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Supplemental information

A novel GPI-anchored dominant-negative

TGF- β receptor II renders T cells

unresponsive to TGF- β signaling

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Fig.S1: GPI-ecto-TGFβRII does interfere with TGFβ induced SMAD2/3 signalling in ATCs and Jurkat cells



Fig. S1: A) Whole WB of pSMAD2/3 from Fig. 2. B) WB of strongly inhibited TGF β -dependent SMAD and ERK signalling in Jurkat cells expressing TGF β -decoy receptors. No changes in Src signaling was observed. C) Flow cytometric analysis of p-SMAD2/3 and 1/8 and ERK1/2 in Jurkat cells gated for either high or low expression of TGF β R2 (decoy or WT). D+E) Readings of calcium flux in the presence of TGF β in either stimulated GFP-ctrl. or HA-GPI-ecto-T β RII expressing Jurkat cells.

Fig. S2: GPI-ecto-TGFbRII in ATCs does not affect stimulation, exhaustion or differentiation in TGFβ low culture conditions



Fig. S2: Expression of TGFβ-decoy receptor does not affect primary T cell marker expression for stimulation, exhaustion or differentiation per se. A) Line graphs representing the expression of CD69 as a marker for T cell stimulation, B) PD1 as a marker for exhaustion, and C) CD57 and D) KLRG1 as marker for T cell differentiation. E) the ratio of CD4+ T cells as well as NKT cells is not affected either. Line graphs display the mean and SEM of at least 6 biological replicates.

Fig. S3: HA-GPI-ecto-TGFbRII has no effect on stimulation or exhaustion or the expression of the chosen marker for differentiation in ATCs in the presence of recombinant TGFb



Fig. S3: Expression of TGF β -decoy receptor does not affect primary T cell marker expression for stimulation, exhaustion or differentiation in the presence of different concentrations of TGF β . A) Line graphs representing the expression of CD69 as a marker for T cell stimulation, B) PD1 as a marker for exhaustion, and C) CD57 and D) KLRG1 as marker for T cell differentiation. E) the ratio of CD4+ T cells is not affected whereas the proliferation of NKT cells seems slightly enhanced in cells expressing HA-GPI-ecto-T β RII. Line graphs display the mean and SEM of at least 6 biological replicates.



