

The regulatory role of interferon- γ producing gamma delta T cells via the suppression of T helper 17 cell activity in bleomycin-induced pulmonary fibrosis

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Introduction

Interstitial pneumonia (IP) is a chronic progressive interstitial lung disease associated with poor prognosis and high mortality [1]. IP is an intractable disease induced by various factors such as autoimmune diseases, drugs, exposure to certain occupational hazards and environmental factors [2]. For example, the chemotherapy with bleomycin and busulphan is reported to cause lung fibrosis in some patients [3]. Following the interstitial inflammation, fibroblast proliferation within both the interstitium and alveolar space is often observed. However, the pathogenesis of IP remains to be elucidated.

In mice, pulmonary fibrosis can be induced by the administration of bleomycin, silica particles or *Bacillus subtilis*. Bleomycin-induced IP mouse is a typical murine model of pulmonary fibrosis [4]. In this model, exposure to bleomycin induces an acute inflammatory response that lasts up to 8 days and is followed by fibrogenic changes,

Summary

Interstitial pneumonia (IP) is a chronic progressive interstitial lung disease associated with poor prognosis and high mortality. However, the pathogenesis of IP remains to be elucidated. The aim of this study was to clarify the role of pulmonary $\gamma\delta$ T cells in IP. In wild-type (WT) mice exposed to bleomycin, pulmonary $\gamma\delta$ T cells were expanded and produced large amounts of interferon (IFN)- γ and interleukin (IL)-17A. Histological and biochemical analyses showed that bleomycin-induced IP was more severe in T cell receptor (TCR)- δ -deficient (TCR $\delta^{-/-}$) mice than WT mice. In TCR $\delta^{-/-}$ mice, pulmonary IL-17A $^{+}$ CD4 $^{+}$ T cells expanded at days 7 and 14 after bleomycin exposure. In TCR $\delta^{-/-}$ mice infused with $\gamma\delta$ T cells from WT mice, the number of pulmonary IL-17A $^{+}$ CD4 $^{+}$ T cells was lower than in TCR $\delta^{-/-}$ mice. The examination of IL-17A $^{-/-}$ TCR $\delta^{-/-}$ mice indicated that $\gamma\delta$ T cells suppressed pulmonary fibrosis through the suppression of IL-17A $^{+}$ CD4 $^{+}$ T cells. The differentiation of T helper (Th)17 cells was determined *in vitro*, and CD4 $^{+}$ cells isolated from TCR $\delta^{-/-}$ mice showed normal differentiation of Th17 cells compared with WT mice. Th17 cell differentiation was suppressed in the presence of IFN- γ producing $\gamma\delta$ T cells *in vitro*. Pulmonary fibrosis was attenuated by IFN- γ -producing $\gamma\delta$ T cells through the suppression of pulmonary IL-17A $^{+}$ CD4 $^{+}$ T cells. These results suggested that pulmonary $\gamma\delta$ T cells seem to play a regulatory role in the development of bleomycin-induced IP mouse model via the suppression of IL-17A production.

Keywords: gamma delta T cell, IL-17A, interferon- γ , interstitial pneumonia

resulting in the deposition of collagen and distortion of the lung structure within 28 days. Thus, the acute inflammation phase switches to the fibrosis phase at approximately day 10 after bleomycin exposure.

Recently, the functional role of gamma delta ($\gamma\delta$) T cells was reported in several IP mouse models [5–9]. The $\gamma\delta$ T cells can produce a wide variety of cytokines, chemokines and growth factors and play an important role in the regulation of the initial immune response to several pathogens by influencing the migration and activity of neutrophils, macrophages, natural killer (NK) and T helper (Th) cells [10]. In fact, $\gamma\delta$ T cells were considered to have regulatory roles in bleomycin-induced IP [5,11]. However, the regulatory function of $\gamma\delta$ T cells in the development of pulmonary fibrosis is not yet clear. Previous studies showed that $\gamma\delta$ T cells were able to produce both interferon (IFN)- γ and interleukin (IL)-17A [12]. IFN- γ is reported to inhibit fibroblast proliferation and production of collagen *in vitro*

as well as the development of pulmonary fibrosis *in vivo* [13–18]. Conversely, IL-17A promotes fibroblast proliferation and production of several fibrotic factors *in vitro* [19]. IL-17A-deficient mice attenuated bleomycin-induced airway inflammation and pulmonary fibrosis [20,21]. Thus, the roles of IFN- γ and IL-17A are thought to have opposite effects in the progression of pulmonary fibrosis.

