Psoriasis is a common immune-mediated chronic inflammatory skin disorder, but the mechanisms of pathogenesis are still poorly understood. IL-23 is expressed in psoriatic skin, and IL-23 injection produces IL-22–dependent psoriasiform changes in mouse skin. Th17 cells produce IL-22 and display CCR6, the CCL20 receptor; CCR6+ T cells and CCL20 are abundant in psoriatic skin. We investigated a possible role for CCR6 in recruiting Th17 cells and producing psoriasiform pathology by injecting IL-23 into the skin of WT and Cer6−/− mice. Unlike for WT mice, IL-23–injected ears of Cer6−/− mice showed neither substantial epidermal/dermal changes nor increased Il22 mRNA expression. However, injection of IL-22 yielded equivalent psoriasiform changes in WT and Cer6−/− mice. Surprisingly, IL-23–injected ears of WT and Cer6−/− mice contained similar numbers of Th cells able to make IL-17A and/or IL-22. Furthermore, in ears of Rag1−/− mice, IL-23 initially induced skin changes and levels of Il22 mRNA that were indistinguishable from WT mice, revealing at least one non–T cell source for IL-22. We conclude that CCR6 is essential in a model of IL-23–induced, IL-22–mediated dermatitis, which develops in sequential T cell–independent and T cell–dependent phases. These findings reveal an expanded role for CCR6 in IL-23–related responses and identify CCR6 as a potential therapeutic target in psoriasis.

Introduction

Psoriasis is one of the most common immune-mediated chronic inflammatory skin disorders in adults, affecting 1%–2% of white individuals, with a considerable impact on quality of life (1). Defining histological features of psoriatic skin lesions include epidermal hyperplasia (acanthosis), caused by aberrant terminal differentiation and hyperproliferation of epidermal keratinocytes, as well as marked infiltration of mononuclear leukocytes into the dermis (2). While an underlying genetic component to the disease has been acknowledged for many years (1–3), important aspects of the complex autoimmune inflammatory mechanisms involved in the pathogenesis of psoriasis are still poorly understood.

Experimental data from various animal models of psoriasis, as well as the clinical effectiveness of newer systemic therapeutics targeting activated T lymphocytes, strongly implicate T cells in the pathogenesis of this disease (2). Psoriasis has traditionally been classified as a Th1-mediated disease. One reason is the abundance of IFN-γ-producing, Th1-polarized T cells within psoriatic skin. A second reason is that TNF-α, another type 1 cytokine produced by several skin immune cells, is highly upregulated in psoriatic lesions (3, 4). TNF-α undoubtedly plays a pathogenic role in the development of psoriasis, as TNF-α inhibitors, such as etanercept and infliximab, are highly effective therapies (5–8).

In spite of the data implicating Th1 cells in psoriasis, recent research has drawn attention to the significance of Th17 cells, a newly identified Th cell subset, in the pathogenesis of psoriasis and other autoimmune inflammatory disorders (9–13). Th17 cells and their downstream effector molecules, which include IL-17A, IL-17F, TNF-α, and IL-22 (14, 15), have been found at increased levels in human psoriatic skin (16–20). In support of a role for Th17 cells in psoriasis, Zaba et al. demonstrated that reduced Th17 responses in patients treated with etanercept correlated with clinical improvement (21). Moreover, psoriatic skin lesions contain high levels of IL-23 compared with nonlesional and normal skin (20, 22, 23). IL-23 is a heterodimeric cytokine that shares its p40 subunit, which includes IL-17A, IL-17F, TNF-α, and IL-22 (14, 15), with other proinflammatory cytokines, such as IL-23. Genome-wide screening has revealed associations between the development of psoriasis and polymorphisms in the genes not only for the p40 subunit but also for the p19 subunit and the IL–23–specific chain of the IL-23 receptor (29–32); moreover, ustekinumab, an antibody that blocks the activity of p40, has produced dramatic clinical responses in patients with chronic plaque psoriasis (33–35).

In support of a role for the IL-23/Th17 axis in psoriasis, it was recently reported that injection of IL-23 into the skin of WT mice induces some psoriasiform changes, such as acanthosis, hyperparakeratosis, and dermal inflammatory infiltrates, yet fails to induce other aspects of the human disease, such as epidermis-infiltrating CD8+ T cells (36, 37). IL-22, a member of the IL-10 family of cytokines and a product of activated Th17 cells, is a critical mediator of the psoriasiform changes, since IL-22 deficiency abrogates much of the acanthosis and dermal inflammation induced by IL-23 (38). In addition, in a model using adoptive transfer of CD4+CD45RBhiCD25- cells into scid mice, neutralizing IL-22 pre-
vented the psoriasis-like skin disease (39). In human studies, IL-22 has been found to produce psoriasiform changes in reconstituted epidermis (40), increased expression of IL-22 has been found in psoriatic skin, and serum levels of IL-22 are elevated in patients with chronic plaque psoriasis—correlating positively with severity of disease and negatively with response to therapy (41–43). Until now, infiltrating Th17 cells have been suggested to be the source of the IL-22 in skin (38, 43). These recent data strongly implicate Th17 cells along with their upstream cytokine, IL-23, and their downstream effector cytokine, IL-22, in the pathogenesis of psoriasis. Such advances notwithstanding, the factors important for migration, maintenance, and function of Th17 cells at the site of cutaneous inflammation are not well understood. Chemokines and their receptors are essential components in leukocyte migration (44–46), and studies of the differential expression of chemokine receptors have been helpful in characterizing subsets of human T cells with distinct migratory and effector profiles (47, 48). For example, the chemokine receptor CCR6 is found on virtually all human CD4+ T cells capable of producing IL-17A, IL-17F, and IL-22 (refs. 13, 49–52, and S.P. Singh and J.M. Farber, unpublished observations), and CCL20, the sole chemokine ligand for CCR6, is produced by Th17/CCR6+ cells (refs. 13, 23, and S.P. Singh and J.M. Farber, unpublished observations). These relationships make observations on CCR6 and CCL20 in psoriasis of particular interest. CCL20 is highly expressed in the epithelium of psoriatic skin and is inducible by IL-1β, IFN-γ, and IL-17A in multiple skin cell types (53). CCR6 is found on a significant fraction of CLA+ (skin homing) CD4+ T cells (53, 54) and on approximately 75% of CD4+ T cells resident in normal skin (55) as well as on T cells in psoriatic epidermis (56).

