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Commensal-dependent expression of IL-25 regulates the IL-23–IL-17 axis in the intestine

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Alterations in the composition of intestinal commensal bacteria are associated with enhanced susceptibility to multiple inflammatory diseases, including those conditions associated with interleukin (IL)-17–producing CD4⁺ T helper (Th17) cells. However, the relationship between commensal bacteria and the expression of proinflammatory cytokines remains unclear. Using germ-free mice, we show that the frequency of Th17 cells in the large intestine is significantly elevated in the absence of commensal bacteria. Commensal-dependent expression of the IL-17 family member IL-25 (IL-17E) by intestinal epithelial cells limits the expansion of Th17 cells in the intestine by inhibiting expression of macrophage-derived IL-23. We propose that acquisition of, or alterations in, commensal bacteria influences intestinal immune homeostasis via direct regulation of the IL-25–IL-23–IL-17 axis.

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The intestinal tract is a major site of colonization by commensal bacteria (1, 2). This bacterial population is acquired shortly after birth, is estimated to number 10¹⁴ organisms, and is remarkably diverse, being composed of at least 500–1,000 individual species (3). The host–commensal relationship is the product of millions of years of coevolution, and it can be influenced by the number and composition of bacterial species within the intestine. For example, studies using germ-free (GF) animals demonstrated that commensal bacteria are required for normal development of the immune system (2). Commensal bacteria have also been implicated in the pathogenesis of inflammatory bowel disease (IBD), as colitis-prone mice fail to develop intestinal inflammation if reared under GF conditions and patients suffering from IBD exhibit dysregulated immune responses against commensal bacteria (4, 5).

IL-17–producing CD4⁺ T (Th17) cells play a significant role in the pathogenesis of IBD (6–8).

The differentiation of Th17 cells is promoted by IL-6– and TGF- β –dependent expression of the transcription factor retinoic acid–related orphan nuclear receptor- γ t (ROR γ t [Rorc]), whereas IL-23 controls their expansion or survival (7, 9). Th17 cells are found constitutively in the small intestine of naive mice housed under conventional conditions (9), suggesting the presence of commensal bacteria may promote their development or maintenance in the intestine (9–11). An alternative model proposes that rather than promoting the presence of effector CD4⁺ T cell populations and cytokines in the intestine, commensal bacteria are recognized by Toll-like receptors, and MyD88-dependent NF- κ B activation results in the maintenance of epithelial integrity that is associated with decreased proinflammatory cytokine and chemokine gene expression, leading to a state of immune hyporesponsiveness in the intestine (12–14). Consistent with this hypothesis, mice depleted of commensal bacteria by antibiotic treatment are highly

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susceptible to chemically induced colitis (12), and disruption of the NF-κB pathway in IECs results in exaggerated expression of proinflammatory cytokines and the development of spontaneous (15) and infection-induced intestinal inflammation before disruption of the epithelial barrier (16).

In this study, we show that intestinal commensal bacteria regulate expression of the IL-17 family of cytokines in the intestine. The frequency of Th17 cells was significantly increased in the large intestine of GF mice compared with conventionally reared (CNV) mice and was associated with heightened levels of IL-23. Expression of the IL-17 family member IL-25 (IL-17E) by intestinal epithelial cells (IECs) was dependent on the presence of commensal bacteria and administration of IL-25 to GF mice reduced expression of IL-23, and the frequency of Th17 cells in the large intestine. Consistent with a role for IL-25 in limiting IL-23 and Th17 cells in the large intestine, neutralization of IL-23 also resulted in decreased frequencies of Th17 cells in the large intestine of GF mice. Further, IL-25 directly inhibited LPS-induced IL-23 expression

by macrophages in a STAT6-independent manner. Thus, these results demonstrate a functional link between commensal bacteria and the IL-25–IL-23–IL-17 axis and identify a novel commensal-dependent mechanism in the regulation of Th17 cells in the intestinal microenvironment.

RESULTS AND DISCUSSION

The role of commensal bacteria in intestinal immune homeostasis remains unclear. One study identified the constitutive presence of Th17 cells primarily in the small intestinal lamina propria (LP) and hypothesized that commensal bacteria may be required for their presence in the intestine (9), whereas another reported that Th17 cells are found only in the large intestinal LP of CNV mice (17). We sought to test whether signals derived from commensal bacteria promote or inhibit the presence of Th17 cells in the intestine. We examined the frequency of Th17 cells in tissues isolated from CNV BALB/c or GF BALB/c mice by flow cytometry. In CNV mice, Th17 cells were present primarily in the LP of the small intestine

