

Introduction

Sapience Therapeutics, Inc. is a clinical-stage biotechnology company focused on the discovery and development of peptide therapeutics to address oncogenic and immune dysregulation that drive cancer. Our first-in-class β -catenin antagonist, ST316, targets the interaction between β -catenin and its co-activator BCL9, a complex that drives oncogene expression in cancers where aberrant Wnt/ β -catenin pathway signaling is observed. Sapience has received clearance from U.S. Food and Drug Administration (FDA) to proceed with a ST316 Phase 1-2 clinical trial for the treatment of solid tumors.

Abstract

The transcription factor β -catenin is a key player in many cellular processes, including stem cell renewal, cellular homeostasis and inflammation. Deregulation of β -catenin occurs frequently in several cancer types by mutation or overexpression of the β -catenin gene or mutation of negative regulators such as Adenomatous Polyposis Coli (APC). ST316 is a novel peptide antagonist that specifically interferes with the association of β -catenin with BCL9, disrupting β -catenin nuclear localization and attenuating target gene expression. In an orthotopic triple-negative breast cancer (TNBC) model, ST316 demonstrates potent anti-tumor activity, resulting in 84% tumor growth inhibition (TGI) ($p < 0.001$ vs. vehicle control). In addition to its cell-autonomous mechanisms, deregulated Wnt/ β -catenin signaling can promote tumorigenesis by impacting the tumor immune microenvironment (TIME). Reprogramming of immunosuppressive M2-like Tumor Associated Macrophages (TAMs) toward an immune-promoting program (M1-like) is an attractive cancer immunotherapeutic strategy. Here we explored ST316 potential for macrophage repolarization toward the M1-like phenotype, activation of cytotoxic T-cells in macrophage/T-cell co-culture assays and cooperation of ST316 with anti-PD-1 to enhance anti-tumor activity in vivo. Initial studies demonstrate that ST316 reprograms human macrophages derived from Peripheral Blood Mononuclear Cells (hPBMCs) and subsequently committed to the M2-like identity, toward an M1-like phenotype. ST316 treatment dose-dependently suppressed expression of the M2 marker CD163 by flow cytometry and quantitative PCR, resulting in 100-fold increase in the M1/M2 ratio without substantial impact on cell viability. Importantly, activation markers associated with M1-state (CD80, CD86), and T-cell viability were not affected by ST316 at the concentrations used in these studies. Further, in co-cultures of M2 macrophages with T cells, ST316 exposure resulted in a three-fold increase in T-cell activation compared to untreated M2/T cell co-cultures, as measured by intracellular IFN- γ staining. Finally, in an orthotopic TNBC model in vivo, subtherapeutic ST316 enhanced the anti-tumor activity of anti-PD-1 by increasing the M1/M2 ratio, increasing the TGI (observed 85% vs. 51% with anti-PD-1 alone, $p < 0.01$; vs. -9% with subtherapeutic ST316 alone, $p < 0.001$). Overall, these results support the immuno-therapeutic potential of ST316 and extend the application range of ST316 to include Wnt-driven cancers with poor clinical response to immune checkpoint blockade and other immunotherapeutic agents.

