

In vivo equilibrium of proinflammatory IL-17⁺ and regulatory IL-10⁺ Foxp3⁺ ROR γ t⁺ T cells

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The nuclear hormone receptor retinoic acid receptor–related orphan receptor γ t (ROR γ t) is required for the generation of T helper 17 cells expressing the proinflammatory cytokine interleukin (IL)–17. In vivo, however, less than half of ROR γ t⁺ T cells express IL–17. We report here that ROR γ t⁺ T α β cells include Foxp3⁺ cells that coexist with IL–17–producing ROR γ t⁺ T α β cells in all tissues examined. The Foxp3⁺ ROR γ t⁺ T α β express IL–10 and CCL20, and function as regulatory T cells. Furthermore, the ratio of Foxp3⁺ to IL–17–producing ROR γ t⁺ T α β cells remains remarkably constant in mice enduring infection and inflammation. This equilibrium is tuned in favor of IL–10 production by Foxp3 and CCL20, and in favor of IL–17 production by IL–6 and IL–23. In the lung and skin, the largest population of ROR γ t⁺ T cells express the γ δ T cell receptor and produce the highest levels of IL–17 independently of IL–6. Thus, potentially antagonistic proinflammatory IL–17–producing and regulatory Foxp3⁺ ROR γ t⁺ T cells coexist and are tightly controlled, suggesting that a perturbed equilibrium in ROR γ t⁺ T cells might lead to decreased immunoreactivity or, in contrast, to pathological inflammation.

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Abbreviations used: BAC, bacterial artificial chromosome; CIA, collagen-induced arthritis; DSS, dextran sodium sulfate; EGFP, enhanced GFP; LTi, lymphoid tissue inducer; RA, retinoic acid; ROR γ t, RA receptor–related orphan receptor γ t; T reg, T regulatory.

T cells adapt immune responses not only to the antigens, but also to the type of pathogen they face. Cell-based pathogens, such as viruses, intracellular bacteria, and tumors evoke a CD4⁺ Th1 type of response characterized by the production of IFN- γ . In contrast, extracellular-based pathogens, such as worms and allergens, evoke a Th2 type of response characterized by the production of IL-4, IL-5, and IL-13 (1). Recently, some bacteria and fungi have been found to evoke a Th17 type of response characterized by the production of IL-17 (or IL-17A), IL-17F, IL-21, IL-22, IL-6, and TNF- α , as well as IL-23 by APCs (2–6). The proinflammatory Th17 response induces the recruitment

of neutrophils and is involved in autoimmune inflammatory diseases. However, the reactivity of T cells is modulated by T regulatory (T reg) cells. These include so-called natural Foxp3⁺ T reg cells that are induced in the thymus (7) as well as adaptive Foxp3⁺ or Foxp3⁻ T reg cells that are induced in the periphery and produce TGF- β and/or IL-10 (8, 9).

The generation of these distinct T cell effector types is under control of specific cytokines and transcription factors (2). In mice, TGF- β is required for the generation of both Th17 and T reg cells. The switch to the Th17 pathway is induced by IL-6 and/or IL-21 (4–6, 10–12), and requires IL-23 for full effector maturation (2, 3). However, TGF- β together with IL-6 induces CD4⁺ T cells to produce both IL-17 and IL-10, a population that is not able to induce inflammation in mice (13).

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In contrast, IL-23 generates proinflammatory cells that produce only IL-17. In accordance with these findings, spontaneous colitis is induced in mice deficient for IL-10 (14), and the inflammatory disease is prevented by concomitant deficiency in IL-23 (15). The immunosuppressive IL-10 can be produced by all types of T cells, including a subset of T reg cells (16), and is induced by IL-27 in Th1 and Th2 cells (17–19). TGF- β alone induces the T reg cell pathway (20) and is strongly promoted in the gut-associated lymphoid tissues by all-trans retinoic acid (RA) (21, 22). The generation of T reg cells requires the forkhead/winged-helix transcription factor Foxp3 (23–25). Mice deficient in Foxp3 lack natural T reg cells (24) and are highly susceptible to inflammatory disease (26).

The development of inflammatory pathologies, such as experimental autoimmune encephalomyelitis (27, 28), an animal model of multiple sclerosis, collagen-induced arthritis (CIA) (29), and inflammatory bowel disease (15, 30), depends on Th17 cells and IL-23. In humans, a coding variant of *Il23* confers protection to Crohn's disease (31). The Th17–IL-23 pathway in also involved in resistance to a growing list of bacterial and fungal pathogens in mice (11, 32–35), and Th17 cells are readily isolated from patients with *Candida* infection (36). However, a comprehensive picture of the class of pathogens targeted by this pathway remains to be drawn. T reg cells, on the other hand, and IL-10 in particular, have been shown to limit inflammation and adaptive immunity to a variety of pathogens (37), as well as to promote long-term T cell memory (38). Collectively, these studies demonstrate the essential role of the Th17 and the T reg cells in immunity, immunoregulation, and immunopathology. They also suggest that a balanced expansion of these Th subsets during immune responses might ensure efficient yet restrained immunity that limits damage to self.

The nuclear hormone receptor RA receptor-related orphan receptor γ t (ROR γ t) is a marker for Th17 cells and is required for their generation (39). Using an enhanced GFP (EGFP) reporter mouse, we found that ROR γ t⁺ T α β cells included IL-10–producing Foxp3⁺ T reg cells. These cells coexisted with IL-17–producing ROR γ t⁺ T α β cells in all tissues examined in an equilibrium regulated by Foxp3 and CCL20 in favor of Foxp3⁺ cells, and by IL-6 and IL-23 in favor of IL-17–producing cells. Foxp3 was shown to bind to ROR γ t, suggesting that it directly regulates ROR γ t activity. The equilibrium within ROR γ t⁺ T α β cells was maintained during massive inflammation of the intestine or the lungs, but it was perturbed in cancer. We suggest that the balanced expansion of ROR γ t⁺ T α β cell subsets during inflammation might ensure efficient yet regulated immunity, and that perturbation of this equilibrium might lead to inadequate immune reactivity to tumors, pathogens, or self.

RESULTS

Diversity and distribution of ROR γ t⁺ T cells in vivo

To optimize the visualization of ROR γ t⁺ cells, we have generated transgenic *Rorc*(γ t)-*Gfp*^{TG} mice expressing EGFP under control of the *Rorc*(γ t) locus on a bacterial artificial

chromosome (BAC) (40). An estimated 5–10 copies of the BAC transgene were integrated, as assessed by quantitative PCR. The expression pattern of EGFP in *Rorc*(γ t)-*Gfp*^{TG} mice was identical to the one described previously in *Rorc*(γ t)^{+/GFP} mice carrying one allele of *Rorc*(γ t) inserted with EGFP (39, 41). Faithful expression of GFP was further confirmed by immunofluorescence histology of GFP and endogenous ROR γ t in various organs from *Rorc*(γ t)-*Gfp*^{TG} mice. In all tissues analyzed, we found a strict correlation between GFP and ROR γ t protein expression (not depicted). High expression of GFP was found in fetal lymphoid tissue inducer (LTi) cells, LTi-like cells in the adult intestine, in immature CD4⁺CD8⁺ thymocytes (not depicted), and in a subpopulation of peripheral T cells (Fig. 1 A). The highest frequency of ROR γ t⁺ T cells was observed in the intestine, as previously reported (39), which were localized in the lamina propria of the villi, whereas CD3[−] ROR γ t⁺ cells, i.e., LTi-like cells, were clustered in cryptopatches that are precursor structures to isolated lymphoid follicles (Fig. 1 C) (42). ROR γ t⁺ T cells were also found in all organs or tissues tested, including the LNs, spleen, lung, skin, and bone marrow (Fig. 1 A). Assessing the homogeneity of these populations within different organs, we observed that up to 50% of ROR γ t⁺ T cells in the lung and skin expressed the γ δ TCR instead of the α β TCR (Fig. 1 B). ROR γ t⁺ T γ δ cells expressed the highest levels of EGFP (Fig. 1 A) and none of the CD4 or CD8 coreceptors (Fig. 1 D).

ROR γ t⁺ T cells include both Th17 and IL-10–producing Foxp3⁺ T reg cells

We assessed whether ROR γ t was a marker for IL-17–producing cells in both T α β and T γ δ populations, and stained ROR γ t⁺ T cells for IL-17, IFN- γ , and Foxp3. Surprisingly, even though a majority of IL-17–producing cells expressed ROR γ t (Fig. 2 A), only 15–50% of ROR γ t⁺ T α β cells expressed IL-17, depending on the organ examined, and few but detectable numbers of ROR γ t⁺ T α β cells produced both IL-17 and IFN- γ (Fig. 2 B). In contrast, a majority (50–90%) of ROR γ t⁺ T γ δ cells expressed high levels of IL-17. Even more surprising, 15–50% of IL-17[−] ROR γ t⁺ T α β cells expressed the T reg cell markers Foxp3 and CD25 (Fig. 2, B and C), with the highest incidence in LNs, even though Foxp3⁺ ROR γ t⁺ T α β cells represented only a minority of the total population of Foxp3⁺ T cells (Fig. 2 A). No ROR γ t⁺ T α β cells were found to express both IL-17 and Foxp3, and no ROR γ t⁺ T γ δ cells expressed Foxp3 (not depicted). The coexpression of ROR γ t and Foxp3 protein at the single cell level was confirmed by immunofluorescence histology of the LNs (Fig. 2 D). Furthermore, Foxp3⁺ ROR γ t⁺ T α β cells were functionally T reg cells, as they suppressed in vitro proliferation of activated CD4⁺ T cells (Fig. 2 E).

