Bleomycin and IL-1β–mediated Pulmonary Fibrosis is IL-17A Dependent

The Harvard community has made this article openly available. Please share how this access benefits you. Your story matters

Citation

Published Version
doi:10.1084/jem.20092121

Citable link
http://nrs.harvard.edu/urn-3:HUL.InstRepos:5343416

Terms of Use
This article was downloaded from Harvard University’s DASH repository, and is made available under the terms and conditions applicable to Other Posted Material, as set forth at http://nrs.harvard.edu/urn-3:HUL.InstRepos:dash.current.terms-of-use#LAA
Bleomycin and IL-1β–mediated pulmonary fibrosis is IL-17A dependent


Idiopathic pulmonary fibrosis (IPF) is a destructive inflammatory disease with limited therapeutic options. To better understand the inflammatory responses that precede and concur with collagen deposition, we used three models of pulmonary fibrosis and identify a critical mechanistic role for IL-17A. After exposure to bleomycin (BLM), but not Schistosoma mansoni eggs, IL-17A produced by CD4+ and γδ+ T cells induced significant neutrophilia and pulmonary fibrosis. Studies conducted with C57BL/6 il17a−/− mice confirmed an essential role for IL-17A. Mechanistically, using ifnγ−/−, il10−/−, il10−/−il12p40−/−, and il10−/−il17a−/− mice and TGF-β blockade, we demonstrate that IL-17A–driven fibrosis is suppressed by IL-10 and facilitated by IFN-γ and IL-12/23p40. BLM-induced IL-17A production was also TGF-β dependent, and recombinant IL-17A–mediated fibrosis required TGF-β, suggesting cooperative roles for IL-17A and TGF-β in the development of fibrosis. Finally, we show that fibrosis induced by IL-1β, which mimics BLM–induced fibrosis, is also highly dependent on IL-17A. IL-17A and IL-1β were also increased in the bronchoalveolar lavage fluid of patients with IPF. Together, these studies identify a critical role for IL-17A in fibrosis, illustrating the potential utility of targeting IL-17A in the treatment of drug and inflammation–induced fibrosis.

Despite distinct etiological and clinical features, most chronic fibrotic disorders have in common a persistent irritant that sustains the production of growth factors, proteolytic enzymes, angiogenic factors, and fibrogenic cytokines (Wilson and Wynn, 2009). Together, these factors stimulate the deposition of connective tissue elements that progressively remodel normal tissue architecture. Although initially beneficial, tissue repair processes become pathogenic when they are not regulated, resulting in substantial deposition of extracellular matrix (ECM) components and development of scar tissue. In some diseases, like idiopathic pulmonary fibrosis (IPF), aberrant healing may lead to organ failure and death (Meltzer and Noble, 2008). Indeed, IPF and other chronic fibrotic lung diseases are associated with high morbidity and mortality and are generally refractory to existing pharmacological therapy (Shah et al., 2005). Therefore, better characterization of the molecular and immunological mechanisms of fibrosis is needed to identify new therapeutic modalities for these diseases.

Although a variety of cytokines, chemokines, and growth factors are important regulators of fibrosis, we identified a critical role for IL-13 in the development of fibrosis in schistosomiasis, a chronic liver disease caused by the parasitic helminth Schistosoma mansoni (Chiaramonte et al., 1999). Since then, IL-13 has been shown to exhibit fibrotic activity in a variety of diseases and tissues, including models of chronic asthma (Blease et al., 2001), skin fibrosis (Aliprantis et al., 2007), and bronchiolitis obliterans (Keane et al., 2007). A few recent studies have also suggested a role for IL-13 in bleomycin (BLM)–induced pulmonary fibrosis, a well-studied model of IPF (Jakubzick et al., 2003; Fichtner-Feigl et al., 2006). It has been suggested that IL-13 triggers fibrosis by inducing and activating TGF-β (Lee et al., 2001). Nevertheless, the mechanism of action of TGF-β in the development of pulmonary fibrosis is not clear.

**Abbreviations used:** BAL, bronchoalveolar lavage; BFA, brefeldin A; BLM, bleomycin; CBC, complete blood count; dKO, double KO; ECM, extracellular matrix; IPF, idiopathic pulmonary fibrosis; MMP, matrix metalloproteinase; mRNA, messenger RNA; NV, normal volunteer; TIMP, tissue inhibitor of MMP.
fibrosis remains controversial (Kaviratne et al., 2004; Varga and Pasche, 2008). Although it has been suggested that TGF-β contributes to BLM-induced inflammation and fibrosis by stimulating fibroblast proliferation and collagen-producing myofibroblasts (Cutroneo et al., 2007), recent studies also identified a critical role for TGF-β in the development of IL-17A–producing CD4+ T cells (Bettelli et al., 2006; Veldhoen et al., 2006), which regulate the pathogenesis of a variety of autoimmune and inflammatory diseases (Bettelli et al., 2008). Similarly, IL-1β can stimulate IL-17A production (Sutton et al., 2009), and IL-1β is a critical mediator of pulmonary fibrosis (Gasse et al., 2007). To date, however, a link between IL-17A–driven inflammation and pulmonary fibrosis has not been established.

The aim of the current study was to characterize the mechanisms of pulmonary fibrosis and to determine whether IL-17A in particular plays an important regulatory role. To do this, three distinct model systems were used, including S. mansoni egg-induced pulmonary fibrosis, BLM-induced pulmonary fibrosis, and the recently described IL-1β–driven fibrosis (Gasse et al., 2007). We report here that S. mansoni egg-mediated fibrosis is IL-13 dependent, as il13-/- mice developed minimal fibrosis compared with WT mice. In marked contrast, BLM-induced pulmonary fibrosis was independent of IL-13 at early time points. Instead, studies with il17a-/- mice revealed a critical role for IL-17A. Using IL-10 reporter mice and newly generated IL-10 and IL-17A double cytokine-deficient animals, we determined that CD4+ cell-derived IL-10 is required to limit the production and frequency of IL-17A+CD4+ and IL-17A+γδ T cells, thus preventing the development of severe IL-17A–driven fibrosis. We also show that IL-17A is essential for the development of fibrosis in response to IL-1β, thus extending recent studies that described an early innate role for IL-1β in pulmonary fibrosis (Gasse et al., 2007). Together, these studies demonstrate that fibrotic tissue remodeling is induced by distinct cytokine-dependent mechanisms, with the effector cytokines IL-13 and IL-17A playing central roles. Moreover, these findings suggest that TGF-β and proinflammatory mediators like IL-1β promote fibrosis by up-regulating the production of IL-17A, thus identifying IL-17A blockade as a potential treatment for fibrotic diseases like IPF.