Wnt/APC/ β -catenin in cancer and TIME

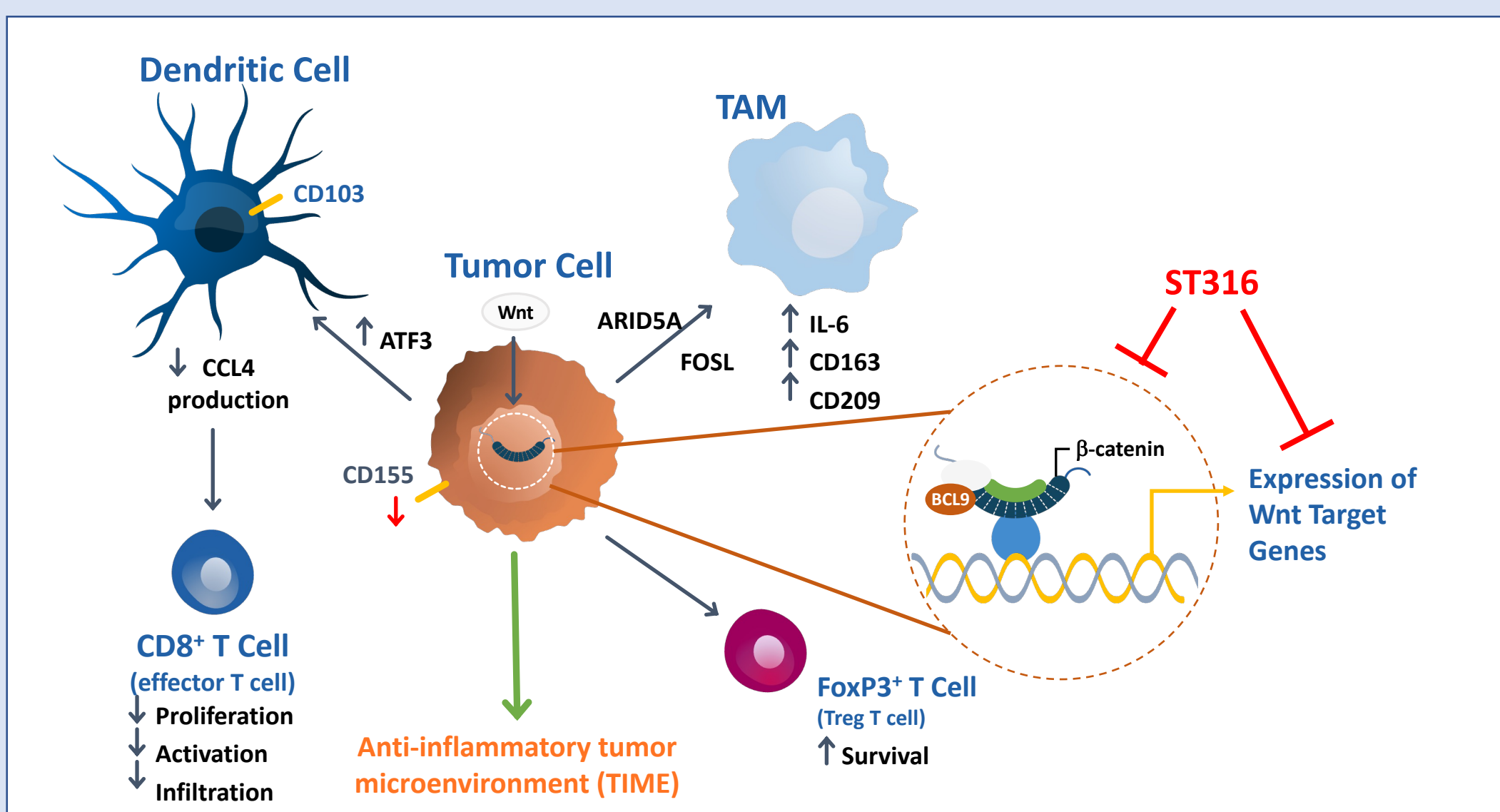


Figure 1: The Wnt/APC/ β -catenin axis is a master regulator of both oncogenesis and the tumor immune microenvironment (TIME). In cancer cells, this pathway is frequently activated by inactivating APC mutations or β -catenin mutations leading to its stabilization. In the TIME, Wnt signaling promotes an immunosuppressive program in Tumor Associated Macrophages (TAM) and Dendritic Cells (DC) leading to upregulation of M2 markers (CD163, DC-SIGN CD209, IL-6) and decrease of immunogenic chemokines (CCL4). This program results in suppression of cytotoxic CD8+ T cell activity and supports survival of immunosuppressive Treg populations. Additionally, Wnt can modulate surface expression of checkpoint molecules such as CD155. References: Pai et al., J of Hemat and Onc 2017; Sarode et al., Sci Adv 2020; Ruiz de Galarreta et al., Cancer Discov 2019; Feng et al., Sci. Adv. 2021).