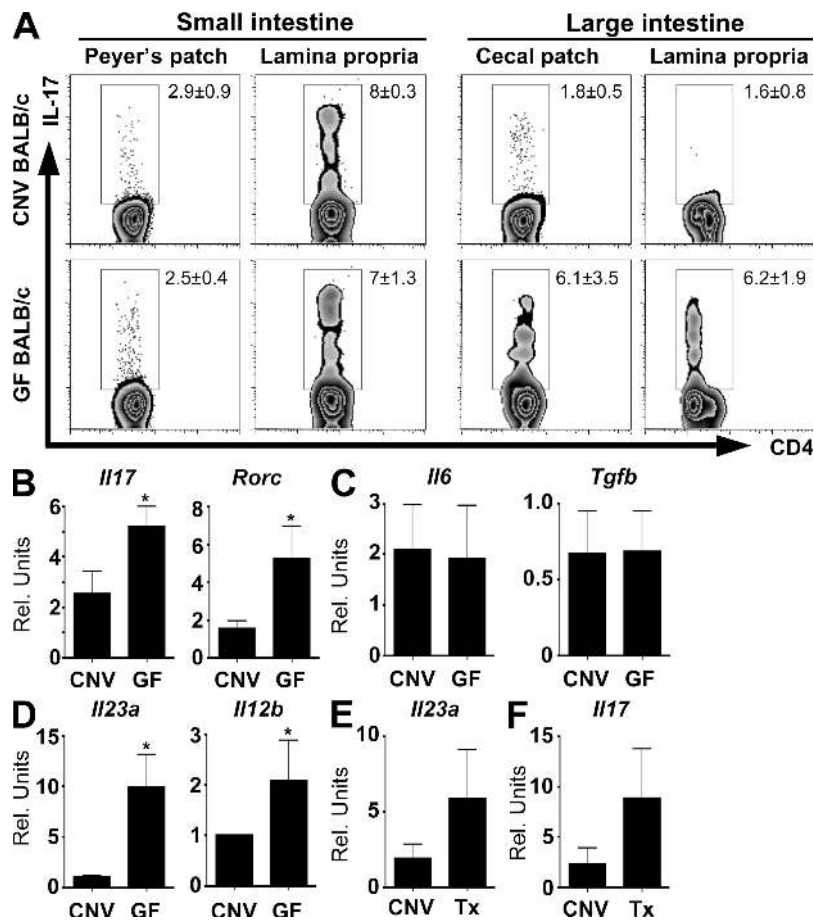


Figure 1. Enteric commensal bacteria are required to limit the frequency of Th17 cells in the intestine. (A) Expression of IL-17 by CD4⁺ T cells in the Peyer's patches, cecal patch, and LP of the small or large intestine of CNV and GF mice was analyzed by flow cytometry. Flow cytometry plots depict log10 fluorescence. (B–D) Expression of mRNA for *Il17* and *Rorc* (B), *Il6* and *Tgfb* (C), or *Il23a* and *Il12b* (D) in the large intestine of CNV and GF mice was analyzed by quantitative real-time PCR. (E and F) Expression of mRNA for *Il23a* (E) and *Il17* (F) in the large intestine of CNV and antibiotic-treated (Tx) CNV mice was analyzed by quantitative real-time PCR. Results are from 3 experiments ($n = 6–9$). Rel. units, relative units. *, $P < 0.05$. Error bars indicate the SEM.

(Fig. 1 A). Comparison of CNV and GF mice revealed equivalent frequencies of Th17 cells in the Peyer's patches and small intestinal LP of CNV and GF mice (Fig. 1 A). In contrast, the frequency of Th17 cells in the cecal patch (a lymphoid follicle associated with the cecum that is analogous to the human appendix) and large intestinal LP was three- to fourfold higher in GF mice compared with CNV animals (Fig. 1 A). Analysis of messenger RNA (mRNA) expression in whole sections of large intestinal tissue isolated from CNV or GF mice demonstrated elevated expression levels of mRNA for *Il17* and *Rorc* (Fig. 1 B). Elevated levels of *Il17* mRNA were independent of genetic background, as GF C57BL/6 mice also expressed significantly higher levels of *Il17* than CNV C57BL/6 mice (Fig. S1, available at <http://www.jem.org/cgi/content/full/jem.20080720/DC1>). These results implicate signals derived from commensal bacteria in limiting the frequency of intestinal Th17 cells.

