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# IL-1 signaling is critical for expansion but not generation of autoreactive GM-CSF<sup>+</sup> Th17 cells

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# Abstract

Interleukin-1 (IL-1) is implicated in numerous pathologies, including multiple sclerosis and its animal model experimental autoimmune encephalomyelitis (EAE). However, the exact mechanism by which IL-1 is involved in the generation of pathogenic T cells and in disease development remains largely unknown. We found that following EAE induction, pertussis toxin administration leads to IL-1 receptor type 1 (IL-1R1)-dependent IL-1 $\beta$  expression by myeloid cells in the draining lymph nodes. This myeloid-derived IL-1β did not vitally contribute to the generation and plasticity of Th17 cells, but rather promoted the expansion of a GM-CSF<sup>+</sup> Th17 cell subset, thereby enhancing its encephalitogenic potential. Lack of expansion of GM-CSF-producing Th17 cells led to ameliorated disease in mice deficient for IL-1R1 specifically in T cells. Importantly, pathogenicity of IL-1R1-deficient T cells was fully restored by IL-23 polarization and expansion in vitro. Therefore, our data demonstrate that IL-1 functions as a mitogenic mediator of encephalitogenic Th17 cells rather than qualitative inducer of their generation.

Keywords autoimmunity; EAE; GM-CSF; IL-1; Th17
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# Introduction

Differentiated CD4 T cells play an essential role in defense against pathogens and are culprits in autoimmune disorders. Their implication among other diseases is well documented in multiple sclerosis (MS), an autoimmune dysfunction affecting the human central nervous system (CNS), and in its animal model experimental autoimmune encephalomyelitis (EAE) (Croxford *et al*, 2011). CD4 T-cell involvement in CNS autoimmunity was initially associated with T helper cells producing IFN $\gamma$  (Th1 cells) (Ando *et al*, 1989). However, with the discovery of cells expressing IL-17 (Th17 cells) (Harrington *et al*, 2005; Park *et al*, 2005), this paradigm was challenged. Th17 cells represent a heterogeneous population of cells expressing not only IL-17 but also additional cytokines, depending on the conditions available during their differentiation. Thus, Th17 cells were found to co-express IFN $\gamma$  (Ivanov *et al*, 2006), GM-CSF (Codarri *et al*, 2011; El-Behi *et al*, 2011), IL-21 (Korn *et al*, 2007), IL-22 (Kreymborg *et al*, 2007), and even the suppressive cytokine IL-10 (McGeachy *et al*, 2007; Ghoreschi *et al*, 2010). Moreover, while the impact of IL-17 itself for EAE pathogenesis is still controversial (Haak *et al*, 2009; Ishigame *et al*, 2009), GM-CSF is indispensable for disease development (Codarri *et al*, 2011; El-Behi *et al*, 2011). In line with that, several recent reports indicated GM-CSF as an essential effector molecule also in MS (Hartmann *et al*, 2014; Li *et al*, 2015).

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Th17 cell differentiation from naïve T cells can be initiated by TGFB, the cytokine that is shared with the T regulatory (Treg) cell developmental program. However, Th17 cell commitment occurs only after subsequent inhibition of the Treg-specific transcription factor FoxP3 by IL-6 (Bettelli et al, 2006; Veldhoen et al, 2006) or IL-21 (Korn et al, 2007). Th17 cells generated only in the presence of TGF<sub>β1</sub> and IL-6 cannot transmit EAE (McGeachy et al, 2007) and require additional exposure to IL-23 to gain pathogenic characteristics (Lee et al, 2012). Another pathway of Th17 cell development is independent of TGF $\beta$  and driven by a combination of IL-6, IL-23, and IL-1ß (Chung et al, 2009; Ghoreschi et al, 2010). Cells generated under these conditions express high levels of GM-CSF (El-Behi et al, 2011) and are highly pathogenic in EAE. Of note, the individual ablation of each aforementioned cytokine-mediated signaling results in complete resistance of mice to EAE induction (Eugster et al, 1998; Cua et al, 2003; Matsuki et al, 2006; Sutton et al, 2006). Whereas deletion of IL-6 prevented Th17 cell differentiation (Korn et al, 2007), IL-23 showed a greater impact on expansion and stabilization of the Th17 cell phenotype (Langrish et al, 2005; Veldhoen et al, 2006). The contribution of IL-1 signaling to the pathogenicity of T cells is still controversial with evidences supporting a role in both de novo generation and expansion of Th17 cells (Sutton et al, 2006; Chung et al, 2009; El-Behi et al, 2011). The mechanism by

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which IL-1 mediates pathogenicity *in vivo* could not yet be fully addressed, mainly due to the lack of suitable genetic tools, namely the conditional knockout of the IL-1 receptor. There is only one known signaling receptor—IL-1 receptor type 1 (IL-1R1)—that is, however, broadly expressed by many cell types of immune and non-immune origin (Boraschi & Tagliabue, 2013).

The induction of active EAE is achieved by the immunization with myelin oligodendrocyte glycoprotein (MOG), emulsified in complete Freund's adjuvant (CFA) and injections of pertussis toxin (PTx) (Mendel et al, 1995). PTx is commonly used in many autoimmune models to enhance disease susceptibility; however, its mode of action is not fully understood. In EAE, PTx application is associated with transient Treg cell suppression (Cassan et al, 2006; Chen et al, 2006), promotion of Th17 cell development (Chen et al, 2007), and contribution to blood-brain barrier disruption (Millward et al, 2007). On the other hand, PTx is also able to block leukocyte migration (Su et al, 2001), demonstrating its complex and diverse involvement in disease progression. Despite these proposed effects of PTx, many of them most likely are secondary events with largely unknown key intermediate factor(s). Importantly, PTx also was shown to stimulate expression and inflammasome-mediated maturation of IL-1 $\beta$  in myeloid cells (Dumas *et al*, 2014), suggesting that IL-1ß could be a potential candidate molecule through which PTx would, at least partially, promote EAE development.

In the present study, we aimed to investigate the requirement of IL-1 signaling for T-cell-mediated autoimmunity. By using novel models of conditional IL-1R1 deletion, we demonstrate that PTx treatment stimulates recruitment and expansion of myeloid cells expressing IL-1 $\beta$  in the draining lymph nodes (dLN), in an IL-1R1-dependent manner. This myeloid cell-derived IL-1 stimulated expansion of GM-CSF<sup>+</sup> Th17 cells. The latter cells also developed in the absence of IL-1 signaling but did not proliferate properly and therefore could not cause severe EAE. However, isolation and expansion of T cells devoid of IL-1 signaling in the presence of IL-23 resulted in recovery of pathogenicity. Thus, we could show that IL-1 signaling in CD4 T cells is critical for their expansion, but not for differentiation to encephalitogenic Th17 cells.

