

# Supplementary Information

## The induction and function of the anti-inflammatory fate of T<sub>H</sub>17 cells

Xu, H., Agaloti, T. *et al.*, 2020

**Supplementary Figure 1: Evaluation of T<sub>H</sub>17-specific IL-10 depletion mouse model**

**Supplementary Figure 2: No immune abnormality observed in the periphery**

**Supplementary Figure 3: T<sub>H</sub>17 cells express higher levels of IL-10R compared to T<sub>H</sub>1 cells**

**Supplementary Figure 4: TGF- $\beta$  promotes IL-10 production in mature T<sub>H</sub>17 cells**

**Supplementary Figure 5: Tgfbr2 is required for T<sub>H</sub>17 cells to produce IL-10**

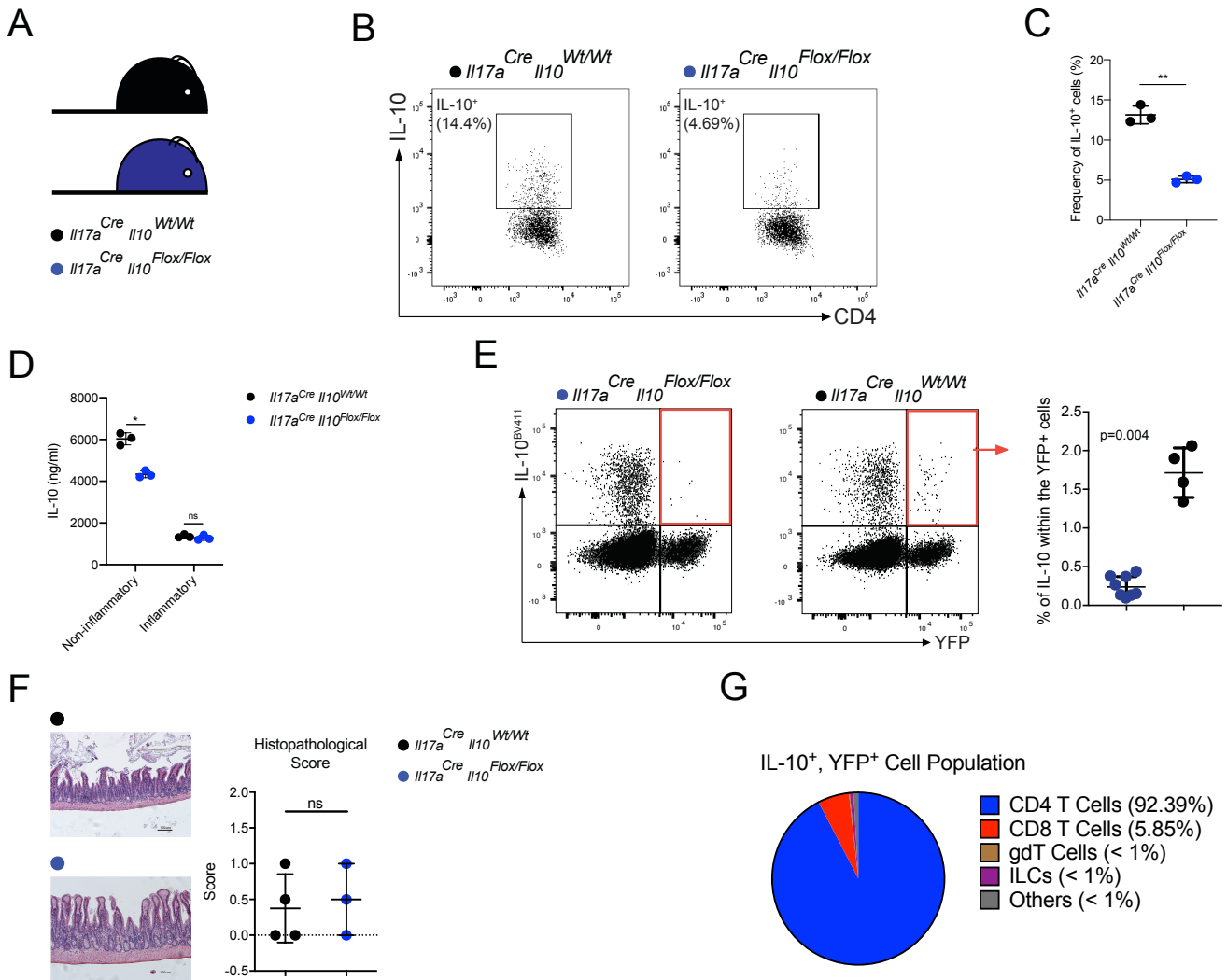
**Supplementary Figure 6: Tgfbr2 is required for T<sub>H</sub>17 cells to produce IL-10 during intestinal inflammation**