CCR6 is a chemokine receptor that was discovered initially by its expression on subsets of dendritic cells and T cells (57). CCR6 has been reported to be of particular importance at mucosal sites. Compared with WT mice, Ccr6−/− mice have smaller Peyer patches (58), fewer associated M cells (58), and defective formation of isolated lymphoid follicles (59). Abnormalities in intestinal immune responses have been reported for Ccr6−/− mice in models of inflammatory bowel disease (60) and during infection with rotavirus (61) and Salmonella typhimurium (62). In the skin, Ccr6−/− mice have been reported to develop either more (58) or less (63) severe contact hypersensitivity as well as diminished delayed-type hypersensitiv-

**Figure 1**

*Ccr6−/−* mice are resistant to IL-23–induced acanthosis and dermal inflammation. Ears of WT and *Ccr6−/−* mice were injected intradermally every other day for 16 days with 20 μl PBS, either alone or containing 500 ng IL-23. (A) Ear thickness was measured on days between injections. Data are from 5 experiments in at least 30 mice/group. **P < 0.01 versus all other groups. (B) H&E-stained sections of PBS- or IL-23–injected ears from WT and *Ccr6−/−* mice at day 15. k, hyperparakeratosis with intraconal neutrophil pustule; a, acanthosis; b, increased proliferative activity of basal layer epidermal keratinocytes; d, dermal mononuclear cell infiltrate; v, telangiectasia of dermal blood vessels. Sections are representative of 2 experiments. (C) Frozen sections of ears from WT PBS-injected or WT and *Ccr6−/−* IL-23–injected mice on day 15 stained with rat anti-mouse CCL20 or isotype-matched control antibody followed by Alexa Fluor 488–conjugated goat anti-rat IgG (green). Nuclei were counterstained with DAPI. Staining is representative of 2 experiments. (D) mRNA for *Ccl20* and *Gapdh* were measured by real-time RT-PCR at day 15 from ears of WT PBS-injected or WT and *Ccr6−/−* IL-23–injected mice. Fold changes normalized for *Gapdh* mRNA are shown versus the PBS-injected ears. Data are from 3 experiments in at least 5 mice/group. *P < 0.05 versus PBS control. Original magnification, ×400.
Results

CCR6 is required for IL-23–induced psoriasiform inflammation. Consistent with previous reports (36–38), intradermal injection of ears of WT mice with IL-23 induced marked ear swelling. Ear thickness increased by more than 100 μm compared with PBS-injected controls after injecting IL-23 every other day over the 16-day experiment (Figure 1A). To test the role of CCR6 in this response, we used Ccr6−/− mice that we generated by gene targeting (Supplemental Figure 1; supplemental material available online with this article; doi:10.1172/JCI37378DS1). As shown in Figure 1A, the ears of the Ccr6−/− mice did not swell after injections of IL-23. H&E staining of sections from injected WT ears on day 15 revealed the epidermal changes described previously, namely acanthosis (thickening of the spinous layer), hyperparakeratosis (retention of nuclei within the stratum corneum), dermal inflammatory infiltrates, and telangiectasis (dilated capillaries and venules; Figure 1B). Occasional intracorneal neutrophils were found, but were not conspicuous (see below). Conversely, IL-23 injection had a minimal effect on Ccr6−/− ears, comparable to that on the PBS-injected controls.

If the role of CCR6 in this model is a direct one at the site of injection, then CCL20, the sole chemokine ligand for CCR6, should be induced in the injected ears. CCL20 protein and Ccl20 mRNA were indeed detected by immunohistochemistry (Figure 1C) and RT-PCR (Figure 1D), respectively. Prominent upregulation of CCL20 was seen in acanthotic mouse epidermis, as previously described for psoriatic skin (53). CCL20 protein and Ccl20 mRNA were induced similarly in WT and Ccr6−/− ears, which suggests that CCR6 itself is not required for their upregulation.

Ccr6−/− mice do not accumulate CD4+ T cells or CD11c+CD11b+ dendritic cells in response to intradermal IL-23 injection. To determine the composition of the IL-23–induced inflammatory infiltrate and quantify differences between WT and Ccr6−/− ears, we prepared single-cell suspensions from treated skin and analyzed them by flow cytometry. Injections with IL-23 led to an approximately 2-fold increase in total CD4+ T cells compared with control PBS injection in WT ears, but no such increase was observed in Ccr6−/− ears (Figure 2A). Analysis of CCR6+CD4+ T cells in WT mice showed that this subset also increased in the IL-23–injected ears (Figure 2B). We found no significant increases in NK1.1+ cells, γδ T cells, macrophages, or neutrophils in the IL-23–injected ears and no difference in the number of these cells between WT and Ccr6−/− ears (data not shown). Trivial numbers of CD8+ T cells were present in the ears, again not increasing after IL-23 injections and not different between WT and Ccr6−/− mice (data not shown). We also found no differences in the number of CD4+ T cells in the noninjected ears between WT and Ccr6−/− mice (data not shown). Differences in cell numbers in the IL-23–injected ears of WT and Ccr6−/− mice appeared to be limited to the site of injection, because the number of CD4+ T cells was similar in the draining cervical lymph nodes and was not affected by IL-23 injections (Figure 2C). For the CD11c+CD11b+ subset of dendritic cells, results were similar to that seen for the CD4+ T cells: a 2-fold increase in WT ears, but not Ccr6−/− ears, after IL-23 injections (Figure 2D). In contrast, no difference between Ccr6−/− and WT mice was detected with regard to the number of CD11c+CD11b+ cells in noninjected ears or in the cervical lymph nodes of IL-23–injected or PBS-injected animals (data not shown).