We next measured the expression, by the different subsets of ROR γ t⁺ T cells isolated by FACS (Fig. S1, available at <http://www.jem.org/cgi/content/full/jem.20080034/DC1>), of a restricted panel of genes involved in the Th17 and

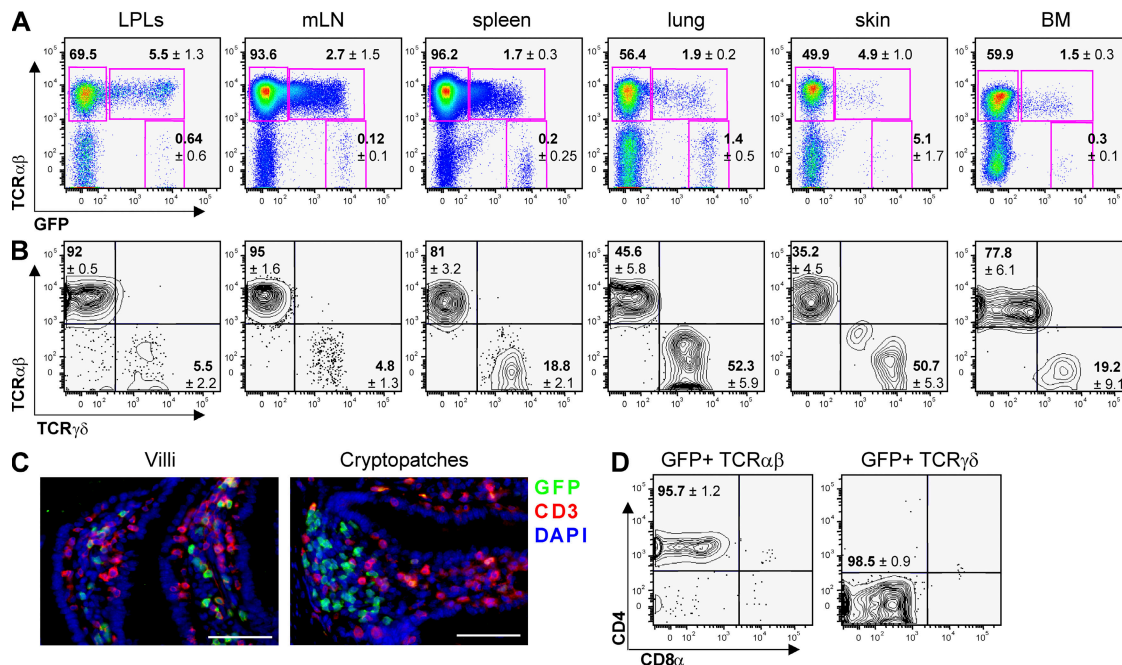


Figure 1. Diversity and distribution of ROR γ t⁺ T cells in vivo. (A and B) Flow cytometry analysis of cells isolated from the organs of 8–12-wk-old *Rorc*(γ t)-*Gfp*^{TG} mice. Plots are gated on CD3⁺ cells (A) or GFP⁺ CD3⁺ cells (B). Numbers indicate mean percent cells in quadrants \pm SD obtained with at least three *Rorc*(γ t)-*Gfp*^{TG} mice. LPLs, lamina propria lymphocytes isolated from small intestine; mLN, mesenteric LN; BM, bone marrow. (C) Immunofluorescence histology of ROR γ t⁺ cells in the small intestine of *Rorc*(γ t)-*Gfp*^{TG} mice. Most ROR γ t⁺ cells in villi are T cells, whereas ROR γ t⁺ cells in cryptopatches located between crypts are CD3⁻ LTi cells. Bar, 50 μ m. (D) Expression of CD4 and CD8 α by spleen GFP⁺TCR- β ⁺ and lung or GFP⁺TCR- δ ⁺ cells.

T reg cell pathways using quantitative qRT-PCR arrays. As compared with ROR γ t⁻ T cells, ROR γ t⁺ T cells expressed higher levels of *Rorc*, *Il17a*, *Il17f*, *il22*, and *Il23r*, as expected, as well as of *Il10*, *Icos*, *Cd20*, *Cx3d1*, and *Mmp9* (Fig. S2). Expression of these genes was further examined in the individual ROR γ t⁺ T cell subsets. As cells could not be fixed and stained for Foxp3 expression before isolation for RNA recovery, Foxp3⁺ ROR γ t⁺ T cells were isolated on the basis of CD25 expression. As expected, CD25⁻ ROR γ t⁺ T α β cells enriched in IL-17-producing cells (Fig. 2 C), as well as ROR γ t⁺ T γ δ cells, expressed high levels of *Il17a*, *Il17f*, and *Il22* transcripts (Fig. 3 A and not depicted). CD25⁺ ROR γ t⁺ T α β cells expressed detectable levels of *Il17a* and *Il17f* transcripts, probably due to the presence of a small population of IL-17-producing CD25⁺ cells (Fig. 2 C). In addition, ROR γ t⁺ T γ δ cells, but not ROR γ t⁻ T γ δ cells, expressed *Csf2*, which codes for GM-CSF involved in the mobilization of granulocytes and monocytes.

CD25⁺ (Foxp3⁺) ROR γ t⁺ T α β cells specifically expressed high levels of the expected *Foxp3* transcript, as well as *Il10*, *Icos*, *Cd20*, *Cx3d1*, and *Mmp9*. IL-10 expression was confirmed at the protein level, and Foxp3⁺, but not Foxp3⁻ ROR γ t⁺ T α β , cells represented a major fraction of total IL-10-producing cells in LNs (Fig. 3 B), and to a lesser extent in the gut (Fig. S3, available at <http://www.jem.org/cgi/content/full/jem.20080034/DC1>). The transcript coding for CCL20 was specifically expressed by Foxp3⁺ ROR γ t⁺ T α β cells and barely detected in other T cell populations. Further-

more, CCR6, the receptor for CCL20, was expressed by both CD25⁺ and CD25⁻ ROR γ t⁺ T α β cells (Fig. S4), suggesting that CD25⁺ ROR γ t⁺ T α β cells can regulate the recruitment of ROR γ t⁺ T α β cells. Interestingly, ICOS (43), CCL20 (13, 44), and MMP-9 (45, 46) have been associated with Th17 cells previously, but not with T reg cells, whereas CX₃CL1 is normally expressed by activated endothelial cells (47).

Generation and regulation of IL-17⁺ or Foxp3⁺ ROR γ t⁺ cells

We further assessed how the generation of IL-17-producing or Foxp3⁺ ROR γ t⁺ T cells in vivo was affected by the absence of factors shown to be required for the generation of Th17 or T reg cells, such as IL-6 (4–6, 10–12) and Foxp3 (24, 26). In IL-6-deficient mice, the number and proportion of ROR γ t⁺ T α β cells, in particular of IL-17-producing ROR γ t⁺ T α β cells, were significantly decreased, whereas the proportion of IL-10-producing Foxp3⁺ ROR γ t⁺ T cells was unaffected (Fig. 4, A, C, and D). This confirms the requirement of IL-6 for the generation of Th17 cells, but also shows that IL-6 is not required for the generation of IL-10-producing ROR γ t⁺ T cells, even though in vitro it acts in synergy with TGF- β to induce IL-10-producing CD4⁺ T cells (13, 17). In contrast, IL-17-producing ROR γ t⁺ T γ δ cells were unaffected by the absence of IL-6 (Fig. 4 B). Furthermore, in IL-12R β 1-deficient mice, unresponsive to both IL-12 and IL-23 (2, 3), the proportion of IL-17-producing ROR γ t⁺ T α β cells was slightly decreased (Fig. 4, C and D),

