

Fc-silent Anti-TIGIT Antibodies Potentiate Anti-tumor Immunity Without Depleting Regulatory T Cells

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Background & Summary

- TIGIT is an inhibitory checkpoint receptor primarily expressed on T and NK cell subsets.
- TIGIT suppresses cell activation through intrinsic mechanisms and by outcompeting a key activating receptor, CD226/DNAM-1, for the shared ligand CD155/PVR¹⁻⁵.
- Analogous to the TCR and CD28, PD-1 can de-phosphorylate CD226 to suppress cell activation^{6,7}.
- To achieve maximal effector cell activation and functionality, it is hypothesized that simultaneous blockade of TIGIT and PD-1 is necessary, particularly to target cell subsets that co-express TIGIT and PD-1, namely pre-exhausted T cells (T_{pe}).
- The TIGIT field has largely focused on elucidating mechanisms of action related to blocking antibodies that retain Fc functionality. In mouse models, Fc-enabled anti-TIGIT can deplete regulatory T cells (T_{reg}), an activity associated with single-agent efficacy^{8,9}; however, whether such mechanisms will translate to humans is unclear.
- In contrast, Fc-silent anti-TIGIT can enhance efficacy of anti-PD-1 without T_{reg} depletion in mice¹⁰.

Here, we report pharmacology and mechanisms of action associated with Fc-silent anti-TIGIT antibodies relative to Fc-enabled counterparts. We found:

- Human and mouse tumor-infiltrating lymphocytes express TIGIT and CD226 on T_{reg} , CD4⁺ non- T_{reg} , and on CD8⁺ T cells with tumor-reactive or exhausted/dysfunctional phenotypes (Figure 2).
- In mice, in combination with anti-PD-1, Fc-silent anti-TIGIT enhances activation, differentiation, and effector function of tumor-specific CD8⁺ T cells in a manner dependent upon T cell egress from the tumor-draining lymph node (Figure 3).
- In vitro*, Fc-enabled human TIGIT-specific antibodies facilitated antibody-dependent cell-mediated cytotoxicity (ADCC) against TIGIT-expressing human T_{reg} , with preferential depletion of a Helios⁺ effector T_{reg} subset (Figure 4 & Figure 5).
- In cancer patients, significant and stable decreases in T_{reg} were measured in the peripheral blood of cancer patients treated with an Fc-enabled (AB308), but not an Fc-silent (domvanalimab), anti-TIGIT antibody (Figure 6).
- Cancer patients treated with domvanalimab in combination with anti-PD-1 antibody, zimberelimab, experienced partial responses while maintaining stable peripheral T_{reg} frequencies on treatment¹⁰.

Silencing the Fc domain of anti-TIGIT antibodies prevents depletion of peripheral T_{reg} , potentially critical for an optimal safety-efficacy profile in cancer patients.

Some images made with BioRender. Referenced DOIs: ¹10.1158/2326-6066.CIR-19-0877, ²10.1038/s41467-023-40755-3, ³10.4049/jimmunol.1003081, ⁴10.1172/JCI81187, ⁵10.1038/nr.1674, ⁶10.1016/j.immuni.2022.02.005, ⁷10.1126/sciimmunol.aat7061, ⁸10.1158/1535-7163.MCT-20-0464, ⁹10.3389/fimmu.2022.828319, ¹⁰10.4049/JIMMUNOL.208.SUPP.120.13; download at: <https://arcusbio.com/our-science/publications/>, ¹¹10.1146/annurev-immunol-042718-041717.