RESULTS

Fibrosis is mediated by IL-13–dependent and IL-13–independent mechanisms

S. mansoni eggs delivered i.v. into S. mansoni egg-sensitized mice induce eosinophil-rich pulmonary granulomas around eggs trapped in the pulmonary microvasculature. The lesions are also associated with significant pulmonary fibrosis, as determined by hydroxyproline assay of lung tissue (Fig. 1 A). As observed in the liver during S. mansoni infection (Chiaramonte et al., 1999), studies conducted with il13-/- mice showed that pulmonary fibrosis is highly dependent on IL-13 (Fig. 1 A). However, unlike chronic liver fibrosis, hydroxyproline assays and Masson's trichrome staining of lung tissue suggested that the IL-13 decoy receptor (IL-13Rα2) was not significantly involved in the regulation of collagen deposition in the lung (Fig. 1, A and B). In contrast to S. mansoni egg-induced pulmonary fibrosis, BLM-induced fibrosis appeared to be both IL-13 and IL-13Rα2 independent. Indeed, C57BL/6 WT, il13-/-, and il13Rα2-/- mice all displayed indistinguishable levels of interstitial fibrosis after intratracheal delivery of BLM (Fig. 1, C and D).

BLM–induced pulmonary fibrosis is characterized by IFN–γ and IL-17A production

To elucidate the mechanisms involved in acute pulmonary inflammation and fibrosis after BLM administration, we characterized the immune and pathological responses in mice over a 21-d period. Significant weight loss occurred during the first week after intratracheal BLM (Fig. 2 A). Marked increases in collagen deposition were also seen by day 7, with peak pulmonary fibrosis developing between days 14 and 21 (Fig. 2, B, D, and F). Cachexia was associated with significant pneumonia and perivascular and peribronchial inflammation as early as day 4 after BLM and persisting through day 21 (Fig. 2 D). The peak in fibrosis was preceded by marked increases in matrix metalloproteinase (MMP) 12, tissue inhibitor of MMP (TIMP) 1 (Manoury et al., 2006), and COL-3 messenger RNA (mRNA) expression (Fig. 2 C), confirming activation of the collagen-producing machinery by BLM. Expression of IFN-γ, IL-13, and IL-17A was also monitored from restimulated bronchoalveolar lavage (BAL) cells, lung leukocytes, and local draining LN cells (thoracic LN) to determine whether BLM-induced fibrosis was associated with a Th1, Th2, and/or Th17-type response. Marked increases in IFN-γ, followed by IL-17A, were detected between days 2 and 7 after BLM, whereas significant increases in IL-13 were not observed before day 7. In fact, peak IL-13 production appeared to occur quite late when compared with IFN-γ and IL-17A (Fig. 2 E). The kinetics of these three mediators suggested that BLM-induced inflammatory and fibrotic responses are associated with a Th1/Th17 response at early time points and a mixed Th1/Th17/Th2-type pattern at later times. As such, they illustrate that BLM triggers significant collagen deposition before the development of the Th2 response (Fig. 2, B and E), thus contrasting with the S. mansoni egg–induced fibrosis, which is highly dependent on IL-4 and IL-13 (Chiaramonte et al., 1999).

IL-10 limits IL-17A/IFN–γ production and pulmonary fibrosis

Exogenous IL-10 treatment suppresses BLM-induced fibrosis (Arai et al., 2000; Kitani et al., 2003; Kradin et al., 2004; Nakagome et al., 2006); however, the mechanism of action and role of endogenous IL-10 remains much less clear. Using IL-10 reporter mice, we identified the major source and pattern of expression of IL-10 after BLM challenge. IL-10 was detected in CD4+ and CD4- cells; however, the major increase in IL-10 was observed in the CD4+ T cell population with minor increases in CD8+, CD19+, and CD11c+ cells (unpublished data). The percentage of CD4+ lymphocytes producing IL-10 increased within 2 d after administration of...
BLM and remained elevated in the BAL and lung at all time points (Fig. 3 A). CD4+IL-10+ and CD4+IL-10− cells were FACS sorted from the lung of BLM-treated mice and analyzed for mRNA transcripts. IL-10 gene transcripts were highly up-regulated in CD4+IL-10+ sorted cells, confirming a role for IL-10 in the pathogenesis of BLM-induced pulmonary fibrosis (Fig. 3 B). These observations are also consistent with severe degradation of the basement membrane, enhanced alveolar permeability, and extensive lung injury in IL-10−/− deficient animals.

The exaggerated pathologies observed in il10−/− mice were strongly correlated with increased production of IL-17A and IFN-γ (Fig. 4 B). FACS analysis identified IFN-γ-producing γδTCR+ cells in the lungs 7 d after BLM treatment. A 10-fold increase in the percentage of CD4+IL-10−/− cells was observed in il10−/− mice (Fig. 4 C). In contrast, only a minor increase in the percentage of γδTCR+ cells was noted. Nevertheless, a significant increase in the total number of both CD4+ and γδTCR+ populations in the lung (8.5-fold increase in CD4+ and 2.5-fold increase in γδTCR+) was observed in BLM-treated il10−/− mice, suggesting that IL-10 regulates the development and influx of both CD4+ and γδTCR+ populations.

Analyses of whole lung homogenates revealed significantly elevated levels of several IL-17A–promoting cytokines in BLM treated il10−/− mice. These mediators included IL-6, IL-12/23p40, the neutrophil chemokine KC (CXCL1), and, perhaps most strikingly, IL-1β, which was recently implicated in BLM-induced fibrosis (Gasse et al., 2007) and in amplifying Th17 responses (Sutton et al., 2006, 2009; Fig. 4 D).
IL-10 may, therefore, operate to regulate the development, in addition to the recruitment, of CD4$^+$ and γδ$^+$IL-17A$^+$ cells. In vitro Th17 cell differentiation was not directly influenced by rIL-10 or anti-IL-10R treatment (Fig. S2), indicating that IL-10 does not directly influence Th17 cells, which is consistent with a recent study examining the in vitro differentiation of human Th17 cells (Naundorf et al., 2009).

Neutrophilia is a common feature of pulmonary fibrosis (Hunninghake et al., 1981; Kinder et al., 2008), and IL-17A has been shown to be critically involved in the recruitment of IL-10 may, therefore, operate to regulate the development, in addition to the recruitment, of CD4$^+$ and γδ$^+$IL-17A$^+$ cells. In vitro Th17 cell differentiation was not directly influenced by rIL-10 or anti-IL-10R treatment (Fig. S2), indicating that IL-10 does not directly influence Th17 cells, which is consistent with a recent study examining the in vitro differentiation of human Th17 cells (Naundorf et al., 2009).