Results

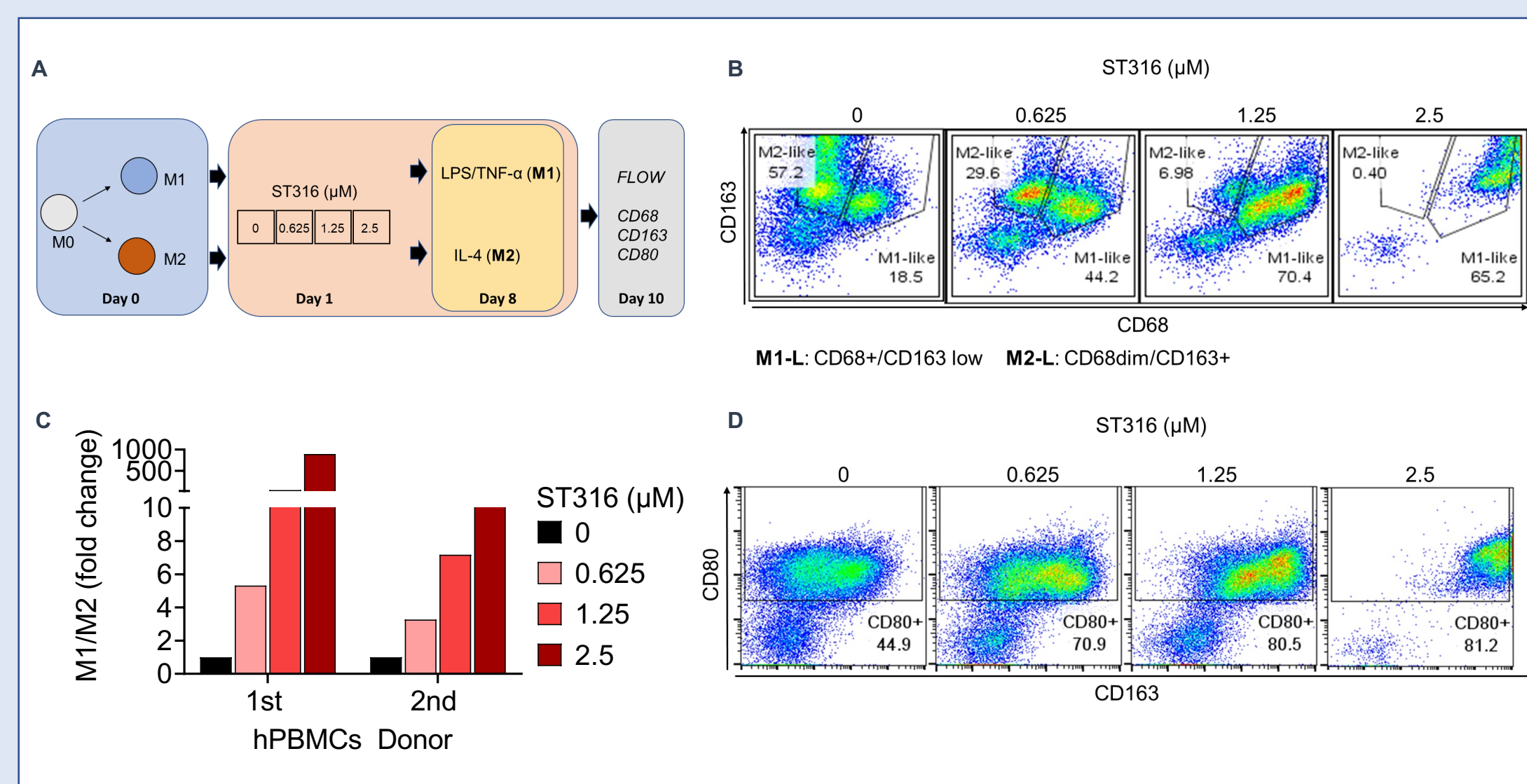


Figure 2: ST316 converts M2-like macrophages to M1-like phenotype. A) Experimental outline of M1 and M2 establishment from hPBMCs. B) Flow cytometry plots at the indicated ST316 concentrations of M1-like (CD68+CD163low) and M2-like (CD68dim/CD163+) populations in M2 conditions at Day 10. C) Normalized hPBMCs M1/M2 population ratios for two unrelated donors at the indicated ST316 concentrations. D) Flow Cytometry plots for the M1 Marker CD80 and CD163 in hPBMC-derived M2 Macrophages treated with the indicated ST316 concentrations