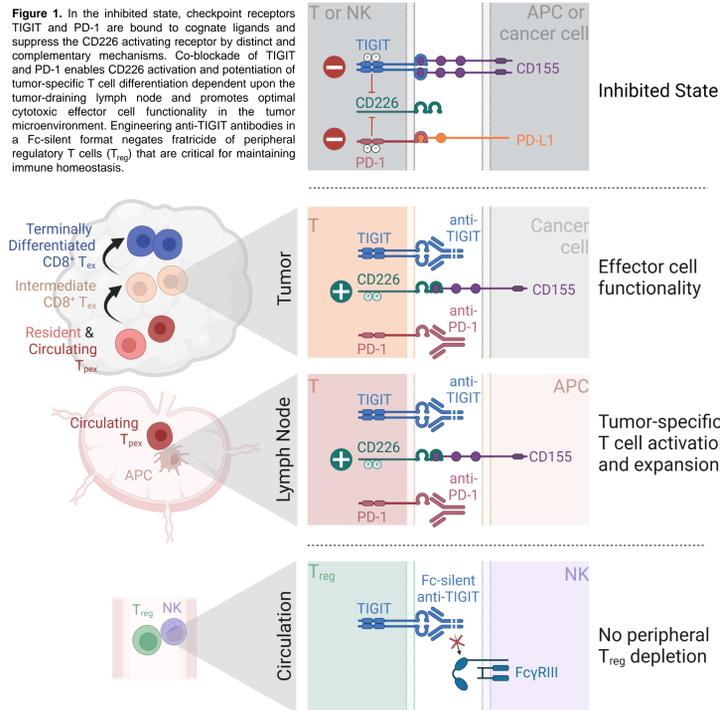


Figure 1. In the inhibited state, checkpoint receptors TIGIT and PD-1 are bound to cognate ligands and suppress the CD226 activating receptor by distinct and complementary mechanisms. Co-blockade of TIGIT and PD-1 enables CD226 activation and potentiation of tumor-specific T cell differentiation dependent upon the tumor-draining lymph node and promotes optimal cytotoxic effector cell functionality in the tumor microenvironment. Engineering anti-TIGIT antibodies in a Fc-silent format negates fratricide of peripheral regulatory T cells (T_{reg}) that are critical for maintaining immune homeostasis.