Neutrophilia is a common feature of pulmonary fibrosis (Hunninghake et al., 1981; Kinder et al., 2008), and IL-17A has been shown to be critically involved in the recruitment of

Figure 2. IL-17A and IFN-$\gamma$ production during BLM-induced fibrosis. 0.15 U BLM was given to WT mice via intratracheal route, as in Fig. 1, with local immune profiling and assessment of tissue pathology analyzed from days 0 to 21, as indicated. One of three independent experiments is shown, with five animals per group. Data shown are mean ± SEM. (A) Weight loss, assessed at several time points after BLM, as indicated. (B) Pulmonary collagen score from histology sections. (C) RNA was extracted from lung tissue, with timp-1, mmp12, and pro–COL-3 mRNA quantified by quantitative RT-PCR. (D) 5-µm sections of paraffin-embedded lung tissue obtained from mice at the indicated day after BLM and stained with Masson’s Trichrome. Images are shown at a 5x magnification. Bars, 60 µm. (E) BAL, lung, and lung-draining thoracic LN (t:LN) cells were isolated and stimulated with anti-CD3ε for 4 d. IL-17A, IFN-γ, and IL-13 were measured in culture supernatants by ELISA. (F) 5-µm sections of paraffin-embedded lung tissue obtained from mice at day 7 after BLM stained with Giemsa or Picrosirius red (inset) shown under polarized light. Images are shown at 10x magnification. Bars, 60 µm.
neutrophils to sites of inflammation via induction of CXC chemokines (Laan et al., 1999; Witowski et al., 2000; Miyamoto et al., 2003). Consistent with the marked IL-17A response, we detected a significant number of neutrophils in the lungs in BLM-treated mice. Indeed, a substantial increase in neutrophils was observed in both the lung and BAL (not depicted), as well as in the peripheral circulation (Fig. S1 B). The neutrophil response was also exacerbated in il10−/− mice, which was consistent with enhanced IL-17A responses (Fig. 4 B). Collectively, these data indicate that IL-10 limits lung injury, neutrophilia, and pulmonary fibrosis by suppressing proinflammatory mediator release and the production of IL-17A from CD4αβ+ and γδTCR+ cells.

**IL-12/23p40 is required for IL-17A production and pulmonary fibrosis**

IL-12/23p40 promotes the differentiation and expansion of both IL-17A and IFN-γ-producing lymphocytes. Given that all of these mediators were elevated in WT mice after BLM treatment, and to a greater extent in il10−/− mice, we investigated whether IL-12/23p40 was regulating the development of IL-17A–producing cells and pulmonary fibrosis.

![Figure 3. IL-10–producing CD4 cells accumulate in the lung.](image-url)

0.15 U BLM was given to C57BL/6 IL-10<sup>gfp</sup> mice via intratracheal route, as in Fig. 1. One of two independent experiments shown with five animals per group. Data shown are mean ± SEM. (A) Lung, BAL, or thoracic LN cells were isolated from IL-10<sup>gfp</sup> reporter mice and stained with anti–mouse CD3 and CD4. Horizontal bars show the mean. (B) CD4<sup>+</sup> IL-10<sup>gfp−</sup> and CD4<sup>+</sup> IL-10<sup>gfp+</sup> cells were FACS sorted (>98% pure) from the lung of BLM-treated mice at day 7. RNA was extracted and analyzed for mRNA transcripts. The dotted line refers to naive CD4<sup>+</sup> cells.
Using mice deficient in the p40 subunit of IL-12/23, we determined that IL-12/23p40 is critically involved in orchestrating the lung inflammatory response after BLM administration (Fig. 5 A). The reduction in BLM-induced lung damage in p40−/− mice was characterized by significantly decreased collagen in the BAL (Fig. 5 B) and lung (Fig. 5 C), which is consistent with related studies (Maeyama et al., 2001; Huaux et al., 2002; Sakamoto et al., 2002). Circulating neutrophilia followed a similar pattern (Fig. 5 D). More importantly, however, we show that the exacerbated disease in il10−/− mice was highly dependent on IL-12/23p40 production because il10−/−il12/23p40−/− mice were almost completely protected from BLM-induced fibrosis (Fig. 5, B and C). We also extend previous studies by demonstrating mechanistically that deletion of IL-12/23p40 is associated with reduced IFN-γ production (Fig. 5 F) but markedly reduced expression of...
IL-17A (Fig. 5 E). Our data, however, suggested a particularly critical role for IL-17A because only IL-17A was significantly reduced in the double il10−/− il12/23p40−/− mice (Fig. 5 F), correlating with markedly reduced disease (Fig. 5, A and B). Similar results were obtained by ELISA (Fig. 5, E and F) and by intracellular cytokine staining (Fig. 5 G). Nevertheless, previous studies identified an important role for IFN-γ in BLM-induced fibrosis (Chen et al., 2001). Consequently, we revisited this work but investigated whether there was a possible mechanistic link between IFN-γ and IL-17A. Interestingly, we discovered that the reduced fibrotic response in ifnγ−/− mice (Fig. S3, C and D) was associated with decreased IL-17A expression (Fig. S3, B and E) and reduced circulating neutrophils (Fig. S3 A). Together, these data suggest that IFN-γ may facilitate an IL-17A response, as suggested in psoriatic lesions (Kryczek et al., 2008). This notion is also supported by an earlier induction of IFN-γ in the lung compared with IL-17A (Fig. 2 E).

**IL-17A is required for BLM-induced pulmonary fibrosis**

To formally investigate the role of IL-17A in BLM-induced pulmonary fibrosis, WT and il17a−/− mice were challenged with BLM. We also generated il10−/− il17a−/− mice (double KO [dKO]) mice to determine whether the exacerbated lung