### Results

#### PTx administration promotes IL-1β expression by myeloid cells during the induction phase of EAE in an IL-1R1dependent manner

Recently, we generated a new mouse line that allows for the conditional deletion of the IL-1 receptor type 1 (Abdulaal *et al*, 2016), the obligatory receptor for IL-1 $\alpha$ - and IL-1 $\beta$ -mediated signaling. By deleting the receptor in the germ line, using CMV-Cre, we obtained mice that globally lack IL-1R1 (hereafter termed IL-1R1<sup>-/-</sup>). These mice differ from previously reported full knockout mice by the deletion of exon 5, which results in inactivation of all known isoforms for IL-1R1 (Glaccum *et al*, 1997; Labow *et al*, 1997; Qian *et al*, 2012). As a result of complete inactivation of IL-1 signaling, IL-1R1<sup>-/-</sup> mice displayed impaired development of CD11b<sup>+</sup> cells in secondary lymphoid organs under steady state conditions, including myeloid cells producing IL-1 $\beta$ , as compared to wild-type (WT) controls (Fig 1A–C). Experimental autoimmune encephalomyelitis can be induced in mice by immunization with the MOG peptide p35-55 emulsified in CFA, along with PTx injections. To study the role of IL-1 signaling in EAE induction, we immunized mice with MOG/CFA with or without addition of PTx. We found dramatically reduced frequencies and numbers of CD11b<sup>+</sup> myeloid cells in the spleen of IL-1R1-deficient animals compared to wild-type controls (Fig 1D). These experiments suggest that IL-1 signaling might be involved in the recruitment of myeloid cells to the secondary lymphoid organs and their subsequent expansion after immunization.

To investigate whether mobilized myeloid cells not only respond to IL-1 but also enhance the immune response by producing this cytokine, we measured the expression of IL-1β upon immunization. Following stimulation with GM-CSF, LPS, or PMA/ionomycin to trigger ex vivo isolated cells, we found that the vast majority of IL-1 $\beta$  originated from CD11b<sup>+</sup> cells (Fig EV1). Moreover, we noted a robust enhancement of IL-1ß expression by myeloid cells when WT animals were additionally treated with PTx, an effect that was completely absent in IL-1R1-deficient animals (Figs 1E and F, and EV1). Further analysis of the myeloid cell populations revealed that treatment of the mice with PTx resulted in increased frequencies of neutrophils and monocytes/macrophages among the cells expressing IL-1 $\beta$  in the WT group, whereas it had a very limited effect on the same cell populations in  $IL-1R1^{-/-}$  mice (Fig 1G and H). In contrast to IL-1 $\beta$ , the expression of IL-1 $\alpha$  in myeloid cells was not affected by PTx treatment (Fig EV2). However, in line with the IL-1 $\beta$  data, IL-1 $\alpha$ -expressing CD11b<sup>+</sup> cells were dramatically reduced in mice deficient for IL-1R1 (Fig EV2).

Collectively, these data show that PTx activates myeloid cells to produce IL-1 $\beta$  in an IL-1R1-dependent manner after MOG/CFA immunization. In addition, our data suggest that IL-1 $\beta$  production by myeloid cells is dependent on an autocrine stimulatory feedback loop.

#### Mouse and human Th17 cells express high levels of IL-1R1

Development of EAE requires the generation of pathogenic CD4 T cells that initiate a coordinated immune attack of the CNS. To study the role of IL-1 signaling in the development of a T-cell-mediated immune response, we investigated the expression of IL-1R1 by different populations of CD4 T cells following immunization with MOG/CFA/PTx (Appendix Fig S1). We found that Th17 cells expressed the highest levels of IL-1R1 compared to conventional Th1 or GM-CSF-producing T cells in both the dLN and the inflamed CNS of EAE-diseased mice (Fig 2A and C). We also noted that FoxP3<sup>+</sup> Treg cells hardly expressed any IL-1R1 (Fig 2A and C). Importantly, IL-17A-expressing cells regardless of their IFN $\gamma$  and/or GM-CSF co-expression equally displayed high levels of IL-1R1 (Fig 2B and D).

Also in humans, CD4 T-cell development was shown to be dependent on IL-1 signaling (Acosta-Rodriguez *et al*, 2007; Zielinski *et al*, 2012). Although mouse models represent useful tools in investigating human pathologies, possible differences in cell biology can result in misleading translation of the findings to the clinic. In line with our mouse data, we found that a large proportion of human Th17 cells also expressed IL-1R1, but very few Th1- and GM-CSFpositive cells do so (Fig 2E). When we further analyzed the different subpopulations of IL-17A-producing human CD4 T cells, we found



Figure 1. Pertussis toxin induces IL-1β-expressing myeloid cells after MOG/CFA immunization dependent on IL-1R1 signaling.

A Frequencies and total numbers (mean + SEM) of CD11b<sup>+</sup> myeloid cells isolated from the spleen and peripheral LN of naïve non-immunized mice.

- B Total numbers (mean + SEM) of CD11b<sup>+</sup> cells expressing IL-1 $\beta$  shown in (A) and restimulated with indicated stimuli for 4 h.
- C Quantification (mean + SEM) of myeloid cell populations expressing IL-1 $\beta$  isolated from LN and restimulated with GM-CSF shown in (B).
- D Frequencies and total numbers (mean + SEM) of CD11b<sup>+</sup> myeloid cells isolated from the spleen of immunized mice.
- E Analysis of IL-1β expression by CD11b<sup>+</sup> cells isolated from dLN of mice treated as in (D). Data are representative FACS histogram overlays, gated on VD<sup>-</sup>CD11b<sup>+</sup> cells with mean frequencies per group.
- F Quantification (mean + SEM) of IL-1 $\beta$  expression by CD11b<sup>+</sup> cells shown in (E).
- $\mathsf{G} \quad \mathsf{Analysis of } \mathsf{IL-1}\beta-\mathsf{expressing } \mathsf{CD11b^+} \ \mathsf{cells } \mathsf{shown in (E)}. \ \mathsf{Data are representative } \mathsf{FACS } \mathsf{plots, gated } \mathsf{on } \mathsf{VD^-CD11b^+} \mathsf{IL-1}\beta^+ \ \mathsf{cells } \mathsf{with } \mathsf{mean frequencies } \mathsf{per group.}$
- H  $\,$  Quantification (mean + SEM) of myeloid cell populations expressing IL-1  $\!\beta$  shown in (G).

Data information: Neutrophils were defined as CD11b<sup>+</sup>Ly6C<sup>-</sup>Ly6C<sup>low</sup> cells, monocytes/macrophages as CD11b<sup>+</sup>Ly6C<sup>-</sup>Ly6C<sup>high</sup> cells. Cells were isolated (A–C) from naïve non-immunized mice n = 5 of each genotype; (D–H) at day 7 after immunization from n = 4 WT PBS-, n = 3 IL-1R1<sup>-/-</sup> PBS-, n = 4 WT PTx-, n = 3 IL-1R1<sup>-/-</sup> PTX MOG/ CFA-immunized mice. Cells (A, C, D–H) were restimulated with 20 ng/ml GM-CSF and (B) with GM-CSF or 500 ng/ml LPS in the presence of monensin for 4 h. \*P < 0.05, \*\*P < 0.01, N.S., not significant; two-tailed unpaired *t*-test. Experiments were performed twice with similar results.

that all of them, regardless of co-expression of IFN $\gamma$  and/or GM-CSF, expressed similar levels of IL-1R1 (Fig 2E and F). We conclude that both mouse and human subpopulations of Th17 cells, including the most autoaggressive GM-CSF<sup>+</sup> Th17 cells, can potentially respond to IL-1 and therefore could be dependent on IL-1 signaling.