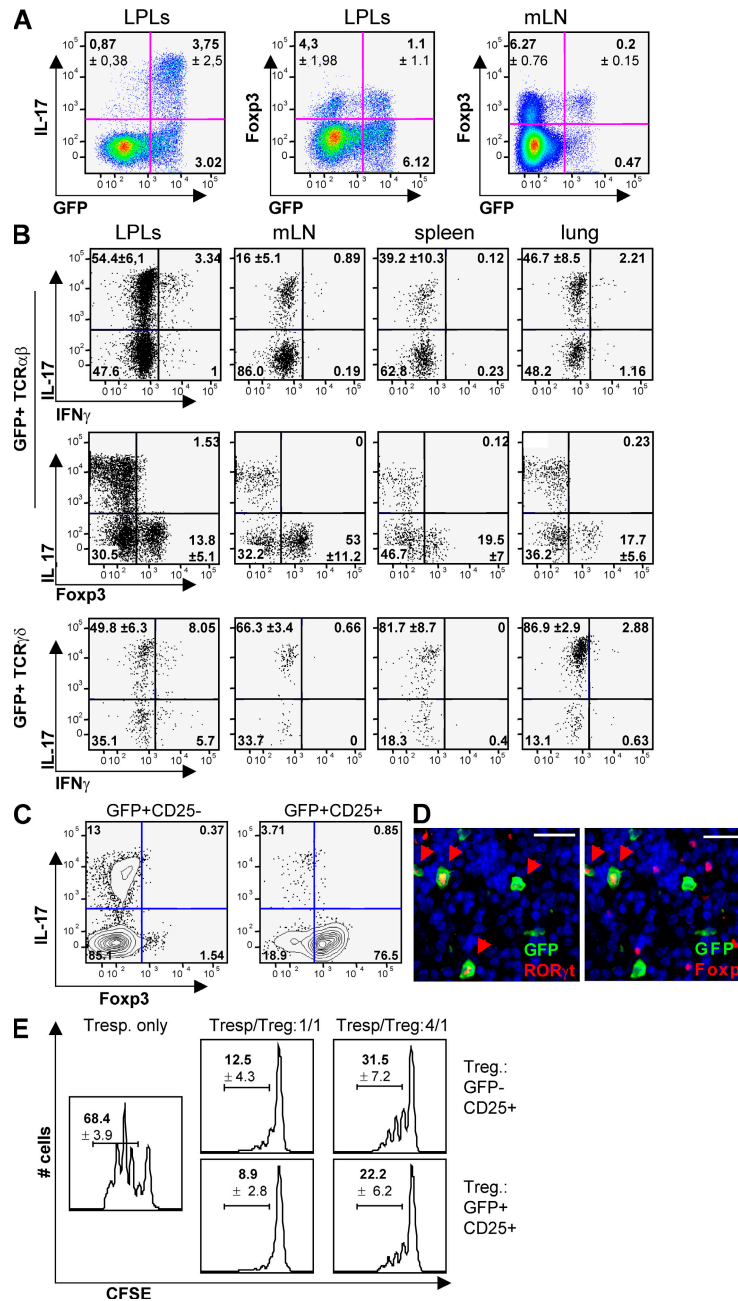


Figure 2. ROR γ t⁺ T cells include Th17 and Foxp3⁺ T reg cells. (A and B) Flow cytometry analysis of cells isolated from *Rorc*(γ t)-*Gfp*^{TG} mice. Cells were restimulated in vitro with PMA/ionomycin for 5 h and subjected to intracellular staining for GFP, IL-17, Foxp3, and/or IFN- γ . Plots are gated on TCR- β ⁺ cells (A) and GFP⁺TCR- β ⁺ or GFP⁺TCR- δ ⁺ cells (B). Numbers indicate mean percent cells in quadrants \pm SD obtained with at least three *Rorc*(γ t)-*Gfp*^{TG} mice. (C) CD4⁺ T cells isolated from the spleen and mesenteric LNs of *Rorc*(γ t)-*Gfp*^{TG} mice were sorted into two populations based on their expression of GFP and CD25. Sorted cells were subsequently restimulated in vitro and stained for IL-17 and Foxp3. Results are representative of three independent experiments. (D) Immunofluorescence histology staining for GFP, ROR γ t, Foxp3, and nuclei (DAPI) of a mesenteric LN isolated from a *Rorc*(γ t)-*Gfp*^{TG} mouse. Arrowheads indicate cells expressing ROR γ t and/or Foxp3. Bar, 20 μ m. (E) Proliferation assay of CD4⁺ T cells isolated from the spleen and mesenteric LNs of *Rorc*(γ t)-*Gfp*^{TG} mice and sorted into two populations based on their expression of GFP and CD25. CFSE-labeled CD25⁻ responder T cells (2.5×10^4) were cultured alone (left) or together with either GFP⁻CD25⁺ T cells or GFP⁺CD25⁺ T cells (2.5×10^4 or 0.625×10^4 cells). Cells were cultured in duplicates for 3 d in the presence of 10^5 irradiated spleen cells and anti-CD3 ϵ antibody. Numbers indicate frequency of proliferating cells (\pm SD) obtained from three independent experiments.

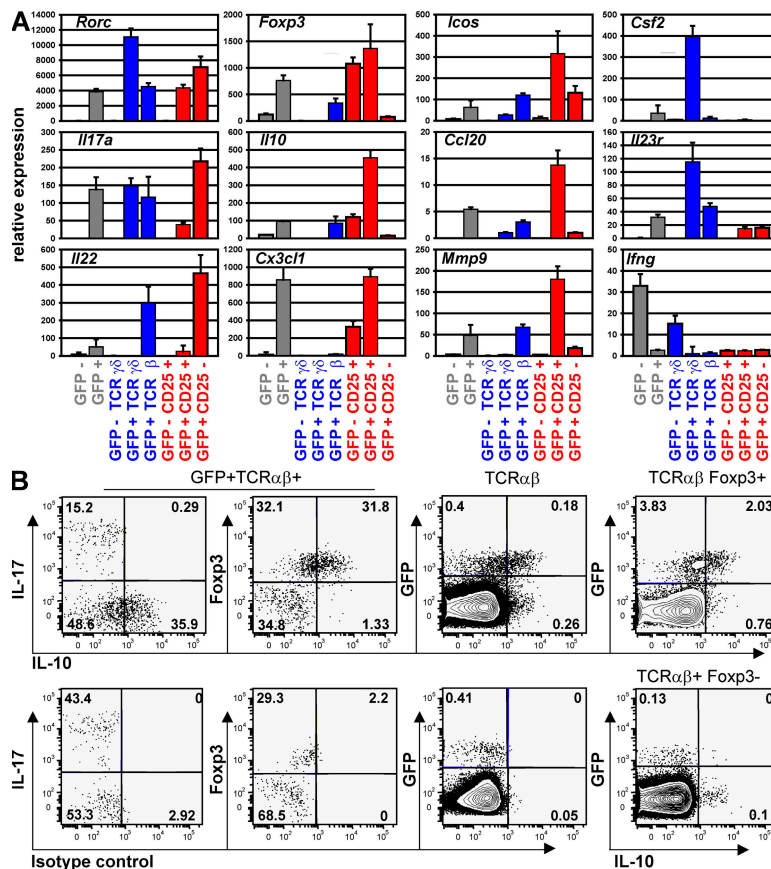


Figure 3. ROR γ t⁺ T cells express genes involved in the Th17 or T reg cell pathway. (A) Cells isolated from the spleen and mesenteric LNs of *Rorc*(γ t)-*Gfp*^{TG} mice were sorted into eight distinct populations based on their expression of GFP, CD3, TCR- β , TCR- δ , CD4, and CD25 (Fig. S1), and gene expression was assessed using real-time PCR. Ct values were normalized to the mean Ct of five housekeeping genes. Data are the mean of two or three independent experiments. (B) Foxp3⁺ ROR γ t⁺ T cells express IL-10. Cells isolated from LNs of *Rorc*(γ t)-*Gfp*^{TG} mice were restimulated in vitro with PMA/ionomycin for 5 h and subjected to intracellular staining for GFP, IL-17, Foxp3, and IL-10 or an isotype control. Numbers indicate percent cells in quadrants. Results are representative of at least three individual experiments.

in accordance with the important but not essential role of IL-23 in the generation of Th17 cells (2, 3).

It has recently been shown that Th17 cells isolated from SKG mice, which spontaneously develop arthritis, as well as human Th17 cells, express CCR6 (44, 48, 49). We therefore assessed the impact of CCR6 on the generation of IL-17-producing or Foxp3⁺ ROR γ t⁺ T cells. In CCR6-deficient mice (50), contrary to IL6- or IL12R β 1-deficient mice, the proportion of IL-17-producing ROR γ t⁺ T α β cells was significantly increased (Fig. 4, C and D). As CD25⁺ (Foxp3⁺) ROR γ t⁺ T α β cells specifically express *Cd20* transcripts (Fig. 3 A), these cells might regulate the generation of IL-17-producing ROR γ t⁺ T α β cells through CCR6.