In humans, several reports showed that $\gamma\delta$ T cells play a role in the development of pulmonary fibrosis [22,23]. We also reported that CD161-expressing $\gamma\delta$ T cells play a regulatory role in IP in patients with systemic sclerosis [24]. To our knowledge, however, there is only little or no information about the properties and functional characteristics of $\gamma\delta$ T cells in pulmonary fibrosis.

The present study was designed to clarify the regulatory role of $\gamma\delta$ T cells in the development of bleomycin-induced IP, with a special focus on the roles of IFN- γ and IL-17A.

Materials and methods

Mice and exposure to bleomycin

C57BL/6 [wild-type (WT)] mice were purchased from Charles River Japan Inc. (Tokyo, Japan). T cell receptor (TCR) δ -deficient (TCR $\delta^{-/-}$) mice [25] with a C57BL/6 background were provided by Riken BRC, a participant in the National Bio-Resource Project of the MEXT, Japan. IFN- γ -deficient (IFN- $\gamma^{-/-}$) mice were obtained from The Jackson Laboratory. IL-17A-deficient (IL-17A $^{-/-}$) mice were kindly provided by Professor Y. Iwakura (Tokyo University of Science). Only female mice were used in this study. The animals were kept under specific pathogen-free conditions and studied at 8–10 weeks of age. The Committee on Institutional Animal Care and Use at Tsukuba University approved all the experimental protocols. Bleomycin (1.25 mg/kg; Nippon Kayaku, Tokyo, Japan) dissolved in phosphate-buffered saline (PBS) was administered intratracheally in this study. Briefly, mice were sedated with isoflurane, attached to a tilting table. Then, the trachea was cannulated with auriscope (WelchAllyn, Skaneateles Falls, NY, USA) and bleomycin was infused directly into the lungs.

Staining and flow cytometry

Pulmonary lymphocytes were isolated from the lungs using the following method. The lungs were perfused thoroughly with PBS to remove circulating blood cells. The dissected lungs were incubated with 1% collagenase D (Roche, Basel, Switzerland) in RPMI-1640 medium containing 10% fetal bovine serum (FBS; BioWest, Miami, FL, USA), 100 units/ml penicillin, 100 μ g/ml streptomycin and 50 μ M 2-mercaptoethanol at 37°C for 1 h. Then, the lungs were minced and passed through nylon mesh to remove debris. The washed and recovered cells were subjected to Percoll

PLUS (GE Healthcare, Uppsala, Sweden) at 800 \times g at room temperature for 20 min. The resultant interface containing pulmonary lymphocytes was recovered and washed with RPMI-1640 medium. Cells were stained with the following monoclonal antibodies (mAbs): fluorescein isothiocyanate (FITC)-conjugated anti-TCR $\gamma\delta$ (clone: GL3), anti-CD11c (N418), anti-TCR β (H57-593), R-phycoerythrin (PE)-conjugated anti-Gr-1 (RB6-8C5), anti-TCR $\gamma\delta$ (GL3), allophycocyanin (APC)-conjugated anti-CD3 ϵ (145-2C11), anti-CD11b (M1/70) and peridinin chlorophyll (PerCP)/cyanin (Cy)5.5-conjugated CD4 (GK1.5) mAbs [all from Biolegend (San Diego, CA, USA)]. Intracellular staining for forkhead box protein 3 (FoxP3) was performed after fixation and permeabilization according to the protocol supplied by the manufacturer (eBioscience, San Diego, CA, USA). Dead cells were stained with Fixable Viability Dye eFluor 780 (eBioscience). The stained cells were analysed on fluorescence activated cell sorter (FACS) Verse flow cytometer (Becton Dickinson, Mountain View, CA, USA) and data were processed using FlowJo software (TreeStar, Ashland, OR, USA).

Intracellular cytokine staining

Pulmonary lymphocytes from WT and TCR $\delta^{-/-}$ mice were prepared as described above. Cells were stimulated with phorbol 12-myristate 13-acetate (PMA, 50 ng/ml; Sigma, St Louis, MO, USA), ionomycin (1 μ g/ml; Sigma) and GolgiStop (BD Pharmingen, Franklin Lakes, NJ, USA) in 96-well round-bottomed plates for 6 h. Cells were first stained for a cell surface marker, then fixed and permeabilized with Cytofix/Cytoperm solution (BD Pharmingen), before further immunostaining with PerCP/Cy5.5-conjugated anti-IFN- γ (XMG1.2) and PE/Cy7-conjugated IL-17A (TC11-18H10.1). Cells were analysed with FACS Verse flow cytometer (Becton Dickinson) and data were analysed with FlowJo software (TreeStar).