# PTx enhances the development of MOG-specific GM-CSF<sup>+</sup> Th17 cells in an IL-1R1-dependent manner

To investigate the role of PTx-induced IL-1 $\beta$  expression on the development of autoreactive T cells, we compared cytokine production by MOG-specific CD4 T cells generated after MOG/CFA immunization of mice with or without PTx injections. To visualize antigen-specific T cells, we re-activated spleen- and dLN-derived lymphocytes with MOG peptide for 6 h and addressed the expression of CD40 ligand

(CD40L) together with the cytokine staining. CD40L is transiently expressed by T cells after their activation and can be used to identify antigen-specific T cells (Chattopadhyay *et al*, 2005; Frentsch *et al*, 2005). In WT mice, addition of PTx to the immunization protocol doubled numbers of MOG-reactive Th17 cells (Fig 3A and B). This effect of PTx was strongly dependent on IL-1R1, since we did not observe it in mice lacking the IL-1 receptor. Instead, IL-1R1<sup>-/-</sup> mice showed a dramatic reduction of antigen-specific Th17 cells compared to WT animals, regardless of the addition of PTx (Fig 3A and B). As a negative control, we restimulated cells with the non-relevant OVA peptide and could not detect OVA-specific cells neither in WT nor in IL-1R1-deficient animals (Fig EV3).

Importantly, the stimulatory effect of PTx observed in WT mice was not restricted to Th17 cells but was also seen in CD4 T cells expressing IFN $\gamma$ , GM-CSF, or both (Fig EV4). These data are in



#### Figure 2. IL-1R1 expression by murine and human Th17 cells.

A–D Analysis of IL-1R1 expression by different populations of murine (A) CD4 T cells and (B) Th17 cells isolated from the dLN 9 days after immunization, and (C) CD4 T cells and (D) Th17 cells isolated from the CNS at the peak of EAE.

E, F Analysis of IL-1R1 expression by different populations of human (E) CD4 T cells and (F) Th17 cells isolated from the peripheral blood.

Data information: Data (A, C) are representative FACS histogram overlay of VD<sup>-</sup>TCRβ<sup>+</sup>CD4<sup>+</sup> cells gated on IL-17A<sup>+</sup>, IL-17A<sup>-</sup>IFNγ<sup>+</sup> (IFNγ<sup>+</sup>), IL-17A<sup>-</sup>GM-CSF<sup>+</sup> (GM-CSF<sup>+</sup>), or IL-17A<sup>-</sup>FNP3<sup>+</sup> (FoxP3<sup>+</sup>) cells (also see Appendix Fig S1) and quantifications shown as bar diagram (mean + SEM). Data (E) are FACS plots of cells gated on CD14<sup>-</sup>CD3<sup>+</sup>CD4<sup>+</sup> cells with mean frequencies per group and quantifications shown as bar diagram (mean + SEM). Data (B, D, F) are FACS plots of Th17 cells and histograms of IL-1R1 staining of indicated Th17-cell subpopulations with mean fluorescent intensity (MFI)  $\pm$  SEM or mean frequencies per group. Cells (A–D) were restimulated in the presence of MOG for 6 h. Data consist of *n* = 4 wild-type mice immunized with MOG/CFA/PTx. Cells (E, F) were restimulated with PMA/ionomycin for 4 h. Data consist of PBMC isolated from *n* = 4 healthy individuals. \**P* < 0.01, \*\*\**P* < 0.001; two-tailed unpaired *t*-test. Experiments were performed twice with similar results.

agreement with our findings that IL-1R1<sup>-/-</sup> mice displayed very limited numbers of such cells regardless of PTx administration (Fig EV4). Previously, our group and others have shown that in the context of EAE, IFN $\gamma$ -, and GM-CSF-positive cells are mainly (or even exclusively) originating from Th17 cells (Kurschus *et al*, 2010; Hirota *et al*, 2011; Brucklacher-Waldert *et al*, 2016). Taken together, we conclude that deficiency in IL-1 signaling (either by IL-1R1

deletion or absence of PTx during immunization) results in reduced Th17 cells and, as expected, their IFN $\gamma$ - and GM-CSF-expressing progenies, during priming of EAE.

As mentioned above, following EAE induction, Th17 cells go through a process of differentiation that results in the co-expression of IFN $\gamma$  and GM-CSF, which are critical for the pathogenicity of CD4 T cells (Kurschus *et al*, 2010; Codarri *et al*, 2011; El-Behi *et al*, 2011;



Figure 3. GM-CSF-expressing Th17 cells are expanded by PTx in response to IL-1 signaling.

A Analysis of cytokine expression by CD4 T cells isolated from the spleen. Data are representative FACS plots, gated on VD<sup>-</sup>TCR $\beta$ <sup>+</sup>CD4<sup>+</sup>CD44<sup>+</sup> cells (upper row) with mean frequencies among CD4 T cells per group  $\pm$  SEM and (lower row) with mean frequencies per group.

B Frequencies and total numbers (mean + SEM) of MOG-specific Th17 cells isolated from the dLN.

C, D Frequencies and total numbers (mean + SEM) of MOG-specific GM-CSF<sup>+</sup> Th17 cells isolated from the spleen (C) and dLN (D).

Data information: Cells (A–D) were isolated at day 9 after immunization and restimulated in the presence of MOG for 6 h. Data consist of n = 5 WT PBS-, n = 4 WT PTx-, n = 4 IL-1R1<sup>-/-</sup> PBS-, n = 4 IL-1R1<sup>-/-</sup> PTx MOG/CFA-immunized mice. \*P < 0.05, \*\*P < 0.01, N.S., not significant; two-tailed unpaired *t*-test. Experiments were performed at least twice with similar results.

Hirota *et al*, 2011). Therefore, we further analyzed antigen-specific IL-17A-producing cells for the production of IFN $\gamma$  and GM-CSF. In WT mice, we found that frequencies and numbers of MOG-specific Th17 cells co-expressing GM-CSF were significantly higher when mice were injected with PTx compared to PBS-treated animals (Fig 3A, C and D). The vast majority of IFN $\gamma$ -positive Th17 cells also expressed GM-CSF, while IFN $\gamma$  single positive Th17 cells represented only a minor subpopulation regardless of the PTx administration (Fig 3A). Importantly, the stimulatory effect of PTx was not observed in mice deficient for IL-1R1, in which autoreactive IL-17A<sup>+</sup> cells displayed impaired co-expression of GM-CSF in both the spleen and dLN (Fig 3C and D). Collectively, these findings suggest that PTx enhances the generation of pathogenic Th17 cells and particularly its GM-CSF-positive subset, by promoting their expansion via IL-1 signaling.

Regulatory T cells are generated early after EAE induction (O'Connor & Anderton, 2008; Yogev *et al*, 2012), and their development was shown to be transiently suppressed by PTx (Cassan *et al*, 2006; Chen *et al*, 2006). However, we did not observe any effects of PTx administration or IL-1R1 deletion on the numbers of FoxP3<sup>+</sup> CD4 T cells in dLN and spleen when mice were analyzed 9 days after immunization (Appendix Fig S2A–C). Moreover, expression levels of Treg co-stimulatory molecules GITR, ICOS, and CTLA-4 were also unchanged in all groups of animals (Appendix Fig S2D and E). We conclude that neither IL-1 signaling, nor PTx administration is critically influencing Treg cell development following EAE induction.