The differentiation of Th17 cells is promoted by IL-6 and TGF- β , whereas IL-23 is required for the subsequent expansion or survival of committed Th17 cells (7). To test whether the absence of commensal bacteria influenced the development or survival/expansion of Th17 cells, we examined ex-

pression of IL-6, TGF- β , and IL-23 in the intestine of CNV and GF mice. Although the levels of *Il6* and *Tgfb* mRNA were equivalent in the large intestine of mice housed under both conditions (Fig. 1 C), the levels of IL-23p19 (*Il23a*) and IL-12/23p40 (*Il12b*) mRNA were significantly higher in the large intestine of GF mice compared with CNV mice (Fig. 1 D). Secretion of IL-6 and IL-12/23p40 protein in large intestinal explant cultures confirmed mRNA expression patterns, with equivalent production of IL-6 (Fig. S2 A, available at <http://www.jem.org/cgi/content/full/jem.20080720/DC1>) by intestinal tissue of CNV and GF mice and increased production of IL-12/23p40 by GF intestinal tissue (Fig. S2 B). Expression of *Il23a* was also significantly increased in the large intestine of GF C57BL/6 mice (Fig. S3). These findings suggest that the commensal bacteria do not influence expression of cytokines that govern Th17 cell differentiation from naive precursors, but rather signals derived from commensal bacteria inhibit expression of IL-23 and the subsequent expansion and/or survival of Th17 cells in the large intestine.

An alternative approach was adopted to determine whether there was a link between commensal bacteria and expression of IL-23 and -17 in CNV mice. Oral administration of antibiotics

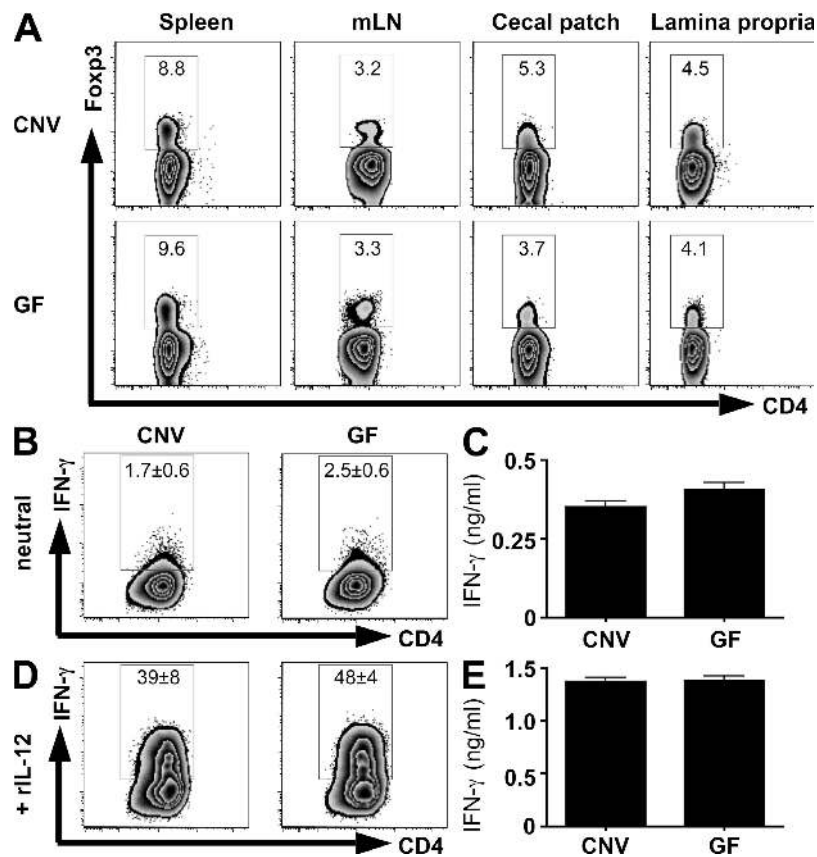


Figure 2. Equivalent regulatory T cell and Th1 cell responses in CNV and GF mice. (A) Cells isolated from the spleen, mLN, cecal patch, or large intestinal LP were stained for expression of CD4 and Fc γ 3 and analyzed by flow cytometry. (B–E) Splenocytes from CNV or GF mice were stimulated with anti-CD3 and -CD28 in the absence (B and C) or presence (D and E) of 1 ng/ml IL-12 for 72 h, followed by incubation with PMA, ionomycin, and brefeldin A for the final 5 h. Harvested cells were stained for expression of CD4 and IFN- γ by flow cytometry (B and D), and cell-free supernatants were analyzed by ELISA (C and E). Flow cytometry plots depict log₁₀ fluorescence. Data are representative of 2 experiments ($n = 8$). Error bars indicate the SEM.