---

**Figure 5.** IL-12/23p40 deficiency significantly impacts IL-17A and IFN-γ and curtails pulmonary fibrosis. 0.15 U BLM was given to WT, il10−/−, il12/23p40−/−, or il10−/− il12/23p40−/− mice via the intratracheal route as in Fig. 1. *, P < 0.05 using a Mann-Whitney test. One of two independent experiments is shown, with five animals per group. Data shown are mean ± SEM. (A) 5-µm sections of paraffin-embedded lung tissue taken from WT or il10−/− mice at day 7 or 21 after BLM and stained with Masson’s Trichrome. Images are shown at 5× magnification with dotted squares magnified at 40× in the insets. Bars: 60 µm; (inset) 7 µm. (B) Lung injury, measured as BAL collagen, was quantified from BAL fluid using Sircol assay. (C) Pulmonary collagen deposition, expressed as micromoles of hydroxyproline per lung. (D) Absolute counts of circulating polymorphonuclear cells (PMNs) were obtained from CBC counts. (E and F) Thoracic LN cells were stimulated with anti-CD3ε, with IL-17A (E) and IFN-γ (F) measured in cultures supernatants. (G) Thoracic LN cells were stained with anti–mouse CD4, CD8, B220, γδTCR, NK1.1, IL-17A, and IFN-γ after a brief stimulation with PMA and ionomycin in the presence of BFA. Data shown is gated on CD4+ cells.
fibrosis in IL-10–deficient animals was attributed to the enhanced Th17 response. IL-17A deficiency significantly reduced BLM-induced fibrosis in WT mice, as shown by analysis of collagen expression in the BAL (Fig. 6 A) and collagen deposition in the lung (Fig. 6 B and E). The significant increase in IL-17A in il10−/− mice (Fig. 4, B and C; and Fig. 6 H), which correlated with increased fibrosis, was also reduced to WT levels by deleting IL-17A (Fig. 6, A, B, and E). The reduction in fibrosis observed in il17a−/− mice and il10−/−il17a−/− mice also correlated with a similar decrease in circulating neutrophils (Fig. 6 C) and degree of weight loss (Fig. 6 D), further illustrating an important role for IL-17A in BLM-induced pathology and morbidity. The findings with il10−/−il12/23p40−/− mice and il10−/−il17a−/− dKO mice also point to IL-17A, rather than IFN-γ, as the critical downstream mediator of fibrosis because the il10−/−il12/23p40−/− mice developed markedly reduced IL-17A and fibrotic responses yet maintained near normal IFN-γ production. These observations also suggest that IL-10 inhibits the development of BLM-induced fibrosis by targeting the proinflammatory IL-23–IL-17A pathway rather than the IL-12–IFN-γ axis. IL17f, il22, and tgfα were elevated in the lungs of BLM-treated il10−/− mice and remained elevated in il10−/−il17a−/− mice, even though the dKO animals developed much less fibrosis (Fig. 6 F). Consequently, it seems unlikely that these Th17-associated mediators are contributing to the exaggerated fibrotic phenotype in il10−/− mice. Interestingly, ifnar mRNA (Fig. 6 F) and protein expression (Fig. 6 G) were significantly reduced in il10−/−il17a−/− mice compared with il10−/− mice, suggesting that IL-17A may facilitate the recruitment of IFN-γ-producing cells via lymphocyte-recruiting chemokines (Zrioual et al., 2008), cell survival (Sergeeva et al., 2005), or direct induction of IL-12 and IFN-γ by macrophages (Lin et al., 2009). It is interesting to note that IL-17A and IFN-γ synergistically promote chemokine responses in a variety of inflamed tissues (Albanesi et al., 1999; Andoh et al., 2001; Eid et al., 2009), suggesting a coordinated response between IL-17A and IFN-γ.

In contrast to il17f, il22, and tgfα, we observed a strong correlation between il17b expression and the degree of fibrosis in il10−/− and il10−/−il17a−/− mice. These observations were revealing because IL-1β was recently implicated in BLM-induced fibrosis in both humans and mice (Gasse et al., 1999; Ortiz et al., 2007; Hoshino et al., 2009). il17b expression was also significantly reduced in il17a−/− mice compared with WT animals, suggesting that either IL-17A or IL-17A–dependent responses promotes IL-1β production. IL-1β can indeed promote IL-17A responses (Sutton et al., 2009); however, whether IL-17A directly or indirectly feeds back to enhance IL-1β has not been reported. Expression of several ECM-associated genes, including col3, timp1, and mmp12, was also reduced in il17a−/− mice (Fig. 6 F). Furthermore, using zymography we observed reduced MMP-2 and MMP-9 activity in BAL fluid of il17a−/− mice compared with WT and in il10−/−il17a−/− mice compared with il10−/− mice (Fig. 6 I). Collectively, these data clearly identify IL-17A as a dominant and critical mediator in the pathogenesis of BLM-induced pulmonary fibrosis.

**IL-1β–driven fibrosis is dependent on IL-17A** IL-1β mRNA expression (Fig. 6 F) and cytokine production (Fig. 4 D) correlated with the Th17 response and degree of fibrosis after BLM administration. Several recent studies have also identified an important role for IL-1β in the development of BLM-induced fibrosis. Indeed, administration of IL-1β alone mimics much of the pulmonary pathology caused by BLM (Wang et al., 2000; Gasse et al., 2007; Ortiz et al., 2007; Hoshino et al., 2009). We confirmed and extended these observations by showing that intratracheal administration of IL-1β induces a marked IL-17A response (Fig. 7 C), as well as significant pulmonary fibrosis (Fig. 7, A and B). The administration of 0.5 µg each of both IL-1β and IL-17A had a mild additive effect with increased collagen deposition and, most strikingly, increased MMP2 bioactivity (Fig. 7 D) compared with IL-1β, IL-17A, or BLM treatment alone, suggesting increased turnover of the ECM by IL-1β and IL-17A.

To test the requirement of IL-17A on IL-1β–induced fibrosis, we administered IL-1β to either WT or il17a−/− mice. As observed earlier, BLM- induced fibrosis was significantly reduced in il17a−/− mice (Fig. 6), and, furthermore, IL-1β–mediated inflammation and fibrosis was also significantly reduced, almost to background levels, in the absence of IL-17A (Fig. 7, E and F). As observed with BLM, intratracheal IL-1β induced significant neutrophilia (Fig. 7 G), TIMP-1 in the BAL (Fig. 7 H), and IFN-γ and IL-17A production in the local LNs (Fig. 7 I). IFN-γ was only slightly elevated in the BAL after IL-1β treatment (Fig. 7 J). These mediators were all reduced in IL-1β–treated il17a−/− mice, indicating that IL-1β–induced pulmonary disease, like BLM, is dependent on IL-17A for airway inflammation and pulmonary fibrosis.