**Supplementary Figure 7: TIF1 $\gamma$  is dispensable for IL-10 production in T<sub>H</sub>17 cells**

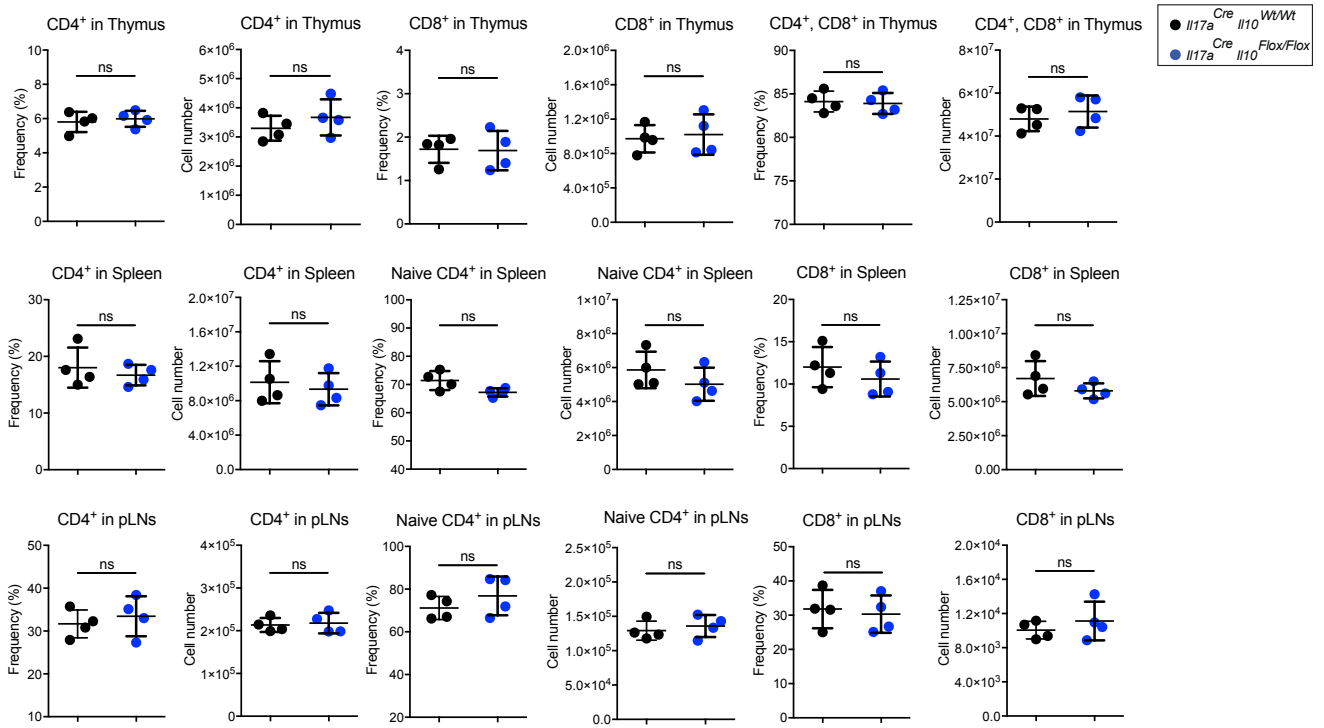
**Supplementary Figure 8: Smad4 is required for T<sub>H</sub>17 cells to produce IL-10**

**Supplementary Figure 9: Smad3 is important for IL-10 production in T<sub>H</sub>17 cells**

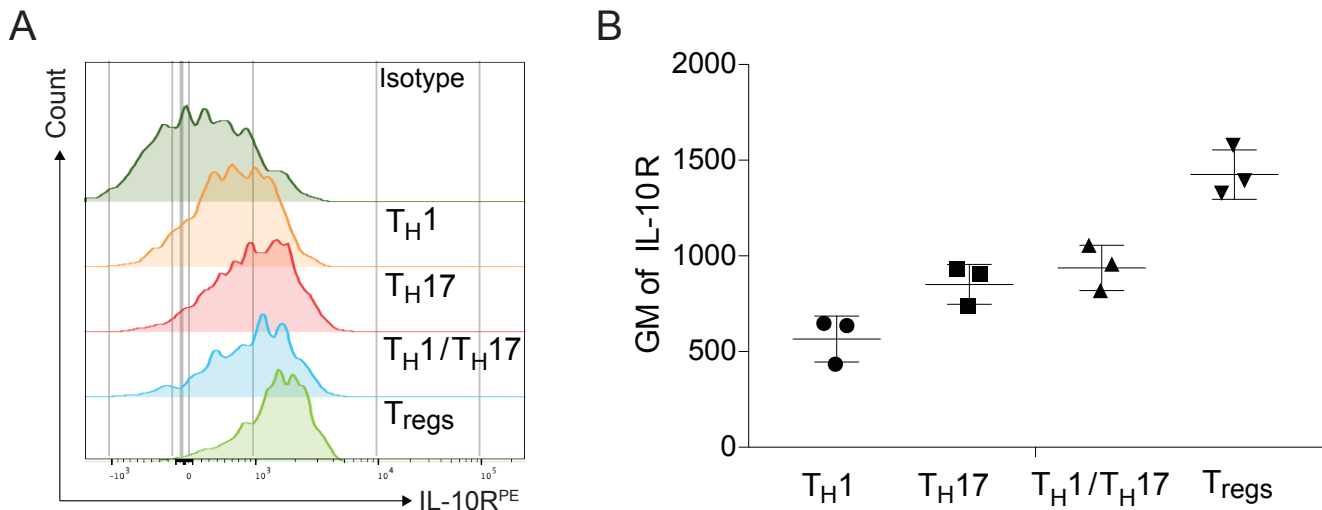
**Supplementary Figure 10: Human intestinal T<sub>H</sub>17 cells produce IL-10; and TGF- $\beta$  further promotes IL-10 expression**



