Rorγt+ innate lymphocytes and γδ T cells initiate psoriasiform plaque formation in mice

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Psoriasis is a common, relapsing inflammatory skin disease characterized by erythematous scaly plaques. Histological manifestations of psoriasis include keratinocyte dysregulation and hyperproliferation, elongated rete ridges, and inflammatory infiltrates consisting of T cells, macrophages, dendritic cells, and neutrophils. Despite the availability of new effective drugs to treat psoriasis, the underlying mechanisms of pathogenesis are still poorly understood. Recent studies have shown that Aldara cream, used to treat benign skin abnormalities, triggers psoriasis-like disease in humans and mice and have implicated Th17 cells in disease initiation. Using this as a model, we found a predominant role for the Th17 signature cytokines IL-17A, IL-17F, and IL-22 in psoriasiform plaque formation in mice. Using gene-targeted mice, we observed that loss of Il17a, Il17f, or Il22 strongly reduced disease severity of psoriasis. However, we found that Th17 cells were not the primary source of these pathogenic cytokines. Rather, IL-17A, IL-17F, and IL-22 were produced by a skin-invading population of γδ T cells and RORγt+ innate lymphocytes. Furthermore, our findings establish that RORγt+ innate lymphocytes and γδ T cells are necessary and sufficient for psoriatic plaque formation in an experimental disease model that closely resembles human psoriatic plaque formation.

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Introduction
Psoriasis is a complex, multifactorial disease, manifesting in an inflammatory response in the skin. The exact mechanistic underpinnings of psoriatic plaque formation are yet to be defined (1). It has been reported that psoriasis occurrences and relapses were inadvertently induced in patients that received topical treatment with Aldara, a cream containing imiquimod (IMQ) (2). In mice, the application of Aldara induces pathology that largely resembles the human disease phenotype of psoriasis (3). Moreover, it was shown to be similarly dependent on IL-23, and IL-17A, signaling, suggesting a role of Th17 effector cells (3–5). Recent studies have suggested a role for IL-17–producing γδ T cells in human psoriasis (6, 7). Here we report that initiation of plaque formation in the Aldara psoriasis model is dependent on RORγt+, skin-infiltrating γδ T cells, and innate lymphoid cells (ILCs). Our observations favor the notion that rather than Th cells, Vγ4+ γδ T cells and ILCs are the dominant and critical source of IL-17A, IL-17F, and IL-22 in the formation of acute psoriasiform lesions.

Results and Discussion
IL-17A, IL-17F, and IL-22 are critical for psoriatic plaque formation. After daily topical application of Aldara, we observed significant thickening, reddening, and scaling of the skin after 3 days (Supplemental Figure 1A; supplemental material available online with this article; doi:10.1172/JCI61862DS1). The clinical course of plaque formation and histopathology for the ear and the back skin were similar, with a slightly different kinetics (Supplemental Figure 1B). Aldara treatment resulted in hyperproliferation of keratinocytes and disturbed epidermal differentiation, as indicated by acanthosis and hyperparakeratosis. An additional feature, reminiscent of Munro’s microabscesses in human psoriasis, was the accumulation of terminal neutrophils (myeloperoxidase [MPO] staining) in the stratum corneum. Additionally, in the dermis extensive infiltration of leukocytes was observed (Supplemental Figure 1C).

Responsiveness to conventional antipsoriatic therapies is a hallmark for a valid psoriasis model (8). Ustekinumab (IL-12/23p40 monoclonal antibody) is one of the most efficacious treatments for psoriasis (4). Mice that were injected with a single dose of neutralizing murine IL-12/23p40 antibody displayed significantly less skin inflammation compared with control mice (Figure 1, A and B). Together with the recent finding that Aldara-induced psoriasiform plaques are responsive to UVB therapy (9), this disease model very closely resembles human disease visually and histologically and responds to known antipsoriatic therapies (1).