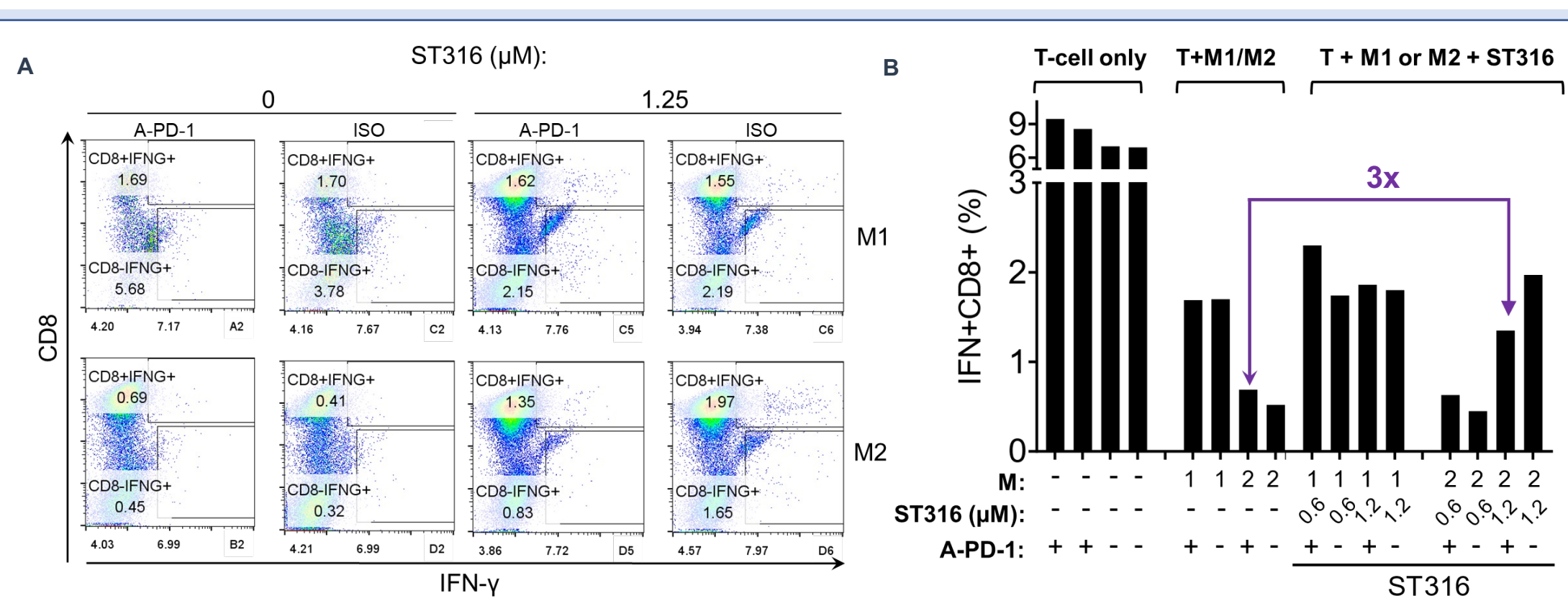


Figure 3: ST316 increases macrophage-induced CD8+ T cell activation in M2 suppressive conditions. A) CD8+ T cell were co-cultured for 3 days with M1 or M2 macrophage and left untreated (0) or treated with 0.625 or 1.25 μ M ST316 and anti-PD-1 (A-PD-1, 20 ng/mL) or isotype IgG4 Control (ISO). Flow cytometry plots for CD8 and intracellular IFN- γ . B) Frequency of IFN-gamma producing CD8+ cells at the indicated conditions. Arrows point to the comparison between M2 condition in the absence or presence of 1.25 μ M ST316, indicating a three-fold increase in IFN- γ (+) cells. T-cell only conditions were assessed in duplicate.