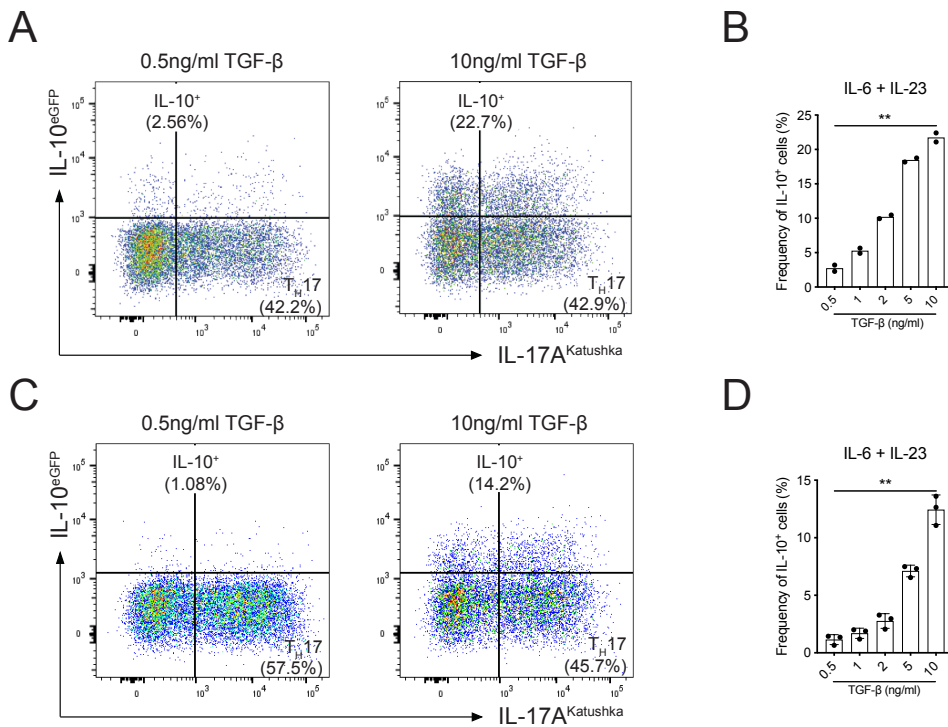
**Supplementary Figure 1. Evaluation of  $T_H17$ -specific IL-10 depletion mouse model.** A. Mouse strains employed to deplete IL-10 production in  $T_H17$  cells. B. Flow cytometric analysis of intestinal lymphocytes isolated from  $Il17a^{Cre} Il10^{Wt/Wt}$  as well as  $Il17a^{Cre} Il10^{FloxFlox}$  mice after anti-CD3 treatment. Intracellular staining for IL-10 was then performed. A pre-gate on  $T_H17$  cells ( $CD4^+$ ,  $TCR\beta^+$ ,  $IL-17A^+$ ) was applied. C. Statistical analysis of frequencies of IL-10<sup>+</sup> cells among  $T_H17$  cells. Each dot represents one mouse ( $n_{wild\ type}=3$ ,  $n_{KO}=3$ ). Mean  $\pm$  S.D.; \*\* $P < 0.01$  by Welch's t-test. D. ELISA measurement of IL-10 after *in vitro* culture. Naive  $CD4^+$  T cells were isolated from the spleens of  $Il17a^{Cre} Il10^{Wt/Wt}$  as well as  $Il17a^{Cre} Il10^{FloxFlox}$  mice and were cultured under non-inflammatory (IL-6 + TGF- $\beta$ ) or inflammatory (IL-6 + IL-23 + IL-1 $\beta$ )  $T_H17$  conditions. IL-10 in the supernatant was measured by ELISA on day 5. Each dot represents one replicate ( $n=3$ ). Mean  $\pm$  S.D.; ns, not significant; \* $P < 0.05$  by Mann-Whitney U test. E. Flow cytometric analysis of *in vitro* cultured  $T_H17$  cells. Naive  $CD4^+$  T cells were isolated from the spleens of  $Il17a^{Cre} Il10^{Wt/Wt}$  as well as  $Il17a^{Cre} Il10^{FloxFlox}$  mice and were cultured under non-inflammatory (IL-6 + TGF- $\beta$ )  $T_H17$  conditions. Cells were re-stimulated and intracellularly stained for IL-10 on Day 5. On the right the statistics of IL-10<sup>+</sup> cells within YFP<sup>+</sup> cells are reported. Each dot represents one separate  $T_H17$  cell culture ( $n_{wild\ type}=4$ ,  $n_{KO}=8$ ) obtained from pooled naïve T cells coming from two mice. Mean  $\pm$  S.D.; Mann-Whitney U-test. F. Representative histological pictures of H&E stained small intestines under steady states. One representative experiment of two is shown. Each dot represents one mouse ( $n_{wild\ type}=4$ ,  $n_{KO}=3$ ). Mean  $\pm$  S.D.; ns, not significant by Welch's t-test. G. Frequencies of different IL-10-producing lymphocyte populations that are derived from IL-17A producers. Intestinal lymphocytes were harvested from steady state *Fate*<sup>+</sup> mice (Gagliani et al., *Nature* 523, 224-224, 2015). Different IL-10-producing immune cell populations that have activated IL-17A<sup>CRE</sup> (indicated by YFP expression) were analyzed by flow cytometry.



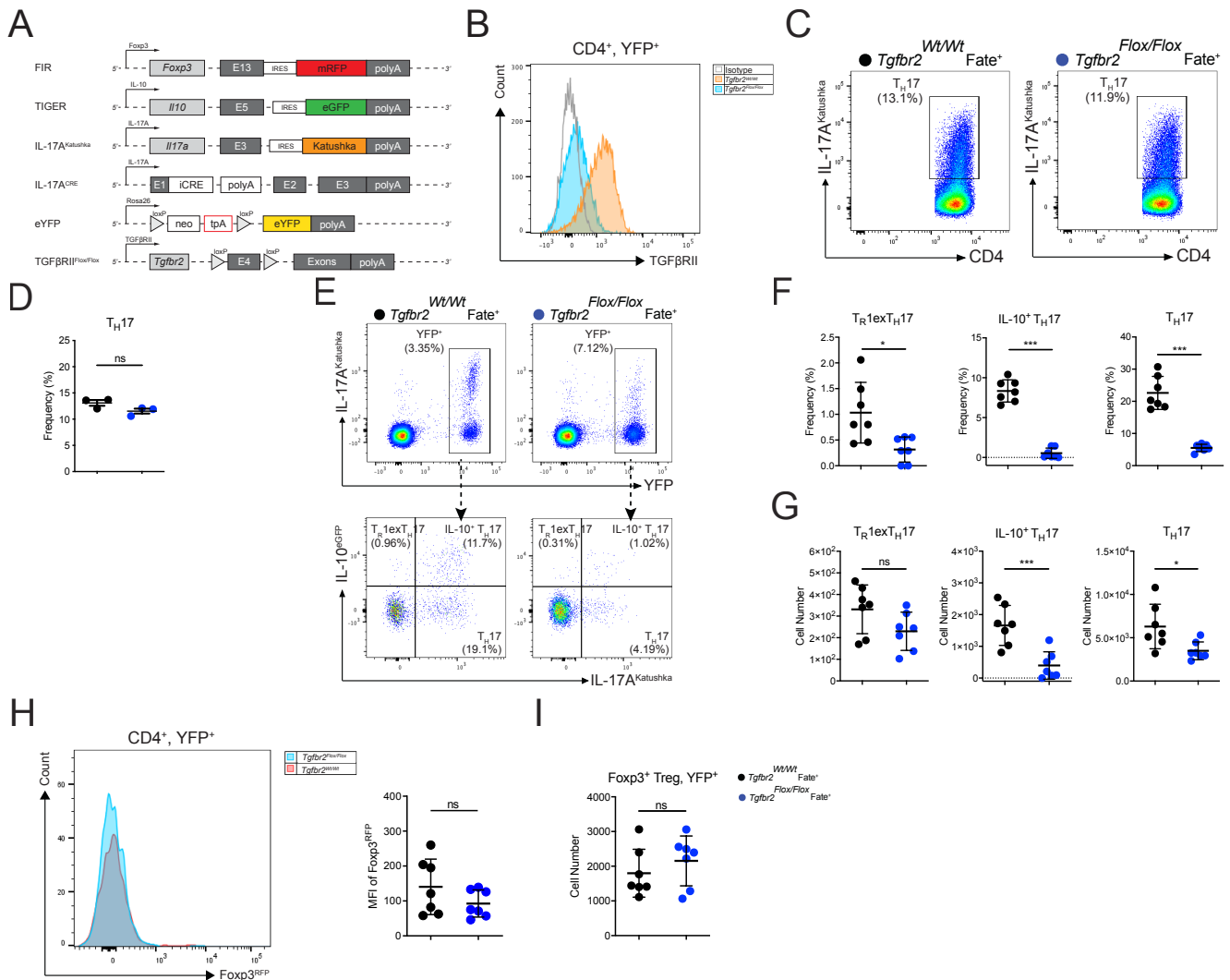
**Supplementary Figure 2. No immune abnormality observed in the periphery.** Frequencies and cell numbers of different T cell subsets in different immune organs. Lymphocytes from thymus, spleen and peripheral lymph nodes (pLNs) were harvested from steady state *Il17a<sup>Cre</sup> Il10<sup>Flox/Flox</sup>* mice as well as their littermate wild-type control mice. Different T cell subsets were then analyzed by flow cytometry. Each dot represents one mouse ( $n_{wild\ type}=4$ ,  $n_{KO}=4$ ). Mean  $\pm$  S.D.; ns, not significant by Mean  $\pm$  S.D.; ns, not significant by Mann-Whitney *U* test.



