Supplementary figures

Ndfip1 restricts Th17 cell potency by limiting lineage stability and proinflammatory cytokine production.

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C IL-17A







Ndfip1 fl/fl CD4 Cre- (WT)

Ndfip1 fl/fl CD4 Cre+ (cKO)











a





rorc

IL-4KO DKO

0.003-

0.002-

0.001

0.000-

















relative expression

Supplementary Figure S1. Ndfip1 limits the abundance of Th17 cells.

Flow cytometric analysis of CD3+ CD4+ T cells among cells isolated from lungs of the wild type (Cre-) or Ndfip1fl/fl CD4-Cre + (cKO) animals shown in Figure 1, but focused only on litter-matched Cre- and Cre+ controls (i.e. litters in which Cre- and Cre+ mice came from the same mothers). (a-b) Percentages of lung CD4+ T cells that are IL-17A+ (a) or IFN γ + (b). (c-d) Percentages of CD44+ CD4+ cells that are IL-17A+ (c) or IFN γ + (d). n= 4 Cre- and n=5 Cre+ in two independent experiments. *p<0.05, **p<0.01,****p<0.0001. p values were calculated by unpaired two-tailed T tests. All error bars represent mean +/- SEM

Supplementary Figure S2. Ndfip1 does not restrain IFN γ secretion from restimulated Th1 cells. (a) Representative plot showing Tbet+ IFN γ + Th1 cells after Th1 polarization and IL-2 expansion of IL-4-/- or Ndfip1 IL-4 DKO CD4+ T cells. (b) Summary of IFN γ ELISAs from plated Th1 cells. Data is pooled for n=2 Ndfip1 IL-4 DKO and n=3 IL-4-/- animals and is analyzed by 2-way ANOVA. *p<0.05, **p<0.01,***p<0.001. Error bars represent mean +/- SEM

Supplementary Figure S3. Loss of Ndfip1 in T cells drives a spontaneous colitis in mice (a) Colon histology of ten week old Ndfip1 fl/fl CD4 Cre+ (cKO) or WT (Ndfip1 fl/fl Cre- or Ndfip1 fl/+ Cre-) animals showing evidence of spontaneous colon inflammation in cKO animals. Bars represent 100um and inset shows enlarged version of the boxed section of the image. (b) Ndfip1 fl/fl CD4 Cre+ (cKO) and WT animals do not show differences in colon length. (c) Flow cytometry data indicating a trend towards increased CD11b+ CD11c- Ly6G+ neutrophils in the colons of Ndfip1 fl/fl CD4 Cre+ (cKO) versus WT animals. Data is shown for an n= 3 Ndfip1 fl/fl CD4 Cre + and n=3 WT (Ndfip1 fl/fl Cre- or Ndfip1 fl/+ Cre-) animals at ten weeks old. All error bars represent mean +/- SEM

Supplementary Figure S4. Th17 cells lacking Ndfip1 show a gene expression profile associated with Th17 pathogenicity.

Naïve sorted CD4+ T cells were differentiated into Th17 cells for 5 days. Cells were harvested and expanded in IL-2 for 3 days and then used for qPCR gene expression analysis shown here and also used in the ELISA studies in Figure 4. Data is shown for n=4 mice per genotype in 2 independent differentiation experiments. All error bars represent mean +/- SEM