We next assessed whether Foxp3 regulates the generation of IL-17- or IL-10-producing ROR γ t⁺ T α β cells. Mice bearing the *Scurfy* mutation in the *Foxp3* gene (*Foxp3*^{scf} mice) lack T reg cells (24, 26). In these mice, however, the analysis of ROR γ t⁺ T cells was complicated by generalized autoimmunity, and the development of immature ROR γ t⁺ CD4⁻CD8⁻ thymocytes was abnormal (not depicted). Therefore, mixed bone

marrow chimeras were generated by adoptive transfer of bone marrow from *Foxp3*^{scf} \times *Rorc*(γ t)-*Gfp*^{TG} and WT mice into irradiated lymphopenic hosts (Fig. S5, available at <http://www.jem.org/cgi/content/full/jem.20080034/DC1>). In such chimeras, ROR γ t⁺ T cells do not express Foxp3, but cells from WT mice protected the host from a scurfy phenotype (not depicted). *Foxp3*^{scf} ROR γ t⁺ T α β cells failed to generate IL-10-producing cells and generated a reduced proportion of IL-17-producing cells as compared with *Foxp3*^{wt} ROR γ t⁺ T α β cells in control chimeras (Fig. 5, A and B). This shows that Foxp3 protein or Foxp3⁺ cells are required for the generation of IL-10-producing ROR γ t⁺ T α β cells and, surprisingly, also favor the differentiation of IL-17-producing ROR γ t⁺ T α β cells.

To test whether Foxp3 regulates ROR γ t⁺ T α β cells through direct interaction with ROR γ t, N-terminally FLAG-tagged ROR γ t was expressed in HEK293T cells together with Foxp3. Foxp3 was specifically coimmunoprecipitated with FLAG-ROR γ t, indicating that the two factors interact (Fig. 5 C). Collectively, these data suggest that Foxp3

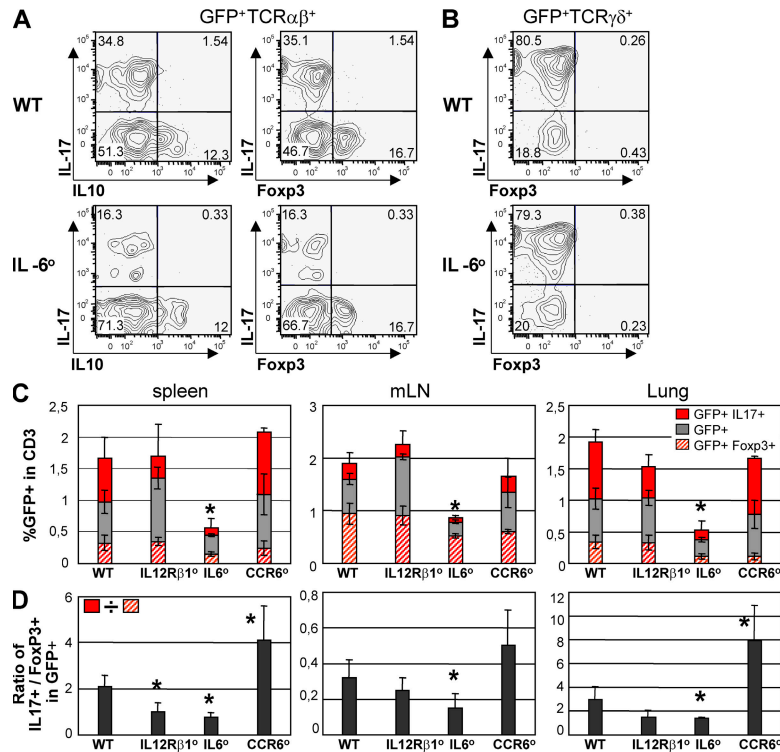


Figure 4. The role of IL-6, IL-12R β 1, and CCR6 in the generation of ROR γ t⁺ T cell subsets. (A and B) Cells were isolated from the spleen of *Rorc*(γ t)-*Gfp*^{TG} (WT) or *Il6*^{-/-} \times *Rorc*(γ t)-*Gfp*^{TG} (IL-6^{-/-}) mice, restimulated in vitro with PMA/ionomycin for 5 h and subjected to intracellular staining for GFP, IL-17, Foxp3, and IL-10. Plots are gated on GFP⁺TCR- β ⁺ cells (A) or GFP⁺TCR- δ ⁺ cells (B). Numbers indicate percent cells in quadrants and are representative of two independent experiments. (C) Cells were isolated from the spleen, mesenteric LN, and lung of *Rorc*(γ t)-*Gfp*^{TG} (WT), *Il12rb1*^{-/-} \times *Rorc*(γ t)-*Gfp*^{TG} (IL12R β 1^{-/-}), *Ccr6*^{-/-} \times *Rorc*(γ t)-*Gfp*^{TG} (CCR6^{-/-}), or *Il6*^{-/-} \times *Rorc*(γ t)-*Gfp*^{TG} mice (IL-6^{-/-}) and processed as in A and B. Histograms report percent GFP+TCR- β ⁺ cell subsets in total T cells. Each bar also shows the percentages of IL-17⁺ and Foxp3⁺ cells within the GFP+TCR- β ⁺ cell population. (D) Histograms report the ratio of the frequencies of IL-17-producing GFP+TCR- β ⁺ cells to Foxp3⁺ GFP+TCR- β ⁺ cells as indicated in the figure. Three to five mice were analyzed per group. *, $P < 0.05$ as compared with WT.

induces the generation of IL-10-producing ROR γ t⁺ T cells and regulates ROR γ t activity directly. On the other hand, ROR γ t is required for the generation of Th17 cells (39), but not for the generation of “mainstream” Foxp3⁺ and/or IL-10-producing T cells. In *Rorc*(γ t)-*Cre*^{TG} \times *Rosa-Stop*^{fl-Dta} mice, which lack ROR γ t⁺ T cells due to their specific expression of the lethal diphtheria toxin subunit A (51), no or few IL-17-producing cells were detected (not depicted), and the proportion of total Foxp3⁺ or IL-10-producing cells was unaffected (Fig. S6, available at <http://www.jem.org/cgi/content/full/jem.20080034/DC1>).

Equilibrium between IL-17⁺ or Foxp3⁺ ROR γ t⁺ cells during inflammation

Given the prominent role of IL-17-producing cells in inflammation and of IL-10-producing cells in immune regulation, we assessed if and how the relative frequencies of the different ROR γ t⁺ T cell subsets were modified in mice enduring severe intestinal inflammation, virus infection, or cancer. Dextran sodium sulfate (DSS) was administered orally to *Rorc*(γ t)-*Gfp*^{TG} mice until the development of chronic intestinal inflammation (Fig. 6 A). A high number of T cells was re-

covered in DSS-treated mice and both the frequency and the number of ROR γ t⁺ T $\alpha\beta$ cells increased, but the ratio of IL-17-producing to Foxp3⁺ ROR γ t⁺ T cells remained constant. Similar results were obtained in mice exposed to intranasal infection with influenza A virus, even though their lungs sustained heavy leukocyte infiltration (Fig. 6 B). In contrast, in *Rorc*(γ t)-*Gfp*^{TG} \times *Apc*^{min/+} mice bearing colonic and ileal polyps and tumors (52), no significant infiltration of leukocyte and ROR γ t⁺ T $\alpha\beta$ cells was observed in the colon and the terminal ileum (Fig. S7, available at <http://www.jem.org/cgi/content/full/jem.20080034/DC1>), whereas the mesenteric LNs were largely hyperplastic. The ratio of IL-17-producing to Foxp3⁺ ROR γ t⁺ T cell in these LNs was significantly decreased, even though the frequency and total number of ROR γ t⁺ T $\alpha\beta$ cells were increased (Fig. 6 C). Similar results have recently been obtained comparing Th17 and Foxp3 cells in mice injected with B16 melanoma (53). These data indicate that the relative frequency of IL-17-producing or Foxp3⁺ ROR γ t⁺ T $\alpha\beta$ cells can remain constant during severe inflammation or infection. However, chronic tumors can alter this balance in favor of the generation or recruitment of Foxp3⁺ ROR γ t⁺ T $\alpha\beta$ cells producing the immunosuppressive IL-10.