has been shown to effectively reduce the numbers of bacteria in the intestine and leads to increased susceptibility to inflammation and food allergy (12, 18). To test whether heightened susceptibility to inflammation after antibiotic treatment was associated with altered expression of the IL-23–IL-17 axis, mice were treated orally with antibiotics for 6 wk, and segments of small and large intestine were analyzed for expression of *Il23a* and *Il17*. Antibiotic treatment of adult mice resulted in increased levels of *Il23a* and *Il17* in the large intestine (Fig. 1, E and F). Critically, this effect was confined to the large intestine, as there were no significant changes in expression of *Il23a*, *Il12b*, or *Il17* in the small intestine (Fig. S4, A–C, available at <http://www.jem.org/cgi/content/full/jem.20080720/DC1>). Thus, reduction in total numbers of commensal bacteria in adult mice results in heightened expression of IL-23 and -17 in the large intestine, suggesting that commensal bacteria are actively promoting inhibition of IL-23 and -17.

Recent studies identified a reciprocal developmental pathway between Th17 cells and Foxp3-expressing regulatory T cells (19). Therefore, we examined whether the elevated frequency of Th17 cells in GF mice was associated with dysregulation of regulatory T cells. No significant differences in the frequency of CD4⁺ Foxp3⁺ regulatory T cells (Fig. 2 A) or IL-10–producing CD4⁺ T cells in the mesenteric LN (mLN), cecal patch, or large intestinal LP were observed between mice housed under CNV or GF conditions (Fig. S5, A–C, available at <http://www.jem.org/cgi/content/full/jem.20080720/DC1>). Additionally, there were no differences in the expression levels of *Il10* mRNA in the large intestine between CNV and GF mice (Fig. S5 D). Consistent with normal frequencies of CD4⁺ Foxp3⁺ T cells in GF mice, there was no evidence of global dysregulation in expression of proinflammatory cytokines. Splenocytes from both CNV and GF mice produced equivalent levels of IFN- γ under neutral conditions (Fig. 2, B and C) or in the presence of exogenous IL-12 (Fig. 2, D and E). Further, there were no significant differences in the expression of IL-12p35 (*Il12a*) mRNA (Fig. S6 A) or the frequencies of CD4⁺ T cells producing IFN- γ in the mLN, cecal patch, and large intestinal LP of CNV and GF mice (Fig. S6 B). Collectively, these results suggest that the absence of commensal bacteria is associated with the selective overexpression of IL-23 and enhanced persistence, survival, or recruitment of Th17 cells in the intestinal microenvironment, rather than a general loss of regulatory T cell- and IL-10–dependent immune regulation.

The absence of commensal bacteria resulted in elevated expression of IL-23 and an increased frequency of Th17 cells in the large intestine. Based on these findings, we sought to test the hypothesis that commensal bacteria promote expression of cytokines that could negatively regulate the frequency of Th17 cells in the intestine. IL-27 and -25 have both been shown to limit Th17 cell responses (20–23). IL-27 acts directly on CD4⁺ T cells to inhibit their differentiation into the Th17 cell lineage (24), whereas IL-25, which can be produced by a variety of cell types, including epithelial cells, mast cells, macrophages, and T cells, limits Th17 cell development by

promoting expression of IL-13 that acts on DCs (22, 25–28). We observed no significant difference in mRNA expression levels of the IL-27 subunits p28 (*Il27p28*) and Epstein-Barr virus–induced (EBI)-3 (*Ebi3*) in the large intestine between CNV and GF mice (Fig. 3 A). In contrast, there was a significant reduction in the levels of *Il25* mRNA in large intestinal tissue isolated from GF mice compared with CNV mice (Fig. 3 B). Consistent with a role for commensal bacteria in the expression of *Il25*, antibiotic treatment of CNV mice resulted in decreased expression of *Il25* in the large intestine (Fig. S7, available at <http://www.jem.org/cgi/content/full/jem.20080720/DC1>). Expression of *Il25* was detected in purified IECs from CNV but not GF mice (Fig. 3 C), suggesting that signals from commensal bacteria may be acting directly on IECs to induce expression of IL-25. Although we did not detect *Il25* expression in the spleen, LNs, Peyer's patches, or LP in CNV or GF mice (unpublished data), we cannot rule out the possibility that other resident or infiltrating LP cell populations could also express *Il25* in the intestine. Thus, absence of commensal bacteria was associated with