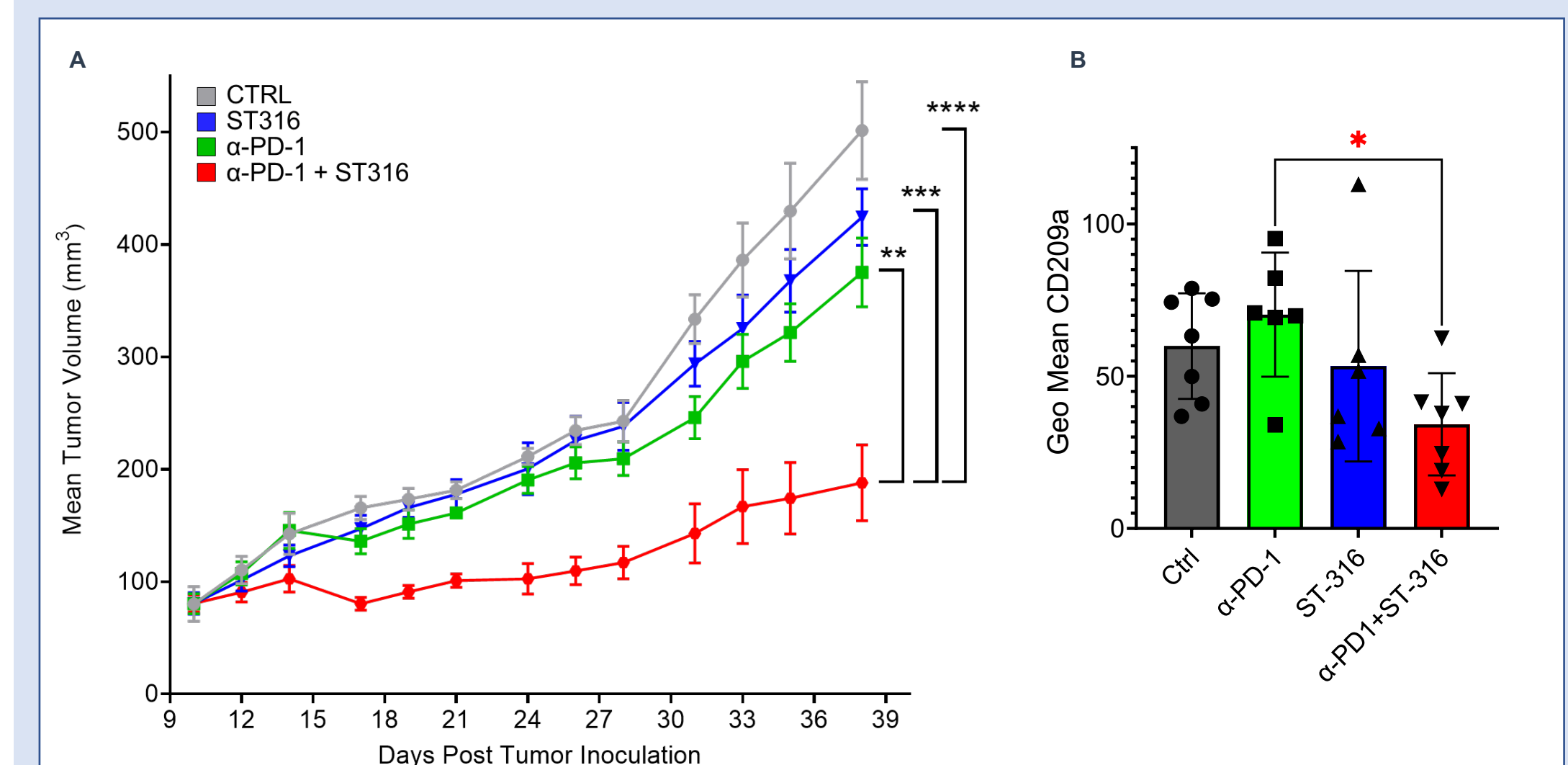


Figure 4: Subtherapeutic ST316 potentiates response to anti-PD-1 therapy in vivo in the orthotopic 4T1 Triple Negative Breast Cancer (TNBC) model. A) Balb/c mice harboring 4T1 tumors implanted into the 4th mammary fatpad were administered subtherapeutic ST316 (2mg/kg once weekly) and/or anti-PD-1 (BioXCell #BE0146, clone 29F.1A12; 12.5 mg/kg/wk). Mean Tumor volumes are shown for the indicated treatment cohorts (n=5/group). Error bars represent SEM. P-value statistics are one-way ANOVA for the indicated comparisons at Day 38 (n=5/group; **** $p < 0.001$; *** $p < 0.01$; ** $p < 0.05$). B) Geometric means for expression of the M2 marker CD209a in TAMs (gated as CD45⁺Gr1^{int}F4/80⁺) for the indicated cohorts (n=5/group; *, $p < 0.05$; one-way ANOVA).