#### IL-1 signaling in T cells is essential for expansion of antigenspecific GM-CSF<sup>+</sup> Th17 cells

To investigate the exact role of IL-1 in the development of autoreactive T cells and their pathogenicity, we specifically deleted the IL-1 receptor type 1 in  $\alpha\beta\text{-}TCR^+$  T cells by crossing conditional IL-1R1<sup>fl/fl</sup> mice to CD4-Cre mice, resulting in IL-1R1<sup> $\Delta T$ </sup> mice. These mice developed normally and did not display abnormalities in the immune system when kept under SPF conditions. However, CD4 T cells isolated from mutant mice failed to proliferate in response to IL-1β administration in vitro (Mufazalov et al, 2016). Previously, it was reported that IL-1R1 fully deficient mice show an impaired CD4 T-cell-mediated antigen response and that these mice are resistant to induction of EAE (Matsuki et al, 2006; Sutton et al, 2006; Chung et al, 2009). To test whether EAE resistance specifically stems from the lack of T-cell responses to IL-1, we immunized WT, IL-1R1 $^{-/-}\!\!\!\!$  , and IL-1R1 $^{\Delta T}$  mice with MOG/CFA along with two injections of PTx and investigated priming of CD4 T cells 9 days postimmunization. As expected, we could not detect IL-1R1 on the mutant T cells isolated from either  $\text{IL-1}\text{R1}^{\Delta T}$  or  $\text{IL-1}\text{R1}^{-/-}$  mice (Fig 4A). We observed that not only global deletion of IL-1R1, but also T-cell-specific deletion of the receptor resulted in a reduction of IL-17A-expressing T cells (Fig 4A).

When we further analyzed antigen-specific CD4 T cells, we found that germ line or T-cell-specific deletion of IL-1R1 resulted in a similar decrease of autoreactive CD4 T cells compared to WT controls (Appendix Fig S3). Moreover, frequencies and numbers of MOG-specific Th17 cells and Th17 cells co-expressing GM-CSF were equally reduced in both groups of IL-1R1-deficient animals (Fig 4B–D), suggesting that this is a T-cell-intrinsic phenomenon, since antigen-presenting cells should not be targeted by CD4-Cre-mediated recombination. Importantly, although numbers of GM-CSF<sup>+</sup> Th17 cells were significantly reduced in both groups of mutant mice compared to WT controls, they were still detectable (Fig 4B and D). This observation is in favor of impaired proliferation or survival,



#### Figure 4. T-cell-specific deletion of IL-1R1 impairs Th17 cell expansion.

- A Analysis of CD4 T cells isolated from the dLN and restimulated with PMA/ ionomycin for 4 h. Data are representative FACS plots, gated on VD<sup>-</sup>CD4<sup>+</sup> cells with mean frequencies per group.
- B Analysis of cytokine expression by CD4 T cells isolated from the spleen and restimulated with MOG for 6 h. Data are representative FACS plots, gated on VD<sup>-</sup>TCR $\beta^+$ CD4<sup>+</sup>CD4<sup>+</sup> cells (upper row) with mean frequencies among CD4 T cells per group  $\pm$  SEM and (lower row) with mean frequencies per group.
- C Total cell numbers (mean + SEM) of MOG-specific Th17 cells isolated from the spleen shown in (B, upper row).
- D Frequencies and total numbers (mean + SEM) of MOG-specific GM-CSF<sup>+</sup> Th17 cells isolated from the spleen shown in (B, lower row).

Data information: Cells (A–D) were isolated at day 9 after MOG/CFA/PTx immunization. Data consist of n = 5 mice of each genotype. \*P < 0.05, \*\*P < 0.01, N.S., not significant; two-tailed unpaired t-test. Experiments were performed twice with similar results.

rather than differentiation failure, of MOG-specific Th17 cells devoid of IL-1 signaling.

#### Development of neuroinflammation is dependent on a T-cellintrinsic response to IL-1

To investigate the consequences of impaired Th17 cell development for autoimmunity, we immunized IL-1R1-deficient mice with MOG/CFA along with PTx injections and followed the progression of EAE. We found that T-cell-specific deletion of IL-1R1 resulted in a mild disease, which was nevertheless aggravated compared to mice fully deficient for IL-1 signaling (Fig 5A–C). Of note, mice with global IL-1R1 deletion were resistant to EAE over 28 days of examination, recapitulating the reported phenotype of previously published IL-1R1-deficient mice (Matsuki *et al*, 2006; Sutton *et al*, 2006).

To investigate which cell types are responsible for the (mild) EAE symptoms seen in the  $IL-1R1^{\Delta T}$  mice, we analyzed immune cell populations in the secondary lymphoid organs and the CNS infiltrates of diseased animals depicted in Fig 5D. Sixteen days after immunization, when WT mice were at the peak of disease, we found no differences in T-cell numbers in the spleen of immunized animals regardless of the genotype (Appendix Fig S4). However, when we analyzed the CNS samples, we found that  $IL-1R1^{\Delta T}$  mice had similar numbers of infiltrating leukocytes but a reduction of CD4 T cells compared to WT controls, while the mice with complete IL-1R1 deficiency had almost no cell infiltrates in the CNS (Fig 5E). Furthermore, when compared to WT mice, we found a strong reduction in antigen-specific Th17 cells in the CNS of  $IL-1R1^{\Delta T}$  animals and a complete absence of these cells in  $IL-1R1^{-/-}$  mice (Fig 5F). The presence of autoreactive Th17 cells in diseased  $IL-1R1^{\Delta T}$  mice, although in reduced numbers, suggests that also without IL-1 signaling, encephalitogenic T cells can develop, as long as other cells in their environment are able to respond to IL-1. In addition to reduced Th17 cell numbers, also the mean fluorescence intensity for IL-17A staining was significantly reduced in cells isolated from  $IL-1R1^{\Delta T}$ animals, suggesting impaired IL-17A expression on the single cell level (Fig 5G). When antigen-specific Th17 cells were further analyzed, we found reduced frequencies and numbers of GM-CSF<sup>+</sup> IL-17A<sup>+</sup> cells in the IL-1R1<sup> $\Delta$ T</sup> mice (Fig 5F and H), correlating with the reduced disease severity of these mice.

In contrast to the effect on the development of pathogenic Th17 cells, T-cell-specific deletion of IL-1R1 had no impact on the development of regulatory T cells (Fig 5I). Therefore, the ratio of FoxP3<sup>+</sup> cells to Th17 cells shifted toward Treg cells in the CNS of IL-1R1<sup> $\Delta$ T</sup> mice (Fig 5J and K), reflecting impaired cytokine production by mutant cells. We conclude that IL-1 signaling is not critical for Treg cell homeostasis.

Migration of inflammatory Th17 cells from the secondary lymphoid organs to the CNS was shown to be dependent on the expression of the chemokine receptor CCR6 (Reboldi *et al*, 2009; Comerford *et al*, 2010). To test whether the reduction in EAE severity observed in IL-1R1<sup> $\Delta$ T</sup> mice stems from altered trafficking of IL-1R1-deficient Th17 cells, we analyzed CCR6 expression in the dLN and CNS of EAE-diseased mice. We found no differences in the expression of this chemokine receptor by IL-1R1-sufficient and IL-1R1-deficient Th17 cells (Fig 5L–N), suggesting normal CCR6dependant migratory capacities of IL-1R1-deficient T cells. Taken together, our data demonstrate that encephalitogenic T cells can develop in the absence of IL-1 signaling, although in much lower numbers, which ultimately leads to the reduced EAE severity in IL-1R1<sup> $\Delta$ T</sup> mice.

# *Ex vivo* expansion of Th17 cells in the presence of IL-23 restores the pathogenic potential of IL-1R1-deficient T cells

To study the role of IL-1 signaling in expansion of MOG-specific Th17 cells, we isolated cells from MOG/CFA-immunized WT mice and cultured them in the presence of MOG peptide and anti-IFN $\gamma$ .