Results

Human and Mouse Tumor Infiltrating Lymphocytes Co-express TIGIT, PD-1, and CD226

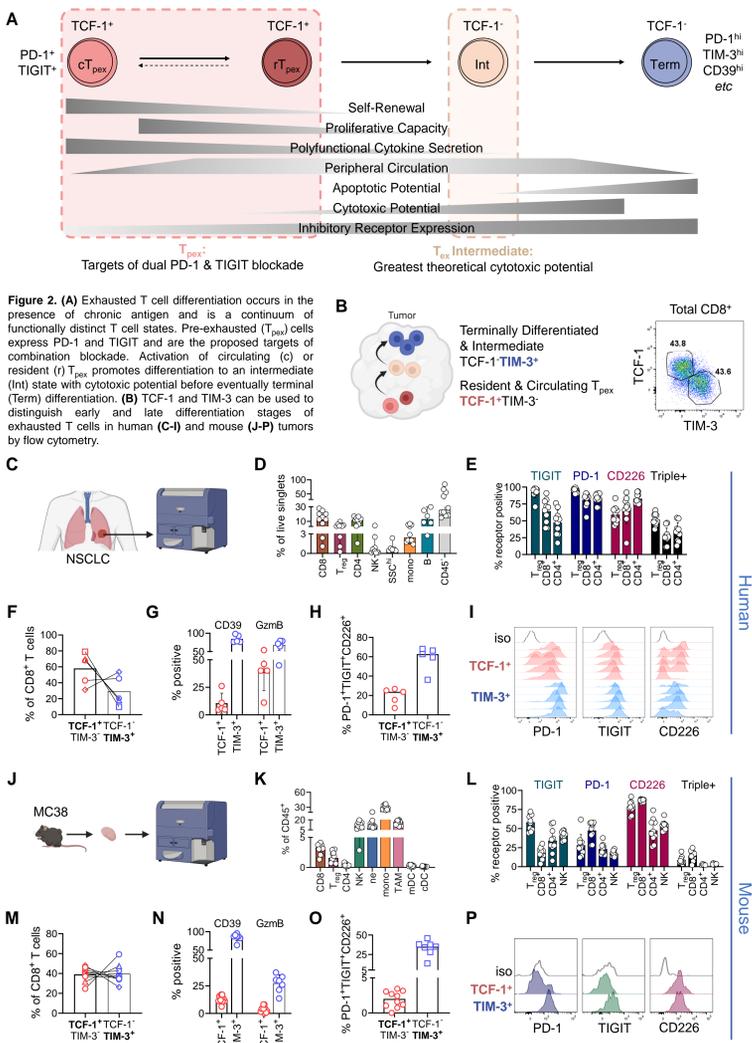


Figure 2. (A) Exhausted T cell differentiation occurs in the presence of chronic antigen and is a continuum of functionally distinct T cell states. Pre-exhausted (T_{pe}) cells express PD-1 and TIGIT and are the proposed targets of combination blockade. Activation of circulating (c) or resident (r) T_{pe} promotes differentiation to an intermediate (Int) state with cytotoxic potential before eventually terminal (Term) differentiation. (B) TCF-1 and TIM-3 can be used to distinguish early and late differentiation stages of exhausted T cells in human (C-I) and mouse (J-P) tumors by flow cytometry.

Results

Fc-silent Anti-TIGIT Potentiates Tumor-specific T Cell Activation and Differentiation that is Dependent on the Tumor Draining Lymph Node