Histological examination

The lung tissues were perfused with formalin. The trachea was cannulated, and formalin (0.5 ml) was infused into the lungs. Fixed lung tissues were removed and embedded in paraffin. Sections (4 μ m) were stained with Masson's trichrome.

Quantitative image analysis (QIA)

The percentage area of fibrosis was quantified as described previously [4]. Briefly, five randomly fields per slide were selected and detected blue-stained collagen within each field. The fraction of collagen areas for each field was averaged for each animal.

Bronchoalveolar lavage fluid (BALF) analysis

The trachea was cannulated, and the lung was lavaged five times with 0.8 ml of cold PBS each. The BALF was centrifuged at $800 \times g$ for 5 min at 4°C.

Determination of soluble collagen by Sircol assay

The left lungs were minced and incubated with 1 ml of pepsin (0.1 mg/ml in 0.5 M acetic acid) at 4°C. After 24 h, samples were centrifuged at $10\,000 \times g$ and these supernatant recovered. Collagen content of the left lung and BALF was performed by Sircol assay according to the protocol supplied by the manufacturer (Biocolor, County Antrim, UK).

Cell proliferation and adoptive transfer of $\gamma\delta$ T cells

Pulmonary lymphocytes were harvested from WT and IFN- $\gamma^{-/-}$ mice and stained with anti-CD3, anti-TCR δ and anti-NK1.1 mAbs. Total $\gamma\delta$ T, NK1.1 $^{-}$ $\gamma\delta$ T and NK1.1 $^{+}$ $\gamma\delta$ T cells were sorted using Moflo XDP (Beckman Coulter, Brea, CA, USA). The purity of each $\gamma\delta$ T cells in this experiment was greater than 90%. The purified $\gamma\delta$ T cells were cultured with IL-2 (100U/ml), IL-7 (20 ng/ml) and IL-15 (20 ng/ml) for 12 days. Then, the expanded $\gamma\delta$ T cells were sorted using Moflo XDP (Beckman Coulter). One day after bleomycin exposure, purified expanded $\gamma\delta$ T cells were infused at 1×10^5 cells in total volume of 100 μ l via the tail vein into TCR $\delta^{-/-}$ mice.

Measurement of cytokines from $\gamma\delta$ T cells

The expanded NK1.1 $^{-}$ $\gamma\delta$ T cells (1×10^5 /ml) and NK1.1 $^{+}$ $\gamma\delta$ T cells (1×10^5 /ml) were stimulated with 1 μ g/ml of anti-CD3 mAb (Biolegend) and 1 μ g/ml of anti-CD28 mAb (Biolegend). After 96 h, IFN- γ and IL-17A in the culture supernatants were evaluated by enzyme-linked immunosorbent assay (ELISA) (R&D Systems, Minneapolis, MN, USA). For intracellular cytokine staining, the expanded NK1.1 $^{-}$ $\gamma\delta$ T cells (5×10^5 /ml) and NK1.1 $^{+}$ $\gamma\delta$ T cells (5×10^5 /ml) were co-cultured with 1 μ g/ml of anti-CD3 mAb (Biolegend) and 1 μ g/ml of anti-CD28 mAb (Biolegend). After 96 h, IFN- γ and IL-17A from NK1.1 $^{-}$ or NK1.1 $^{+}$ fractions were analysed by flow cytometry.

Isolation of CD4 $^{+}$ T cells and *in-vitro* T cell cultures

Splenic CD4 $^{+}$ T cells were isolated by positive selection, using a magnetic-activated cell sorter (MACS) system with anti-CD4 monoclonal antibody (mAb; Miltenyi Biotec, Bergisch Gladbach, Germany). For Th17 cell differentiation, CD4 $^{+}$ cells (1×10^6 /ml) were cultured in medium with 1 μ g/ml of anti-CD3 mAb (Biolegend), 1 μ g/ml of anti-CD28 mAb (Biolegend), 1 ng/ml of human transforming growth factor (TGF)- β (R&D Systems), 20 ng/ml of mouse IL-6 (eBioscience), 10 μ g/ml of anti-IFN- γ mAb (Biolegend) and 10 μ g/ml of anti-IL-4 mAb (Biolegend). On day 4, the cells were restimulated for 6 h with 50 ng/ml

of phorbol myristate acetate (PMA) and 500 ng/ml of ionomycin and used in the experiments.