Figure 5. Mice with T-cell-specific deletion of IL-1R1 are only partially resistant to EAE induction.

- A-C (A) Mean, (B) maximal EAE clinical scores, and (C) area under the curve (AUC) of mice immunized with MOG/CFA/PTx. Data shown as (A) mean (+ SEM) and (B, C) individual plots with mean and are combined from at least four independent experiments and consist of n = 41 WT, n = 24 IL-1R1<sup> $\Delta$ T</sup>, n = 18 IL-1R1<sup>-/-</sup> mice. \*P < 0.05, \*\*\*P < 0.001; two-tailed unpaired t-test. Black asterisks in (A) represent statistical analysis of IL-1R1<sup> $\Delta$ T</sup> vs WT and blue asterisks of IL-1R1<sup> $\Delta$ T</sup> vs IL-1R1<sup> $\Delta$ T</sup>
- D, E (D) Maximal EAE clinical scores (individual plots with mean) of selected mice and (E) total cell numbers (mean + SEM) of infiltrates isolated from CNS and used for further analysis.
- F–H Analysis of cytokine expression by CD4 T cells isolated from the CNS of mice shown in (D). Data (F) are representative FACS plots, gated on VD<sup>-</sup>CD90<sup>+</sup>CD4<sup>+</sup> cells (upper row) with mean frequencies among CD4 T cells per group ± SEM and (lower row) with mean frequencies per group. (G) Mean fluorescent intensity (mean + SEM) of IL-17A staining by MOG-specific Th17 cells shown in (F, upper row). (H) Quantification of MOG-specific GM-CSF<sup>+</sup> Th17 cells (mean + SEM) shown in (F, lower row).
- I-K Analysis of Treg cells isolated from mice shown in (D). (I) Total Treg cell numbers (mean + SEM) isolated from CNS and dLN. (J) Ratio of Treg and Th17 cells isolated from the CNS. Data are representative FACS plots, gated on VD<sup>-</sup>CD90<sup>+</sup>CD4<sup>+</sup> cells with mean frequencies per group. (K) Quantification of Treg/Th17 cell ratio (mean + SEM) within CNS and dLN.
- L-N Analysis of CCR6 expression by CD4 T isolated from the dLN and CNS of EAE-diseased mice. Data (L) are representative FACS histogram overlays of (black) control, (blue) IL-1R1<sup>ΔT</sup> cells, gated on CD3<sup>+</sup>CD4<sup>+</sup>IL-17A<sup>+</sup> cells; and of (black filled) control, (blue filled) IL-1R1<sup>ΔT</sup> cells, gated on CD3<sup>+</sup>CD4<sup>+</sup>IL-17A<sup>-</sup> cells. (M) Mean fluorescent intensity (mean + SEM) of CCR6 staining by CD4 T cells shown in (L). (N) Mean fluorescent intensity (mean + SEM) of CCR6 staining by GM-CSF<sup>+</sup> and GM-CSF<sup>-</sup> Th17 cells isolated from CNS shown in (L).

Data information: Cells (F–H) were restimulated with MOG for 6 h and (I–N) in the presence of PMA/ionomycin for 4 h. Data (D–K) consist of n = 4 WT, n = 3 IL-1R1<sup> $\Delta$ T</sup>, and n = 3 IL-1R1<sup>-I-</sup> mice and (L–N) of n = 5 mice of each genotype. \*P < 0.05, \*\*P < 0.01, \*\*\*P < 0.001, N.S., not significant; two-tailed unpaired t-test. Experiments were performed at least twice with similar results.

We detected a dramatic increase in the frequencies and numbers of Th17 cells in cultures supplemented with IL-1 $\beta$  compared to cytokine-free conditions (Fig 6A). Apart from IL-1 $\beta$ , also IL-23 was shown to play a critical role in the establishment of T-cell-mediated pathogenicity (Cua *et al*, 2003). When we added IL-23 to the cultures, we observed an even stronger stimulatory effect for the



Figure 6. IL-23 increases numbers of encephalitogenic IL-1R1-deficient Th17 cells.

- A–C Analysis of (A) IL-17A expression by wild-type CD4 T cells, followed with the analysis of (B) GM-CSF and (C) GM-CSF and IFNγ co-expression by Th17 cells, shown as mean + SEM.
- D–G Analysis of cytokine expression by cells isolated from WT (black), IL-1R1<sup>ΔT</sup> (blue), and IL-1R1<sup>-/-</sup> (red) mice and cultured with or without IL-23. Data (D) are representative FACS plots, gated on VD<sup>--</sup>TCRβ<sup>+</sup> cells with mean frequencies per group. Quantification (mean + SEM) of (E) MOG-specific Th17 cells, (F) GM-CSF<sup>+</sup> Th17 cells, and (G) GM-CSF<sup>+</sup>IFNγ<sup>+</sup> Th17 cells.
- H, I (H) Mean EAE clinical scores (+ SEM) of Rag1<sup>-/-</sup> mice which received 1 × 10<sup>5</sup> IL-17A<sup>+</sup> cells of indicated genotypes and (I) total cell numbers (mean + SEM) of infiltrates isolated from CNS and used for further analysis.
- J-L Analysis of cytokine expression by CD4 T cells isolated from CNS of mice shown in (H) and restimulated with MOG for 6 h. (J) Data are representative FACS plots, gated on VD<sup>-</sup>TCR $\beta^+$  cells (upper row) with mean frequencies ± SEM, and (lower row) with mean frequencies per group. (K) Total cell numbers (mean + SEM) of Th17 cells shown in (J, upper row). (L) Frequencies and total numbers (mean + SEM) of GM-CSF<sup>+</sup> Th17 cells shown in (J, lower row).

Data information: (A–G) Cells were isolated from dLN and the spleen of MOG/CFA-immunized mice, cultured under indicated conditions for four days and restimulated with MOG for 6 h prior to analysis. Data consist of n = 4 of each genotype. (H–L) Data consist of Rag1<sup>-/-</sup> mice which received WT (n = 6), IL-1R1<sup>AT</sup> (n = 3) and IL-1R1<sup>-/-</sup> (n = 4) cells polarized in the presence of IL-23 prior to transfer. \*P < 0.05, \*\*P < 0.01, \*\*\*P < 0.001, N.S., not significant; two-tailed unpaired *t*-test. All experiments were performed at least twice with similar results.

expansion of Th17 cells, with the greatest outcome when both IL-1 $\beta$  and IL-23 were combined together (Fig 6A). Accordingly, numbers of GM-CSF<sup>+</sup>, as well as GM-CSF<sup>+</sup>IFN $\gamma^+$  Th17 cells, were also increased in cultures supplemented with either IL-1 $\beta$  or IL-23 or both (Fig 6B and C). Therefore, we hypothesized that stimulation with IL-23 may substitute the deficiency in IL-1 signaling. Indeed, cells isolated from both IL-1R1-deficient strains, namely IL-1R1<sup> $\Delta$ T</sup> and IL-1R1<sup>-/-</sup> mice, responded equally to IL-23 and showed robust expression of IL-17A after culturing for four days (Fig 6D and E). Exposure to IL-23 also had a stimulatory effect on numbers of

GM-CSF-producing cells among the MOG-specific IL-1R1-deficient Th17 cells (Fig 6F). Moreover, IFN $\gamma$  co-expression by GM-CSF<sup>+</sup> Th17 cells was comparable between different genotypes when IL-23 was added (Fig 6G).