Skin-invading lymphocytes were analyzed for cytokine production at peak disease. IL-17F+ lymphocytes were substantially more abundant than IL-17A– and IL-22–producing cells in the skin of Aldara-treated mice (Figure 1C). We tested individual contributions of the implicated cytokines using Il17a–/–, Il17f–/–, and Il22–/– mice and found a significant reduction in the percentage change in skin thickness in all of the mutant mouse strains, compared with wild-type mice (Figure 1D). Histological analysis of the inflammation revealed a significant decrease in acanthosis in Il17a–/–, Il17f–/–, Il22–/– mice compared with wild-type mice, with the most pronounced effects in mice lacking IL-17F and IL-22 (Figure 1E). The fact that there were more skin-invading cells secreting IL-17F than IL-17A is in line with the stronger disease resistance in Il17f–/– compared with Il17a–/– mice. IL-17AF heterodimers have been described as sharing the biological properties with IL-17A and IL-17F (10), and we found significantly increased levels of IL-17AF heterodimers in the
draining LNs, as well as slightly elevated levels in the inflamed skin of wild-type mice (Figure 1F).

An IL-17A–, IL-17F–, and IL-22–dependent mechanism of inflammation is in line with the TLR7 agonistic activity of IMQ (11), as skin inflammation could not be induced in Tlr7–/– mice. The induction of keratinocyte hyperproliferation in Tlr7–/– mice, however, revealed TLR7-independent activities of Aldara (Supplemental Figure 1D). Apart from additional active ingredients in the Aldara cream formulation (M. van den Broek, unpublished observations), TLR7-independent pathways of keratinocyte activation have been previously reported by others (12).

RORγt+γδ T cells and innate lymphocytes are the main producers of IL-17A, IL-17F, and IL-22 in psoriasiform plaques. The transcription factor Rorγt is required for IL-17 and IL-22 production (13, 14). To determine the main producers of these cytokines in the Aldara cream formulation (M. van den Broek, unpublished observations), TLR7-independent pathways of keratinocyte activation have been previously reported by others (12).

**RORγt+γδ T cells and innate lymphocytes are the main producers of IL-17A, IL-17F, and IL-22 in psoriasiform plaques.** The transcription factor Rorγt is required for IL-17 and IL-22 production (13, 14). To determine the main producers of these cytokines, we used Rorc-Cre × EYFP fate-mapping mice. Surprisingly, only a small proportion of the cytokine producers were CD4+ Th cells. In contrast, about 80% of IL-17A as well as nearly 60% of IL-22 were produced by γδ T cells. The remaining IL-17A and IL-22 producers were RORγt+CD3– ILCs, accounting for 10% of IL-17A and approximately 40% of IL-22 (Figure 2, A and B).

There is one report suggesting that human γδ T cells can potentially upregulate TLR7 expression and thus are equipped to respond to IMQ (15), but we failed to observe a direct effect of Aldara or IMQ in purified murine Vγ4 T cells (data not shown). There is, however, emerging evidence that conventional DCs initiate the immune response to IMQ (B. Clausen, unpublished observations).

At steady state, γδ T cells in the murine skin are primarily represented by Vγ4′ resident epidermal cells (nomenclature according to ref. 16), termed dendritic epidermal T cells (DETCs), a cell population not found in human skin. Recently, it was shown that the dermis also contains Vγ4′ resident cells expressing Rorγt, CCR6, and IL-17A (17, 18). Staining for the γδ lineage (γ chain) of the lymphocytes responsible for the observed Aldara skin inflammation revealed that there were two distinct populations (Supplemental Figure 2A). DETCs represented the TCRγδhi cells, and the major
Tcrγδ–/– thymus (19). However, Cai and colleagues, using lineage in the altered due to the developmental changes in the γδ T cell pool in Tcrb–/– γδ T cells is critically lower, but still noticeable inflammation in the skin (Figure 2D). It is also possible that the T cell pool in Tcrd–/– γδ T cells and innate lymphocytes are the main producers of IL-17A, IL-17F, and IL-22 in psoriasiform plaques. (A) Intracellular cytokine staining in the skin of Rorc-Cre × EYFP mice after 5 days of Aldara treatment, gated on CD4+ cells (n = 3), with (B) scatter plots showing percent distribution (n = 3). (C) Cytokine staining of Vγ4+ versus Vγ5+ for IL-17F and IL-22, pre-gated on TCRγδ+ cells. (D and E) Kinetics of Aldara-induced skin inflammation in (D) WT versus Tcrb–/– and Tcrd–/– mice (n = 4) and (E) WT versus Il15ra–/– mice (n = 3), shown as the percent increase in skin thickness. Each experiment was performed independently at least 3 times.*P < 0.05, **P < 0.001.

