## TREM1 Agonist Antibody PY159 Promotes Myeloid Cell Reprogramming and Unleashes Anti-tumor Immunity

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Figure 5. (A) Single-cell RNAseq of CD45+ cells from a human ovarian cancer UMAP plots depict distinct latilocula schools (2.5) as impacts

TREM1 expression levels in individual leukocyte subsets are presented using violin plots (right). (B) TREM1 staining on TANs, mMDSCs, TAMs, cDC1, cDC2, and

pDCs by flow cytometry, in 30 dissociated human tumor samples. Tumor types

Background: Tumor-associated myeloid cells can impede productive anti-tumor immunity. One strategy for targeting immunosuppression is myeloid reprogramming, which drives immunosuppressive myeloid cells to acquire an immunostimulatory phenotype. Triggering receptor expressed on myeloid cells-1 (TREM1) is an immunoglobulin superfamily cell surface receptor expressed on neutrophils, subsets of monocytes and tissue macrophages. TREM1 associates with the DAP12 adaptor protein and induces proinflammatory signaling, amplifies inpate immune responses and is implicated in the development of acute and chronic inflammatory diseases. TREM1 is also enriched in tumors, specifically on tumor-associated myeloid cells. To investigate the potential of TREM1 modulation as an anti-cancer therapeutic strategy, we developed PY159, an afucosylated humanized anti-TREM1 monoclonal antibody, and characterized it in the pre-clinical assays described below.

Materials and Methods: An FcyR binding ELISA and a Jurkat TREM1/DAP12 NFAT-luciferase reporter cell line were used to assess PY159 binding to human FcvRs and TREM1 signaling. respectively. PY159 responses in human whole blood in vitro were evaluated by flow cytometry. transcriptional analysis of sorted leukocyte subsets, and measurement of secreted cytokines/chemokines by MSD. A Transwell system was used to evaluate PY159 effects on neutrophil chemotaxis. TREM1 expression in human tumors was validated by scRNAseg and flow cytometry. Anti-tumor efficacy of a surrogate anti-mouse TREM1 antibody, PY159m, was evaluated using syngeneic mouse tumor models, either as a single agent or in combination with anti-PD-1

## **TREM1** Receptor



PY159 is an Afucosylated Anti-human TREM1 Antibody With Enhanced FcγR Binding



Figure 1. (A) PY159 binding to cell surface expressed human TREM1 (hTREM1) was evaluated using HEK 293 cells recombinantly expressing human TREM1 and DAP12. Antibody binding was detected by flow cytometry after staining with an APC-labeled secondary anti-human IgG antibody. (B) PY159 and PY159F (fully fucosylated version of PY159) were tested for binding to immobilized recombinant human FcyRilla by ELISA. Antibody binding was detected using a secondary HRP-conjugated goat anti-human F(ab')2 antibody, followed by the measurement of absorbance (optical density) at 450 nm.





Tumor #1

Tumor #2

 Tumor #3 Tumor #4

## **Results & Conclusions**

Afucosylation of PY159 increased its binding affinity for EcvR and its ability to activate TREM1/DAP12 signaling In human blood assays, PY159 treatment upregulated monocyte activation markers, promoted neutrophil chemotaxis, and induced proinflammatory cytokines and chemokines, which was dependent on PY159 afucosylation. In human tumors, TREM1 was detected on tumor-associated neutrophils, tumor-associated macrophages, and monocytic myeloid-derived suppressive cells. PY159 induced proinflammatory cytokines and chemokines in dissociated human tumors in vitro, demonstrating that PY159 can reprogram tumor-associated myeloid cells. A surrogate anti-mouse TREM1 antibody, PY159m, exhibited anti-tumor efficacy in several syngeneic mouse tumor models, both as single agent and in combination with anti-PD-1. These results show that PY159 reprograms myeloid cells and unleashes anti-tumor immunity. PY159 safety and efficacy are currently being evaluated in first-in-human clinical trial (NCT04682431) involving patients resistant and refractory to standard of care therapies.



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CXCL-10 CCL3 CCL4

Donor 4

foure 4. (A) Whole blood from six healthy hu Figure 4. (A) Whole blood from six healthy human donors was treated for 24 hrs with 25 nM PY159 or isotype control. Plasma cytokines and chemokines were measured using the MSD platform. The graph represents PY159-induced cytokines as fold increases relative to the effects of isotype controls in individual donors. (B) Whole blood from two donors was treated with 10µg/mL of PY159, fully fucosylated PY159F, or corresponding isotype controls. Cytokine induction by PY159 or PY159F is presented as fold BV150E or ( increase rela tive to the effects of corresponding sotype controls in individual donors

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