As addition of IL-23 led to higher numbers of IL-1R1-deficient GM-CSF-producing Th17 cells, we tested whether these cells are pathogenic by using a model of passive EAE by transfer of *ex vivo* reactivated T cells. For that we isolated cells from the spleen and dLN of WT, IL-1R1<sup> $\Delta$ T</sup>, and IL-1R1<sup>-/-</sup> MOG/CFA-immunized mice and polarized them *in vitro* in the presence of MOG peptide,

anti-IFN $\gamma$ , and IL-23, as described above. After four days of culture, the numbers of harvested cells were adjusted to  $1 \times 10^5$  IL-17A<sup>+</sup> cells of each genotype and total cell preparations were transferred into Rag1<sup>-/-</sup> mice. These cells, regardless of the genotype, transmitted disease and caused strong EAE symptoms in recipient mice (Fig 6H), confirming the pathogenicity of IL-1R1-deficient T cells observed upon active immunization. At the peak of disease, we isolated cellular infiltrates from the CNS and found that CD4 T cells represented the major population of immune cells and were equally present in mice that received WT or IL-1R1-deficient cells (Fig 6I). Furthermore, we observed high numbers of IL-17A<sup>+</sup> CD4 T cells within the inflamed CNS in all groups of diseased animals (Fig 6J and K). In line with this, we did not find differences in the percentages and total numbers of GM-CSF-co-expressing cells among the mice of the different groups (Fig 6J and L). CD4 T-cell-derived GM-CSF was shown to activate CNS resident microglia cells (Ponomarev et al, 2007) as well as monocytes (Croxford et al, 2015) and to sustain neuroinflammation by promoting IL-1ß expression. Therefore, we investigated IL-1ß production by myeloid cells isolated from inflamed CNS and indeed detected large populations of CD11b<sup>+</sup> cells expressing IL-1 $\beta$  in a similar manner in all analyzed animals (Appendix Fig S5).

To directly study the role of IL-23 *in vivo*, we induced EAE in WT and IL- $1R1^{\Delta T}$  mice and administered the cytokine via i.p. injections. Constitutive treatment with IL-23 had a mild, but stimulatory effect on EAE severity of both WT and IL- $1R1^{\Delta T}$  mice (Fig EV5). Collectively, we conclude that the necessity of IL-1 signaling for the

expansion of encephalitogenic T cells can be substituted by IL-23 in an adoptive transfer system and partially *in vivo*.

#### Treg cells suppress disease in IL-1R1-deficient mice

Treg cells are the potent suppressors of autoimmunity and can be transiently depleted by injections of PC61 antibodies directed against CD25 (McGeachy et al, 2005). To study whether the small numbers of Th17 cells found after immunization of IL-1R1<sup>-/-</sup> mice can be pathogenic also in active EAE, we depleted Tregs 5 days prior to immunization. This led to the development of strong EAE symptoms in IL-1R1<sup>-/-</sup> mice treated with PC61, although with later disease onset compared to controls (Fig 7A). Nineteen days after immunization, we selected mice with similar clinical scores (Fig 7B) and analyzed them for cytokine-producing T cells in the CNS. As seen in Fig 7C, we found no differences in numbers of Th17 cells deficient or sufficient for IL-1 signaling. These data suggest that IL-1 signaling is not critical for the suppressive capacity of Treg cells, which is in agreement with our findings that these cells hardly express the IL-1R1 (Figs 2A and C, and 4A) and that IL-1R1-deficient Treg cells develop normally after EAE induction (Appendix Fig S2 and Fig 5I). Our data suggest that under IL-1R1-deficient conditions, Th17 cells cannot properly expand due to their impaired IL-1 response and consequent suppression by Tregs.

Importantly, IL-1R1 expression by encephalitogenic Th17 cells was not altered in wild-type mice treated with PC61 in comparison with the non-treated group (Fig 7D and E). Furthermore, we could



Figure 7. Treg depletion results in EAE susceptibility of IL-1R1-deficient mice.

- A Mean EAE clinical scores (+ SEM).
- B Maximal EAE clinical scores (individual plots with mean) of selected mice.
- C Total cell numbers (mean + SEM) of infiltrates isolated from the CNS of mice shown in (B).
- D Analysis of IL-1R1 expression by Th17 cells isolated from CNS of EAE-diseased mice shown in (B). Data are representative FACS histogram overlays of cells isolated from (gray) WT mice treated with PBS, (black) WT, (blue) IL-1R1<sup>ΔT</sup>, and (red) IL-1R1<sup>-/-</sup> mice treated with PC61, gated on VD<sup>-</sup>TCRβ<sup>+</sup>CD4<sup>+</sup>IL-17A<sup>+</sup> cells.
- E Quantification (mean + SEM) of IL-1R1 expression by Th17 cells, shown in (D).

Data information: Cells (C, D, E) were restimulated in the presence of MOG for 6 h. Data consist of (A) n = 10 WT PBS-treated, n = 4 IL-1R1<sup>-/-</sup> PBS-treated, n = 4 WT PC61-treated, n = 4 IL-1R1<sup>-/-</sup> PC61-treated and n = 5 IL-1R1<sup>-/-</sup> PC61-treated mice and of (B–E) n = 4 WT PBS-treated, n = 4 WT PC61-treated, n = 3 IL-1R1<sup>AT</sup> PC61-treated mice and n = 5 IL-1R1<sup>-/-</sup> PC61-treated mice and of (B–E) n = 4 WT PBS-treated, n = 4 WT PC61-treated, n = 3 IL-1R1<sup>AT</sup> PC61-treated mice and n = 5 IL-1R1<sup>-/-</sup> PC61-treated mice and n = 3 IL-1R1<sup>-/-</sup> PC61-tr

show that T cells isolated from the CNS of PC61-treated IL- $1R1^{\Delta T}$  mice expressed no or very little IL-1R1 (Fig 7D and E), indicating that disease in these mice was not driven by cells that could potentially escape CD4-Cre-mediated IL-1R1 deletion.

## Discussion

The development of EAE requires CD4 T-cell maturation in secondary lymphoid organs upon immunization. Here, we demonstrate that T-cell-intrinsic IL-1 signaling is necessary to promote the expansion of encephalitogenic CD4 T cells. We show that PTx, an essential component of the immunization protocol, stimulates IL-1 $\beta$  production by myeloid cells, which leads to proliferation of encephalitogenic GM-CSF-producing Th17 cells. Importantly, our data emphasize that IL-1 $\beta$  signaling is not necessary for primary differentiation of pathogenic Th17 cells *in vivo*.