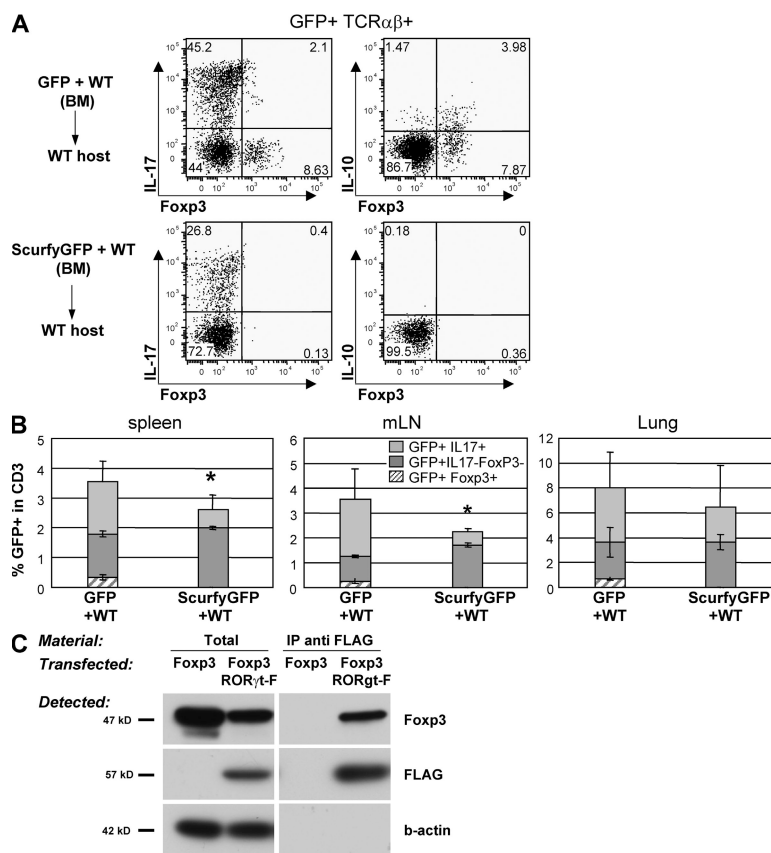


Figure 5. Regulation of ROR γ t⁺ T cells by Foxp3. (A and B) Bone marrow cells isolated from CD45.2⁺ *Rorc*(γ t)-*Gfp*^{TG} (WT) or *Foxp3*^{fl} \times *Rorc*(γ t)-*Gfp*^{TG} (Scurfy) mice were mixed with bone marrow cells isolated from CD45.1⁺ C57BL/6 mice and injected into irradiated lymphopenic mice. 6 wk after transfer, cells were isolated from the spleen, LN, and lung and restimulated in vitro with PMA/ionomycin for 5 h and subjected to intracellular staining for GFP, IL-17, Foxp3, and IL-10. (A) Spleen cells are gated on GFP+TCR- β ⁺ cells. Numbers indicate percent cells in quadrants. Reconstitutions with CD45.1⁺ and CD45.2⁺ cells were comparable in all chimeras. Results are representative of two independent experiments. (B) Histograms report percent GFP+TCR- β ⁺ cell subsets in total CD45.2⁺ T cells (see Fig. 4 C). *, $P < 0.05$ as compared with WT. (C) Immunoprecipitation of ROR γ t and Foxp3. Foxp3 was expressed in HEK293T cells alone or together with N-terminally FLAG-tagged ROR γ t. Total cell lysates or immunoprecipitates (IP) using anti-FLAG antibody were blotted and probed with antibodies to Foxp3, FLAG, or β -actin. Results are representative of two independent experiments.

Generation of Th17, Foxp3⁺, or IL-10–producing ROR γ t⁺ T cells in vitro

Many steps in the differentiation pathway of Th17 cells have been inferred from in vitro studies. We therefore assessed if the distinct subsets of ROR γ t⁺ T $\alpha\beta$ cells we have identified in vivo were derived in vitro from naive CD4⁺ T cells. TGF- β together with IL-6 or IL-21 induced the generation of Th17 cells from naive CD4⁺ T cells (4–6, 10–12) after massive up-regulation of ROR γ t expression after 1–2 d of culture (Fig. 7) (39). In contrast, TGF- β alone induces the generation of T reg cells (20). However, we also observed that most cells up-regulated of ROR γ t after 1–2 d of culture in the presence of TGF- β alone, and all Foxp3⁺ cells co-expressed ROR γ t after 2–3 d. Up-regulation of ROR γ t was specific to these conditions because stimulation under Th1 conditions did not induce ROR γ t⁺ expression (Fig. S8, available at <http://www.jem.org/cgi/content/full/jem.20080034/DC1>). Expression of ROR γ t in the culture of naive CD4⁺ T cells in the presence of TGF- β has been shown before at the

population level (39), but not in individual cells expressing Foxp3. In contrast to their in vivo counterparts, however, few in vitro–derived Foxp3⁺ ROR γ t⁺ T $\alpha\beta$ cells expressed IL-10, which was rather expressed by ROR γ t[−] T $\alpha\beta$ cells. IL-10 was not induced by TGF- β alone but required the addition of IL-6, as found previously (13, 17), or the addition of IL-21. IL-10 production was inhibited in the presence of RA, even though RA favored the generation of Foxp3⁺ ROR γ t⁺ T $\alpha\beta$ cells and decreased the generation of IL-17–producing cells in the presence of TGF- β and IL-6. A combination of TGF- β with IL-6 (with or without RA) or IL-21 generated IL-17–producing as well as IL-10–producing or Foxp3⁺ ROR γ t⁺ T $\alpha\beta$ cells, showing that the engagement of these pathways is not mutually exclusive at the cell population level, in agreement with recent reports (13, 17). ROR γ t⁺ T $\alpha\beta$ cells producing both IL-17 and IL-10 could be generated in the presence of TGF- β and IL-6, but Foxp3⁺ ROR γ t⁺ T cells produced neither IL-10 nor IL-17 (not depicted). Collectively, these data show that even though naive CD4⁺

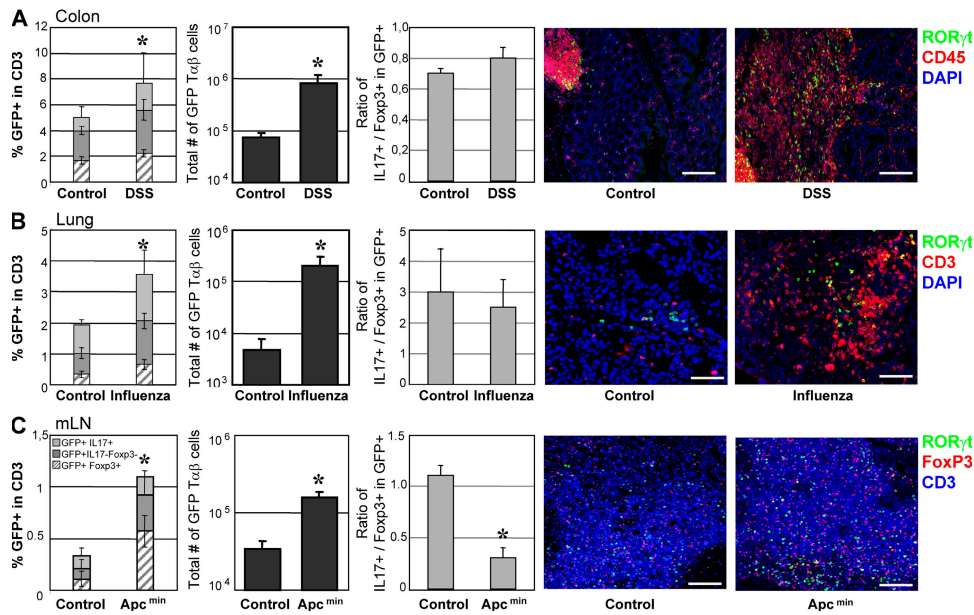


Figure 6. Equilibrium in ROR γ t⁺ T cell subsets during pathology. (A) *Rorc*(γ t)-*Gfp*^{TG} mice were treated with DSS in the drinking water for 6 d, followed by water for 10 d. This protocol was repeated for a total of three cycles. After the last cycle, cells isolated from the colon were restimulated in vitro with PMA/ionomycin for 5 h and subjected to intracellular staining for GFP, IL-17, and Foxp3. Histograms (from left to right) report percent GFP⁺TCR- β ⁺ cell subsets in total T cells (see Fig. 4 C), total numbers of ROR γ t⁺ T α β cells present in the organ, and the ratio of IL-17–producing to Foxp3⁺ cells within ROR γ t⁺ T α β cells (see Fig. 4 D). Right panels show immunofluorescence histology of a colon from a healthy or a treated *Rorc*(γ t)-*Gfp*^{TG} mouse. Bar, 100 μ m. (B) *Rorc*(γ t)-*Gfp*^{TG} mice were infected intranasally with 100 PFUs of influenza A virus for 7 d. Cells were then isolated from the lung and processed as in A. Right panels show immunofluorescence histology of a lung from healthy or an infected *Rorc*(γ t)-*Gfp*^{TG} mouse. Bar, 50 μ m. (C) Cells were isolated from the mesenteric LNs of a 4-mo-old *Rorc*(γ t)-*Gfp*^{TG} \times *Apc*^{min/+} mouse and processed as in A. Right panels show immunofluorescence histology of a mesenteric LN from a normal or a tumor-bearing mouse. Bar, 100 μ m. Data shown are representative of at least three independent experiments. Three to four mice were analyzed per group. *, $P < 0.05$ as compared with control (mock-treated or WT mice).

T cells differentiate in vitro into IL-17–producing, IL-10–producing, or Foxp3⁺ ROR γ t⁺ T α β cells, substantial differences exist with the in vivo ROR γ t⁺ T α β cell subsets in the phenotype of the cells producing IL-10, and the proportion of Foxp3⁺ cells expressing ROR γ t.

DISCUSSION

In this study, we have shown that ROR γ t⁺ T cells include both IL-17– and IL-10–producing Foxp3⁺ T reg cells, which are endowed with proinflammatory and regulatory functions, respectively. In addition, ROR γ t⁺ T cells include T γ δ cells producing high levels of IL-17 in an IL-6–independent pathway. The relative frequency of IL-17– or IL-10–producing ROR γ t⁺ T α β cells is maintained constantly during infection or inflammation, suggesting that a robust mechanism maintains an equilibrium between these two effector arms. However, this equilibrium is perturbed in mice harboring intestinal tumors. IL-6, Foxp3, and CCR6, and to a lesser extent IL-23, are essential regulators of this equilibrium.