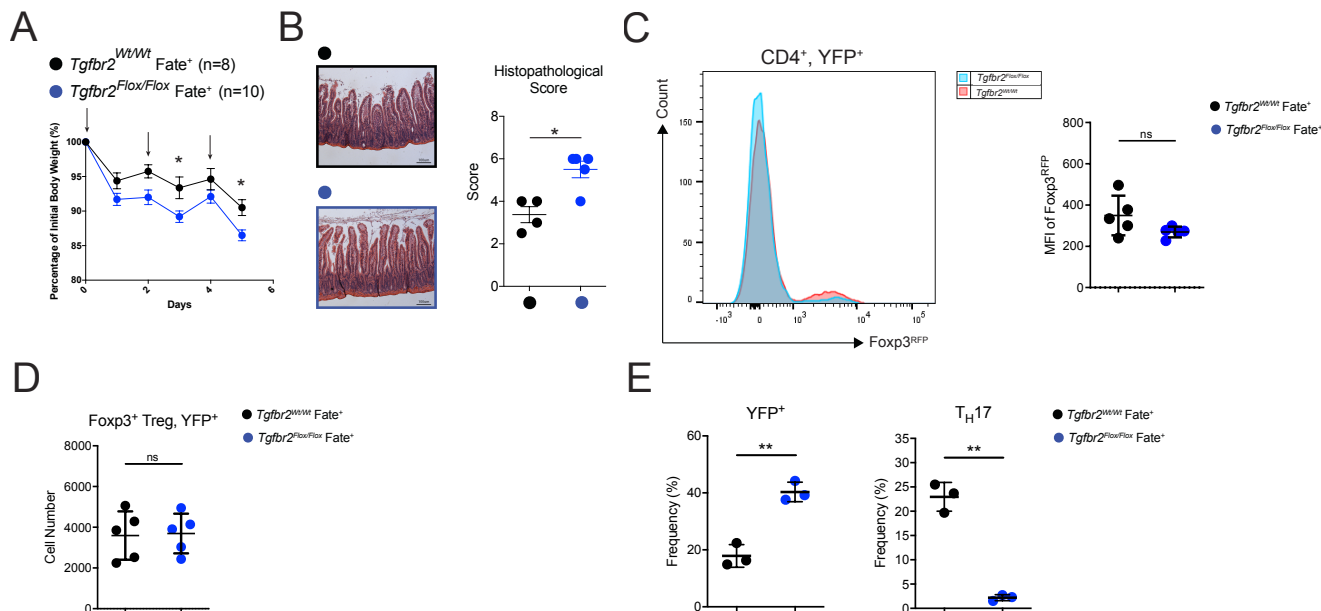
**Supplementary Fig. 3 T<sub>H</sub>17 cells express higher levels of IL-10R compared to T<sub>H</sub>1 cells.** A. Histograms showing the expression of IL-10R for T<sub>H</sub>1 cells (defined as CD4<sup>+</sup> CD44<sup>+</sup> IFN- $\gamma$ <sup>Katushka<sup>+</sup></sup> IL-17a<sup>eGFP<sup>-</sup></sup>), T<sub>H</sub>17 cells (defined as CD4<sup>+</sup> CD44<sup>+</sup> IFN- $\gamma$ <sup>Katushka<sup>-</sup></sup> IL-17a<sup>eGFP<sup>+</sup></sup>), T<sub>H</sub>1/T<sub>H</sub>17 cells (defined as CD4<sup>+</sup> CD44<sup>+</sup> IFN- $\gamma$ <sup>Katushka<sup>+</sup></sup> IL-17a<sup>eGFP<sup>+</sup></sup>) and Tregs (defined as CD4<sup>+</sup> Foxp3<sup>RFP<sup>+</sup></sup> IFN- $\gamma$ <sup>Katushka<sup>-</sup></sup> IL-17a<sup>eGFP<sup>-</sup></sup>) isolated from the intestinal lamina propria of a representative Foxp3<sup>RFP</sup> IL-17a<sup>eGFP</sup> IFN- $\gamma$ <sup>Katushka</sup> reporter mouse injected with anti-CD3 mAb. B. Geometric Mean (GM) values of IL-10R from different CD4 T cell subsets. Each dot represents one mouse ( $n=3$ ). Mean  $\pm$  S.D.



**Supplementary Fig. 4 TGF- $\beta$  promotes IL-10 production in mature T<sub>H</sub>17 cells.** A, C. T<sub>H</sub>17 cells generated *in vitro* under either TGF- $\beta$  condition (IL-6 + IL-23 + TGF- $\beta$ ) (A) or IL-1 $\beta$  condition (IL-6 + IL-23 + IL-1 $\beta$ ) (C) were purified and then re-activated with IL-6, IL-23 as well as different amounts of TGF- $\beta$ . Cells in the above dot plots are pre-gated on viable CD4<sup>+</sup> T cells. B, D. Statistical analysis of the frequencies of IL-10<sup>+</sup> cells in A, C respectively across different TGF- $\beta$  conditions. \*\*P < 0.01, by ordinary one-way ANOVA.