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| IFNg | | GFP | s HA-DN | GFP vs HA-Alkphos | | | | GFP vs HA-DN | | GFP vs HA-Alkphos | | | | GFP vs HA-DN | | GFP vs HA-Alkphos | |
|------|------------|--------------|--------------|-------------------|--------------|---------|------------|--------------|--------------|-------------------|--------------|--------|------------|--------------|--------------|-------------------|--------------|
| | TGFb conc. | p value | significance | p value | significance | GM-CSF | TGFb conc. | p value | significance | p value | significance | | TGFb conc. | p value | significance | p value | significance |
| | untreated | 0.07213 | NA | 0.08900 | NA | | untreated | 0.05978 | NA | 0.36468 | NA | THE | untreated | 0.009872 | ** | 0.07086 | NA |
| | 1ng/ml | 0.00158 | ** | 0.00255 | ** | | 1ng/ml | 0.00026 | *** | 0.00024 | *** | INFU | 1ng/ml | 0.000855 | *** | 0.00287 | ** |
| | 10ng/ml | 0.00350 | ** | 0.00631 | ** | | 10ng/ml | 0.00014 | *** | 0.00034 | *** | | 10ng/ml | 0.000821 | *** | 0.00661 | ** |
| | 100ng/ml | 0.00092 | *** | 0.00649 | ** | | 100ng/ml | 0.00027 | *** | 0.00196 | ** | | 100ng/ml | 0.000332 | *** | 0.00736 | ** |
| | | GFP vs HA-DN | | GFP vs HA-Alkphos | | | | GFP vs HA-DN | | GFP vs HA-Alkphos | | | | GFP vs HA-DN | | GFP vs HA-Alkphos | |
| | TGFb conc. | p value | significance | p value | significance | | TGFb conc. | p value | significance | p value | significance | | TGFb conc. | p value | significance | p value | significance |
| TNEO | untreated | 0.329954 | NA | 0.7425 | NA | MID 1h | untreated | 0.521679 | NA | 0.98554 | NA | | untreated | 0.02701 | * | 0.62017 | NA |
| пигр | 1ng/ml | 0.089237 | NA | 0.2067 | NA | WIIP-10 | 1ng/ml | 0.036379 | • | 0.07303 | NA | IL5 | 1ng/ml | 0.00253 | ** | 0.02117 | • |
| | 10ng/ml | 0.084746 | NA | 0.2439 | NA | | 10ng/ml | 0.050498 | NA | 0.04275 | * | | 10ng/ml | 0.00184 | ** | 0.04388 | • |
| | 100ng/ml | 0.042144 | * | 0.1323 | NA | | 100ng/ml | 0.018459 | • | 0.05823 | NA | | 100ng/ml | 0.00127 | ** | 0.03139 | • |
| IL4 | | GFP vs HA-DN | | GFP vs HA-Alkphos | | | | GFP vs HA-DN | | GFP vs HA-Alkphos | | | | GFP vs HA-DN | | GFP vs HA-Alkphos | |
| | TGFb conc. | p value | significance | p value | significance | | TGFb conc. | p value | significance | p value | significance | | TGFb conc. | p value | significance | p value | significance |
| | untreated | 0.20879 | * | 0.66077 | NA | IL5 | untreated | 0.52152 | NA | 0.37745 | NA | IL6 | untreated | 0.13642 | NA | 0.54948 | NA |
| | 1ng/ml | 0.00367 | ** | 0.01860 | • | | 1ng/ml | 0.03666 | • | 0.04560 | * | | 1ng/ml | 0.00010 | *** | 0.05073 | NA |
| | 10ng/ml | 0.02040 | * | 0.03725 | • | | 10ng/ml | 0.06233 | NA | 0.04387 | * | | 10ng/ml | 0.00001 | *** | 0.00791 | ** |
| | 100ng/ml | 0.00061 | *** | 0.01556 | • | | 100ng/ml | 0.03469 | • | 0.04952 | * | | 100ng/ml | 0.00334 | ** | 0.00378 | ** |
| IL13 | | GFP vs HA-DN | | GFP vs HA-Alkphos | | | | GFP vs HA-DN | | GFP vs HA-Alkphos | | | | GFP vs HA-DN | | GFP vs HA-Alkphos | |
| | TGFb conc. | p value | significance | p value | significance | IL10 | TGFb conc. | p value | significance | p value | significance | cCD401 | TGFb conc. | p value | significance | p value | significance |
| | untreated | 0.98495 | NA | 0.79050 | NA | | untreated | 0.00498 | ** | 0.02654 | * | | untreated | 0.29442 | NA | 0.58031 | NA |
| | 1ng/ml | 0.00016 | *** | 0.00049 | *** | | 1ng/ml | 0.00097 | *** | 0.01471 | * | SCD40L | 1ng/ml | 0.00146 | ** | 0.00138 | ** |
| | 10ng/ml | 0.00023 | *** | 0.00205 | ** | | 10ng/ml | 0.00032 | *** | 0.02407 | + | | 10ng/ml | 0.00171 | ** | 0.03216 | • |
| | 100ng/ml | 0.00013 | *** | 0.00509 | ** | | 100ng/ml | 0.00019 | *** | 0.00593 | ** | | 100ng/ml | 0.00094 | *** | 0.00906 | ** |

Fig. S4A) Heat map presentation of concentrations of IL10 and sCD40L measured in cell culture supernatants after ATC culture in the presence of different concentrations of TGF β . B) Statistical analyses of cytokine concentrations measured in culture supernatants of GFP ctrl. and TGF β -decoy receptor expressing ATCs. Heat maps display the data of 6 biological replicates.