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Supplemental information

Microbiota-mediated skewing of tryptophan

catabolism modulates CD4⁺ T cells

in lupus-prone mice

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Α

в

1616_Fecal_super

414_Fecal_supe Pool5p.mz XMLPeakheight 1717_Fecal_sup 1818_Fecal_sup 1919_Fecal_supe 2121_Fecal_sup

Pool6p.mz XMLPeakheigl

020_Fecal_supr

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r dues

2222_Feca 2424_Fecal_supe I Pool8p 1313_Fecal_sup

323_Fecal_sup

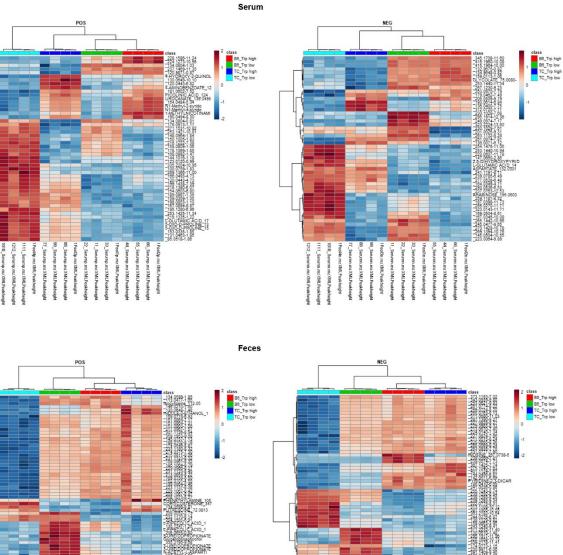


Figure S1. Heat maps of the top 50 metabolites as determined from ANOVA from serum) and feces from B6 and TC mice fed high or low tryptophan for 1 month. Related to Figure 1.

424_Feca

heigh

222 Feca 323_recal_supe 313_Fecal 515 Fecal 414_Fecal_supe 616_Fecal_supern 818_Fecal_sup 717_Fecal_supern

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ool5n.mz XMLPea

1919_Fecal_supern

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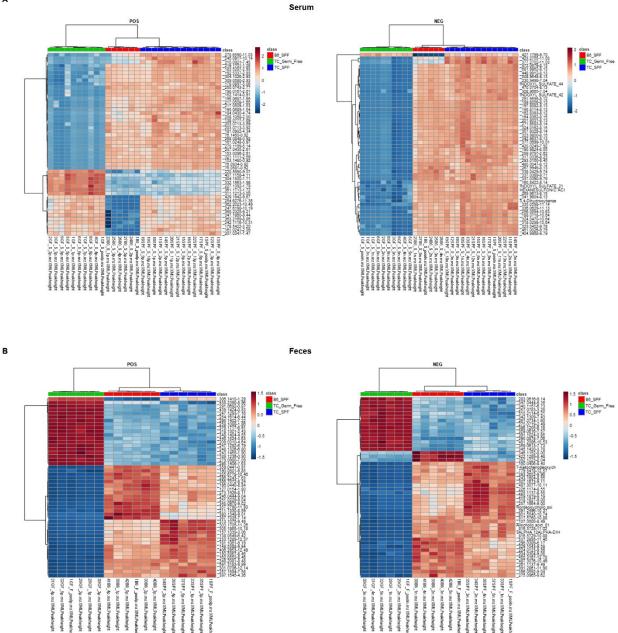


Figure S2. Heat maps of the top 50 metabolites as determined from ANOVA from serum and fecal metabolite profiles in GF TC, SPF TC, and SPF B6 mice. Related to Figure 3.

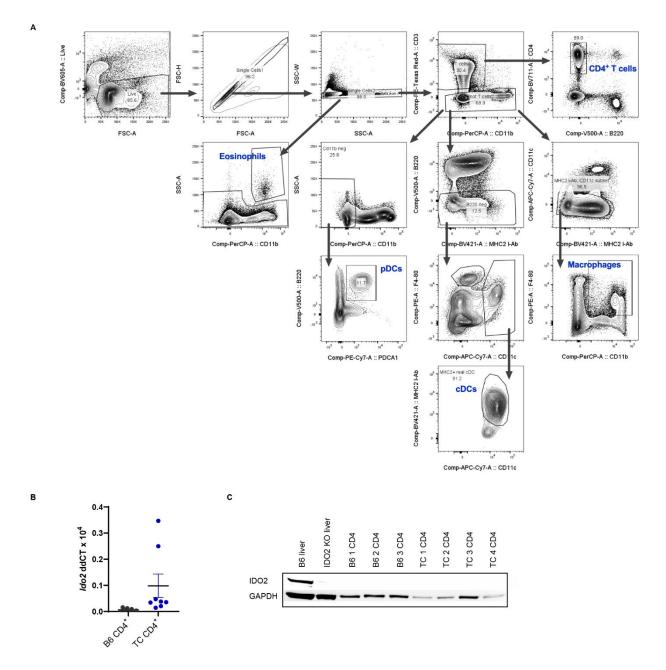


Figure S3. A. Gating strategy for splenocytes populations and *Ido2* gene expression. Related to Figure 4.