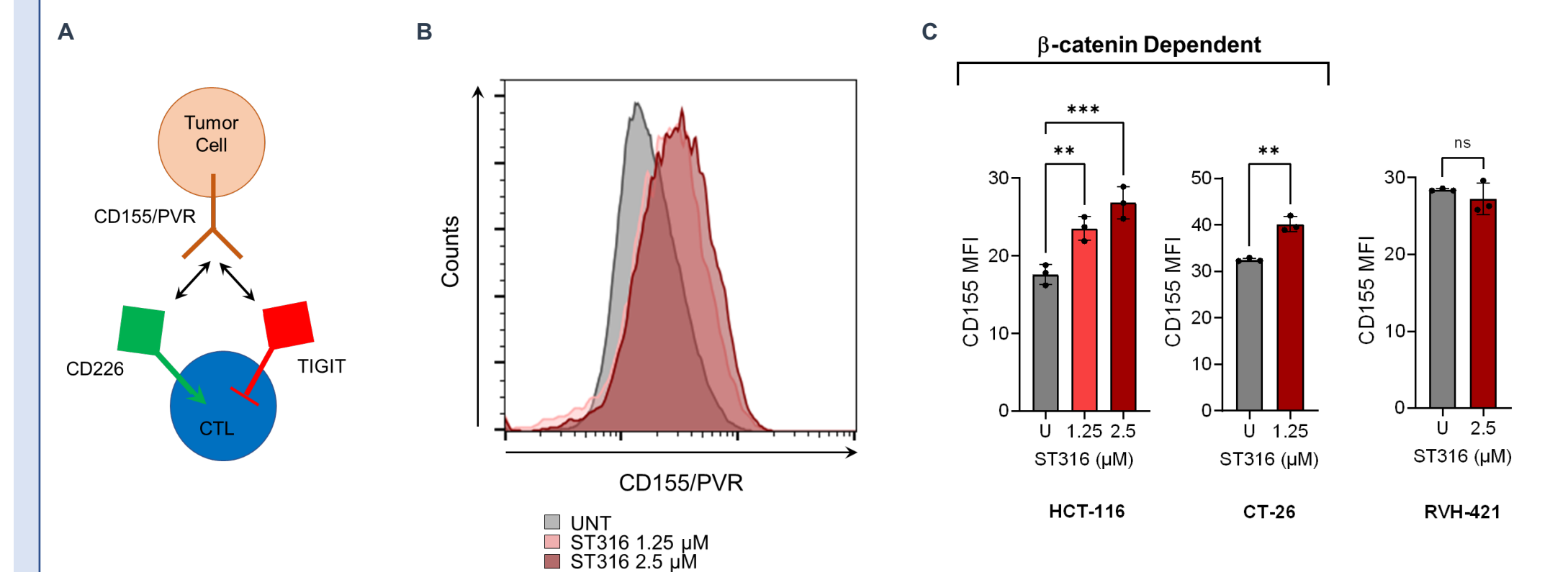


Figure 5: ST316 increases surface expression of checkpoint activator CD155/PVR in β -catenin-driven lines. A) Schematic representation of the CD155/TIGIT checkpoint. CD155/PVR mediates activation of cytotoxic T-cells (CTL) by binding CD226. TIGIT binds competitively CD155 and suppresses T-cell activation. B) Representative histograms (mode-normalized) for HCT116 cells untreated (dark gray) or treated with 1.25 μ M ST316 (orange) or 2.5 μ M ST316 (red) for 48hrs. C) Median Fluorescence Intensity of CD155 in the β -catenin-driven lines HCT116 and CT-26 untreated or treated at the indicated ST316 concentrations (Statistics represent t-test of the indicated pair; *** $p < 0.001$; ** $p < 0.01$; ns, not significant). The melanoma cell line RVH-421 is shown as control.

Conclusions

- ST316 induces a dose-dependent shift in hPBMC-derived M2 macrophage to the M1 identity and induces the M1 markers CD80 in M2 conditions.
- Exposure to ST316 increases CD8 T-cell activation in macrophage/T cell mixed cultures as measured by IFN- γ production by CD8+ T cells.
- Subtherapeutic ST316 enhances the activity of anti-PD-1 antibody in the syngeneic (immunocompetent) orthotopic 4T1 TNBC tumor model; Isolated TAM expressed decreased M2 marker DC-SIGN (CD209a).
- ST316 promotes surface expression of CD155/PVR in β -catenin dependent cell lines.
- These data support a model in which ST316 promotes a shift in the tumor microenvironment via multiple mechanisms, including driving macrophage polarization toward an immune-promoting phenotype, augmenting activity of cytotoxic T cells and increasing expression of checkpoint activators like CD155/PVR in cancer cells.