Ccr6−/− mice produce less IL-22 after IL-23 injections. IL-22 has been previously reported to be induced in ears after IL-23 injection and to be critical for the IL-23–induced changes in the skin (38). Consistent with this report, we found elevated levels of Il22 mRNA in the skin (58) and graft-versus-host–mediated injury (64). In this study, we assessed the role of CCR6 in psoriasiform changes after injection of IL-23 into mouse skin and found that the IL-23–induced pathology was CCR6 dependent. Furthermore, contrary to our expectations, our findings suggest that the requirement for CCR6 is not through a direct effect on T cells, but rather by establishing an inflammatory response that is initially T cell independent and that includes a non–T cell source of IL-22.
the IL-23–injected WT ears. However, induction of IL22 mRNA in Ccr6−/− ears was much reduced (Figure 3A). If lower levels of IL-22 induction in the ears were responsible for the overall failure of the Ccr6−/− mice to respond to IL-23, then injecting IL-22 into the ears would be expected to produce similar changes in Ccr6−/− and WT mice. WT and Ccr6−/− ears swelled in parallel in response to IL-22 (Figure 3B), although IL-22 produced somewhat less swelling in the WT mice than did IL-23 (compare with Figure 1A). H&E staining of sections from the IL-22–injected ears showed acanthosis, dermal inflammation, and dilatation of dermal vasculature that were comparable in the WT and Ccr6−/− mice (Figure 3C), although changes in the WT mice were again more modest than those seen after IL-23 injections (compare with Figure 1B). As done for the ears in the IL-23 experiments, we prepared single-cell suspensions from PBS- and IL-22–injected ears and analyzed them by flow cytometry. IL-22 injection induced an approximately 2-fold increase in the number of CD4+ T cells in both WT and Ccr6−/− mice (Figure 3D), similar to the increases seen after IL-23 injection in the WT mice (compare

Figure 3
IL-22 production deficiency is the critical factor in protection against psoriasis-like changes in Ccr6−/− mice. (A) Ears were injected as in Figure 1, and mRNAs for Il22 and Gapdh were measured by real-time RT-PCR at day 15. mRNA for Il22 was undetectable (UD) in PBS-injected ears; Ct for these samples was set at 37, the limit of detection determined experimentally. Minimal fold changes were calculated for Il22 mRNA normalized for Gapdh mRNA versus the PBS-injected sample with the greatest ΔCt. Data are from 5 experiments in 12 IL-23–injected and 4 PBS-injected mice/group. *P < 0.05 versus all other groups. (B–D) Ears were injected as in Figure 1, except that 500 ng IL-22 was used in place of IL-23. (B) Ear thickness, measured as in Figure 1A. Data are from 2 experiments in 16 mice/group. *P < 0.05, IL-22–injected versus PBS control. (C) H&E-stained sections of PBS- or IL-22–injected ears of WT and Ccr6−/− mice on day 15. Sections are representative of 2 experiments. (D) Total numbers of CD4+ T cells and CD11c+CD11b+ (dendritic) cells from PBS- or IL-22–injected ears from WT and Ccr6−/− mice, measured as in Figure 2. Data are from 5 mice/group in 1 representative experiment of 2. *P < 0.05 versus PBS control. Other pairwise comparisons did not yield significant differences. Original magnification, ×400.
with Figure 2A). However, unlike the IL-23 injections, IL-22 injection did not induce the accumulation of CD11c<sup>+</sup>CD11b<sup>+</sup> dendritic cells (Figure 3D). Moreover, we found no significant increases in CD8<sup>+</sup> T cells, NK1.1<sup>+</sup> cells, γδ T cells, macrophages, or neutrophils in IL-22–injected ears (data not shown).

We presumed that the lack of IL-22 in the ears of Ccr6<sup>−/−</sup> mice was due to a failure of IL-22–producing cells to traffic to the site. However, to rule out an essential defect in the ability of cells from Ccr6<sup>−/−</sup> mice to make IL-22, we performed experiments using splenocytes and CD4<sup>+</sup> T cells ex vivo. Naive CD4<sup>+</sup> T cells purified from WT and Ccr6<sup>−/−</sup> mice were cultured for 7 days under either nonpolarizing or Th17-polarizing conditions. After 7 days, cells were treated with PBS or 100 ng/ml IL-23 for 12 hours, and mRNAs for IL-22 and Gapdh were measured by real-time RT-PCR. Fold changes were calculated for values of IL-22 mRNA normalized for Gapdh mRNA versus the nonpolarized sample with the greatest ΔCt within each plot. Values denote cells from WT or Ccr6<sup>−/−</sup> mice in 1 representative experiment of 3.

Figure 4

CD4<sup>+</sup> T cells from Ccr6<sup>−/−</sup> mice are able to differentiate into Th17/IL-22–producing cells. (A) Naive CD4<sup>+</sup> T cells (CD62L<sup>+</sup>CD44<sup>−</sup>CD25<sup>−</sup>) were purified from WT or Ccr6<sup>−/−</sup> mice and activated and cultured under nonpolarizing (NP) or Th17-polarizing conditions. After 7 days, cells were treated with PMA and ionomycin, and mRNAs for Il17a, Il22, and Gapdh were measured by real-time RT-PCR. Fold changes were calculated for Il17a and Il22 mRNAs normalized for Gapdh mRNA versus the nonpolarized sample with the greatest ΔCt within each plot. Values denote cells from WT or Ccr6<sup>−/−</sup> mice from a single representative experiment of 4. (B) Splenocytes were cultured in medium alone or with 100 ng/ml IL-23 for 12 hours, and mRNAs for Il22 and Gapdh were measured by real-time RT-PCR. Fold changes were calculated for values of Il22 mRNA normalized for Gapdh mRNA versus WT cells in medium alone. Data are from 3 samples from WT or Ccr6<sup>−/−</sup> mice in 1 representative experiment of 2. *P < 0.05 versus medium. Values for IL-23–treated splenocytes from WT and Ccr6<sup>−/−</sup> mice were not significantly different.

IL-23 injection did not induce the accumulation of CD11c<sup>+</sup>CD11b<sup>+</sup> dendritic cells (Figure 3D). Moreover, we found no significant increases in CD8<sup>+</sup> T cells, NK1.1<sup>+</sup> cells, γδ T cells, macrophages, or neutrophils in IL-22–injected ears (data not shown).

