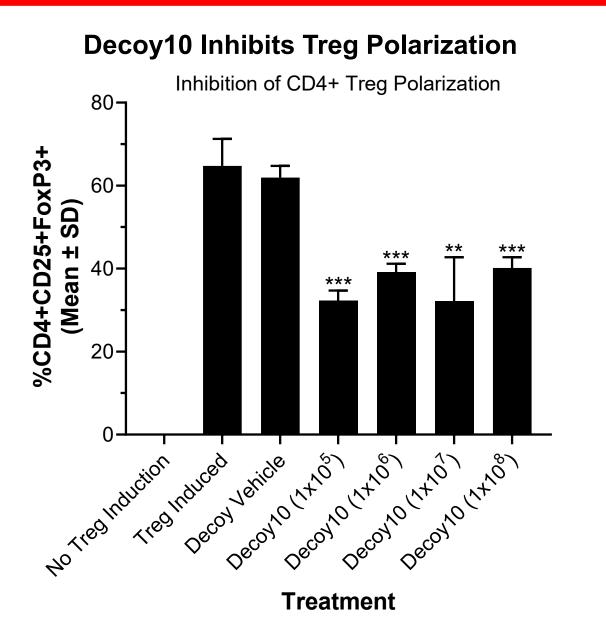


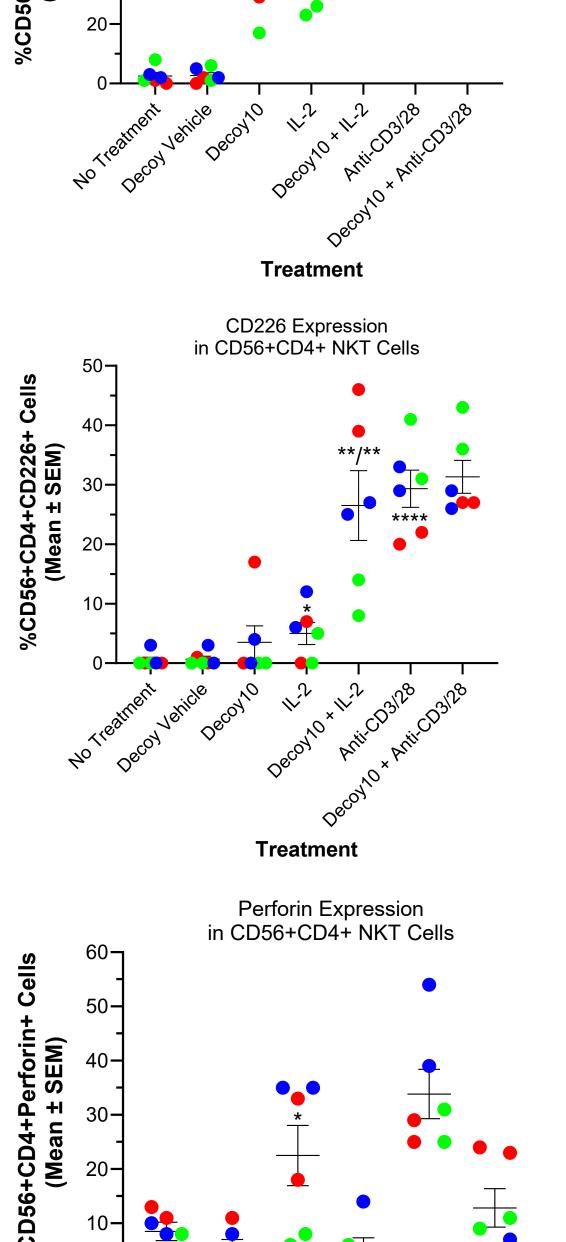


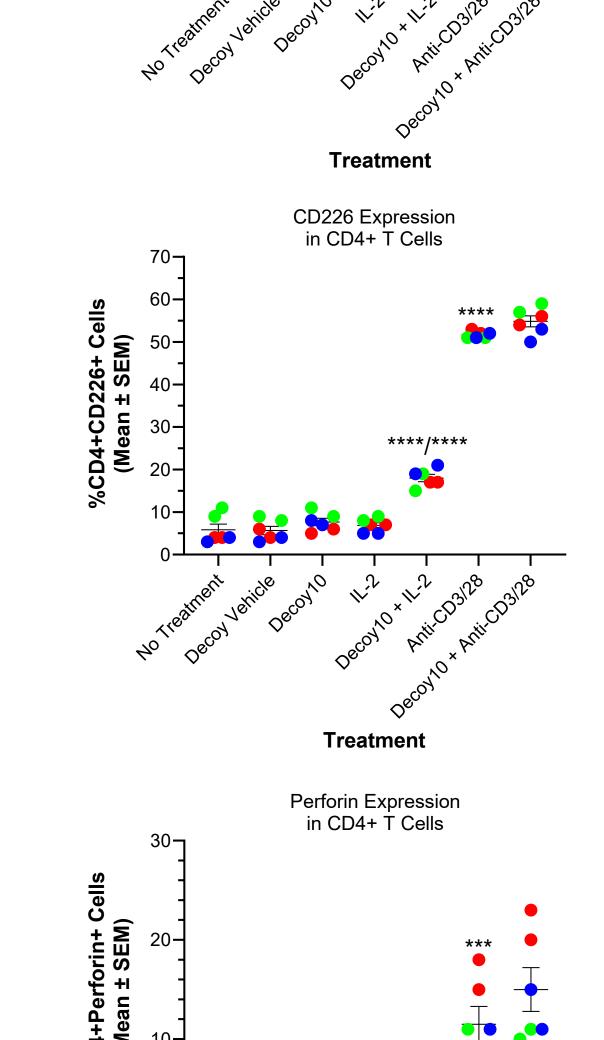


Treatment