Thus far, our findings show the reliance of Aldara-induced psoriasis on psoriasiform plaque formation, with Tcrd–/– mice lacking both Vγ4+ and DETCs. To formally rule out the involvement of DETCs in plaque formation, we treated wild-type and Il15ra–/– mice, in which DETCs do not develop (Supplemental Figure 2E), with Aldara and found inflammation to be indistinguishable (Figure 2E).

We found a similar increase in proliferation of Vγ4+ cells in both the skin and skin-draining lymph nodes, suggesting that cytokine-secreting Vγ4+ cells are not predominantly skin resident (Supplemental Figure 2F). A recent study on psoriasis in humans has demonstrated that a subset of γδ T cells seems to play a role in disease development and that these cells express high levels of CCR6 and cutaneous leukocyte antigen (CLA) (7). Correspondingly, we found Vγ4+ cells to express the relevant skin-homing receptors CLA and CCR6, which could facilitate their recruitment to the skin (Supplemental Figure 2G).

Analysis of Aldara-treated skin revealed a small population of Analysis of Aldara-treated skin revealed a small population of Lin NKp46− cells that were Vγ4+ (Supplemental Figure 2B). Upon disease induction, only the frequency (Supplemental Figure 2C) and total number of dermal Vγ4+ cells increased, whereas DETC numbers were unaltered (Supplemental Figure 2D). Intracellular cytokine staining of γδ subpopulations in the skin at peak disease revealed that Vγ4+ cells produced IL-17F and IL-22, whereas DETCs did not (Figure 2C), suggesting that Vγ4+ cells drive the inflammatory skin alteration in this model.

To verify this, we compared the responses to Aldara in Tcrb−/−, Tcrd−/−, and wild-type mice. Tcrb−/− mice developed inflammation similar to that in wild-type mice. In contrast, Tcrd−/− mice had drastically lower, but still noticeable inflammation in the skin (Figure 2D). It is also possible that the γδ T cell pool in Tcrb−/− mice is altered due to the developmental changes in the γδ lineage in the thymus (19). However, Cai and colleagues, using Tcrα−/− mice, also found them to be fully susceptible to IL-23–induced plaque forma-
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The production (13, 14). Consequently, as in Rag2−/−Il2rg−/− mice, no skin inflammation or thickening was observed in Rorc−/− mice (Figure 3E). Moreover, IL-17A, IL-17F, and IL-22 were virtually absent in the skin of these mice compared with wild-type mice (Figure 3F and data not shown). Therefore, Aldara-induced psoriasiform plaque formation seems to be completely dependent on the transcription factor RORγt in Vγ4γδ T cells and RAG-independent ILCs.

We set out to dissect the cellular underpinnings of psoriatic plaque formation and employed a recently described mouse model (3). Aldara-induced inflammatory lesions bear a substantial number of major hallmark histopathological and molecular features of human psoriatic plaques (1). The model is, however, limited to the simulation of psoriatic plaque formation, whereas the underlying natural pathogenic trigger of psoriasis, or any aspect of chronicization of the inflammation, cannot be addressed. In a study parallel to ours, Cai and colleagues reported the pivotal role of dermal γδ T cells and their production of IL-17A in IL-23–induced skin inflammation in mice (6) and found evidence for their involvement in human psoriasis (6, 7). Some supporting evidence has also been reported in studies using the Aldara-induced psoriasis model (6, 20), which our study links to with an in-depth analysis of the pathophysiologic events during psoriatic plaque formation.

In summary, our data show that RORγt-dependent ILCs and γδ T cells are necessary and sufficient to drive psoriasiform plaque formation in mice through the collective delivery of IL-17A, IL-17F, and IL-22 to the skin. The current notion that IL-23 induces Th17 cells stems from the observation that activated T cells are a major part of the skin-infiltrating immune cells and are a known source of these cytokines. Only recently have innate lymphocytes been acknowledged to be highly effective producers of these mediators. The fact that OKT3 antibody depleted human T cells was interpreted to indicate that psoriasis is dependent on CD4+ and CD8+ T cells (21), even though it also depletes γδ T cells (22). Without dismissing adaptive immune processes in the etiology of psoriasis, our study does establish the sufficiency of a dysregulated innate immune compartment for psoriatic plaque formation. Thus, our proposed paradigm of lesion development not only provides a new basis for understanding the therapeutic efficacy of new biological drugs to treat human psoriasis, but may also lead to more in depth research on γδ T cell and ILC involvement in human disease.