Ivanov et al. (39) reported that ROR γ t⁺ T cells or IL-17–producing cells were only present in the intestinal lamina propria. These authors therefore suggested that the intestinal flora induces the differentiation of ROR γ t⁺ IL-17–producing T cells. We show now that although these cells are more abundant in the intestine, sizeable populations of ROR γ t⁺ T

cells were found in all organs examined, including the LNs, spleen, lung, skin, and bone marrow. This extended distribution of ROR γ t⁺ T cells was not due to infection, as specific pathogen-free mice were used in all experiments. This discrepancy might be explained by the sensitivity of the reporter mice that have been used. Ivanov et al. (39) used a knock-in mouse encoding one copy of the *Egfp* reporter gene inserted into the *Rorc*(γ t) locus (41), whereas we used a transgenic mouse encoding 5–10 copies of a BAC carrying a similar *Egfp* insertion and therefore induces a brighter reporter expression. The more extended distribution of ROR γ t⁺ IL-17–producing T cells, in particular in the LNs and spleen, might better explain the occurrence of Th17-dependent inflammatory disease in the brain or joints of mice enduring experimental autoimmune encephalomyelitis or CIA (27–29, 39).

Foxp3⁺ ROR γ t⁺ T α β cells were an important fraction of ROR γ t⁺ T cells. They produced the immunosuppressive IL-10 and functioned as genuine T reg cells in vitro. Furthermore, these cells, but not IL-17–producing ROR γ t⁺ T cells, also expressed high levels of the transcript coding for CCL20. This transcript was undetected in ROR γ t[–] T cells, suggesting that Foxp3⁺ ROR γ t⁺ T cells were the main source of CCL20 in T cells, even though it might be expressed by a small subset of ROR γ t[–] T cells. In accordance with our results, McGeachy et al. (13) have reported that CD4⁺ T cells

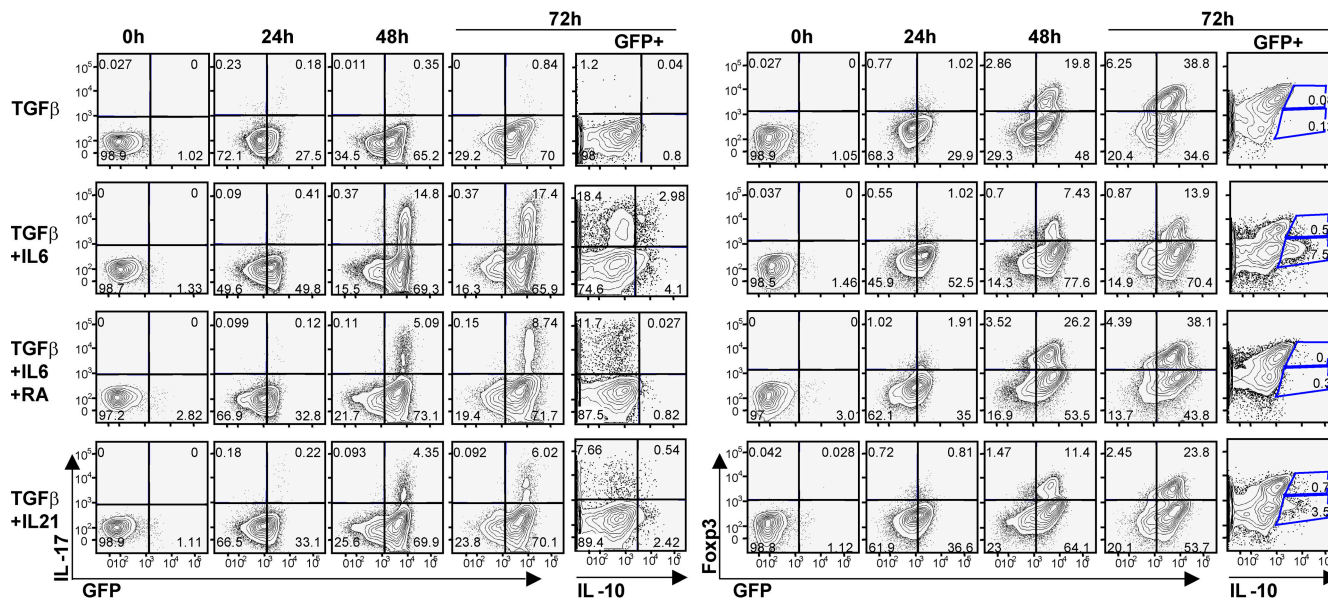


Figure 7. Generation and differentiation of ROR γ t⁺ T α β cells in vitro. MACS-sorted naive (CD62L⁺) CD4⁺ T cells from the spleens of *Rorc*(γ t)-*Gfp*^{fl} mice were stimulated in duplicates with anti-CD3 ϵ and anti-CD28 in the presence of blocking anti-IFN- γ and anti-IL-4 antibodies and the indicated cytokines or RA. After different periods of time, cells were restimulated with PMA/ionomycin for 5 h and analyzed by flow cytometry for the expression of GFP, Foxp3, IL-17, and IL-10. All plots are gated on TCR- β ⁺ cells, except plots for IL-10 that are gated on GFP⁺TCR- β ⁺ cells. Numbers indicate percent cells in quadrants. Data are representative of three independent experiments.

cultured in the presence of TGF- β and IL-6, which produce both IL-17 and IL-10, express *Ccl20*, in contrast to cells cultured with IL-23 alone, which produce only IL-17. CCL20 is the ligand for CCR6, suggesting that Foxp3⁺ ROR γ t⁺ T cells regulate the recruitment of CCR6⁺ T cells. Th17 cells isolated from SKG mice, which spontaneously develop arthritis, as well as human Th17 cells, express CCR6 (44, 48, 49). CCR6 is also expressed by a subset of “memory-like” CD25⁺ T reg cells that produce elevated levels of *Il10* transcripts (54). We found that 35–50% of ROR γ t⁺ T α β cells expressed CCR6, regardless of their expression of CD25 (or Foxp3). Importantly, CCR6-deficient mice generated an increased ratio of IL-17- to IL-10-producing ROR γ t⁺ T α β cells as compared with WT mice. These results suggest that Foxp3⁺ ROR γ t⁺ T cells producing IL-10 and CCL20 recruit CCR6⁺ ROR γ t⁺ T cells and negatively regulate their differentiation into IL-17-producing cells.

Foxp3⁺ ROR γ t⁺ T cells also expressed elevated levels of the costimulatory receptor gene *Icos*, shown to be essential for the induction of CIA and the optimal generation of Th17 cells (43, 55). The expression of ICOS is also induced by IL-23 in human cells (56). Furthermore, Foxp3⁺ ROR γ t⁺ T cells specifically expressed the metalloproteinase gene *Mmp9*, induced by IL-23 in a mouse tumor model (57), by IL-17 in the lung (45) and by IL-17R in inflamed joints (46), as well as elevated levels of the fractalkine gene *Cx3cl1*. The functional consequences of elevated expression of *Icos*, *Mmp9*, and *Cx3cl1* by Foxp3⁺ ROR γ t⁺ remain to be explored, but this expression pattern appears paradoxical as the product of these genes might promote rather than modulate inflammation

through the recruitment and activation of effectors, as well as through tissue remodeling through MMP-9. It might be suggested that the role of Foxp3⁺ ROR γ t⁺ T cells is not merely to regulate inflammation through IL-10, but also to participate in the recruitment of inflammatory cells such as Th17 cells to an immunosuppressive environment. On the other hand, we cannot formally exclude that Foxp3⁺ ROR γ t⁺ T cells contain distinct functional subsets of IL-10-producing and/or CCL20-, ICOS-, MMP-9-, and CX₃CL1-expressing cells.

Of note, 30–50% of ROR γ t⁺ T α β cells did not express IL-17, IL-10, or Foxp3. We could not purify these cells from IL-17 producers for transcript analysis, as no markers were found to distinguish these cells before intracellular cytokine staining. It is possible that non-IL-17 and non-IL-10 producers contained uncommitted ROR γ t⁺ T cells, i.e., not yet committed to either the Th17 or the T reg cell pathway. They could also represent a third committed subpopulation that remains to be characterized. In favor of the first hypothesis, only few IFN- γ - and no IL-4-producing cells (not depicted) could be detected in ROR γ t⁺ T cells.