We presumed that the lack of IL-22 in the ears of Ccr6<sup>−/−</sup> mice was due to a failure of IL-22–producing cells to traffic to the site. However, to rule out an essential defect in the ability of cells from Ccr6<sup>−/−</sup> mice to make IL-22, we performed experiments using splenocytes and CD4<sup>+</sup> T cells ex vivo. Naive CD4<sup>+</sup> T cells purified from WT and Ccr6<sup>−/−</sup> mice were cultured for 7 days under either nonpolarizing or Th17-polarizing conditions, and Il17a and Il22 mRNA levels were measured after restimulation. Il17a and Il22 mRNAs were highly induced in cells cultured under Th17-polarizing conditions, and these levels did not differ between cells from WT and Ccr6<sup>−/−</sup> mice (Figure 4A). Similarly, IL-23 induced Il22 mRNA to equal levels in unfractionated splenocytes from WT and Ccr6<sup>−/−</sup> mice (Figure 4B). Most likely, then, the low levels of IL-22 in the IL-23–injected Ccr6<sup>−/−</sup> ears were not caused by an inherent inability of Ccr6<sup>−/−</sup> cells to produce IL-22, but rather by differences between Ccr6<sup>−/−</sup> and WT mice in an IL-22–producing population of cells resident in the ear skin are γδ T cells, although these cells are not able to produce IL-22 (65). As noted above, we saw no difference in the number of γδ T cells after IL-23 injection or between Ccr6<sup>−/−</sup> and WT mice. We did detect CD3<sup>+</sup>CD4<sup>+</sup>IL-17<sup>+</sup> cells from the ears after activation ex vivo, but again, the numbers did not differ between PBS- and IL-23–injected or between Ccr6<sup>−/−</sup> and WT mice (data not shown).

We detected small numbers of IL-22–producing CD4<sup>+</sup> T cells after activation ex vivo, but only in IL-23–injected ears (Figure 5C). IL-22–producing cells were highly enriched within the IL-17A–producing subset. Cells were also identified within the IL-22<sup>+</sup>IL-17A<sup>−</sup> quadrant, although their tendency to sit just to the left of the IL-17A gate suggests they might be IL-17A<sup>+</sup>. The number of cells capable of producing IL-22 was similar in WT and Ccr6<sup>−/−</sup> ears (Figure 5D). The findings from intracellular staining for IL-22 were supported by measuring the IL-22 produced by ear T cells after stimulating with anti-CD3 ex vivo. We detected secretion of IL-22 only by cells from IL-23–injected ears, and similar amounts of IL-22 were produced by cells from WT and Ccr6<sup>−/−</sup> mice (Figure 5E). In addition to analyzing ears for Il22, Il17a, and Il17f, we also performed RT-PCR for Tnfα and Ifng mRNA. Ifng mRNA was not detectable, and Tnfα mRNA
IL-23–injected ears of WT and Ccr6−/− mice have similar *IL17a* and *IL17f* mRNA, Th17 cells, and IL-22–producing cells at day 15. Ears were injected as in Figure 1 and harvested on day 15. (A) Cells from IL-23–injected WT ears were activated with PMA and ionomycin, stained, and analyzed. Shown are CD3+CD4+ cells only, from 1 mouse in 1 representative experiment of 6. (B) mRNAs were measured by real-time RT-PCR. Data are Gapdh-normalized fold changes versus the PBS-injected sample with the greatest ΔCt, from 5 experiments in 12 (IL-23) and 4 (PBS) mice/group. (C) Cells were activated with PMA and ionomycin, stained, and analyzed. Shown are CD3+CD4+ cells only, from a sample pooled from 3 ears in 1 representative experiment of 3. (D) Total IL-17A– or IL-22–expressing CD3+CD4+ T cells analyzed as in C. Data are from at least 14 mice/group in 8 experiments (IL-17A) or from 3 pooled samples of 3 ears/group in 1 representative experiment of 3 (IL-22). (E) Cells were cultured in medium alone or with 10 μg/ml soluble anti-CD3 for 12 hours, and IL-22 in the culture media was measured. Data are from duplicate wells in 1 representative experiment of 2. *P < 0.05 versus respective PBS control; #P < 0.05 versus WT PBS. (A and C) Percent cells in the respective quadrant is indicated. (B, D, and E) Values for IL-23–injected WT and Ccr6−/− mice (E, anti-CD3–stimulated only) were not significantly different.
was not increased, after IL-23 injections; moreover, expression did not differ between Ccr6−/− and WT mice (data not shown). Together, these data demonstrate that although Ccr6−/− ears had more Th17 cells at baseline, there were no significant differences between day-15, IL-23–injected Ccr6−/− and WT ears with respect to Il17a and Il17f mRNA levels or the number of T cells capable of producing IL-17A and IL-22.

Rag1−/− mice initially show WT responses to IL-23 injection, including IL-22 induction. The discordance between Il22 mRNA level, on the one hand, and Th17/IL-22–producing cell numbers and levels of Il17a and Il17f mRNA, on the other, suggested that CD4+ T cells might not be the principal source of IL-22 in the IL-23–injected ears. We investigated this possibility by comparing the responses of Rag1−/− and WT mice. As shown in Figure 6A, the IL-23–injected Rag1−/− and WT ears swelled similarly until day 5, after which ear thickness in the Rag1−/− mice decreased toward the baseline values of PBS-injected control and Ccr6−/− ears. At day 5, tissue sections from IL-23–injected ears of both WT and Rag1−/− mice showed focal hyperparakeratosis and dilated dermal capillaries with mild acanthosis and dermal inflammation (Figure 6B). Particularly striking were the multiple intracorneal pustules, which were much reduced in day-15 ears and rarely seen in Ccr6−/− ears (Figure 6B and Supplemental Figure 2). By day 15, acanthosis and dermal inflammation increased in IL-23–injected WT ears, with less pronounced changes observed in Rag1−/− ears (Figure 6B).