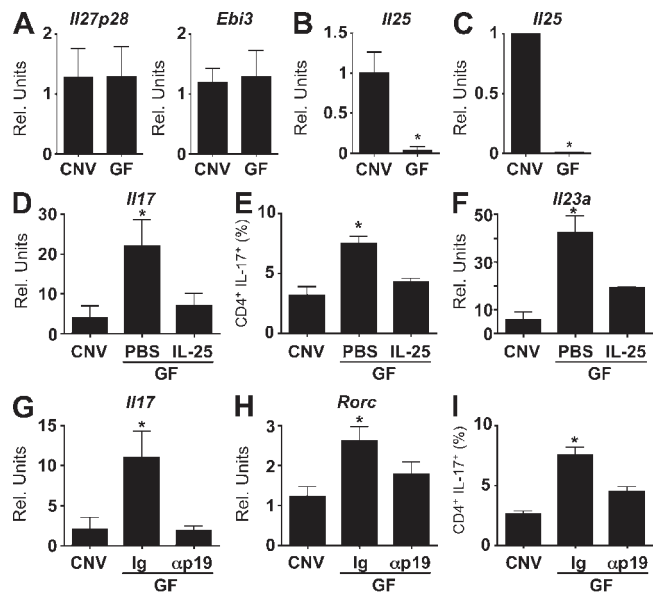


Figure 3. Commensal-dependent expression of IL-25 inhibits IL-23 and the frequency of Th17 cells in the large intestine. (A) Expression of mRNA for *Il27p28* and *Ebi3* in the large intestine of CNV and GF mice was analyzed by quantitative real-time PCR. (B and C) Expression of mRNA for *Il25* in the large intestine (B) and purified IECs (C) of CNV and GF mice was analyzed by quantitative real-time PCR. Data are from 3 experiments ($n = 12$). (D–F) Expression of mRNA for *Il17* (D), the frequency of IL-17⁺ CD4⁺ cells (E), and expression of mRNA for *Il23a* (F) in the large intestine of CNV mice, GF mice, or GF mice treated with IL-25 (0.5 μ g daily for 7 d) was analyzed by quantitative real-time PCR. Data are from 2 experiments ($n = 6$). (G–I) Expression of mRNA for *Il17* (G) and *Rorc* (H) and the frequency of IL-17⁺ CD4⁺ cells (I) in the large intestine of CNV mice, GF mice, or GF mice treated with a neutralizing monoclonal antibody against IL-23p19 (α p19; 1 mg daily for 7 d) was analyzed by quantitative real-time PCR. Data are from 2 experiments ($n = 4–6$). Rel. units, relative units. *, $P < 0.05$. Error bars indicate the SEM.

decreased expression of IL-25 in IECs, heightened IL-23, and increased frequencies of Th17 cells in the large intestine.

Based on these findings, we next sought to examine whether there was a functional interaction between reduced expression of IL-25 and the exaggerated production of IL-23 and -17 in the intestine. To test this, exogenous IL-25 was administered to GF mice. IL-25 treatment resulted in decreased expression of *Il17* (Fig. 3 D) and lower frequencies of Th17 cells in the LP of the large intestine (Fig. 3 E). IL-25 treatment also resulted in significantly decreased expression of *Il23a* mRNA (Fig. 3 F). The association between administration of IL-25 and reduced levels of IL-23 and -17 in GF mice suggested that the effects of IL-25 were mediated through the inhibition of IL-23. To directly address whether dysregulated expression of IL-23 in GF mice resulted in the increased frequency of Th17 cells observed, GF mice were treated with a neutralizing monoclonal antibody against IL-23 (α p19). Consistent with results obtained from administration of IL-25, neutralization of IL-23p19 in GF mice resulted in significantly decreased expression of *Il17* and *Rorc* in the large intestine (Fig. 3, G and H), as well as a significant decrease in the frequency of Th17 cells in the large intestinal LP (Fig. 3 I). These results suggest that commensal-dependent expression of IL-25 is a component of the pathways that regulate expression of IL-23 and the frequency of Th17 cells in the intestinal microenvironment.

As GF mice and antibiotic-treated CNV mice expressed lower levels of IL-25, we hypothesized that the absence of IL-25 in CNV mice would recapitulate the heightened expression of IL-23 and increased frequency of Th17 cells observed in GF mice. To test this, expression of IL-17 and -23 in the large intestine of naive WT and IL-25-deficient (*Il25*^{-/-}) mice housed under conventional conditions was examined. Expression of *Il23a*, *Il12b*, and *Il17* was increased in naive *Il25*^{-/-} mice compared with WT controls (Fig. 4, A–C), similar to results obtained in GF mice. Moreover, flow cytometric analysis of lymphocytes isolated from the large intestinal LP demonstrated an increased frequency of Th17 cells in *Il25*^{-/-} mice under steady-state conditions (Fig. 4 D). The consequences of IL-25 deficiency on IL-17 expression were apparent primarily in the intestinal tissue, as the frequencies of Th17 cells in the mLN or spleen were similar in WT and *Il25*^{-/-} mice (Fig. 4 D). Thus, the absence of endogenous IL-25 in CNV mice is associated with increased levels of IL-23 and heightened frequencies of Th17 cells in the large intestine of naive *Il25*^{-/-} mice.

