

Figure S1. Sunitinib impairs the host defense against intestinal C. rodentium infection.

(A) 8 weeks old wild type mice were orally inoculated with 2×10^9 CFU of *C. rodentium*. Sunitinib (4 mg/Kg, n=6 or 40 mg/Kg, n=8) or control (n=5) was administered orally from day 0 to day 7, once daily. Histological analysis of representative colons from naïve, control or Sunitinib (40 mg/Kg) treated mice at day 11 after infection were shown. H&E staining shows severely disruption of the colonic epithelial layer and massive cell infiltration in Sunitinib treated mice. Scale bars, 50 µm. Data are representative of two independent experiments. Figure S1, related to Figure 1.



Figure S2. STAT3 signaling from RORyt⁺ lymphocytes controls *C. rodentium* infection.

(A-C) Conditional deletion of STAT3 in ROR γ t⁺ lymphocytes. (A) Colonic LPLs were isolated from 6 to 8 week-old *Stat3*^{fl/fl} and *Rorc-cre-Stat3*^{fl/fl} littermate mice and intracellular stained with STAT3 and ROR γ t. STAT3 expression was analyzed by flow cytometry after gating on CD3⁻ ROR γ t⁺ CD45⁺, CD3⁻ ROR γ t⁻ CD45⁺ and CD3⁺ CD45⁺ cells. Data are representative of two experiments with 2-3 mice for each genotype. (B) The STAT3 protein from the colon, liver, CD4⁺ and CD4⁻ splenocytes of WT (*Stat3*^{fl/fl}) or KO (*Rorc-cre-Stat3*^{fl/fl}) were examined by western blot. GAPDH was as internal control. Data are representative of two independent experiments. (C) Different ILCs were sorted from the intestine LPLs of *Rorc-cre-Stat3*^{fl/fl} *Rorc*^{gfp/+} and *Rorc*^{gfp/+} mice. ILC1, CD90⁺CD3⁻NK1.1⁺ROR γ t⁻; ILC2, CD90⁺CD3⁻NK1.1⁺ROR γ t⁻ KLRG1⁺; ILC3, CD90⁺CD3⁻ROR γ t⁺; other ILCs, CD90⁺CD3⁻NK1.1⁺ROR γ t⁻ KLRG1⁻. The mRNA levels of STAT3 were examined by realtime PCR. Data are representative of two independent experiments.

(**D**, **E**) 7 weeks old *Rorc-cre-Stat3*^{fl/fl}, wild type *Stat3*^{fl/fl} and *Rorc-cre-Stat3*^{fl/+} heterozygous mice (n = 5 per group) were orally inoculated with *C. rodentium* (2 × 10⁹ CFU) and body weight was measured at the indicated time points. (**D**) Body weight change and survival rates are shown. (**E**) Bacterial titers from fecal homogenate cultures and blood at indicated day post infection in the mice in **D**. *P<0.05, **P<0.01, ***P<0.001; ns, no significant difference (Student's *t*-test); nd, nondetectable. Data are representative of two independent experiments (mean \pm s.e.m.).

(**F**) Histological analysis of representative colons from $Stat3^{fl/fl}$ and $Rorc-cre-Stat3^{fl/fl}$ mice at day 8 after infection. *Rorc-cre-Stat3*^{fl/fl} and their littermate wild type $Stat3^{fl/fl}$ mice were orally inoculated with high dose (2 × 10⁹ CFU) of *C. rodentium*. Naïve $Stat3^{fl/fl}$ was as a control. H&E

staining shows severely disruption of the colonic epithelial layer and massive cell infiltration in *Rorc-cre-Stat3*^{fl/fl} mice. Scale bars, 50 μ m. Figure S2, related to Figure 2.



Figure S3. STAT3 signaling is required for IL-22 and IL-17 production from both innate and adaptive lymphocytes.

(A-D) IL-22 and IL-17 expression in CD3⁻ and CD3⁺ cells were analyzed by intracellular cytokine staining followed by flow cytometry. Intestinal lamina proprial lymphocytes (LPL) were isolated from the colons of *Stat3*^{fl/fl} or *Rorc-cre-Stat3*^{fl/fl} mice at day 5 post-infection and stimulated with IL-23 (25 ng/ml), PMA (50 ng/ml) and ionomycin (750 ng/ml) for 4 hours. Naïve mice were used as control.

(A, C) Percentage of IL-22⁺ or IL-17⁺ cells in the CD3⁻ Thy1⁺ ILCs population, as well as the absolute numbers of IL-22⁺ or IL-17⁺ ILCs in the colons of naïve or *C. rodentium* infected *Stat3*^{fl/fl} and *Rorc-cre-Stat3*^{fl/fl} mice are shown.