Consistent with the importance of IL-22 in the skin pathology, levels of IL-22 in the ears correlated with ear thickness and histology. At day 5, Il22 mRNA was elevated in IL-23–injected Rag1−/− ears, close to the level seen in WT ears and significantly higher than that in Ccr6−/− ears, while by day 15, Il22 mRNA levels in Rag1−/− ears had fallen substantially (Figure 7A). These data demonstrate that T cells are not required initially for either psoriasiform inflammation or local production of IL-22 induced by IL-23 injections. In the absence of T cells, however, both the inflammatory changes and the elevated levels of IL-22 subsequently diminished, despite continued injections of IL-23.

For Il17a and Il17f mRNA, Rag1−/− ears showed low levels at days 5 and 15 (Figure 7, B and C). Similar to our results shown above, Il17a and Il17f mRNA levels were comparable in IL-23–injected WT and Ccr6−/− ears at day 15. At day 5, however, IL-23–injected Ccr6−/− ears contained amounts of Il17a and Il17f mRNA that were similar to the level in Rag1−/− ears and much less than that in WT ears. We conclude that T cells are the principal source of IL-17A and IL-17F in the IL-23–injected ears, and that despite Th17 cell numbers similar to or higher than WT levels, the Ccr6−/− mice show a delay in the induction of IL-17A and IL-17F — along with a consistent defect in induction of IL-22 — in response to the IL-23 injections. Unlike Il22 mRNA levels, Il17a and Il17f mRNA levels did not correlate with psoriasiform pathology, since they could be low in ears with substan-
†tial pathology (e.g., Rag1−/− mice at day 5) and high in ears with few changes (e.g., Ccr6−/− mice at day 15).

Discussion

Our principal finding is that CCR6 was required for the changes induced in mouse skin by intradermal injection of IL-23. The importance of our findings depends on the relevance of the model for understanding at least some components of human disease. An increasing body of data from both mouse (9–11) and human studies (66) suggests a role for IL-23 in autoimmune disorders. In psoriasis, the most compelling evidence includes genetic data that link both IL-23 and its receptor to disease risk (29–32) and clinical trials suggesting that neutralizing the p40 chain of IL-23 and IL-12 is an effective treatment (33–35). Th17 cells constitute the effector population driven by IL-23, and evidence from human studies suggests that IL-22 is a key Th17 cytokine in psoriasis (40, 42, 43). Moreover, experiments in mice have established a direct connection between IL-23 and the production of IL-22 by both T cells and non–T cells (38, 67–69). Together, the data provide strong support for using the IL-23 injection model in mice, in spite of some clear limitations (37, 70).

Figure 7

Il22 mRNA correlates with IL-23–induced inflammation, but not with Il17a and Il17f mRNA, and is produced in T cell–independent and T cell–dependent phases. (A–C) Ears were injected as in Figure 1, and mRNAs for Il22 (A), Il17a (B), Il17f (C), and Gapdh were measured at days 5 and 15 by real-time RT-PCR. Data are Gapdh-normalized fold changes versus the PBS-injected sample with the greatest ΔCt, from 3 mice/group in 1 representative experiment of 2. Note that values cannot be compared between plots. *P < 0.05 vs. IL-23 Ccr6−/−; **P < 0.05 vs. IL-23 Rag1−/−. Other pairwise comparisons between the IL-23–injected groups did not yield significant differences.
A possible link between CCR6 and psoriatic inflammation was suggested by data on expression of CCR6 and CCL20 in psoriatic skin (53, 56, 71) and by the observation that IL-17A- and IL-22–producing cells cloned and expanded from psoriatic skin expressed CCR6 (13). The latter finding is consistent with the importance of CCR6-deficient mice. While deficiency in CCR6 prevented almost all of the histological changes induced by IL-23 in WT skin, the absence of IL-22 was not sufficient to block the effects of IL-23 completely (38), nor were injections of IL-22 alone able to recapitulate fully the magnitude of the IL-23 effects (present study).

The lack of IL22 expression in IL-23–injected Ccr6–/– ears was not due to any inherent defect in the ability of CCR6-deficient cells to differentiate appropriately, and we presumed that any role for CCR6 in this model would be through effects on Th17 cell migration at the inflammatory site. However, evaluation of the day-15 ears for Th17/IL-22–producing T cells, which were analyzed after activation ex vivo, also provided no explanation for the low levels of Il22 mRNA in Ccr6–/– mice. Although Th17 cells in IL-23–injected WT ears expressed CCR6, we found no differences in the IL-23–injected Ccr6–/– versus WT ears with regard to Il17a or Il17f mRNA level, number of cells that could be activated to produce IL-17A or IL-22, or IL-22 secretion after stimulation with anti-CD3 ex vivo.

Our failure to find deficits in the Th17/IL-22–producing T cells in the Ccr6–/– mice led us to examine the role of T cells in the model by using Rag1–/– mice. At day 5, IL-23–injected Rag1–/– and WT ears showed equivalent degrees of swelling, with focal hyperparakeratosis, dilated dermal capillaries, mild acanthosis and dermal inflammation, and intracorneal pustules. After day 5, while changes in the WT mice progressed, those in the Rag1–/– mice began to resolve. Relative Il22 mRNA levels in the ears correlated well with pathology: in WT mice, persistently high; in Ccr6–/– mice, persistently low; in Rag1–/– mice, high at day 5 and diminished at day 15. The Rag1–/– data provided a way to reconcile, at least in part, our findings that the Ccr6–/– mice showed little induction of Il22 despite WT-level numbers of Th17/IL-22–producing T cells. A critical component of the IL-23–induced pathology was an early, CCR6-dependent increase in IL-22 from a non–T cell source.