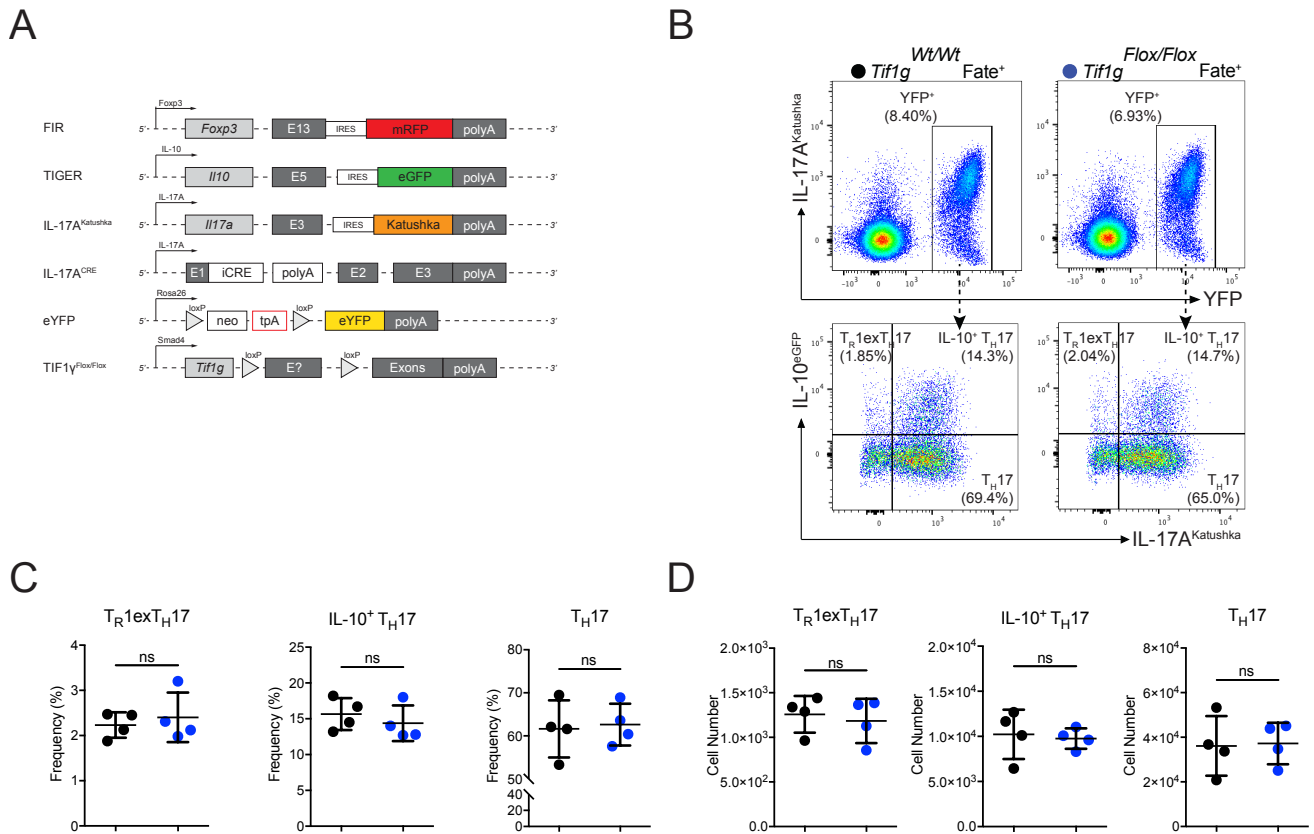


**Supplementary Fig. 5 Tgfr2 is required for TH17 cells to produce IL-10.** A. Constructs contained in Tgfr2<sup>fl/fl</sup> Fate<sup>+</sup> mouse line. B. Flow cytometry analysis of TGF- $\beta$ RII expression in intestinal CD4<sup>+</sup> T cells expressing YFP (indicating CRE activation). One representative of two independent experiments is shown. C. Flow cytometry analysis of naive CD4<sup>+</sup> T cells polarized under TH17 condition *in vitro*. Cells are pre-gated on viable CD4<sup>+</sup> T cells. D. Frequency of TH17 cells shown in C. One representative of two independent experiments is shown. Mean  $\pm$  S.D.; ns, not significant by Welch's t-test. E. Flow cytometry analysis of small intestinal CD4<sup>+</sup> T cells under steady state. Top panel is pre-gated on CD4<sup>+</sup> Fcpx3<sup>-</sup> T cells and bottom panel on CD4<sup>+</sup> Fcpx3<sup>-</sup> YFP<sup>+</sup> T cells. The cell populations indicated on the FACS dot plots are identified by different combination of reporter expression (T<sub>R</sub>1exTH17: IL-10<sup>eGFP+</sup> YFP<sup>+</sup> IL17a<sup>Kata-</sup>; IL-10<sup>+</sup> TH17: IL-10<sup>eGFP+</sup> YFP<sup>+</sup> IL17a<sup>Kata+</sup>; TH17: IL-10<sup>eGFP-</sup> YFP<sup>+</sup> IL17a<sup>Kata+</sup>). F, G. Frequencies (F) and numbers (G) of T<sub>R</sub>1<sup>exTH17</sup>, IL-10<sup>+</sup> TH17 and TH17 cells. Data are pooled from three independent experiments. Each dot represents one mouse ( $n_{wild\ type}=7$ ,  $n_{KO}=7$ ). Mean  $\pm$  S.D.; ns, not significant; \*P < 0.05, \*\*\*P < 0.001 by Mann-Whitney U test. H. Left, Flow cytometric analysis of the Fcpx3 expression level among the intestinal YFP<sup>+</sup> CD4<sup>+</sup> T cells under steady state conditions. Fcpx3 expression was determined by RFP reporter fluorescence. A pre-gate on YFP<sup>+</sup> CD4<sup>+</sup> T cells is applied. Right panel shows the statistical analysis of Fcpx3 Mean Fluorescence Intensity (MFI) among YFP<sup>+</sup> CD4<sup>+</sup> T cells. I. Numbers of YFP<sup>+</sup> Fcpx3<sup>+</sup> Treg cells in the small intestine under steady state conditions in the two indicated mouse lines. Each dot represents one mouse ( $n_{wild\ type}=7$ ,  $n_{KO}=7$ ). Mean  $\pm$  S.D.; ns, not significant by Mann-Whitney U test.