Methods

Further information is available in Supplemental Methods.

Mice. C57BL/6 (wild-type) mice were obtained from Janvier. RorcGFPGFP (in text referred to as Rorc−/−), Tcrδ−/−, Rag1−/−, and Rag2−/−Il2rg−/− mice were purchased from The Jackson Laboratory. Terb−/− and Tcrδ−/− mice were bred from Tcrδ−/− mice. Rorc-Cre crossed with Rosa26-stop-eYFP mice (called

Figure 3

RORγt+ innate lymphocytes are essential for psoriasiform plaque formation. (A) Kinetics of Aldara-induced skin inflammation in WT versus Tcrδ−/− versus Rag1−/− mice. (B) Cytokine staining in ILCs in the skin of Aldara- versus control-treated mice, pre-gated on Lin–CD45+ cells (n = 3). (C) Fate map analysis and IL-22 staining in the Aldara-treated skin of Rorc-Cre X EYFP and Rorc-Cre X EYFP x Rag1−/− mice gated on CD45+ cells. (D) WT versus Rag1−/− versus Rag2−/−Il2rg−/− mice and (E) WT versus Rorc−/− mice showing percent change in skin thickness (n = 3). (F) Staining for IL-17F and IL-22 in CD45+ cells from the skin of control and Aldara-treated WT versus Rorc−/− mice (n = 3). Each experiment was performed independently at least 3 times. *P < 0.001.
Rorc-Cre x EYFP mice here) were provided by A. Diefenbach (University of Freiburg, Freiburg, Germany), Il17a−/− mice by Y. Iwakura (University of Tokyo, Tokyo, Japan), Il17f−/− mice by Merck Serono S.A., Il22−/− mice by J.C. Renaudel (Université Catholique de Louvain, Yvoir, Belgium), and Il15ra−/− mice by S. Belfone-Paus (University of Giessen, Giessen, Germany).

**Treatment.** The mouse back was shaved, and a daily dose of 50 mg of Aldara of 5% IMQ cream; 3M Pharmaceuticals) or control vehicle cream (Soft Kreme KA, Kantonsapotheke Zurich) was applied on the back and 5 mg to each ear for 5–7 days. Anti–IL-12/23p40 treatment was performed by injection of the mice with 200 µg/mouse rat anti–mouse IL-12/23p40 mAb (C17.8) and respective IgG2A isotype control antibody (2A3, both Bio X Cell) on day 2 of Aldara application.

**Cell preparation.** Skin and ears were cut into small pieces and digested with 1 mg/ml collagenase type IA and 100 mg/ml DNase (Sigma-Aldrich) for 60 minutes at 37°C. Isolation of leukocytes from the LN involved teasing the organs apart. Both were followed by filtering through 70-µm cell strainers to obtain single-cell suspensions.

**Antibodies.** Cells were incubated with antibodies for 20 minutes at 4°C. For intracellular cytokine staining, cells were stimulated with PMA (Appli-chem) and ionomycin (Invitrogen) and treated with GolgiStop (BD) for 3 hours. After surface staining, cells were permeabilized according to the manufacturer’s (BD) recommendations and stained intracellularly. Stainings were analyzed with a FACS LSRII Fortessa (BD). Post-acquisition analysis was performed with FlowJo (Tree Star) software.

**Histology.** Tissue samples were fixed in 4% paraformaldehyde and embed-ded in paraffin. Deparaffinized sections were stained with H&E. Immunohistochemical staining for MPO (1:100, rabbit, Thermo Scientific) was performed by a Ventana Benchmark XT automated staining system according to the manufacturer’s guidelines.

**Statistics.** For disease severity, differences between groups were evaluated by 2-way ANOVA with Bonferroni’s post hoc test. For analysis of scatter plots of maximum thickness comparing ≥3 groups of mice, 1-way ANOVA with Bonferroni’s post-test was used. Differences between two sets of data were evaluated by 2-tailed Student’s t test. Data represent mean ± SEM. P ≤ 0.05 was considered statistically significant. All statistics were done using GraphPad Prism (GraphPad Software).

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