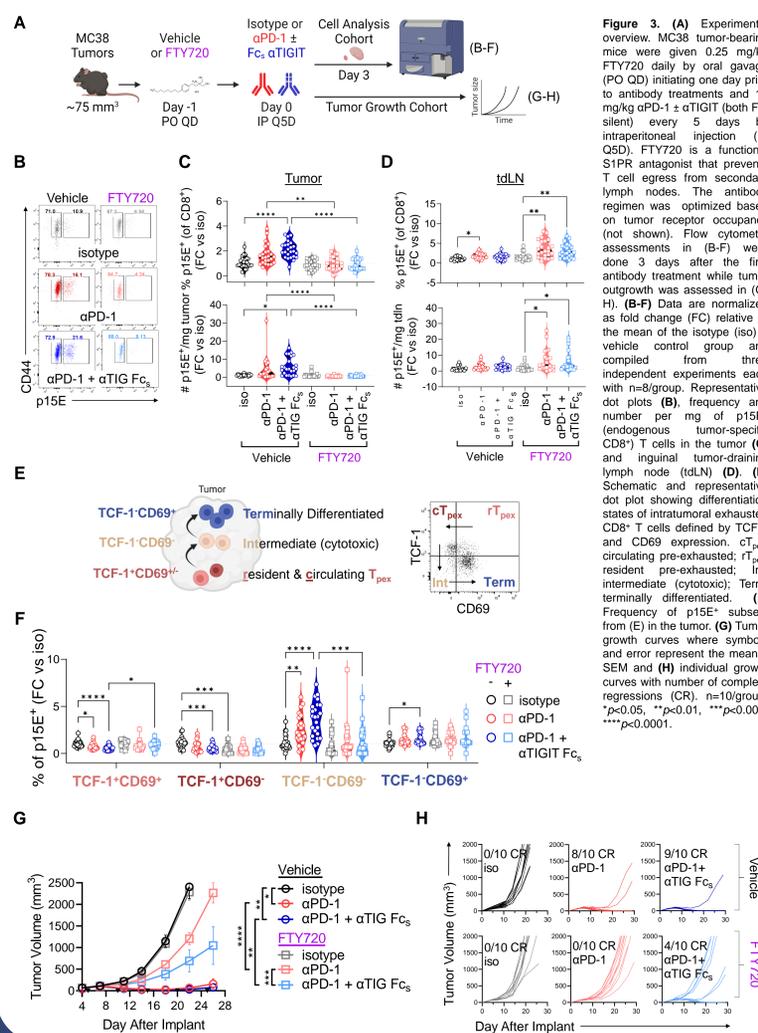


Figure 3. (A) Experimental overview. MC38 tumor-bearing mice were given 0.25 mg/kg FTY720 daily by oral gavage (PO QD) initiating one day prior to antibody treatments and 10 mg/kg aPD-1 ± aTIGIT (both Fc-silent) every 5 days by intraperitoneal injection (IP Q5D). FTY720 is a functional S1PR antagonist that prevents T cell egress from secondary lymph nodes. The antibody regimen was optimized based on tumor receptor occupancy (not shown). Flow cytometry assessments in (B-F) were done 3 days after the first antibody treatment while tumor outgrowth was assessed in (G-H). (B-F) Data are normalized as fold change (FC) relative to the mean of the isotype (iso) + vehicle control group and compiled from three independent experiments each with n=8/group. Representative dot plots (B), frequency and number per mg of p15E⁺ (endogenous tumor-specific CD8⁺) T cells in the tumor (C) and inguinal tumor-draining lymph node (tdLN) (D). (E) Schematic and representative dot plot showing differentiation states of intratumoral exhausted CD8⁺ T cells defined by TCF-1 and CD69 expression. CT_{pe}: circulating pre-exhausted; T_{pe}: resident pre-exhausted; Int: intermediate (cytotoxic); Term: terminally differentiated. (F) Frequency of p15E⁺ subsets from (E) in the tumor. (G) Tumor growth curves where symbols and error represent the mean ± SEM and (H) individual growth curves with number of complete regressions (CR). n=10/group. *p<0.05, **p<0.01, ***p<0.001, ****p<0.0001.

Results

Fc-enabled Anti-TIGIT Directs NK-mediated ADCC Against TIGIT-expressing Target Cells

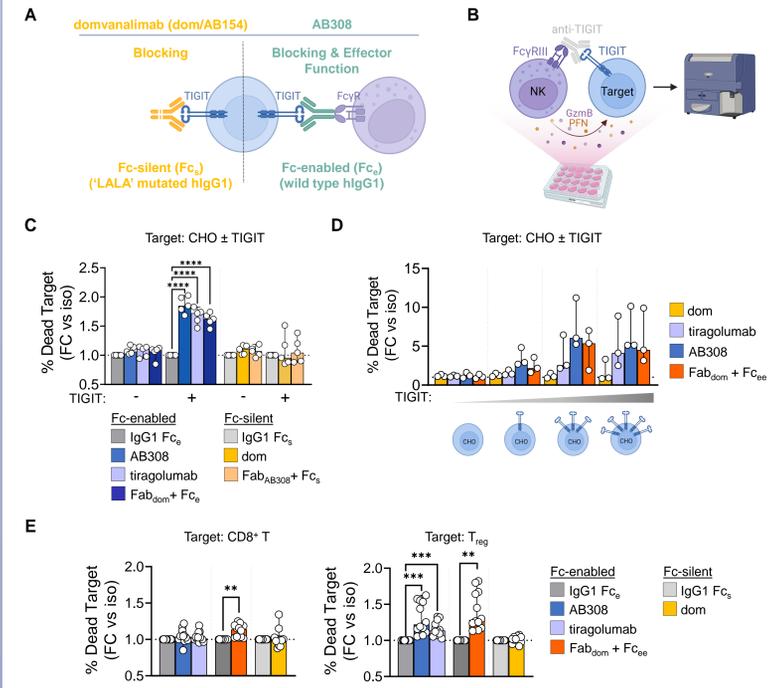


Figure 4. (A) Overview of clinical human TIGIT-specific antibodies. (B) Schematic *in vitro* human NK-mediated ADCC assay using TIGIT-expressing target cells in (C-E). Cell death was measured at 22 hours by flow cytometry and plotted as fold change (FC) relative to the respective isotype (iso) control for each NK donor. (C & D) ADCC assay with human TIGIT-expressing CHO target cells treated with Fc-enabled (Fc_e) or Fc-silent (Fc_s) antibodies. (E) ADCC assay with enriched primary human allogeneic CD8⁺ T or T_{reg} target cells treated with Fc_e or Fc_s antibodies. Fc_{pe}: Fc-effector enhanced (DLE: mutated hlgG1). Tiragolumab was manufactured by Arcus using sequences disclosed in the WHO Drug Information proposed INN publication (List 117; Vol 31 No. 2, 2017). **p<0.01, ***p<0.001, ****p<0.0001.