The experiments with Rag1–/– mice demonstrated not only the early, T cell–dependent component to the IL-23–induced changes, but also a later, T cell–dependent component. An associated defect in T (Th17) cell responses within the Ccr6–/– ears was in fact revealed in the cytokine profiles. Rag1–/– mice showed consistently low levels of Il17a and Il17f mRNA compared with WT mice, in line with T cells being an important source of these cytokines (73) and suggesting that levels of Il17a and Il17f mRNA in the ears served as a read-out of Th17 cell activity. Although by day 15, levels of these mRNAs were not markedly lower in the IL-23–injected ears of Ccr6–/– versus WT mice, at day 5, Il17a and Il17f mRNA induction in Ccr6–/– mice was much reduced compared with WT mice. These findings suggest that although we were unable to detect deficiencies in Th17/IL-22–producing T cells from IL-23–injected Ccr6–/– ears when analyzed ex vivo, Th17 cells in the Ccr6–/– and WT ears were not responding to IL-23 injections equivalently in situ. Taken together, our data suggest that direct effects of IL-23 on Th17/IL-22–producing T cells in the skin are not sufficient to produce psoriasis-like pathology—otherwise, such changes would have been seen in the ears of Ccr6–/– mice, which have normal (or higher) numbers of Th17 cells. Rather, IL-23 initiated a T cell–independent inflammatory process, which resulted in the production of IL-22 and early changes in the skin and was required for optimal activation of Th17 cells (and perhaps other T cells) and the sustained, T cell–dependent inflammation observed in the WT mice. A parsimonious explanation for our findings is that, by preventing the early, T cell–dependent events, the absence of CCR6 also blocked subsequent T cell activation and the full development of skin pathology, even though the relevant T cell subsets were present.

Although Th17 cells were identified in control and IL-23–injected ears, in both WT and Ccr6–/– mice, we only found T cells capable of making IL-22 in IL-23–injected ears. Together with the fact that the IL-23–injected Ccr6–/– ears showed little change in the numbers of T/Th17 cells, this observation raises the possibility that one effect of IL-23 was to make Th17 cells within the ear IL-22 capable. Regardless of whether these T cells were a significant source of IL-22 in WT ears, injected IL-23 was clearly not sufficient to induce enough IL-22 from T cell or non–T cell sources to produce pathology in the Ccr6–/– mice.

Although Il17a and Il17f mRNA levels were much lower in the IL-23–injected ears of Ccr6–/– compared with WT mice at day 5, we have no evidence that the failure to produce IL-17A and/or IL-17F early on was important in preventing inflammatory changes. IL-17A and/or IL-17F might have been presumed to have a role particularly in forming the early intracorneal pustules, given their roles in recruiting neutrophils and the ability of IL-17A to induce neutrophil-attracting chemokines in human keratinocytes (74). However,
we found no simple correlation between the pustules and these cytokines, as Il17a and Il17f mRNA levels were not markedly different between the Rag1–/– (pustules prominent) and Ccr6–/– (pustules uncommon) mice at day 5. In fact, we generally found that Il17a and Il17f mRNA levels did not correlate well with pathology, consistent with earlier work showing that blocking IL-17A had no effect on epidermal hyperplasia induced by 4 daily injections of IL-23 and that intradermal injections of IL-17A failed to cause the epidermal hyperplasia seen with IL-23 (37). An auxiliary role for IL-17A and IL-17F in the IL-23–induced changes cannot be excluded.

CCR6 is expressed not only on Th17 cells, but also on a subset of mouse and human CD25+ Tregs that produce IL-10 (75). Since the Ccr6–/– mice showed an absence of, rather than an exacerbation of, inflammation after IL-23 injection, a defect in Treg recruitment could not explain our findings. However, we considered the possibility that CCL20 might act through CCR6 as an inhibitor of Treg function. In fact, CCL20 showed no such activity when analyzed in CCR6–/– mice compared with WT mice. Higher numbers of Th17 cells in Ccr6–/– mice were also recently reported by others, who speculated that this increase was the result of a failure of CCL20-mediated colocalization of CCR6–Tregs and differentiating CCR6+ effectors, leading to diminished suppression of Th17 differentiation (65).

Important unresolved and interrelated issues are the identities of the cells mediating early inflammation and IL-22 production, the precise ways in which their roles in the response depend on CCR6, and the relationships between the early and late phases of the response. Principal candidates for non–T cells mediating the early response are subsets of dendritic cells and/or NK cells. Dendritic cells are proposed as important contributors to psoriasis (2, 76) and have been found to produce IL-22 in response to IL-23 (69). Moreover, there is a demonstrated role for CCR6 in dendritic cell recruitment to the inflamed skin (77, 78). It is noteworthy that the reports of CCR6-dependent trafficking of skin dendritic cells (77, 78) did not demonstrate CCR6 expression on these cells. Although we found increased numbers of dendritic cells in the IL-23–injected ears of WT mice, but not Ccr6–/– mice, we were also unable to detect CCR6 on the CD11c+CD11b+ cells from the ears (data not shown); thus, the role of CCR6 in recruitment of dendritic cells to the IL-23–injected ears may have been indirect. Whatever the role for CCR6 on dendritic cells under inflammatory conditions, we found no difference in the number of CD11c+CD11b+ cells in nonmanipulated ears in Ccr6–/– and WT mice, nor did we find differences between these mice in the number of Langerhans cells in epidermal sheets (data not shown), consistent with published data (58, 78). Very recent reports have described cells with NK cell markers that are an innate source of IL-22, so-called “NK-22” cells, which express — and whose development depends on — RORγt (79–82). In humans, these cells express CCR6 and have been found in inflamed skin (79). In mice, at least under homeostatic conditions, the equivalent subset appears confined to the gut (80), and we did not detect these cells in the noninjected ears (data not shown). It will be of interest to investigate whether these NK-22 cells are found in extraintestinal sites in mice during inflammatory responses.

Our present findings showed an expanded role for CCR6 in IL-23–related responses, one that was not limited to effects on Th17 cell trafficking. Such a broad role makes CCR6 an appealing target for therapies against the lengthening list of diseases mediated by Th17 cells. The recent approval of a chemokine receptor antagonist for clinical use (83) provides an encouraging precedent and suggests that developing drugs against additional chemokine receptors is a realistic goal. The accumulated mouse and human data suggest that blocking a chemokine receptor such as CCR6 is likely to cause less overall immunosuppression than therapies that inhibit pleiotropic immune modulators.