A recent study identified an important role of IL-1 produced by neutrophils and monocyte-derived myeloid cells during their transmigration into the CNS for EAE development (Levesque et al, 2016). We speculate that IL-1 does not only activate the endothelial cells of the blood-spinal cord barrier, but promotes the secondary activation and expansion of CD4 T cells during the disease onset within the CNS. Here, we investigated IL-1 production in secondary lymphoid organs during CD4 T-cell priming and found that also at this stage, the myeloid cells represent the main source of IL-1ß in draining lymph nodes and spleen. Addition of PTx to the immunization protocol greatly increased the numbers of autoreactive CD4 T cells, although the primary effects of PTx on T cells were described to be rather negative due to suppressed proliferation and migration (Robbinson et al, 1996; Su et al, 2001). On the other hand, EAE cannot be induced using most protocols of active immunization without the addition of PTx (Stromnes & Goverman, 2006). Together, these observations suggested that PTx might promote EAE development through intermediate factor(s). In line with that, PTx-induced IL-6 was suggested as one of the factors that stimulates Th17 cell development (Chen et al, 2007). Later, it was found that IL-6 in fact stimulates IL-1R1 expression on Th17 cells rendering them responsive to IL-1 (Chung et al, 2009). In agreement with our data, neutrophils, macrophages, and monocyte-derived dendritic cells were shown to produce IL-1 in response to PTx administration, thereby promoting CD4 T-cell priming (Dumas et al, 2014; Lin et al, 2016; Ronchi et al, 2016). It was also reported that deletion of IL-1 receptor antagonist, a key negative regulator of IL-1 signaling, allows EAE development even without PTx (Matsuki et al, 2006). Together, these findings suggest that PTx promotes EAE development via IL-1 signaling.

In line with previous reports (Chung *et al*, 2009; Guo *et al*, 2009; Martin *et al*, 2016), we found that Th17 cells, including its IFN $\gamma$ and GM-CSF-positive subpopulations, express the highest levels of IL-1R1 among other CD4 T cells. Therefore, the stimulatory effect of myeloid cell-derived IL-1 $\beta$  was not only seen in the general expansion of MOG-specific Th17 cells but was also observed in IL-17A-positive cells that co-express GM-CSF. The importance of IL-1 signaling for GM-CSF production by CD4 T cells *in vivo* has been reported previously (Lukens *et al*, 2012), but the question of its T-cell-intrinsic function remained open. Using a novel mouse strain with IL-1R1 conditionally deleted in  $\alpha\beta$ -TCR<sup>+</sup> T cells, we could show that T cells, which have developed in the absence of IL-1 signaling but in an otherwise IL-1R1-sufficient environment, also displayed reduced GM-CSF expression in the Th17 subset. This notion is important in light of broad IL-1R1 expression by different cell types and restricts its necessity directly to T cells. Of note, previous studies reported that T-cell-derived GM-CSF is a potent inducer of IL-1 $\beta$  expression by myeloid cells (Croxford *et al*, 2015), which then stimulates further expansion of GM-CSF<sup>+</sup> Th17 cells in a positive feedback loop, as we show here. Our data contribute in elucidating such GM-CSF/IL-1 proinflammatory axis (reviewed in Paré *et al* (2016)).

Interestingly, some  $IL-1R1^{\Delta T}$  mice developed a mild paralysis after EAE induction, showing for the first time that CD4 T cells indeed gain pathogenic potential also in the absence of IL-1 signaling. This in vivo data suggested that IL-1 is not an indispensible factor for pathogenicity and our following transfer experiments confirmed this notion. Cells, which differentiated independently of IL-1 signaling, successfully transmitted EAE after polarization in the presence of IL-23. It is broadly accepted that IL-23 does not stimulate de novo Th17 cell generation, but rather stabilizes this phenotype and promotes Th17 cell expansion (Bettelli et al, 2006; Veldhoen et al, 2006; McGeachy et al, 2007). This led us to speculate that in our cultures, supplemented with a very limited cytokine cocktail, the main effect was the expansion of already existing pathogenic cells rather than their *de novo* generation. The observed pathogenicity of the transferred cells is in contrast to the reported inability of IL-1R1-deficient cells to transmit EAE into Rag1<sup>-/-</sup> mice (Chung et al, 2009). However, we believe that differences in the experimental setting may explain this discrepancy. Chung and coauthors did not expand cells prior to transfer but rather immunized recipient animals, while we enriched transferred cell populations for IL-17A-positive cells after in vitro expansion. Nevertheless, their independent experiments with chimeric mice, and experiments reported by Croxford et al (2015), which both consisted of WT and IL-1R1-deficient bone marrow-derived progenitors, showed that neuroinflammation driven by WT cells resulted also in recruitment of IL-1R1-deficient CD4 T cells into the CNS. Although IL-1R1deficient CD4 T cells were reduced in numbers and showed impaired cytokine production, this finding is in full agreement with our data showing the presence of small but sufficient quantities of IL-1R1-deficient cells in the CNS of diseased IL-1R1<sup> $\Delta T$ </sup> mice.

The balance between Tregs and Th17 cells is crucial for development of autoimmunity. Analyzing both the preclinical and the effector phase of EAE, we did not find effects of disrupted IL-1 signaling neither on peripheral nor on CNS infiltrated Tregs. However, we did find increased frequencies of Tregs among CD4 T-cell CNS infiltrates in IL-1R1<sup> $\Delta$ T</sup> mice, reflecting the impaired effector T-cell development in these animals. One might speculate that the shift of the Treg-Th17 ratio in IL-1R1<sup> $\Delta$ T</sup> mice may even disguise the real encephalitogenic potential of effector T cells devoid of IL-1 signaling. Our data are in line with an impaired Th17 response combined with increased frequencies of Treg cells seen in mice with T cells lacking MyD88, the adaptor molecule for Toll-like receptor and IL-1 receptor family signaling (Schenten *et al*, 2014). This study and our data together support the notion that IL-1 signaling acts on precursor cells of Tregs, rather than on already developed regulatory T cells.

Importantly, mice completely lacking IL-1 signaling did not show EAE symptoms or CNS infiltrates, unless regulatory T cells were ablated. This indicates that IL-1 has multiple roles in the induction of EAE beyond expansion of encephalitogenic T cells. One of probably many roles might be the activation and recruitment of myeloid cell populations as shown here. Only under very restrictive conditions, when the suppressive arm was removed by Treg cell depletion,  $IL-1R1^{-/-}$  mice developed EAE. Together, our data clearly show that the main role of IL-1 in terms of T helper cell-mediated pathogenicity in mice is not primarily the differentiation of Th17 cells but rather the expansion or boosting of GM-CSF-producing encephalitogenic Th17 cells.

Using mouse models greatly contributed to the development of therapeutic interventions for human diseases. Suppression of IL-1 signaling has beneficial effects in several pathologies, including those with autoimmune and inflammatory components (reviewed in Schett *et al* (2016)). Human Th17 cells were found to express high levels of IL-1R1 (Lee *et al*, 2010) and IL-1 was proposed as a critical factor in pathogenic Th17 cell development (Zielinski *et al*, 2012). Our data demonstrate that like the mouse counterpart, also human GM-CSF<sup>+</sup> Th17 cells express IL-1R1, making them responsive to IL-1. Importantly, IL-1 might be essential not only for the proliferation of human Th17 cells but also for their differentiation (Acosta-Rodriguez *et al*, 2007). Our data provide the mechanistic basis for better characterization of T cells recovered from MS patients, studies that are needed in order to explore the involvement of IL-1 signaling in this human disease.

# **Materials and Methods**

#### Mice

The *ll1r1*<sup>fl/fl</sup> mouse strain with exon 5 of *ll1r1* gene flanked by loxP sites was crossed to *Cd4-Cre* and *CMV-Cre* mice resulting in IL-1R1<sup> $\Delta$ T</sup> and IL-1R1<sup>-/-</sup> mouse strains, respectively (Mufazalov *et al*, 2016). Rag1<sup>-/-</sup> mice were obtained from in-house breeding.