ROR γ t⁺ T γ δ cells have been reported previously to represent 50% of total T γ δ cells in the lamina propria of the small intestine, a majority of them expressing IL-17 (39). In the CIA model, 60–80% of V γ 4⁺ T γ δ cells and an equivalent number of CD4⁺ T α β cells produce IL-17 in the draining LNs (58). Furthermore, T γ δ cells are a primary source of IL-17 in the lung before and during infection with *Mycobacterium tuberculosis* (59) or *Mycobacterium bovis* (60), and V δ 1⁺ T γ δ cells produce IL-17 during intraperitoneal infection with *Escherichia coli* (61). In accordance with these data, we found

that ROR γ t⁺ T γ δ cells were a major source of IL-17 in the lung and skin, as >80% of them produced IL-17, representing half of the total ROR γ t⁺ T cell population in these compartments. ROR γ t⁺ T γ δ cells expressed no Foxp3, but produced the highest levels of ROR γ t, as well of *Il23r* and *csf2* coding for GM-CSF. The latter induces the differentiation of granulocytes and monocytes, as well as the production of proinflammatory cytokines such as TNF- α and IL-6 (62). In contrast to ROR γ t⁺ T α β cells, IL-17-producing ROR γ t⁺ T γ δ cells were not affected by the absence of IL-6 in deficient mice, indicating that ROR γ t⁺ T γ δ cells and IL-17-producing ROR γ t⁺ T α β cells are generated by distinct pathways. Collectively, these data suggest that ROR γ t⁺ T γ δ cells play an important role in inflammation to pathogens through the production of IL-17 and GM-CSF; however, their precise role in vivo remains to be assessed.

ROR γ t is required for the generation of Th17 cells and has been shown to induce Th17 cells when transduced into mouse naive CD4⁺ T cells (39). Furthermore, naive CD4⁺ T cells cultured in the presence of TGF- β and IL-6 or IL-21 generate Th17 cells (4–6, 10–12) and IL-17-producing ROR γ t⁺ T cells (Fig. 4). We find that a large fraction of ROR γ t⁺ T α β cells expressed Foxp3, in particular in the LNs, as well as the immunosuppressive IL-10, and functioned as T reg cells. It has been shown previously that culture conditions that favor the generation of Th17 cells from naive CD4⁺ T cells, i.e., in the presence of TGF- β and IL-6, also induce the generation of IL-10-producing cells (13, 17). But in contrast to the in vivo situation, IL-10 was produced by Foxp3⁻ rather than Foxp3⁺ ROR γ t⁺ T cells, indicating that these culture systems might not faithfully replicate in vivo pathways. Interestingly, RA blocked the generation of IL-10-producing cells, whereas it favored the generation of Foxp3⁺ ROR γ t⁺ T cells. Furthermore, cultures of CD4⁺ T cells in the presence of TGF- β and IL-6 or IL-21 generated ROR γ t⁺ T cells expressing both IL-17 and IL-10, as observed previously (13, 17). However, ROR γ t⁺ T cells producing both cytokines could not be detected in vivo, exposing potential conflicts between in vivo and in vitro differentiation pathways.

Most cells cultured in the presence of TGF- β alone expressed ROR γ t after 1–2 d, but then differentiated into Foxp3⁺ ROR γ t⁺ T cells that did not express IL-17 after 2–3 d. It has been previously reported that TGF- β alone induces significant levels of *Rorc*(γ t) transcripts but no detectable *Il17* or *Il23r* transcripts (4–6). These data indicate that ROR γ t is necessary but not sufficient for the generation of Th17 cells from naive CD4⁺ T cells, in accordance with the observation that cells cultured in the presence of TGF- β and IL-6 express ROR γ t after 1–2 d, but express IL-17 only after 2–3 d of culture (Fig. 4) (39). Collectively, these data indicate that TGF- β is opening a ROR γ t-dependent differentiation pathway in CD4⁺ T cells that leads either to Th17 or T reg cells, depending on the presence of maturation and polarizing factors such as IL-6, IL-21, RA, IL-23, and possibly IL-10.

The ratio of Th17 to T reg cells within ROR γ t⁺ T cells varied between organs, with the highest proportions of Th17 cells in the intestine and lung, and of Foxp3⁺ ROR γ t⁺ T cells in the LNs. This ratio was tuned in favor of Th17 cells by IL-6 and IL-23, as the ratio of IL-17-producing to Foxp3⁺ ROR γ t⁺ T cells decreased in IL-6- or IL-12R β 1-deficient mice. However, in IL-6-deficient mice, the total number of ROR γ t⁺ T α β cells was also significantly diminished, indicating that IL-6 is involved more generally in the generation of ROR γ t⁺ T α β cells, as observed previously (6, 39). Conversely, the ratio of Th17 to T reg cells within ROR γ t⁺ T cells was tuned in favor of T reg cells by Foxp3 and CCL20, as inferred from scurfy mice or CCR6-deficient mice. Foxp3 might regulate the fate of “uncommitted” ROR γ t⁺ T cells into IL-17- or IL-10-producing cells by directly modifying the molecular function of ROR γ t, as Foxp3 was coimmunoprecipitated with ROR γ t. Foxp3 might also act through Foxp3⁺ ROR γ t⁺ T cells or “mainstream” Foxp3⁺ T cells regulating ROR γ t⁺ T cells. Interestingly, cells derived from scurfy mice generated a decreased proportion of ROR γ t⁺ T α β cells and IL-17-producing ROR γ t⁺ T α β cells, indicating a more complex role for Foxp3 in the generation and regulation of ROR γ t⁺ T α β cells that remains to be elucidated.

A robust mechanism maintains an equilibrium between Th17 to T reg cells within ROR γ t⁺ T cells. During massive inflammation induced by DSS in the intestine, the total number of ROR γ t⁺ T cells increased 10-fold in the intestine and >10-fold in influenza A virus-infected lungs. However, the numbers of IL-17- and IL-10-producing (Foxp3⁺) ROR γ t⁺ T cells were increased in comparable proportions. Regulating IL-17 versus IL-10 production might promote inflammation while limiting “collateral” damage, a necessary compromise between effective immunity and tissue integrity. IL-10-deficient mice, which lack one of these effectors arms, generate increased proportions of IL-17-producing CD4⁺ T cells and develop spontaneous colitis in an IL-23-dependent pathway (15). In contrast, *Apc*^{min/+} mice that developed large colon and ileal polyps and tumors by 3–4 mo of age generated an increased proportion of IL-10-producing ROR γ t⁺ T cells. Similar observations have been made in mice injected with the B16 melanoma, which develop an increased ratio of Foxp3⁺ cells versus IL-17-producing cells as early as 8 d after tumor injection (53). The mechanisms used by tumors to alter the equilibrium between IL-17- and IL-10-producing ROR γ t⁺ T cells remain to be investigated, but this phenomenon is reminiscent of the IL-10-mediated immunosuppression induced by diverse viruses and bacteria (63). In contrast, the equilibrium between IL-17- and IL-10-producing ROR γ t⁺ T cells might be disrupted in favor of Th17 cells in chronic inflammatory diseases such as rheumatoid arthritis, multiple sclerosis, or inflammatory bowel disease. Tools to control the fate of ROR γ t⁺ T cells might therefore prove useful either to boost or to limit IL-17-producing cells. Potential targets include ROR γ t as well as any factors that regulate the differentiation of ROR γ t⁺ T cells into IL-17- or IL-10-producing effectors.

MATERIALS AND METHODS

Mice. BAC-transgenic *Rorc*(γ)-*Gfp*^{TG} or *Rorc*(γ)-*Cre*^{TG} mice were generated as described previously (40). The coding sequence for EGFP or Cre, including the stop codon, was inserted into exon 1 of *Rorc*(γ) in place of the endogenous ATG translation start codon, on a 200-kb BAC (Invitrogen) carrying at least 70 kb of sequence upstream of the *Rorc*(γ) translation start site. Transgenic mice used in transfer experiments were backcrossed eight times to C57BL/6 mice. CCR6-deficient (50), *Rosa-Stop^{fl}-Dta* (51), and *Rag2^{-/-}Il2rb^{-/-}* (64) mice have been described previously. *Foxp3^{fl}*, *Il12rb1^{-/-}*, *Il6^{-/-}*, and *Apc^{min/+}* mice were obtained from The Jackson Laboratory. All mice were kept in specific pathogen-free conditions, and all animal experiments were approved by the committee on animal experimentation of the Institut Pasteur and by the French Ministry of Agriculture.

Antibodies. The following mAbs were purchased from BD Biosciences: PE-conjugated anti-IL-17 (TC11-18H10), PerCP-conjugated anti-CD4 (RM4-5) and purified anti-CD3 ϵ (145-2C11), and anti-IFN- γ (XMG1.2). The following were purchased from eBioscience: PE-conjugated anti-IL-10 (JES5-16E3), TCR- δ (GL3), TCR- β (H57-597), and rat IgG1 isotype control; APC-conjugated anti-IL-10 (JES5-16E3), CD45.2 (104), TCR- β (H57-597), and rat IgG2b isotype control; PE-Cy5-conjugated anti-Foxp3 (FJK-16S); PE-Cy7-conjugated anti-CD4 (GK1.5), CD8 α (53-6.7), and streptavidin; biotinylated anti-TCR- δ (GL3), CD25 (PC61.5), and CD45.2 (104); APC-Alexa750-conjugated anti-CD3 ϵ (17A2); Alexa647-conjugated anti-IFN- γ (XMG1.2); and purified anti-CD28 (37.51) and anti-IL-4 (11B11). PE-conjugated anti-CCR6 was from R&D Systems. Purified anti-GFP (A-11122) and FITC-conjugated anti-rabbit polyclonal antibodies were from Invitrogen. Anti-ROR γ was described previously (41).