Methods

Mice. C57BL/6 WT mice were purchased from the Division of Cancer Therapy, NCI, NIH; from Taconic; or from The Jackson Laboratory. Rag1–/– mice were purchased from The Jackson Laboratory. C57BL/6 Ccr6–/– mice were derived as described below. All mice were housed in sterile cages in pathogen-free conditions and used at 8–20 weeks of age. Animal protocols were approved by the Animal Care and Use Committee of the NIAID, NIH.

Targeting of the mouse Ccr6 gene and creation of mouse lines. The mouse Ccr6 gene was isolated by using a human CCR6 cDNA fragment to probe a 129/SvJ mouse genomic DNA library in Lambda Fix II (Stratagene), generated and provided by S.-J. Lee (Johns Hopkins University School of Medicine, Baltimore, Maryland, USA). A BamHI fragment of the mouse Ccr6 gene corresponding to positions 73,2125–73,8664 in GenBank record NT_039638—which includes the entire Ccr6 ORF—was sequenced and used for producing fragments for gene targeting. Details of Ccr6–/– mouse production are shown in Supplemental Figure 1. The targeting DNA was constructed on the pPNT backbone (84) with the Ccr6 fragments flanking an insertion cassette containing EGFP (taken from pEGFP; Clontech) downstream of an internal ribosome entry site [taken from pCITE-2a(+)]; Novagen) and followed by the small t antigen intron and early region polyadenylation sequence of SV40 (taken from pMSXND; provided by S.-J. Lee; ref. 85) and a neomycin resistance gene flanked by loxP sites (taken from pL2-neo; provided by H. Gu, NIAID, NIH). The pPNT-muCCR6 targeting vector was linearized using the unique NotI site and transfected into R1 ES cells, followed by selection with G418 and ganciclovir. DNAs from 220 G418- and ganciclovir-resistant colonies were digested with BamHI and screened by Southern blot analysis using a 515-bp fragment from sequences upstream of the s’ recombination fragment. Of these 220 ES cell colonies, 8 yielded a BamHI fragment of approximately 3 kb, the predicted size for the targeted allele. Following additional analysis by Southern blotting to rule out multiple insertions and assure proper structure at the targeted site, ES cells were chosen for injection into C57BL/6 blastocysts. The neomycin resistance gene and its associated promoter were removed by crossing chimeras with mice containing the ElaCre transgene in the C57BL/6 background, and offspring were screened for loss of both neo and ElaCre. Mice were backcrossed to C57BL/6 at least 8 times and intercrossed to yield breeders that were used to produce mice for experiments.

Analysis of CCR6 expression and function in Ccr6–/– mice. Resting B cells were purified from mouse splenocytes by negative magnetic selection using CD43 MicroBeads per the manufacturer’s instructions (Miltenyi Biotec). B cells were either used fresh or stimulated for 48 hours with 10 μg/ml goat anti-mouse IgM F(ab’)2; from ICN Biomedicals. Analysis of CCR6 surface expression was analyzed by flow cytometry, as described below. For assaying chemotaxis, B cells were loaded into transwell inserts with 5-μm pores (Costar) sitting in wells containing chemotaxis buffer, which consisted of RPMI 1640 (Invitrogen) supplemented with 20 μM HEPES (Invitrogen) and 0.5% BSA (Invitrogen), for 30 minutes at 37°C in 5% CO2. Inserts were moved to wells containing chemotaxis media alone or supplemented with 1 μg/ml mouse CCL20 (PeproTech) or mouse CXCL12 (PeproTech) and...
incubated for 3 hours. Migrated cells were counted using a flow cytometer after adding a known number of fluorescent counting beads (see below).

**Intradermal cytokine injections.** We performed intradermal injection of 20 μl PBS, either alone or containing 500 ng recombinant mouse IL-23 or IL-22 (eBioscience), into the ears of anesthetized mice using a 30-gauge needle every other day for 16 days. Ear thickness was measured before injection on day 0 and thereafter on days without injections. Ear measurements were made at the center of the ears using a G-1A dial thickness gauge (Pecoak). All measurements were performed blinded. At various times, mice were euthanized, and tissue was collected.

**Processing of ear skin, lymph nodes, and splenocytes for flow cytometry and ex vivo stimulation.** Injected ears were collected, and cell suspensions from individual ears were prepared as previously described (86). Briefly, after collection of ears, ventral and dorsal dermal sheets were separated from cartilage and incubated for 45 minutes at 37°C with dermal side down in RPMI 1640 containing 0.28 Wunsch units/ml of Liberase CI blendzyme (Roche Diagnostics). Next, the ear sheets of individual animals were mechanically separated for 5 minutes using a Medimachine tissue disruptor according to the manufacturer's instructions (BD Biosciences — Pharmingen), with complete RPMI supplemented with 0.05% DNase (Sigma-Aldrich) as a buffer. Complete RPMI contained RPMI 1640 supplemented with 10% fetal bovine serum (Invitrogen), then washed in PBS and mounted using ProLong Gold anti-fade reagent (Invitrogen), then washed in PBS and mounted using ProLong Gold anti-fade reagent (Invitrogen). Lymph nodes or spleens were mechanically disrupted using a syringe plunger in complete RPMI. Cells were filtered through 100-μm nylon mesh and washed before activation and/or staining.

**Histopathological analysis and immunofluorescence microscopy.** Sections from paraffin-embedded whole ears (7 μm) were prepared by American Histo-labs or HistoServe Inc. and stained with H&E. Staining for CCL20 was done using frozen 7-μm sections of whole mouse ears; sections were air-dried after 5 minutes of fixation in ice-cold acetone, blocked for 1 hour at room temperature with 5% goat serum in PBS containing 3% skim milk, and incubated with rat anti-mouse CCL20 (clone 114906; R&D Systems) or rat IgG isotype control antibody (eBioscience) for 1 hour at room temperature. After washing in PBS, the sections were incubated for 30 minutes at room temperature with an Alexa Fluor 488-conjugated goat-anti-rat IgG (Invitrogen), then washed in PBS and mounted using ProLong Gold anti-fade reagent with DAPI nuclear stain (Invitrogen). Images were acquired at ×400 magnification using a Carl Zeiss microscope and AxioVision software (release 4.5; Zeiss). H&E images were captured using a DP71 digital camera (Olympus) attached to an Olympus BX41 microscope.