Peripheral Helios⁺ Effector T_{reg} (e T_{reg}) are Preferentially Targeted By TIGIT Directed ADCC

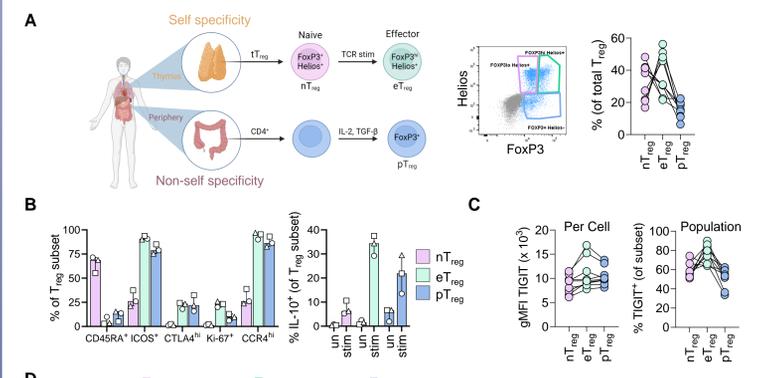


Figure 5. (A) Schematic overview¹¹, example dot plot, and frequency of three T_{reg} subsets detectable in peripheral human blood of healthy donors (n=9). (B) Phenotypic and functional characterization of subsets from (A) (n=3). Stim: overnight TCR stimulation. (C) GMFI and frequency of TIGIT of subsets from (A) (n=9). (D) After completion of ADCC assay outlined in Figure 4B, remaining live event counts for each T_{reg} subset were quantified (n=6). %, median decrease for all donor pairs treated with AB308 (Fc-enabled) relative to isotype control.

Peripheral T_{reg} Significantly Decrease in Patients Treated with Fc-enabled Anti-TIGIT, But Remain Stable in Patients Treated with Fc-silent Anti-TIGIT

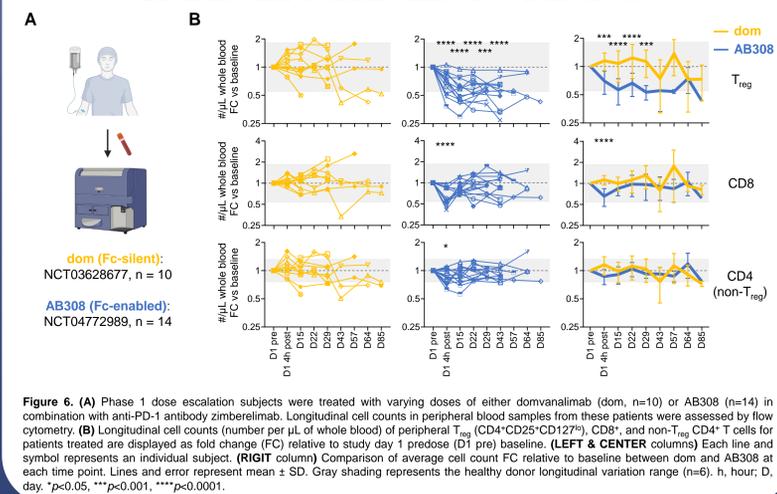


Figure 6. (A) Phase 1 dose escalation subjects were treated with varying doses of either domvanalimab (dom, n=10) or AB308 (n=14) in combination with anti-PD-1 antibody zimberelimab. Longitudinal cell counts in peripheral blood samples from these patients were assessed by flow cytometry. (B) Longitudinal cell counts (number per μ L of whole blood) of peripheral T_{reg} (CD4⁺CD25⁺CD127⁺), CD8⁺, and non- T_{reg} CD4⁺ T cells for patients treated are displayed as fold change (FC) relative to study day 1 predose (D1 pre) baseline. (LEFT & CENTER columns) Each line and symbol represents an individual subject. (RIGHT column) Comparison of average cell count FC relative to baseline between dom and AB308 at each time point. Lines and error represent mean ± SD. Gray shading represents the healthy donor longitudinal variation range (n=6). h, hour; D, day. *p<0.05, **p<0.01, ***p<0.001, ****p<0.0001.