Co-culture with CD4 $^{+}$ and expanded $\gamma\delta$ T cells

CD4 $^{+}$ T cells (2×10^5 /well) were co-cultured with expanded $\gamma\delta$ T cells (1×10^4 /well) in conditions of Th17 cell differentiation as above. On day 4, the cells were restimulated for 6 h with 50 ng/ml of phorbol myristate acetate and 500 ng/ml of ionomycin and used in the experiments.

Quantification of gene expression by reverse transcription–polymerase chain reaction (RT–PCR)

Total RNA was extracted from isolated from lung tissues, and reverse-transcribed into cDNA using RevertAidTM first-strand cDNA Synthesis kit (Fermentas, Burlington, Ontario, Canada), according to the protocol supplied by the manufacturer. The cDNA samples were amplified with specific primers and fluorescence-labelled probes for target genes. Specific primers and probes for TGF- β , collagen type I alpha 1 (Col1a1) and glyceraldehyde 3-phosphate dehydrogenase (GAPDH) were purchased from Applied Biosystems Japan (Tokyo, Japan). The amplified product genes were monitored on an ABI 7700 sequence detector (Applied Biosystems Japan). The quantitative PCR master mix was purchased from Applied Biosystems Japan. The final concentrations of the primers were 200 nM for each of 5' and 3' primers, and the final probe concentration was 100 nM. The thermal cycler conditions were 50°C for 2 min, 95°C for 10 min, then 50 cycles of 95°C for 15 s and 60°C for 1 min. Serial dilutions of a standard sample were incubated in every assay, and standard curves for the genes of interest and GAPDH genes were generated. All measurements were performed in duplicate. The gene expression level was calculated from the standard curve, and expressed relative to GAPDH gene expression.

Statistical analysis

Data were expressed as median and mean \pm standard deviation (s.d.). Differences between groups were examined for statistical significance using Student's *t*-test. A *P*-value less than 0.05 denoted the presence of a statistically significant difference. For multiple group comparisons, one-way analysis of variance (ANOVA) was performed, followed by the Tukey test. The Statistical Package for Social Sciences (SPSS Inc., Chicago, IL, USA) was used for all statistical analyses.

Results

$\gamma\delta$ T cells suppress bleomycin-induced IP using TCR $\delta^{-/-}$ mice

To examine the effects of $\gamma\delta$ T cells in the progression of pulmonary fibrosis, WT and TCR $\delta^{-/-}$ mice were treated