(**B**, **D**) Percentage of IL-22⁺ or IL-17⁺ cells in the CD3⁺ Thy1⁺ T cell population, as well as the absolute numbers of IL-22⁺ or IL-17⁺ T cells in the colons of naïve or *C. rodentium* infected *Stat3*^{fl/fl} and *Rorc-cre-Stat3*^{fl/fl} mice are shown. *P<0.05, **P<0.01, ***P<0.001 (Student's *t*-test). Each dot represents one individual mouse. Data are representative of two independent experiments (**A-D**; mean \pm s.e.m.).

(E) Sunitinib suppresses IL-22 production in innate lymphocytes. Colonic LPLs were isolated from 6 to 8 week-old wild type mice were stimulated with IL-23 or IL-23 and Sunitinib (10 μ M) for 4 hours. IL-22 expression in CD3⁻ Thy1⁺ cells were analyzed by intracellular cytokine staining followed by flow cytometry. Data are representative of two independent experiments.

(F) LPLs were isolated from the colons of $Stat3^{fl/fl}$ or *Rorc-cre-Stat3*^{fl/fl} mice at day 8 postinfection and stimulated with either IL-23 (25 ng/ml), PMA (50 ng/ml) and ionomycin (750 ng/ml) or IL-23 (25 ng/ml), anti-CD3 (plate coated, 1 µg/ml) and anti-CD28 (2 µg/ml) for 4 hours. IL-22 and IL-17 expression in CD3⁺CD4⁺ T cells were analyzed by intracellular cytokine staining. Data are representative of two independent experiments. Figure S3, related to Figure 3.



Figure S4. CD4⁺ T cells but not STAT3 signaling in T cells are required for the production of *C. rodentium* specific IgG.

(**A**, **B**) Wild type *Stat3*^{fl/fl} mice were infected with 5×10^6 CFU of *C. rodentium* and injected intraperitoneally with mAb GK1.5 (n=5) or Rat IgG (n=4) (50 µg per mouse each time) at day -5, 0, 5, 10, 15 post infection for depletion of CD4⁺ T cells. (**A**) The depletion efficiency was examined at day 8 post infection. (**B**) Blood samples were collected at day 12 post-infection and serum *C. rodentium* specific IgG was determined by ELISA.

(**C**, **D**) 7 weeks old *Rorc-cre-Stat3*^{fl/fl} and wild type *Stat3*^{fl/fl} mice (n = 5 per group) were orally inoculated with *C. rodentium* (2 × 10⁹ CFU). One group of *Rorc-cre-Stat3*^{fl/fl} mice were rescued by IL-22-Ig fusion protein injection. At day 60 after first *C. rodentium* infection, IL-22-rescued *Rorc-cre-Stat3*^{fl/fl} and wild type *Stat3*^{fl/fl} mice were challenged with high dose of *C. rodentium* (2 × 10⁹ CFU) again. (**A**) Blood samples were collected at day 14 post-infection and serum *C. rodentium* specific IgG was determined by ELISA. (**B**) Body weight was measured at the indicated time points. ***P<0.001 (Student's *t*-test). Data are representative of two independent experiments.

(**E**, **F**) *Cd4-cre-Stat3*^{fl/fl} and wild type *Stat3*^{fl/fl} mice were infected with 5×10^6 CFU of *C*. *rodentium* and injected intraperitoneally with mAb GK1.5 or Rat IgG (50 µg per mouse each time) at day -5, 0, 5, 10 and 15 post infection for depletion of CD4⁺ T cells. Average body weight change (**E**) and survival rates (**F**) at the indicated time points are shown. Figure S4, related to Figure 4.



Figure S5. Intestinal CD90^{hi}CD45^{lo} LPLs are RORyt⁺ ILCs.

LPLs were isolated from both the large intestine and small intestine of wild type C57BL/6 mice and gated by CD45 and CD90 expression. Two separate populations, CD90^{hi}CD45^{lo} and CD90^{lo}CD45^{hi} were further analyzed with the expression of CD3 and RORyt. Data are representative of two independent experiments. Figure S5, related to Figure 5.



Figure S6. The development of $ROR\gamma t^+$ Th cells, but not $ROR\gamma t^+$ ILCs, requires STAT3 signaling.

(A) Colonic LPLs were isolated from $Stat3^{fl/fl}$ and Cd4-cre-Stat3^{fl/fl} littermate mice. ROR γ t expression were analyzed in colonic LPLs by flow cytometry after gating on CD4⁺ TCR β^+ , CD4⁻ TCR β^+ adaptive lymphocytes and CD4⁺ TCR β^- , CD4⁻ TCR β^- innate lymphoid cells.