**Supplementary Fig. 6 Tgfb2 is required for T<sub>H</sub>17 cells to produce IL-10 during intestinal inflammation.**

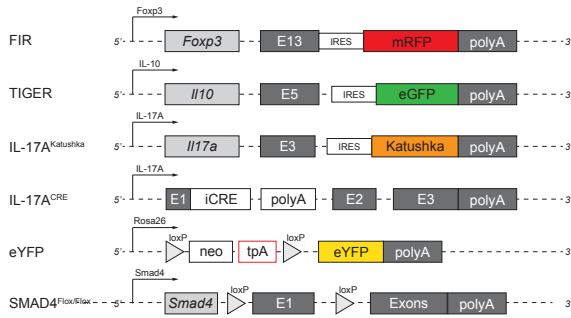
A. Percentages of initial body weight of mice after anti-CD3 mAb treatment. The arrows indicate the injection of anti-CD3 mAb. Data are cumulative of three independent experiments. Mean  $\pm$  SEM.; \*P < 0.05, by 2-way ANOVA with Bonferroni's post-test. B. Representative histological pictures of H&E stained small intestines after anti-CD3 treatment. Statistical analysis is representative of two experiments. Each dot represents one mouse ( $n_{wild\ type}=4$ ,  $n_{KO}=5$ ). Mean  $\pm$  SEM, \*P < 0.05 by Mann-Whitney U test. C. Left, flow cytometric analysis of Foxp3 expression among YFP<sup>+</sup> CD4<sup>+</sup> T cells in the small intestine during anti-CD3 mAb-induced intestinal inflammation. Foxp3 expression was determined by RFP reporter fluorescence. A pre-gate on YFP<sup>+</sup> CD4<sup>+</sup> T cells is applied. Right panel shows the statistical analysis of Foxp3 Mean Fluorescence Intensity (MFI) among YFP<sup>+</sup> CD4<sup>+</sup> T cells between the two indicated mouse lines. Each dot represents one mouse ( $n_{wild\ type}=5$ ,  $n_{KO}=5$ ). D. Numbers of YFP<sup>+</sup> Foxp3<sup>+</sup> Treg cells in the small intestine in the two indicated mouse lines. Each dot represents one mouse ( $n_{wild\ type}=5$ ,  $n_{KO}=5$ ). Mean  $\pm$  S.D.; ns, not significant by Mann-Whitney U test. E. As part of Fig.4G, here it is reported the statistical analysis of YFP<sup>+</sup> and T<sub>H</sub>17 cells after CD4<sup>+</sup> T cells transfer. CD4<sup>+</sup> T cells were isolated from the indicated mouse lines. Each dot represents one mouse ( $n_{wild\ type}=3$ ,  $n_{KO}=3$ ). Mean  $\pm$  S.D.; \*\*P < 0.01 by Welch's t-test.



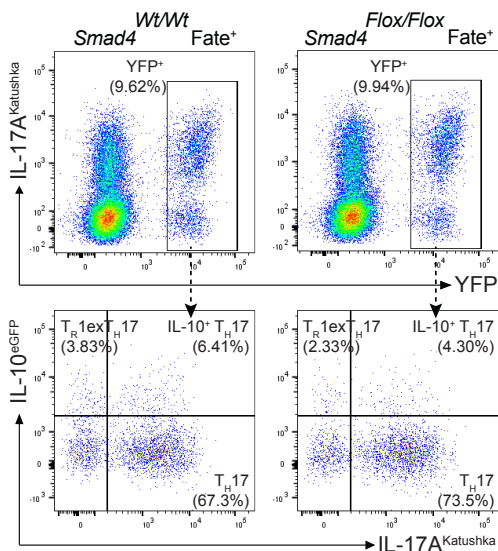
**Supplementary Fig. 7 TIF1 $\gamma$  is dispensable for IL-10 production in T<sub>H</sub>17 cells.** A. Constructs contained in *Tif1g<sup>fl/fl</sup>* *Fate<sup>+</sup>* mice. B. Flow cytometry analysis of small intestinal CD4<sup>+</sup> T cells after anti-CD3-induced intestinal inflammation. Top panel, pre-gated on *Foxp3<sup>-</sup>* CD4<sup>+</sup> T cells; bottom panel, pre-gated on YFP<sup>+</sup> *Foxp3<sup>-</sup>* CD4<sup>+</sup> T cells. The populations indicated on the FACS dot plots are identified by different combination of reporter expression (T<sub>R</sub>1exT<sub>H</sub>17: IL-10<sup>eGFP+</sup> YFP<sup>+</sup> IL17a<sup>Kata-</sup>; IL-10<sup>+</sup>T<sub>H</sub>17: IL-10<sup>eGFP+</sup> YFP<sup>+</sup> IL17a<sup>Kata+</sup>; T<sub>H</sub>17: IL-10<sup>eGFP-</sup> YFP<sup>+</sup> IL17a<sup>Kata+</sup>). C, D. Frequencies (C) and numbers (D) of T<sub>R</sub>1exT<sub>H</sub>17, IL-10<sup>+</sup> T<sub>H</sub>17 and T<sub>H</sub>17 cells. One representative of three independent experiments is shown. Each dot represents one mouse ( $n_{wild\ type}=4$ ,  $n_{KO}=4$ ). Mean  $\pm$  S.D.; ns, not significant by Mann-Whitney U test.