The increased expression of IL-23 in GF mice and the ability of IL-25 to inhibit IL-23 and -17 expression in vivo led to the hypothesis that IL-25 could act on accessory cells to limit expression of IL-23 and subsequent expansion or survival of Th17 cells. To test this, CD11b⁺ cells were isolated from the large intestinal LP of CNV and GF mice and

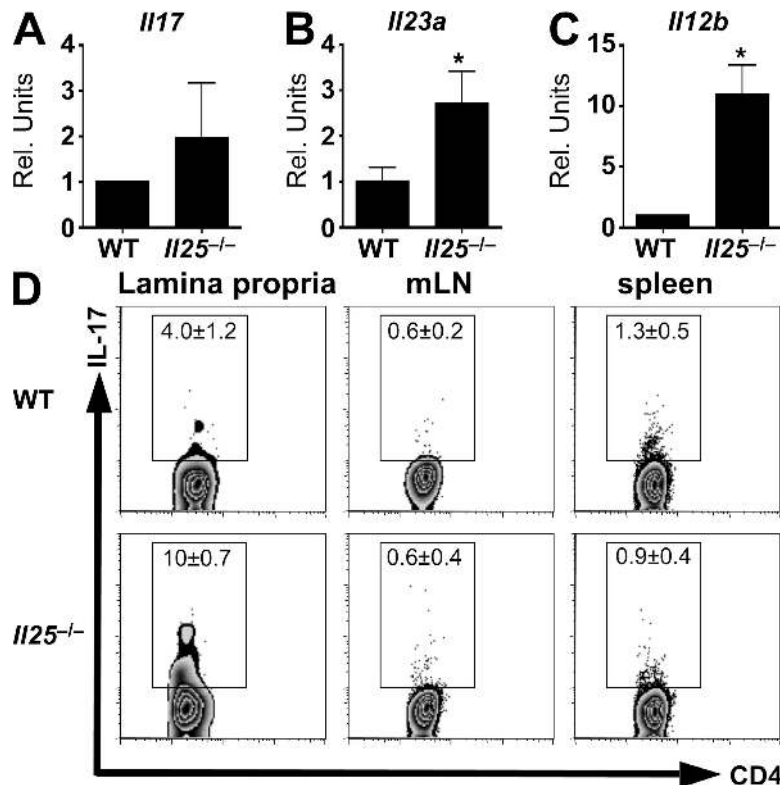


Figure 4. Endogenous IL-25 is required to limit IL-23 and -17 expression in the large intestine. (A–C) Expression of mRNA for *Il17* (A), *Il23a* (B), and *Il12b* (C) in the large intestine of CNV and GF mice was analyzed by quantitative real-time PCR. (D) Expression of IL-17 by CD4⁺ T cells in the large intestinal LP was analyzed by flow cytometry. Flow cytometry plots depict log₁₀ fluorescence. Data are from 3 experiments ($n = 6-9$). Rel. units, relative units. *, $P < 0.05$. Error bars indicate the SEM.

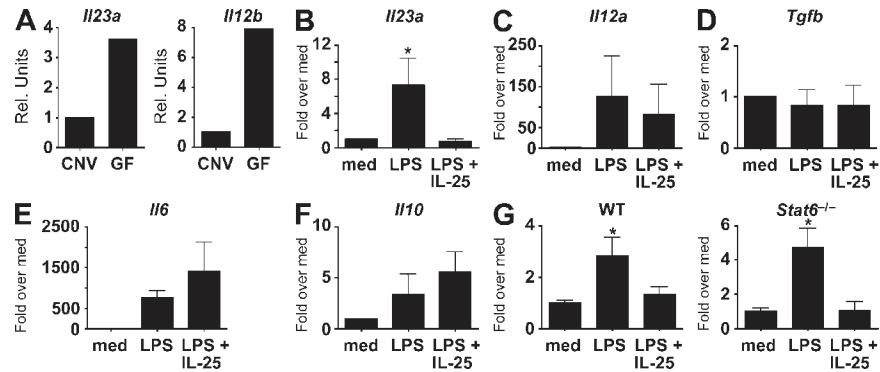


Figure 5. IL-25 limits accessory cell-derived IL-23 independently of STAT6. (A) Increased expression of *Il23a* and *Il12b* by CD11b⁺ cells isolated from the large intestinal LP of GF mice. (B–F) Expression of mRNA for *Il23a* (B), *Il12a* (C), *Tgfb* (D), *Il6* (E), and *Il10* (F) in macrophages 4 h after stimulation with LPS in the absence or presence of IL-25 was analyzed by quantitative real-time PCR. Results are from four experiments. (G) mRNA expression of IL-23p19 (*Il23a*) in WT or *Stat6*^{-/-} macrophages was analyzed by real-time PCR after stimulation with media alone, LPS, or LPS and IL-25. Results are representative of two experiments. *, $P < 0.05$. Error bars indicate the SEM.