(**B**, **C**) LPLs were isolated from both large and small intestine of *Rorc-cre-Stat3*^{fl/fl}*Rorc*^{gfp/+} and *Rorc*^{gfp/+} mice. ROR γ t⁺ T cells, NCR⁺ROR γ t⁺ ILC3, NCR⁻ROR γ t⁺ ILC3, NCR⁺ROR γ t⁻ ILC1 and ROR γ t⁻KLRG1⁺ ILC2 were analyzed by flow cytometry. Figure S6, related to Figure 6.

<i>ll</i> 22 FW	CATGCAGGAGGTGGTACCTT
1122 RV	CAGACGCAAGCATTTCTCAG
1123p19 FW	GGTGGCTCAGGGAAATGT
1123p19 RV	GACAGAGCAGGCAGGTACAG
1123p40 FW	ACATCTACCGAAGTCCAATGCA
1123p40 RV	GGAATTGTAATAGCGATCCTGAGC
ll23r FW	TCAGTGCTACAATCTTCAGAGGACA
ll23r RV	GCCAAGAAGACCATTCCCGA
Rorc FW	TACCTTGGCCAAAACAGAGG
Rorc RV	ATGCCTGGTTTCCTCAAAA
Ahr FW	GGCTTTCAGCAGTCTGATGTC
Ahr RV	CATGAAAGAAGCGTTCTCTGG
Reg3g FW	CAAGGTGAAGTTGCCAAGAA
Reg3g RV	CCTCTGTTGGGTTCATAGCC
Reg3b FW	ATGGCTCCTACTGCTATGCC
Reg3b RV	GTGTCCTCCAGGCCTCTT
Hprt FW	TGAAGAGCTACTGTAATGATCAGTCA
Hprt RV	AGCAAGCTTGCAACCTTAACCA
<i>ll17a</i> FW	CTCCAGAAGGCCCTCAGACTAC
<i>ll17</i> a RV	AGCTTTCCCTCCGCATTGACACAG
Stat3 FW	GAAGACCAAGTTCATCTGTGTGAC
Stat3 RV	GGTTTCAGCTCCTCACATGG
IL22 pro FW	GGGAGATCAAAGGCTGCTCTA
ll22 pro RV	CCACCTTGAGAGATGGGAAGT
II22 Intron FW	AGCTGCATCTCTTTCTCTCCA
II22 Intron RV	TATCCTGAAGGCCAAAATAGGA
ll17 pro FW	GCCTTTGTGATTGTTTCTTGCAG
ll17pro RV	CCTTGCCCAAAGAAACCCACTC
Ltbr FW	CAGTGGCTCCAAGTGGCTTG
Ltbr RV	GCAAACCGTGTCTTGGCTGC

Table S1Primers for realtime RT-PCR and ChIP assays.

SUPPLEMENTAL EXPERIMENTAL PROCEDURES

C. rodentium specific IgG

Blood samples were also collected and serum *C. rodentium* specific IgG was determined by ELISA with *C. rodentium* lysate and anti-mouse IgG (Jackson ImmunoResearch).

Plasmids and Retroviral Transduction of Cell Lines

STAT3C (constitutively active forms of STAT3) (Zhou et al., 2007) was cloned into a DFTC (double flag epitope tagged)-retroviral vector containing an IRES-puromycin resistant gene. Plasmid DNA was confirmed by sequencing. HEK293T cells were transfected with retroviral plasmids and the packaging plasmid 10A1 using Polyethylenimine (PEI). Viral supernatant was collected after transfection. EL4 cells were spun in 2 ml of virus supernatant containing 8 μ g/ml polybrene at 2500 rpm for 1.5 hr at 30 °C. The retroviral transduction was repeated 24 h later. The cells were then selected in the presence of 7 μ g/ml puromycin for 5 days (Qiu et al., 2011).

Western Blot

Colon and liver tissues were washed in PBS and homogenized in RIPA lysis buffer. Protein samples were resolved by SDS–PAGE and transferred to a nitrocellulose membrane. Western blotting was performed with anti-STAT3 rabbit mAb (D3Z2G) (Cell Signaling Technology) and a monoclonal anti-GAPDH antibody (Sigma).

SUPPLEMENTAL REFERENCE

Qiu, J., Heller, J.J., Guo, X., Chen, Z.M., Fish, K., Fu, Y.X., and Zhou, L. (2011). The aryl hydrocarbon receptor regulates gut immunity through modulation of innate lymphoid cells. Immunity *36*, 92-104.

Zhou, L., Ivanov, II, Spolski, R., Min, R., Shenderov, K., Egawa, T., Levy, D.E., Leonard, W.J., and Littman, D.R. (2007). IL-6 programs T(H)-17 cell differentiation by promoting sequential engagement of the IL-21 and IL-23 pathways. Nat Immunol 8, 967-974.