Flow cytometry. Fragments of small intestine or colon were first incubated in PBS (Ca/Mg free) containing 15 ml of 1 mM DTT and 3 mM EDTA, and then for 40 min at 37°C in DMEM medium containing 1 mg/ml collagenase (Roche) and 1 U/ml DNase (Invitrogen). Lung and ear skin were directly incubated for 40 min to 1 h at 37°C in DMEM medium containing collagenase and DNase. Tissue suspensions were then pressed through a 100- μ m mesh, pelleted, resuspended in a 40% isotonic Percoll solution (GE Healthcare), and underlaid with an 80% isotonic Percoll solution. Centrifugation for 20 min at 2,000 rpm yielded the mononuclear cells at the 40–80% interface. Cells were finally washed twice with PBS-F (PBS containing 2% FCS). Single cell suspensions were prepared from the thymus, spleen, and LNs by pressing the tissues through a 100- μ m mesh and from the bone marrow by flushing the long bones with PBS. All cells were first preincubated with mAb 2.4G2 to block Fc γ receptors, and then washed and incubated with the indicated mAb conjugates for 40 min in a total volume of 100 μ l PBS-F. For intracellular cytokine staining, cells were stimulated for 5 h in complete medium in the presence of 50 ng/ml phorbol 12-myristate 13-acetate and 500 ng/ml ionomycin (both from Sigma-Aldrich). For the last 2 h, 10 μ g brefeldin A (Sigma-Aldrich) was added to the cultures. After surface staining for CD3, CD4, TCR- β , and/or TCR- δ , cells were fixed with fresh 2% paraformaldehyde in PBS for 20 min. Intracellular staining was performed in permeabilization-solution (1% saponin in FACS buffer) with anti-IL-17 (BD Biosciences), anti-IFN- γ , anti-Foxp3, and/or anti-IL-10. Polyclonal anti-GFP was used to increase GFP signal during intracellular staining. Cells were analyzed on a FACSCanto (BD Biosciences), followed by analysis with FlowJo software (Tristar).

In vitro T cell differentiation. Spleen and mesenteric LN cells from *Rorc*(γ)-*Gfp*^{TG} mice were pooled and T cells were purified using a CD4⁺ T cell isolation kit (Miltenyi) and further sorted into naive CD4⁺CD62L⁺ cells using CD62L Microbeads (Miltenyi). 3×10^5 cells were stimulated for 1–3 d in complete DMEM with 5 μ g/ml anti-CD3, 5 μ g/ml anti-CD28, 10 μ g/ml of neutralizing anti-IFN- γ , and 10 μ g/ml anti-IL-4 in the presence of 10 ng/ml human TGF- β , 10 ng/ml mouse IL-6, 100 ng/ml mouse IL-21, and/or 100 nM RA (Sigma-Aldrich).

In vitro proliferation assay. 2.5×10^4 FACS-sorted CD4⁺CD25⁻ T cells were labeled with CFSE using CellTrace Cell Proliferation kit (Invitrogen) and cultured alone or in the presence of 2.5×10^4 (or 0.625×10^4) of either GFP⁺ or GFP⁻ CD4⁺CD25⁺ T cells sorted from the spleen and LNs of *Rorc*(γ)-*Gfp*^{TG} mice. T cells were stimulated in duplicates with irradiated splenic feeder cells (10^5 per well) and 1 μ g/ml of soluble anti-CD3 in 96-well U-bottom plates for 96 h.

Gene expression analysis. To obtain RNA for gene expression analysis by real-time RT-PCR, 500–5,000 cells were directly sorted into vials containing RLT buffer (QIAGEN) supplemented with β -mercaptoethanol, and total RNA was extracted using an RNeasy Micro kit (QIAGEN). The quality of total RNA was assessed using the 2100 Bioanalyzer system (Agilent Technologies). 250–500 μ g of high quality total RNA was subjected to one linear mRNA amplification cycle using the MessageBooster kit for quantitative RT-PCR (Epicentre Biotechnologies). 50–100 ng of amplified mRNA was then converted to cDNA using the RT² PCR Array first strand kit (SuperArray Bioscience Corporation). All procedures were performed according to the manufacturer's protocols. The expression of 84 different genes was measured using the mouse Th17 RT² Profiler PCR Array (SuperArray Bioscience Corporation). Real-time PCR was performed on a PTC-200 thermocycler equipped with a Chromo4 detector (Bio-Rad Laboratories). Data were analyzed using Opticon Monitor software (Bio-Rad Laboratories).

Immunofluorescence histology. Tissues were washed and fixed overnight at 4°C in a fresh solution of 4% paraformaldehyde (Sigma-Aldrich) in PBS. The samples were then washed for 1 d in PBS, incubated in a solution of 30% sucrose (Sigma-Aldrich) in PBS until the samples sank, embedded in OCT compound 4583 (Sakura Finetek), frozen in a bath of isopentane cooled with liquid nitrogen, and stocked at -80°C . Frozen blocs were cut at 8- μ m thickness, and sections were collected onto Superfrost/Plus slides (VWR). Slides were dried for 1 h and processed for staining or stocked at -80°C . For staining, slides were first hydrated in PBS-XG (PBS containing 0.1% Triton X-100 and 1% normal goat serum; Sigma-Aldrich) for 5 min and blocked with 10% bovine serum in PBS-XG for 1 h at room temperature. Endogenous biotin was blocked with a biotin blocking kit (Vector Laboratories). Slides were then incubated with primary polyclonal antibody or conjugated mAb (in general 1/100) in PBS-XG overnight at 4°C, washed three times for 5 min with PBS-XG, incubated with secondary conjugated polyclonal antibody or streptavidin for 1 h at room temperature, washed once, incubated with DAPI (Sigma-Aldrich) for 5 min at room temperature, washed three times for 5 min, and mounted with Fluoromount-G (SouthernBiotech). Slides were examined under an AxioImager M1 fluorescence microscope (Carl Zeiss, Inc.) equipped with a CCD camera, and images were processed with AxioVision software (Carl Zeiss, Inc.).

Immunoprecipitation. A FLAG-encoding sequence was added to the 5' end of the *Rorc*(γ) cDNA by PCR and cloned into the pIRES2-EGFP vector (Clontech Laboratories, Inc.), and the *FoxP3* cDNA was cloned into the pIRES-DsRed vector (Clontech Laboratories, Inc.). 4 μ g of each construct was cotransfected into 1.106 293T cells using Lipofectamine 2000 (Invitrogen). After 48 h, immunoprecipitation was performed using the FLAG Tagged Protein Immunoprecipitation kit (Sigma-Aldrich) and resolved on a 10% gel bis-tris acrylamide gel.

DSS and influenza. To induce colitis, *Rorc*(γ)-*Gfp*^{TG} mice were treated with 2.5% DSS (MP Biomedicals) in the drinking water for 6 d, followed by water for 10 d. This protocol was repeated for a total of three cycles. To infect lungs, *Rorc*(γ)-*Gfp*^{TG} mice were infected intranasally with 100 PFUs of influenza A virus (H3N2 strain Scotland/20/74), and mice were killed after 7 d.

Chimeras. Bone marrow cells isolated from CD45.2⁺ *Rorc*(γ)-*Gfp*^{TG} or *Foxp3^{fl} × Rorc*(γ)-*Gfp*^{TG} mice were treated with a lineage depletion kit (Miltenyi). Purified cells were then mixed in a 1:1 ratio with lin⁻ bone marrow cells from CD45.1⁺ C57BL/6 mice, and a total of 4×10^5 was injected

into *Rag2^{-/-}Il2rb^{-/-}* mice (64) irradiated at 600 rads. After 4 wk, reconstitution of the mice was assessed in peripheral blood. Reconstituted mice were killed and analyzed 6 wk after transfer.

Online supplemental material. In Fig. S1, the purification of the different subsets of ROR γ ⁺ T cells is shown. Fig. S2 shows that ROR γ ⁺ T cells express genes involved in the Th17 or T reg cell pathway. In Fig. S3, expression of IL-10 by intestinal ROR γ ⁺ T α β cells is depicted. Fig. S4 shows the expression of CCR6 by ROR γ ⁺ T α β cells. Fig. S5 shows the generation of mixed bone marrow chimeras, and Fig. S6 shows the generation of Foxp3⁺ T cells in mice ablated of ROR γ ⁺ cells. In Fig. S7, immunofluorescence histology of terminal ileum from a WT or a 4-mo-old *Rorc*(γ t)-*Cjfp*^{TG} x *Apc*^{min/+} mouse developing polyps and carcinoma is shown. Figs. S1–S7 are available at <http://www.jem.org/cgi/content/full/jem.20080034/DC1>.

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