analyzed for ex vivo expression of *Il23a* and *Il12b* mRNA. CD11b⁺ cells isolated from GF mice expressed higher levels of *Il23a* and *Il12b* than those isolated from CNV mice (Fig. 5 A). To examine whether IL-25 could directly act on macrophages to limit IL-23 expression, bone marrow-derived macrophages were activated with LPS, a TLR ligand that is known to induce IL-23, in the absence or presence of IL-25. Examination of mRNA levels revealed that IL-25 inhibited LPS-induced *Il23a* gene expression (Fig. 5 B). This effect was specific for IL-23, as no down-regulation of *Il12a*, *Tgfb*, *Il6*, or *Il10* gene expression was observed after exposure to IL-25 (Fig. 5, C–F). A previous study demonstrated that IL-25-induced IL-13 production by DCs that could inhibit IL-23 production (22). However, treatment of macrophages with IL-25 failed to induce *Il4* or *Il13* mRNA (unpublished data), and IL-25 inhibited LPS-induced expression of *Il23a* in macrophages derived from mice deficient in STAT6 (*Stat6*^{-/-} mice), which is the primary signaling molecule downstream of IL-4 and -13 (Fig. 5 G). Consistent with this, despite higher levels of *Il23a* mRNA in GF mice, we did not detect any difference in *Il13* mRNA in the large intestine of mice housed under CNV or GF conditions (Fig. S8, available at <http://www.jem.org/cgi/content/full/jem.20080720/DC1>). Therefore, IL-25 can act directly on macrophages to inhibit TLR-induced IL-23 expression independently of the IL-4–IL-13–STAT6 pathway.

Collectively, these results identify a previously unrecognized commensal-dependent immunoregulatory pathway associated with the maintenance of intestinal immune homeostasis. Specifically, recognition of commensal bacteria promotes expression of IL-25 by IECs that is a component of the cytokine network necessary to limit expression of IL-23 and the size of the Th17 cell pool in the intestine under steady-state conditions (Fig. S9, available at <http://www.jem.org/cgi/content/full/jem.20080720/DC1>). Several studies suggest that commensal bacteria can promote immune cell hyporesponsiveness to maintain intestinal immune homeostasis. For example, commensal bacteria can actively inhibit the activation of

NF- κ B and expression of proinflammatory cytokines by several mechanisms, including sequestration of signaling components such as PPAR- γ and RelA and inhibition of I κ B degradation (13, 14, 29). The absence of these signals in GF mice could contribute to a mechanism that results in increased IL-23 and -17 expression in the absence of commensal bacteria. Supporting a model in which innate recognition of commensal bacteria shapes the local immune environment, a recent study in *Drosophila* highlighted the influence of signals derived from commensal bacteria in the maintenance of intestinal immune homeostasis (30). When NF- κ B-dependent expression of antimicrobial peptides was disrupted in intestinal cells of the fly, there was an outgrowth of a population of pathogenic commensal bacteria that led to fly mortality. In mammals, disruption of innate recognition of commensal bacteria via IEC-specific deletion of the NF- κ B pathway by targeting I κ B kinase- β (IKK- β) or IKK- γ /NEMO, resulted in spontaneous and infection-induced intestinal inflammation (15, 16). Collectively, these studies highlight the growing recognition of the influence of commensal bacteria-derived signals that promote immunoregulatory pathways in the intestine. Thus, in addition to established immunoregulatory cytokines such as IL-10 and TGF- β , the demonstration of a role for commensal bacteria in influencing IL-25 expression identifies a previously unrecognized pathway through which commensal bacteria can regulate intestinal immune responses.

MATERIALS AND METHODS

Mice. BALB/c, C57BL/6, and BALB/c *Stat6*^{-/-} mice were obtained from The Jackson Laboratory. *Il25*^{-/-} mice have been described previously (22, 23). GF BALB/c and C57BL/6 mice were maintained in plastic isolator units and fed autoclaved feed and water. Conventional animals were maintained in a specific pathogen-free facility and routinely tested negative for pathogens. In individual experiments, mice were age matched and used at 6–8 wk of age or 12–14 wk of age. All experiments were performed under the guidelines of the Institutional Animal Care and Use Committee at the University of Pennsylvania.