**RT-PCR, quantification, and display.** Extraction of RNA from whole ears was performed using a Qiagen RNeasy Kit according to the manufacturer's instructions. RNA extraction from cultured cells was performed using TRIzol reagent according to the manufacturer's instructions (Invitrogen). We used Platinum Quantitative RT-PCR Thermoscript One-Step System (Invitrogen) to perform one-step RT-PCR. The following primers and probes were purchased from Applied Biosystems: GAPDH (catalog no. Mm99999915_g1), IL-17A (catalog no. Mm00439619_m1), IL-17F (catalog no. Mm00521423_m1), CCL20 (catalog no. Mm01268754_m1), and IL-22, which was a custom construct using previously described sequences (38). The reactions were run on an Applied Biosystems 7900HT system using the standard protocol provided by Invitrogen. Data were represented as fold differences relative to control/reference samples. Fold difference was calculated by the 2^−ΔΔCt method, where ΔCt = Ct_target_cDNA − Ct_ref_CDNA and ΔΔCt = ΔCt_treated − ΔCt_reference. ΔCt_reference was chosen as the PBS-injected or other reference sample with the greatest ΔCt for each plot.

**Flow cytometry and intracellular staining.** Monoclonal fluorescent antibodies against mouse CD45, CD3, CD4, CD8, γδ TCR, NK1.1, CD11b, B220, CD11c, CD25, CD62L, and CD44 as well as streptavidin-PE were purchased from BD Biosciences. Antibodies against IL-17A, Ly-6G, and F4/80 were purchased from eBioscience. Antibodies against TCRβ, CCR6, and IL-22 were purchased from Biolegend, R&D Systems, and Antigenex, respectively. Propidium iodide (PI) was purchased from Invitrogen. Flow cytometry was performed on single-cell suspensions of ear skin (see above) or lymph node cells. For immunotyping of cells, dead cells were eliminated on the basis of positive PI staining. CD4+ T cells were identified by their scatter profile and as PI CD4+ CD3+CD4+. For some samples, CCR6 expression was also evaluated on CD4+ T cells. Dendritic cells were identified by their scatter profile and as PI CD45+CD3+CD11c+CD11b+. Intracellular staining for IL-17A was done after activating cells for 6 hours with Leukocyte Activation Cocktail with GolgiPlug (BD Biosciences — Pharmingen). Cells were then fixed and permeabilized using the Cytofix/Cytoperm Plus Kit according to the manufacturer's instructions (BD Biosciences). For intracellular staining, the same surface markers were used to identify CD4+ T cells, and PI was omitted. Absolute numbers of cells were calculated by adding a known number of fluorescent counting beads (Spherotech) to each of the samples. The ratio of beads counted on the flow cytometer to the number of beads added to the sample was determined and used to calculate the absolute numbers in the samples for cells of a given immunophenotype. All samples were analyzed on an LSR II System flow cytometer (BD Biosciences), and the data were analyzed using FlowJo software (version 8.2; Tree Star).

**Sorting naïve cells and differentiation of Th17 cells in vitro.** For the sorting of naïve CD4+ T cells, splenocytes and lymph node cells were harvested from nonmanipulated mice, and CD4+ T cells were first enriched by using the CD4+ negative selection kit from Miltenyi Biotec and their autoMACS machine, following the manufacturer’s instructions. Naïve cells were purified by cell sorting based on the criterion CD4+CD25 CD62L+CD45+. The sorted cells were stimulated with 10 μg/ml plate-bound hamster anti-mouse CD3e (clone 145-2C11; R&D Systems) and 1 μg/ml soluble hamster anti-mouse CD28 (BD Biosciences) for 3 days under nonpolarizing or Th17-polarizing conditions in complete RPMI. For nonpolarizing conditions, complete RPMI was supplemented with 10 μg/ml rat anti-mouse IFN-γ (BD Biosciences), 10 μg/ml rat anti-mouse IL-4 (BD Biosciences), 10 ng/ml mouse IL-2 (R&D Systems), and 10 ng/ml mouse TGF-β1 (Sigma-Aldrich). For Th17 conditions, complete RPMI was supplemented with 10 μg/ml rat anti-mouse IFN-γ, 10 μg/ml rat anti-mouse IL-4, 10 ng/ml mouse IL-23 (eBiosciences), 20 ng/ml mouse IL-6 (R&D Systems), 10 ng/ml mouse TGF-β1, 5 ng/ml mouse IL-1β (R&D Systems), and 5 ng/ml mouse TNF-α (R&D Systems). After 3 days of stimulation, the cells were washed and resuspended in fresh media supplemented as above per the respective polarizing condition, but without anti-CD3 or anti-CD28. After a total of 7 days in culture, the cells were harvested and stimulated with 20 ng/ml PMA and 1 μM ionomycin for 6 hours before preparing RNA.

**Measuring IL-22 produced by cells stimulated ex vivo with IL-23 or anti-CD3.** For unfractionated splenocytes, cells were cultured at 2 × 10^6 cells/ml in complete RPMI supplemented with 100 ng/ml mouse IL-23 (eBioscience) for 12 hours at 37°C. For stimulating cells from the ears, single-cell suspensions were pooled from multiple ears and then divided so that cells equivalent to a single ear were cultured per well in a 24-well plate. Whole-ear suspensions were stimulated with 10 μg/ml soluble hamster anti-mouse CD3e for 12 hours at 37°C. Supernatants were harvested and analyzed using a mouse IL-22 ELISA Construction Kit (Antigenex) according to the manufacturer’s instructions. We used 3,3′,5,5′-tetramethylbenzidine as a chromogen, detected with a microplate reader set at 450 nm.

**Statistics.** All quantitative data are shown as mean ± SEM unless otherwise indicated. All samples were compared using 2-tailed, unpaired Student’s t test. A P value less than 0.05 was considered significant.
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