All mice were bred in-house under SPF conditions. Age- and gender-matched genetically modified animals carrying loxP sites without *Cre* transgene were used as controls. Experiments were performed with 6- to 14-week-old mice on C57BL/6 background in accordance with the guidelines of the central animal facility institution (TARC, University of Mainz).

#### Immunization, active EAE induction, and clinical assessment

Mice were immunized subcutaneously at the tail basis with 50 µg MOG<sub>35-55</sub> peptide (Gene Script) emulsified in complete Freund's adjuvant (Difco) supplemented with 1.1 mg of heat-inactivated *Mycobacterium tuberculosis* (Difco). Along with immunization and on day 2 postimmunization, 200 ng of pertussis toxin (List Biological Laboratories) or Dulbecco's phosphate-buffered saline (PBS, Sigma) was administered by intraperitoneal injection, when indicated. Mice were scored daily for clinical signs of EAE as described (Huppert *et al*, 2010) with modifications. The clinical assessment scale was graded from 0 to 6 as follows: 0, no disease; 0.5, partial loss of tail tonicity; 1, complete loss of tail tonicity; 1.5, partially impaired righting reflex; 2.5, partial hindlimb paresis resulted in staggering gait; 3, hindlimb paresis; 3.5, hindlimb paralysis; 4,

unilateral forelimb paresis; 4.5, complete forelimb paresis; 5, forelimb paresis resulted in inability to move the body; 5.5, moribund; and 6, dead animal. Calculation of the area under the curve (AUC) was performed as described elsewhere (Fleming *et al*, 2005).

#### Cytokines and cell culture

For *in vitro* culture, cells were isolated from emulsion-draining lymph nodes (dLN) and spleen at day 10 after MOG/CFA immunization and mixed at 1–3 ratios. Cells were cultured at a concentration of  $0.6 \times 10^6$  cells/well in 200 µl complete medium (RPMI medium containing 10% FCS, 2 mM L-glutamine, 100 units/ml penicillin, 100 mg/ml streptomycin, 1 mM sodium pyruvate, 50 mM 2-mercaptoethanol, 10 mM HEPES, and 1% non-essential amino acids) supplemented with 10 µg/ml MOG<sub>35-55</sub> and 10 µg/ml  $\alpha$ -IFN $\gamma$  antibodies (BioXCell) in 96-well plates as triplicates. When indicated, cultures were additionally supplemented with 15 ng/ml IL-23 (Miltenyi), 25 ng/ml IL-1 $\beta$  (R&D Systems), or both. After culturing for 4 days, triplicates were pooled and restimulated in fresh medium containing MOG/brefeldin A for 6 h prior to cytokine analysis.

#### Transfer EAE

Donor mice were immunized with MOG/CFA as described above. On day 10 postimmunization, cells from dLN and spleen were isolated, mixed at 1 to 4 ratio, and cultured at a concentration of  $5 \times 10^6$  cells/ml in complete medium supplemented with 10 µg/ml MOG<sub>35-55</sub>, 10 µg/ml  $\alpha$ -IFN $\gamma$  antibodies, and 15 ng/ml IL-23 for 4 days. Afterward, total cell preparations containing  $1 \times 10^5$  blasting MOG-specific Th17 cells were washed twice in PBS and injected intraperitoneally into Rag1<sup>-/-</sup> mice. Along with cells transfer and on day 2 posttransfer, 200 ng of PTx was administered into recipient mice. Mice were scored for EAE signs as described above.

#### In vivo Treg depletion and IL-23 treatment

For Treg cell depletion, mice were treated intraperitoneally with 0.5 mg PC61 (anti-CD25 antibody, BioXCell) or with PBS five and three days prior to active EAE immunization. The treatment with 1  $\mu$ g IL-23 (Miltenyi) or with PBS was performed intraperitoneally at day 3, 5, 7, 9, 11, and 13 after active EAE immunization.

#### **Organ preparation**

Single cell suspensions of spleen and dLN were prepared by mechanical dissociation in PBS supplemented with 2% FCS. For the analysis of myeloid cells, spleens and dLN were digested with 2 mg/ml collagenase II (Sigma) and 40  $\mu$ g/ml DNase I (Roche) for 20 min at 37°C and homogenized with syringe and needle (18G). To obtain CNS infiltrating lymphocytes, brain and spinal cord were isolated from transcardially perfused mice, pooled, enzymatically digested, and centrifuged in 30/37/70 Percoll gradient as previously described (Mufazalov & Waisman, 2016).

#### Cells restimulation prior to cytokine staining

For cytokine detection, cells were incubated at (4–20)  $\times$   $10^6$  cells/ ml in complete culture medium. For IL-1 detection, cells were

incubated for 4 h in the presence of 2  $\mu$ M monensin (BioLegend) and 20 ng/ml GM-CSF (Peprotech), 500 ng/ml LPS (Sigma), or 50 ng/ml phorbol-12-myristate-13-acetate (PMA, Sigma) together with 500 ng/ml ionomycin (Invitrogen). For polyclonal stimulation of CD4 T cells, samples were incubated for 4 h in the presence of PMA, ionomycin together with monensin, or 1  $\mu$ g/ml brefeldin A (AppliChem). For antigen-specific CD4 T-cell restimulation, cells were incubated for 6 h using a combination of 20  $\mu$ g/ml MOG<sub>35-55</sub> or 20  $\mu$ g/ml OVA (Calbiochem) in the presence of monensin or brefeldin A.

#### Antibodies and flow cytometry

Single cell suspensions were stained with antibodies (Appendix Table S1) specific to epitopes present at the cell surface together with fixable viability dyes (VD, eBioscience). Afterward, cells were fixed and permeabilized with Cytofix/Cytoperm (BD Biosciences) or FoxP3 staining kit (eBiosciences) and stained intracellularly with antibodies (Appendix Table S2) according to the manufacturer's recommendations. Stained cells were acquired on FACS-Canto II (BD Biosciences) and obtained data were further analyzed with FlowJo software. Gating strategy always considered cell size, excluded duplets, and defined living cells as viability dye negative (VD<sup>–</sup>).

#### **Human PBMCs**

Buffy coats of healthy human donors who gave their informed consent were obtained from the Center for Blood Transfusion University Medical Center Mainz, Germany. PBMCs were isolated by Ficoll-Hypaque gradient centrifugation. Afterward,  $2 \times 10^6$  cells were reactivated in complete medium with PMA, ionomycin, and monensin for 4 h prior to staining. For the surface staining, CD14, CD3, and CD4 (BioLegend) FACS antibodies were used. Anti-IL-17A, IFN $\gamma$ , GM-CSF (BioLegend), IL-1R1, and goat IgG isotype control (R&D Systems) were applied after fixation with the cytokine staining kit (BD Biosciences).

#### Statistical analysis

Graphical and statistical analysis was done using Prism 5 software (GraphPad) and Excel (MS Office). Statistical significance was calculated using the two-tailed unpaired *t*-test. Values of P < 0.05, P < 0.01, and P < 0.001 were marked as \*, \*\*, and \*\*\*, respectively. N.S., not significant.

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#### Author contributions

IAM designed experiments; IAM, CS, TR, JK, FW, LAG, and JH performed experiments; IAM analyzed data; IAM, FCK, and AW interpreted data; WM and EP supplied essential reagents; TR, WM, and EP corrected the manuscript; IAM, FCK, and AW wrote the manuscript; AW supervised study.

#### **Conflict of interest**

The authors declare that they have no conflict of interest.

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