Cytokine and antibody treatments. IL-25 (R&D Systems) was administered daily (0.5 μ g i.p.) to GF mice for 7 d. Control mice received PBS.

Monoclonal anti-IL-23p19 was produced at Schering-Plough Biopharma, and 1 mg was administered i.p. daily. Control mice received 1 mg of control antibody i.p. daily.

Isolation of cells. At necropsy, mLN, spleen, cecal patches, and Peyer's patches were harvested, and single-cell suspensions were prepared. IECs and LP lymphocytes were isolated as previously described (9). Purity of IECs was determined by flow cytometric analysis using anti-Ep-Cam antibody (G8.8, obtained from Developmental Studies Hybridoma Bank, University of Iowa, Iowa City, Iowa) and was routinely >90%.

Macrophage isolation, culture, and stimulation. CD11b⁺ cells were isolated from LP of CNV and GF mice using magnetic beads (Miltenyi Biotech). Bone marrow macrophages were cultured for 7 d, plated overnight at 10⁶ cells/ml, and pulsed with 10 ng/ml LPS (*Salmonella typhimurium*; Sigma-Aldrich) in the absence or presence of 50 ng/ml IL-25 the next day.

Cell stimulations and cytokine assays. Spleen cells were plated in medium alone or in the presence of anti-CD3 and -CD28 (1 µg/ml each) in the presence or absence of 1 ng/ml IL-12 (eBioscience). Large intestinal tissue explant cultures were performed as previously described (12). Cell-free supernatants were harvested after 72 h, and analyzed for cytokine secretion by sandwich ELISA (eBioscience). Cells were stimulated by incubation for 4 h with 50 ng/ml PMA (Sigma-Aldrich) and 750 ng/ml ionomycin (Sigma-Aldrich) in the presence of 10 µg/ml Brefeldin A (Sigma-Aldrich), and then surface stained with fluorochrome-conjugated antibodies against CD4 and CD3 fixed in 2% paraformaldehyde. Fixed cells were permeabilized with 0.5% saponin (Sigma-Aldrich) and stained intracellularly for IL-17, IFN-γ, TNF-α, IL-10 or Foxp3 (eBioscience), acquired on a FACSCalibur using CellQuest Pro software (BD Biosciences) and analyzed with FlowJo software (Version 8.5; Tree Star, Inc.).

RNA isolation and real-time PCR. RNA was isolated from tissues using RNeasy Spin columns (QIAGEN) after disruption in a homogenizer (TissueLyzer; QIAGEN). RNA was reverse transcribed into cDNA, and quantitative real-time PCR was performed on cDNA using primer sets for *Il10*, *Il12a*, *Il12b*, *Il23a*, *Tgfb*, *Il6*, *Il27p28*, and *Ebi3* (QIAGEN) using SYBR Green chemistry. Primers for RORγt (*Ror*) have been previously described (9). *Il17* and *Il25* were analyzed using TaqMan primer/probe pairs (Applied Biosystems). All reactions were run on an ABI 7500 Fast Real-Time PCR System (Applied Biosystems). Samples were normalized to naive controls unless specifically stated.

Statistics. Results represent the mean ± SEM. Statistical significance was determined by Student's *t* test (between two groups or conditions) or analysis of variance with a post-hoc test (three or more groups or conditions) using Prism 4.0 (GraphPad Software).

Online supplemental material. Increased expression of *Il17* in C57BL/6 GF mice is shown in Fig. S1. Production of IL-6 and -12p40 in large intestinal tissue explant cultures is shown in Fig. S2. Increased expression of *Il23a* in C57BL/6 GF mice is shown in Fig. S3. Fig. S4 shows that there are minimal antibiotic-induced changes in gene expression in the small intestine of CNV mice. Equivalent *Il10* mRNA expression and frequencies of IL-10-producing CD4⁺ T cells in CNV and GF mice are shown in Fig. S5. Fig. S6 depicts expression of IL-12 and IFN-γ in CNV and GF mice. Fig. S7 shows that expression of *Il25* mRNA is decreased in the large intestine of antibiotic-treated CNV mice. Similar levels of *Il13* expression in CNV and GF mice are shown in Fig. S8. Fig. S9 depicts a model of how commensal-dependent IL-25 regulates the IL-23-IL-17 axis in the intestine. The online version of this article is available at <http://www.jem.org/cgi/content/full/jem.20080720/DC1>.

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