**IL-17A–dependent pulmonary fibrosis requires TGF-β** Many studies have identified a profibrotic role for TGF-β, particularly in the pathogenesis of pulmonary fibrosis (Cutroneo et al., 2007). More recently, studies have also suggested important inflammatory activities for TGF-β, with TGF-β providing an integral differentiation signal for the development of proinflammatory IL-17A–secreting cells (Bettelli et al., 2006; Veldhoen et al., 2006). We therefore tested whether the profibrotic properties of TGF-β were linked to IL-17A. Intratracheal delivery of BLM or IL-17A induced significant collagen release in the BAL (Fig. 8 A) and deposition in the lung (Fig. 8, B and D). TGF-β blockade significantly reduced BLM and IL-17A–induced fibrosis (Fig. 8, B and D), identifying a critical requirement for TGF-β in both BLM and IL-17A–induced fibrosis. rIL-17A and BLM both increased IL-17A responses in the local LNs (Fig. 8 C), and TGF-β blockade curtailed endogenous IL-17A production, indicating that TGF-β is required for the genesis of endogenous IL-17A. Together, these findings suggest cooperative and amplifying roles TGF-β and IL-17A in the development of fibrosis.
Figure 6. Attenuated pulmonary fibrosis in \( \text{il17a}^{-/-} \) mice. 0.15 U BLM was given to WT, \( \text{il10}^{-/-} \), \( \text{il17a}^{-/-} \), or \( \text{il10}^{-/-}\text{il17a}^{-/-} \) mice, via the intratracheal route, as in Fig. 1. *, \( P < 0.05 \) using a Mann-Whitney test. One of two independent experiments is shown, with five animals per group. Data shown are mean ± SEM. (A) Lung injury, measured as BAL collagen, was quantified from BAL fluid using Sircol assay. (B) Pulmonary collagen deposition, expressed as micromoles of hydroxyproline per lung. (C) Absolute counts of circulating polymorphonuclear cells were obtained from CBC counts. (D) Percentage of weight change, 7 d after BLM. (E) 5-µm sections of paraffin-embedded lung tissue taken from WT or \( \text{il10}^{-/-} \) mice at day 7 or 21 after BLM and stained with Masson’s Trichrome. Images are shown at a 40× magnification. Bar, 7 µm. (F) RNA was extracted from lung tissue, with indicated mRNA quantified by quantitative RT-PCR. (G and H) Thoracic LN cells were stimulated with anti-CD3ε, with IL-17A (G) and IFN-γ (H) measured in culture supernatants. (I) MMP2 and MMP9 bioactivity measured in BAL by zymography.
Figure 7. IL-17A–dependent IL-1β–induced collagen deposition. Mice were given intratracheal 0.15 U BLM, 1 µg IL-1β, 1 µg IL-17A, or 0.5 µg each of both IL-1β and IL-17A with pulmonary collagen deposition assessed on day 7. *, P < 0.05 using a Mann-Whitney test. One of two independent experiments is shown, with five animals per group. Data shown are mean ± SEM. (A and E) 5-µm sections of paraffin-embedded lung tissue were stained with Masson’s Trichrome. Images are shown at a 20× magnification. Bars, 20 µm. (B and F) Pulmonary collagen deposition, expressed as micromoles of hydroxyproline per lung. (C and I) Thoracic LN cells were stimulated with anti-CD3ε, with IL-17A and IFN-γ measured in culture supernatants. (D) MMP2 bioactivity in BAL fluid measured by zymography. (G) Absolute counts of circulating polymorphonuclear cells were obtained from CBC counts. (H) BAL fluid TIMP-1 and IFN-γ was measured by ELISA.
IL-13–dependent pulmonary fibrosis is IL-17A independent

To determine whether IL-17A was functioning as a master regulator of fibrosis, we also exposed il17a−/− and il10−/− il17a−/− mice to another model of pulmonary fibrosis. Previous studies have shown that S. mansoni egg–induced pulmonary fibrosis is highly dependent on IL-13 signaling (Ramalingam et al., 2008). However, IL-17A has been implicated in IL-13–associated airway hyperreactivity (Nakae et al., 2002). The possible connection between IL-17A expression and Th2-dependent fibrosis, however, has not been previously investigated (Wynn, 2008). We therefore assessed the impact of IL-17A on IL-13–dependant pulmonary fibrosis using the S. mansoni egg–induced pulmonary granuloma model. CD4+IL-17A+ (1.1%), CD4+IFN-γ+ (13.1%), and CD4+IL-13+ (22.5%) lymphocytes were observed in WT mice 7 d after i.v. delivery of S. mansoni eggs to the lungs (Fig. 4A), suggesting a potential role for IL-17A in the development of fibrosis in this model. Deletion of il10 resulted in an approximate doubling of IL-17A–producing cells (1.1–2.5%) and a modest increase in IFN-γ–producing cells (13.1–16.3%). Similarly, when whole lung tissue was analyzed by quantitative RT-PCR, significant increases in il17a and ifnγ mRNAs were observed, with only modest changes in il13 (Fig. S4, A and E). To test the importance of IL-17A in this model, WT, il17a−/−, il10−/−, and il10−/− il17a−/− mice were challenged i.v. with S. mansoni eggs. In marked contrast to the BLM model (Fig. 6), BAL collagen (Fig. S4 B), hydroxyproline measurements (Fig. S4 C), and microscopic assessments of lung fibrosis (Fig. S4, D and G) revealed little to no role for IL-17A. Therefore, although IL-17A was significantly upregulated and increased further in the absence of IL-10 (Fig. S4, A and E), IL-13 appears to function as the dominant profibrotic mediator in this model. Production of IL-13 was not significantly affected by the presence or absence of IL-17A, suggesting no obvious regulatory role for IL-17A (Fig. S4, E and F). In fact little modulation of the ECM–related genes (mmp12, timp1, and col6a) was observed in the absence of IL-17A (Fig. S4 E). Moreover, although there was a modest increase in IFN-γ in the IL-10–deficient groups, it had no significant impact on the development of IL-13–dependent fibrosis, perhaps because IL-13 was upregulated ~20-fold more than IFN-γ. Thus, in marked contrast to the BLM model, where IL-17A and, to a lesser extent, IFN-γ appeared to play critical roles in the development of fibrosis, pulmonary fibrosis induced by schistosome eggs appeared to be dependent on Th2 cytokines. Thus, unique cytokine–dependent mechanisms can be exploited to induce pulmonary fibrosis, with IL-17A playing a critical role in BLM–induced fibrosis.

Increased IL-1β and IL-17A in BAL fluid of IPF patients

To determine whether IL-1β and IL-17A are involved in human IPF, we obtained lung biopsies and BAL fluid from normal volunteers (NVs) and IPF patients. Masson’s trichrome–stained lung sections revealed inflammatory foci within the parenchyma, surrounded by dense collagen deposits (Fig. 9 B). Similar to the mouse model, IL-17A (NV, 1.48 ± 0.98 pg/ml;
IPF, 14.31 ± 3.76 pg/ml) and IL-1β (NV, 9.58 ± 1.58 pg/ml; IPF, 21.36 ± 3.97 pg/ml), but not IL-22 (NV, 56.1 ± 2.00 pg/ml; IPF, 50.7 ± 1.47 pg/ml; Whittington et al., 2004), in BAL fluid of IPF patients were increased (Fig. 9A), indicating that an IL-1β–IL-17A pathway may be involved in the development of human IPF.