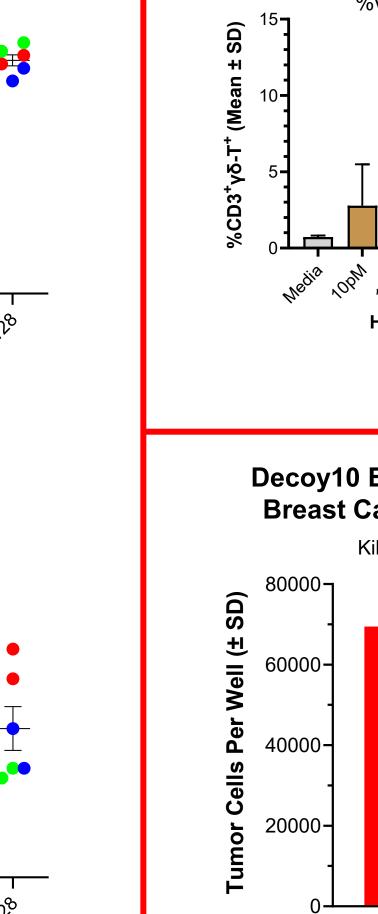


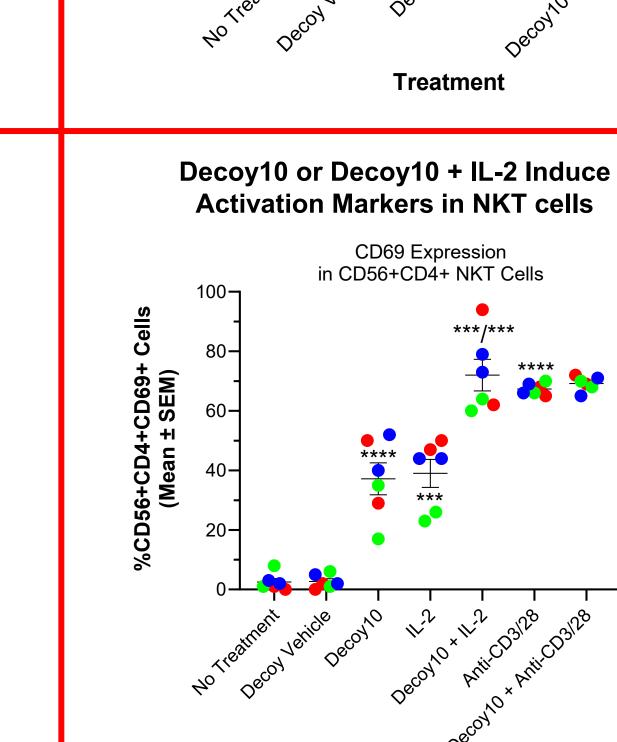






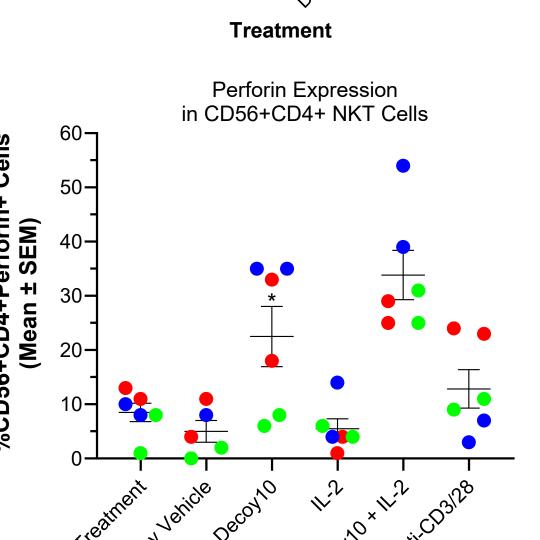
Treatment





orin+ (SEM)

CD56+F (Mea



Treatment