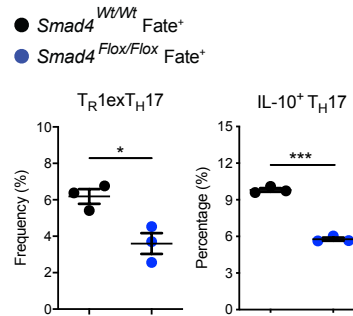
**A**



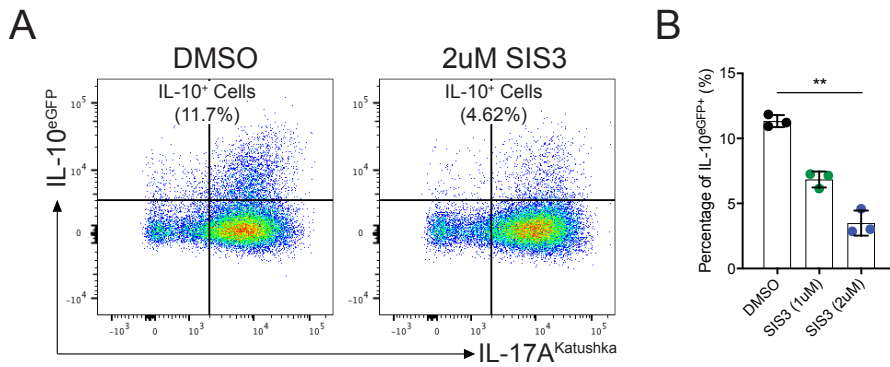
**B**



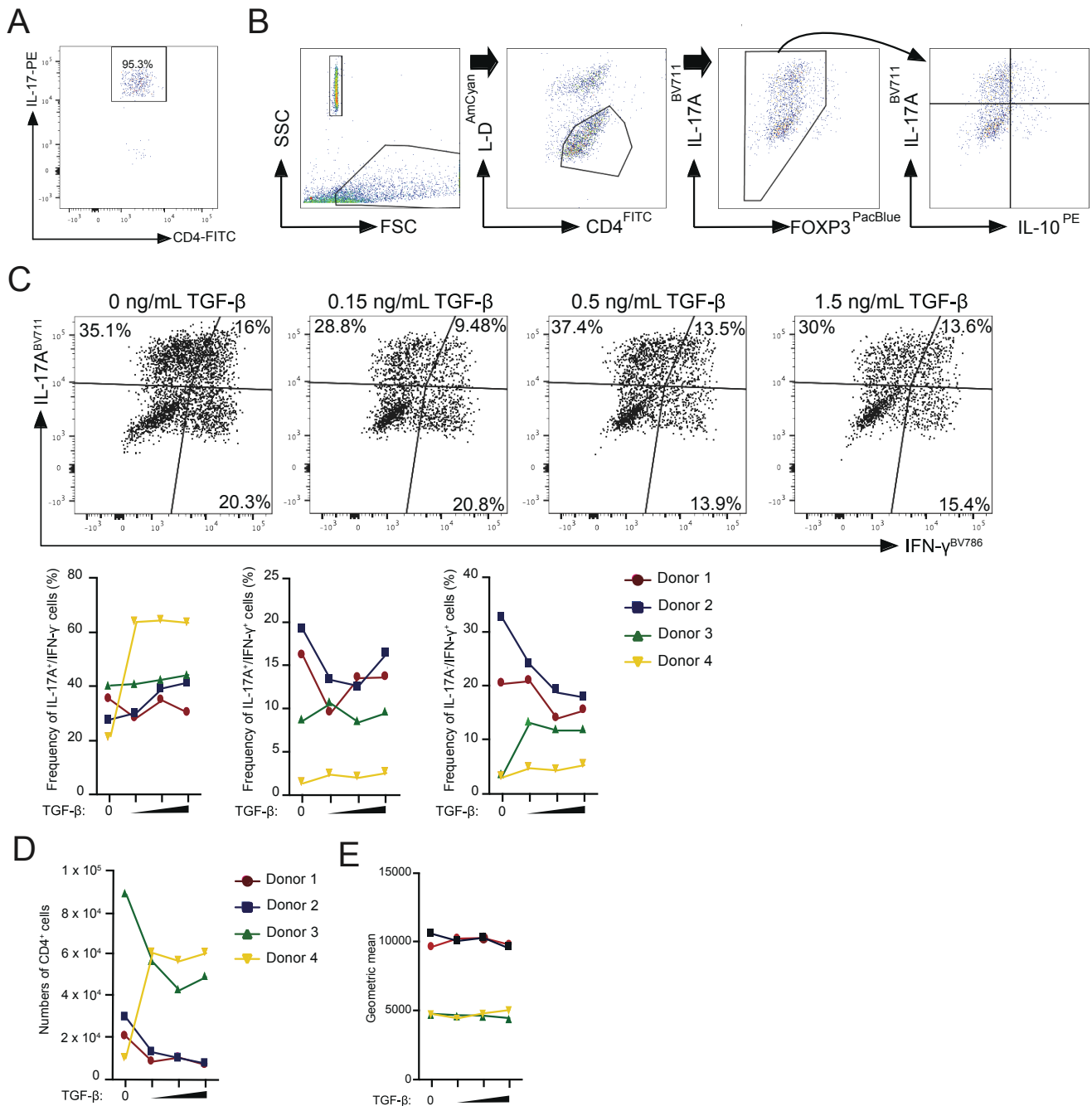
**C**



**Supplementary Fig. 8 Smad4 is required for T<sub>H</sub>17 cells to produce IL-10.** A. Constructs contained in *Smad4*<sup>fl/fl</sup> Fate<sup>+</sup> mice. B. Flow cytometric analysis of *in vitro* cultured naive CD4<sup>+</sup> T cells. Cells were polarized under T<sub>H</sub>17 condition for 5 days. Top panel, pre-gated on Foxp3<sup>-</sup> CD4<sup>+</sup> T cells; bottom panel, pre-gated on YFP<sup>+</sup> Foxp3<sup>-</sup> CD4<sup>+</sup> T cells. The populations indicated on the FACS dot plots are identified by different combination of reporter expression (T<sub>R</sub>1exT<sub>H</sub>17: IL-10<sup>eGFP+</sup> YFP<sup>+</sup> IL17a<sup>Kata-</sup>; IL-10<sup>+</sup>T<sub>H</sub>17: IL-10<sup>eGFP+</sup> YFP<sup>+</sup> IL17a<sup>Kata+</sup>; T<sub>H</sub>17: IL-10<sup>eGFP-</sup> YFP<sup>+</sup> IL17a<sup>Kata+</sup>). B. Statistical analysis of T<sub>R</sub>1exT<sub>H</sub>17, IL-10<sup>+</sup>T<sub>H</sub>17 and T<sub>H</sub>17 cells. Data are representative of one out of three experiments. Each dot represents one mouse ( $n_{wild\ type}=3$ ,  $n_{KO}=3$ ). Mean ± S.D.; \*P < 0.05, \*\*\*P < 0.001 by Welch's t-test.



**Supplementary Fig. 9 Smad3 is important for IL-10 production in T<sub>H</sub>17 cells.** *In vitro* differentiated T<sub>H</sub>17 cells were purified and cultured in the presence 2ng/ml TGF- $\beta$  together with DMSO or different amount of SIS3. A. Representative flow cytometry analysis of DMSO and 2uM SIS3 conditions. Cells were pre-gated on viable CD4<sup>+</sup> T cells. Expression of IL-10<sup>eGFP</sup> was detected by reporter fluorescence. B. Statistical analysis of the frequencies of IL-10<sup>+</sup> cells across different conditions. Each dot represents one replicate ( $n=3$ ). Mean  $\pm$  S.D.; \*\*P < 0.01 by Kruskal-Wallis test.