**DISCUSSION**

We have demonstrated that BLM-induced, but not *S. mansoni* egg-induced, pulmonary fibrosis is dependent on IL-17A produced by γδ+ and CD4+αβ+ T cells. In addition, we show that IL-17A is required for the development of inflammation, neutrophilia, and pulmonary fibrosis after exposure to IL-1β, a recently described initiator of fibrosis (Wang et al., 2000; Ortiz et al., 2007; Gasse et al., 2007; Cassel et al., 2008; Hoshino et al., 2009). Mechanistically, using *il10*−/−, *il10*−/− *il17a*−/−, and *il10*−/−*il12p40*−/− dKO mice and TGF-β blockade, we also show that fibrosis is tightly controlled by IL-10–producing CD4+ lymphocytes, which regulate IL-17A production. Together, these studies identify IL-17A as a critical mediator of pulmonary fibrosis after BLM administration, illustrating the potential utility of targeting IL-17A in the treatment of IPF and other fibrotic diseases.

Previous studies have identified important roles for TGF-β1 and IL-13 in the development of fibrosis in a variety of tissues (Chiaramonte et al., 1999; Lee et al., 2001; Varga and Pasche, 2008), whereas a role for IL-17A has remained unclear. Some studies found that IL-13 induces and activates TGF-β1 (Lee et al., 2001), which then serves as the primary mediator of fibrosis by stimulating collagen synthesis in fibroblasts (Czaja et al., 1989). Nevertheless, other studies demonstrated that IL-13 could function independently of TGF-β1 (Kaviratne et al., 2004). To better characterize both the unique and convergent pathways of fibrosis, we dissected the mechanisms of pulmonary fibrosis in two distinct models in which TGF-β1 and IL-13 have been shown to function as critical mediators. Unexpectedly, although small interfering RNA gene silencing studies implicated IL-13 in the mechanism of BLM-induced pulmonary fibrosis (Fichtner-Feigl et al., 2006), our studies conducted with *C57BL/6 il13−/−* and *il13Rα2−/−* mice failed to reveal a significant role for IL-13 signaling at this early stage of fibrosis in this model. In contrast, *S. mansoni*...
egg–induced pulmonary fibrosis was completely IL-13 dependent. Together, these studies indicate that pulmonary remodeling and fibrosis can be induced by distinct nonoverlapping mechanisms, with IL-13 functioning as the key driver of *S. mansoni* egg–driven fibrosis (Kaviratne et al., 2004) and IL-17A serving as a key mediator of BLM-induced fibrosis. The discovery that distinct immunological mechanisms can regulate fibrosis suggests that the etiology and/or route of the initial insult, as well as the character of the subsequent inflammatory response, may have a significant bearing on the nature of the lung wound healing response. The distinct route of tissue damage observed in both models may contribute to the unique fibrotic mechanisms used. Indeed, *S. mansoni* eggs delivered i.v. primarily disrupt endothelial cells lining the pulmonary vasculature, whereas BLM administered intratracheally is directly toxic to epithelial cells lining the airway. *S. mansoni* eggs are also highly immunogenic, inducing robust antigen–specific CD4+ Th2 responses (Everts et al., 2009; Steinfelder et al., 2009), whereas BLM induces rapid epithelial cell death and destruction of DNA (Hay et al., 1991), leading to acute proinflammatory Th1/Th17-type cytokine production.

Most studies of BLM-induced fibrosis have focused on the hypothesis that TGF-β regulates the development of pulmonary fibrosis by directly promoting the differentiation and activation of collagen-producing myofibroblasts (Varga and Pasche, 2008). However, recent studies demonstrated that TGF-β is also critically involved in the development of IL-17A–producing lymphocytes (Bettelli et al., 2006; Veldhoen et al., 2001), which regulate the development of a variety of inflammatory and autoimmune diseases (Tesmer et al., 2008). Using *il17a−/−* mice, we show that IL-17A is essential for the development of BLM-induced fibrosis. Similarly, blockade of TGF-β also significantly reduces BLM and IL-17A–induced fibrosis and correlates with reduced IL-17A production. These findings suggest that the profibrotic activity of TGF-β may, at least in part, be attributed to the induction of IL-17A. Therefore, by stimulating myofibroblast activation and the production of IL-17A by T cells, TGF-β likely promotes BLM-induced fibrosis through both direct and indirect mechanisms. In contrast, development of *S. mansoni* egg–induced pulmonary fibrosis was completely IL-17A independent. Therefore, IL-17A appears to induce fibrosis in models where TGF-β, but not IL-13, has been shown to play critical roles. IL-17A can directly induce the collagenase MMP-1 (Cortez et al., 2007) and progelatinase, MMP-3 (Beklen et al., 2007) from various fibroblasts, suggesting that IL-17A may facilitate tissue disruption, in addition to its ability to promote granulopoiesis (Schwarzenberger et al., 1998) and inflammation. To this end, many studies have identified the proinflammatory properties of IL-17A. In the context of inflammation and pulmonary fibrosis, IL-17A can stimulate IL-6, IL-8, and MCP-1 from bronchial epithelial cells (Luan et al., 2001) or fibroblasts (Hata et al., 2002; Mahanonda et al., 2008) and, as mentioned in the previous paragraph, is inducible by IL-1β and IL-23, placing IL-17A in a central position, magnifying the inflammatory cascade. One previous study suggested that IL-17A producing γδ+ T cells regulate cellular recruitment with γδ+ T cell–deficient mice developing increased inflammation and collagen deposition (Braun et al., 2008). These data suggest that γδ+ T cells may be heterogeneous and responsible for more than IL-17A, with populations of γδ+ T cells providing a brake on the inflammatory response in addition to the early IL-17A production. A similar scenario could be envisioned for CD4+ T cells, providing both proinflammatory and fibrotic IL-17A in addition to antinflammatory IL-10.

Several studies have identified an important regulatory role for IL-10 in BLM-induced pulmonary fibrosis (Kradin et al., 2004; Nakagome et al., 2006). These studies suggested that IL-10 controls BLM-induced inflammatory and fibrotic response by regulating chemokine and TGF-β1 production (Kradin et al., 2004; Nakagome et al., 2006). However, we also found a marked increase in IL-17A in *il10−/−* mice, which develop markedly exacerbated inflammation, neutrophil mobilization, and pulmonary fibrosis after BLM administration. These observations are consistent with studies that suggested IL-10 could regulate the magnitude and effector function of Th17 responses (Fitzgerald et al., 2007; McGeachy et al., 2007; Chang et al., 2008). Similar to previous studies (McGeachy et al., 2007; Stunhofer et al., 2007), IL-10 was often coexpressed with IL-17A and RoRγt, suggesting that Th17 cells may develop into IL-10–secreting cells to limit Th17 effector function.