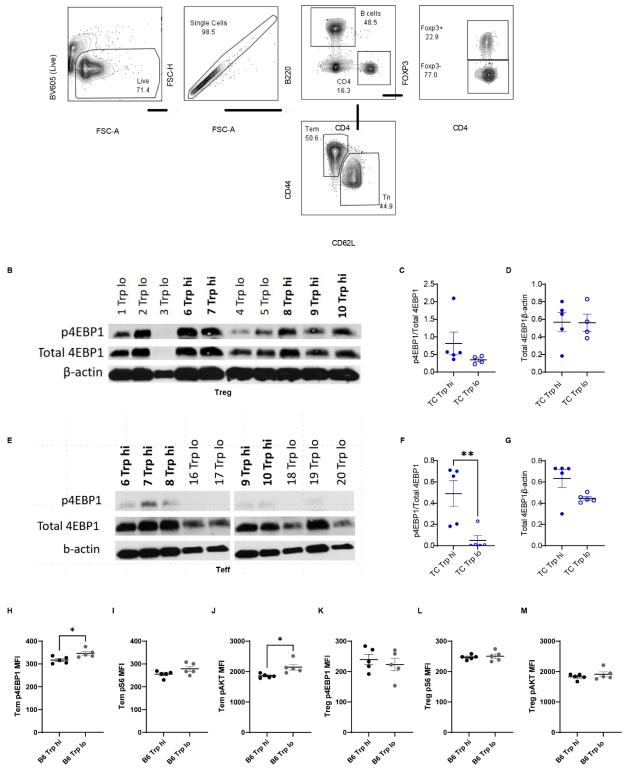


Figure S4. Gating strategy for Treg, Tn and Tem cells and expression of targets of mTOR signaling. Related to Figure 6.

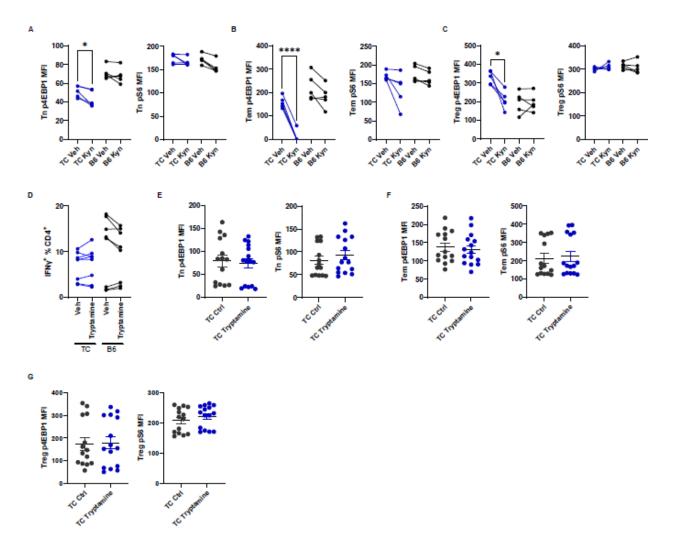


Figure S5. Effect of kynurenine and tryptamine on TC and B6 T cell phenotypes. Related to Figure 7.

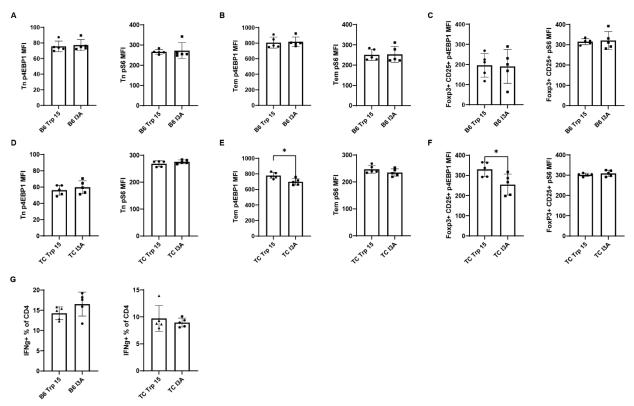


Figure S6. Effect of I3A on TC and B6 T cell phenotypes. Related to Figure 7.