**Supplementary Fig. 10 Human intestinal T<sub>H</sub>17 cells produce IL-10 and TGF-β further promotes IL-10 expression.** A. The dot plot shows the purity of the sorted CD4<sup>+</sup> IL17A<sup>+</sup> cells corresponding to blood donor 1 shown in Fig. 7. B. Gating strategy of human T<sub>H</sub>17 cells isolated from PBMCs of one representative healthy human donor. This strategy applies to the data shown here and in Figure 7B. C. Dot plots of PBMC-sorted T<sub>H</sub>17 cells from the human blood, stimulated and cultured under increasing TGF-β1 concentrations for 5 days and stained for IL-17A and IFN-γ. The graphs below show the percentages of IL-17A<sup>+</sup>, IL-17A<sup>+</sup>/IFN-γ<sup>+</sup>, and IFN-γ<sup>+</sup> cells from four different blood donors. Each colored line represents one donor ( $n=4$ ). D. Absolute total numbers of living CD4<sup>+</sup> T cells corresponding to (C). E. Foxp3 expression (MFI) among the *in vitro* cultured human T<sub>H</sub>17 cells shown in Fig. 7, in the presence of increasing TGF-β concentrations.

**Supplementary Table 1: Table of Antibodies**

<b>Name</b>	<b>Clone</b>	<b>Company</b>	<b>Cat. Number</b>	<b>Use</b>
Anti-Smad3 (phosphoS423+S425)	EP823Y	Abcam	ab52903	WB, PLA
Anti-Smad3	-	Abcam	ab52903	ChIP
Smad4 Monoclonal Antibody	4G1C6	ThermoFischer Scientific	MA5-15682	WB, PLA, ChIP
Anti-Smad4	EP618Y	Abcam	ab40759	ChIP
Anti-acetyl Histone H4	-	Merck	06-866	WB
Pac Blue anti-mouse CD4	GK1.5	Biolegend	100428	FACS
Alexa Fluor® 700 anti- mouse CD4	RM4-5	Biolegend	100536	FACS
PE.Cy5 anti-mouse CD8	53-6.7	Biolegend	100722	FACS
PE.Cy7 anti-mouse NK1.1	PK136	Biolegend	108713	FACS
PE anti-mouse CD19	6D5	Biolegend	115508	FACS
PE.Cy7 anti-mouse CD11b	M1/70	Biolegend	101216	FACS
PE.Cy7 anti-mouse CD11c	N418	Biolegend	117318	FACS
PE.Cy7 anti-mouse γδTCR	GL3	Biolegend	118123	FACS
PE anti-mouse TGFβRII	-	R&D Systems	FAB532P	FACS
BV421 anti-mouse IL- 17A	TC11- 18H10.1	Biolegend	506925	FACS
PE anti-mouse IFN-γ	XMG1.2	Biolegend	554412	FACS
FITC anti-human CD4	OKT4	Biolegend	317408	FACS
Alexa Fluor® 700 anti- human CD45RA	HI100	Biolegend	304120	FACS
PE/Cy7 anti-human CD127	019D5	Biolegend	351320	FACS
Brilliant Violet 650™ anti-human CD25	BC96	Biolegend	302633	FACS
Brilliant Violet 711™ anti-human IL-17A	BL168	Biolegend	512328	FACS
PE anti-human IL-10	JES3- 19F1	Biolegend	506804	FACS
Pacific Blue™ anti- mouse CD4	RM4-5	Biolegend	100531	FACS

WB: Western Blot

PLA: Proximity Ligation Assay

ChIP: Chromatin Immunoprecipitation

FACS: Fluorescent Activation Cell Sorting

**Supplementary Table 2: Table of primers**

<b>Name</b>	<b>Sequence</b>	<b>Amplicon size</b>	<b>Use</b>
<i>Il10pr Fw</i>	5' gttgcttctgctgttggaacg 3'	1447bp	PCR, CL
<i>Il10pr Rev</i>	5' gtagacctcctgttcttggtccc 3'		PCR, CL
<i>Il10pr-pGL2 Fw</i>	5' gcgtcgtagctcgaggttgcttctgctgttggaacg 3'	1477bp	PCR, CL
<i>Il10pr-pGL2 Rev</i>	5' cggaatgccaaagctttagacctcctgttcttggtccc 3'		PCR, CL
<i>Smad3 Fw</i>	5' atgctgcatcctgcccttcac 3'	1278bp	PCR, CL
<i>Smad3 Rev</i>	5' ctaagacacactggaacagcggatg 3'		PCR, CL
<i>Smad3-pT<sub>N</sub>T Fw</i>	5' cttttgcactcgagatgctgcatcctgcccttcac 3'	1308bp	PCR, CL
<i>Smad3-pT<sub>N</sub>T Rev</i>	5' cgggtcgactctagactaagacacactggaacagcggatg 3'		PCR, CL
<i>Smad4 Fw</i>	5' atggacaatatgtctataacaaatacaccaa 3'	1656bp	PCR, CL
<i>Smad4 Rev</i>	5' tcagtctaaaggctgtgggtccg 3'		PCR, CL
<i>Smad4-pTag2 Fw</i>	5' cgggctgcaggaattcatggacaatatgtctataacaaatacaccaa 3'	1687bp	PCR, CL
<i>Smad4-pTag2 Rev</i>	5' cgtatcgataagcttcagtctaaaggctgtgggtccg 3'		PCR, CL
<i>Il10A Fw</i>	5' caggttgagtggaggaaacaa 3'	138bp	ChIP
<i>Il10A Rev</i>	5' ggcagacagctgttctatgt 3'		ChIP
<i>Il10B Fw</i>	5' gccaggtacagaatgaaa 3'	173bp	ChIP
<i>Il10B Rev</i>	5' agctgtgaaggatggagatg 3'		ChIP
<i>Il10C Fw</i>	5' agccattatccacgctatta 3'	94bp	ChIP
<i>Il10C Rev</i>	5' actggtcggaatgaacttctg 3'		ChIP
<i>Gapdh Fw</i>	5' accagggagggtgcagtcc 3'	237bp	ChIP
<i>Gapdh Rev</i>	5' tcagttcggagcccacaggc 3'		ChIP

PCR: Polymerase Chain Reaction

CL: Cloning

ChIP: Chromatin Immunoprecipitation