To determine whether there was a functional connection between IL-10 and IL-17A, we generated *il10−/−*/*il17a−/−* dKO mice and then challenged the animals with BLM. Strikingly, the increase in BLM-induced pulmonary fibrosis observed in *il10−/−* mice was completely reversed in the absence of IL-17A. As such, these studies explain the severe inflammatory response observed in *il10−/−* mice (Kradin et al., 2004). They also identify IL-17A, which is produced predominantly by CD4+ T cells, as a critical target of IL-10. Gene transcripts for the IL-17A–related molecules IL-17F and IL-22 were also increased in the absence of IL-10, correlating with exaggerated inflammation and fibrosis (Fig. 6). However, these molecules were not markedly affected in *il17a−/−*/*il10−/−* mice despite a significant reduction in pulmonary fibrosis, suggesting that IL-17A, but not IL-17F or IL-22, functions as critical mediators of pulmonary inflammation and fibrosis at this stage. Furthermore, IL-22 levels were not significantly different between control and pulmonary fibrosis patients (Whittington et al., 2004). A comprehensive study by Ishigame et al. (2009) highlighted the overlapping but also distinct roles of IL-17A and IL-17F. For example, IL-17A, but not IL-17F, was required for DTH, autoimmune encephalomyelitis, and collagen-induced arthritis. However, IL-17A and IL-17F were both required for host defense against *Staphylococcus aureus* and *Citrobacter rodentium*. Functionally, IL-17F and IL-17A also appear to have opposing roles in the allergic lung, with IL-17F negatively regulating and IL-17A positively regulating Th2-mediated inflammation (Yang et al., 2008). From our data, it
appears that IL-17A, and not IL-17F or IL-22, is the dominant molecule involved in BLM and IL-1β-mediated pulmonary inflammation and fibrosis.

Previous studies identified a role for IFN-γ in BLM-induced fibrosis (Chen et al., 2001), which was consistent with our findings. Surprisingly, we found that IFN-γ deficiency was associated with reduced IL-17A production, suggesting a link between IFN-γ and IL-17A. BLM studies performed with il12p40−/− and il10−/− il12p40−/− dKO mice confirmed this association; however, they indicated that IL-17A was functioning as the dominant inducer of fibrosis. Indeed, reduced BLM-induced fibrosis in il12/23p40−/− mice was associated with a much greater decrease in IL-17A than IFN-γ. A similar pattern was observed in BLM-treated il10−/− il12p40−/− dKO mice. il17a−/− and il10−/− il17a−/− dKO mice also developed relatively normal IFN-γ responses, despite significantly reduced fibrosis. IFN-γ responses in the lungs preceed IL-17A responses (Fig. 2 E), suggesting that the IL-12–IFN-γ axis may be involved in the recruitment of IL-23-dependent IL-17A–producing cells into the lung, as previously reported (Kryczek et al., 2008). Nevertheless, these findings strongly suggest that the IL-23–IL-17A axis, rather than the IL-12–IFN-γ pathway, plays the dominant role in BLM-induced fibrosis, although both mechanisms are clearly involved and connected.

In addition to TGF-β1 and IL-23 (Langrish et al., 2005), recent studies identified an important role for IL-1β in Th17 responses (Sutton et al., 2006). Interestingly, a recent study showed that IL-1R signaling is involved in BLM-induced pulmonary fibrosis, with exogenous IL-1β treatment recapitulating much of the lung pathology seen with BLM (Gasse et al., 2007). We confirmed and extended these observations by showing that IL-1β–mediated inflammation and fibrosis is dependent on IL-17A. These findings suggest that IL-17A functions as a critical downstream mediator of fibrosis. Interestingly, the reduced inflammation and fibrosis observed in BLM-treated il17a−/− animals was associated with reduced IL-1β expression, suggesting that IL-17A promotes IL-1β production, as has been observed in vitro with cultured fibroblasts (Beklen et al., 2007). These data suggest that IL-17A and IL-1β cross-regulate each other, thus providing an explanation for their additive roles in BLM-induced fibrosis.

BLM-induced fibrosis is associated with significant neutrophil recruitment to the lung. Indeed neutrophilia is a common feature of fibrosis and pulmonary alveolitis in IPF patients (Hunninghake et al., 1981; Kinder et al., 2008), with airway neutrophilia identified as a predictor of early mortality in IPF (Kinder et al., 2008). Interestingly, it is now well appreciated that IL-17A is involved in the recruitment of neutrophils to sites of inflammation via induction of IL-8 and CXC chemokines, including MIP-2 (Laan et al., 1999; Witowski et al., 2000; Miyamoto et al., 2003; Lindén et al., 2005). Nevertheless, a direct link between IL-17A, neutrophilia, and IPF has not been previously investigated. In our study, we observed a strong correlation between IL-17A and neutrophil recruitment, with il17a−/− and il10−/− il17a−/− mice displaying significantly fewer neutrophils in the lung than their respective control groups. Finally, IL-17A and IL-1β were also detected in the BAL fluid of IPF patients. When viewed together, these observations make a compelling case for IL-17A in the pathogenesis of pulmonary inflammation and fibrosis.

In conclusion, genetic deletion of IL-17A significantly attenuated lung inflammation, neutrophilia, and fibrosis induced by BLM treatment in both WT and IL-10–deficient mice. In contrast, IL-17A appeared to play little to no role in the development of IL-13–dependent fibrosis. As such, these data identify distinct nonoverlapping immunological roles for IL-13 and IL-17A in the development of fibrosis. They also suggest that the IL-17A pathway might provide a novel therapeutic target for the treatment of pulmonary fibrosis, for which few therapeutic options currently exist.

MATERIALS AND METHODS

Animals. Female 6–10-wk-old C57BL/6 (WT), il10−/−, il12/23p40−/−, gfi1−/−, d133−/−, d13Ra2−/−, il10−/− il12/23p40−/−, and OVA-specific OT-Il [C57BL/6-Tg(TCR-α TCR-β)] (all C57BL/6 background for ≥10 generations) were obtained from National Institute of Allergy and Infectious Disease (NIAID) animal facilities at Taconic. IL-10−/− reporter mice were provided by R. Flavell (Yale University, New Haven, CT; Kamanaka et al., 2006). il17a−/− animals were generated and provided by Regeneron (Leppkes et al., 2009), il10−/− il17a−/− animals were generated by crossing il10−/− with il17a−/−, with gene expression and protein production confirmed by PCR and ELISA. All animals were housed under specific pathogen-free conditions at the National Institutes of Health in an American Association for the Accreditation of Laboratory Animal Care–approved facility. The NIAID animal care and use committee approved all experimental procedures. A minimum of five mice per group were used in each experiment, unless indicated.

Human tissues. Paraffin-embedded lung sections and BAL fluid were obtained from 14 NVs and 7 patients diagnosed with IPF. 5-µm lung sections were stained with Masson’s trichrome as described in Histopathology and fibrosis. Subjects were enrolled in protocols 99-HG-0056 and/or 04-HG-0211, which were approved by the National Human Genome Research Institute Institutional Review Board. Written informed consent was obtained from all subjects. Bronchoscopy with BAL was performed as previously described (Ren et al., 2007).

Reagents and cell culture. For in vitro cell culture, LN cells were isolated, washed, and plated at 5 × 105 cells per well of a 96-well plate and cultured with gene expression and protein production confirmed by PCR and ELISA. Briefly, animals were generated by crossing il10−/− with il17a−/−, with gene expression and protein production confirmed by PCR and ELISA. All animals were housed under specific pathogen-free conditions at the National Institutes of Health in an American Association for the Accreditation of Laboratory Animal Care–approved facility. The NIAID animal care and use committee approved all experimental procedures. A minimum of five mice per group were used in each experiment, unless indicated.

Complete blood count (CBC) analysis. EDTA-treated blood was processed for automated counting using a Vista Analyzer (Siemens).

Pulmonary fibrosis models. For Schistosoma mansoni egg–induced pulmonary fibrosis, mice were immunized with 5,000 S. mansoni eggs by i.p. injection. Tail vein injection of 5,000 eggs was given on day 14, with analysis of fibrotic granuloma development 7 d later on day 21. For BLM-induced pulmonary fibrosis, mice were anesthetized with a xylazine and ketamine...
cocktail and given 0.15 U BLM sulfate (EMD) in saline via the intratracheal route. Weight change, pulmonary inflammation, and fibrosis were assessed daily or 7 d later, as designated in figure legends. 1 µg IL-1β (R&D Systems) or 1 µg IL-17A (R&D Systems) was administered by the intratracheal route to anesthetized mice in the same way as BLM.

BAL fluid and differential cell counts. Mice were terminally anesthetized with sodium pentobarbital. The trachea was cannulated and pulmonary airspaces lavaged with an initial 500 µl of sterile PBS, followed by two 500-µl PBS washes. Fluids were centrifuged at 1,200 g, and pellets recovered from all three lavage washes were pooled into 1 ml PBS for total BAL cell counts and cellular analysis. The supernatants of the initial 500 µl BAL fluid were stored at -80°C for biochemical analyses. BAL collagen was measured using a Sircol assay, according to the manufacturer’s recommendations.

ELISA. Cytokines, chemokines, and TIMP-1 were measured by ELISA using Immulon 2HB plates (Thermo Fisher Scientific) and the manufacturer's guidelines. Paired capture and detection antibodies (R&D Systems) for IL-17A (human and mouse), IFN-γ, IL-13, IL-10, TIMP-1, CXCCL1 (KC), IL-6, IL-12/23p40, and IL-1 (human) were used. Plates were washed with 0.05% Tween 20 in PBS (PBST) and blocked with 5% milk in PBST. Recombinant cytokine standards (R&D Systems) were used to assess quantity using a standard curve, with OD acquired at 405 nm in an ELISA reader.

Flow cytometry. After a 3-h incubation with 10 µg/ml M. pneumoniae (BFA), cells were stained with antibodies diluted in PBS with 0.05% sodium azide (Sigma-Aldrich) for 20 min at 4°C. Surface molecule staining (CD4 [BD], CDF4 [BioLegend], B220 [BD], CD19 [BD], and γδ [BD]), followed by fixation and permeabilization (Cytofix/Cytoperm; BD) and intracellular cytokine staining (IL-17A [BD], IFN-γ [BD], IL-10 [BD], IL-13 [Contecor], and IL-22 [provided by Wtyeh; Ma et al., 2008]), was performed on freshly isolated cells. IL-10gp cells were detected ex vivo, without stimulation, and were only stained for surface molecules. The expression of surface molecules and intracellular cytokines was analyzed on a flow cytometer (LSR II; BD) using Flowjo software (v.8; Tree Star, Inc.).

Histopathology and fibrosis. Pulmonary collagen was measured as hydroxyproline after hydrolysis of a known weight of lung tissue in 5 ml of 6N HCl at 110°C for 18 h. The increase in pulmonary hydroxyproline per mouse was calculated based upon total lung weight and expressed as micrograms of hydroxyproline per milligram of lung tissue.

Statistical analysis. Datasets were compared by a Mann Whitney test or one-way analysis of variance, as specified in the figure legends, using Prism software v5. Differences were considered significant (*) at P < 0.05.

Online supplemental material. Fig. S1 demonstrates increased pulmonary inflammation and circulating neutrophilia in BLM-treated iIL-10−/− mice, compared with BLM-treated WT mice. Fig. S2 shows that neither recombinant IL-10 nor anti–IL-10R blockade has a direct impact on Th17 cell generation in vitro. Fig. S3 corroborates other studies and demonstrates, using iIfnγ−/− mice, that IFN-γ contributes to BLM-induced pulmonary inflammation and fibrosis. Fig. S4 shows that IL-17A is not required for S. mansoni egg-induced pulmonary inflammation and fibrosis. Online supplemental material is available at http://www.jem.org/cgi/content/full/jem.20092121/DC1.

We are very grateful to the Intramural Research Program of the National Institutes of Health, National Institute of Allergy and Infectious Disease, and National Human Genome Research Institute, who supported this research.

The authors acknowledge the meticulous care of animals used in this study by Nicole Reiferf, Joy McFarlane, Jose Encarnacion, Lauren Donato, and SoBrann staff. We thank Sandra D. White, Robert W. Thompson, and clinical center staff for additional animal care, technical assistance, and CBC processing. We also thank Drs. Margaret Karow, Andrew J. Murphy, David M. Valenzuela, and George D. Paty. We thank Sandra D. White, Robert W. Thompson, and clinical center staff for additional animal care, technical assistance, and CBC processing. We also thank Drs. Margaret Karow, Andrew J. Murphy, David M. Valenzuela, and George D. Paty.

The authors have no competing financial interests.

Submitted: 30 September 2009
Accepted: 1 February 2010

REFERENCES


doi:10.1046/j.1365-2249.2001.01588.x