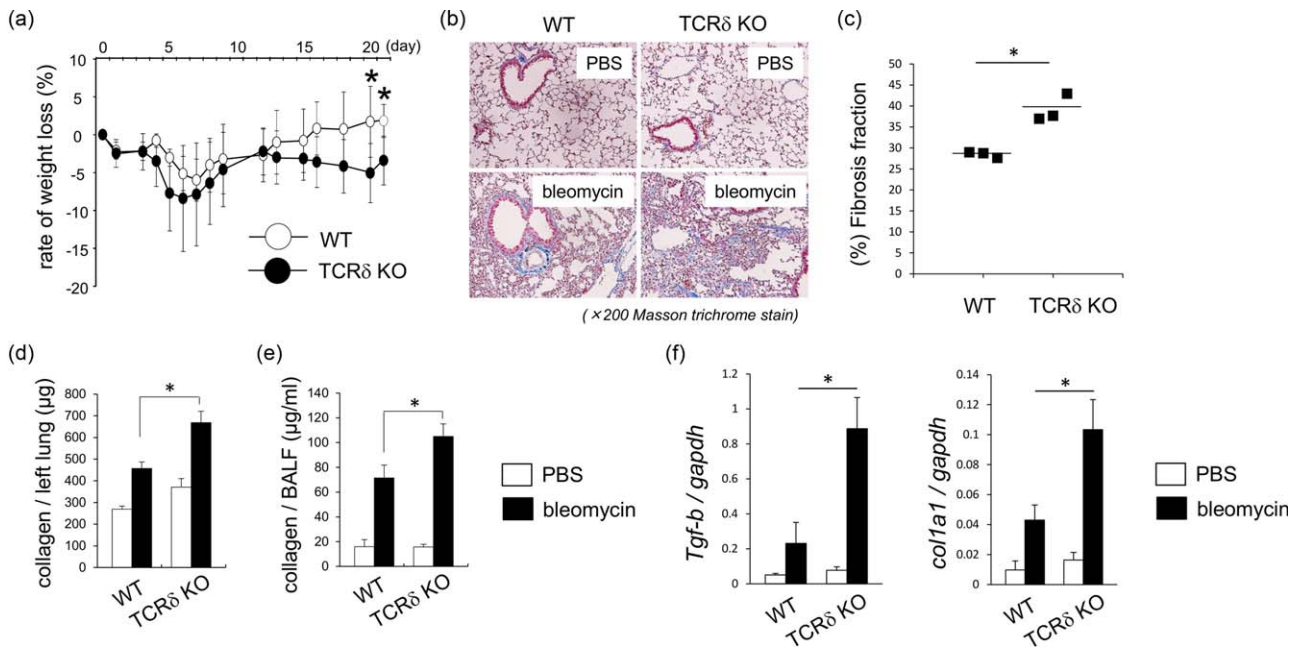


Fig. 1. Histological and biochemical analyses of wild-type (WT) and T cell receptor (TCR) $\delta^{-/-}$ mice exposed to bleomycin. (a) Change in body weight after bleomycin exposure in WT ($n = 4$) and TCR $\delta^{-/-}$ mice ($n = 7$). Data are representative of at least two independent experiments. Data are mean \pm standard deviation (s.d.). * $P < 0.05$. (b) Lung tissues were obtained from WT ($n = 3$) and TCR $\delta^{-/-}$ mice ($n = 4$) on day 21 after phosphate-buffered saline (PBS) or bleomycin exposure. Paraffin sections were stained with Masson's trichrome. Original magnification $\times 200$. (c) Fibrosis fraction in WT ($n = 3$) and TCR $\delta^{-/-}$ ($n = 3$) mice was measured on 21 days after bleomycin exposure by quantitative image analysis. Data are representative of at least three independent experiments. Data are mean \pm s.d. * $P < 0.05$. (d,e) The lung tissues and bronchoalveolar lavage fluid (BALF) were obtained from WT ($n = 4$) and TCR $\delta^{-/-}$ mice ($n = 7$) on day 21 after bleomycin exposure and collagen production was determined. Data are representative of at least three independent experiments. Data are mean \pm s.d. * $P < 0.05$. (f) Lung tissues were harvested from WT ($n = 4$) and TCR $\delta^{-/-}$ mice ($n = 7$) on day 21 after bleomycin exposure. Lung mRNA was extracted and the expression of transforming growth factor (TGF)- β and *col1a1* mRNA was analysed by reverse transcription–polymerase chain reaction (RT–PCR). Data are representative of at least two independent experiments. Data are mean \pm s.d. * $P < 0.05$.

with bleomycin. As shown in Fig. 1a, TCR $\delta^{-/-}$ mice showed significant body weight loss compared with WT mice on days 20 and 21 after bleomycin exposure. Histological analysis showed that TCR $\delta^{-/-}$ mice had thickened alveolar septa and ablation of alveolar space at day 21 compared with WT mice (Fig. 1b). Using the quantitative image analysis (QIA), the fibrosis fraction was significantly larger in TCR $\delta^{-/-}$ mice compared with WT mice 21 days after bleomycin exposure (Fig. 1c). The amounts of collagen in lung tissues and BALF was significantly higher in TCR $\delta^{-/-}$ mice than in WT mice 21 days after bleomycin exposure (Fig. 1d,e). Furthermore, the expression of TGF- β and *Col1a1* mRNA in lung tissues was significantly higher in TCR $\delta^{-/-}$ mice than WT mice (Fig. 1f). These results indicate that the deficiency of $\gamma\delta$ T cells enhances the progression of bleomycin-induced IP.

Expansion of pulmonary $\gamma\delta$ T cells in bleomycin-induced IP

To examine the phenotype of pulmonary $\gamma\delta$ T cells following exposure to bleomycin, pulmonary lymphocytes were ana-

lysed by flow cytometry (FCM). As shown in Fig. 2a, the number of pulmonary $\gamma\delta$ T cells increased at days 3, 7 and 14 after bleomycin exposure. Then, we examined the expression of IFN- γ and IL-17A on pulmonary $\gamma\delta$ T cells. At 3 days after bleomycin exposure, IFN- γ -producing (IFN- γ^+) $\gamma\delta$ T cells expanded in lung tissues (Fig. 2b). In comparison, at 7, 14 and 21 days after bleomycin exposure, the number of IL-17A $^+$ $\gamma\delta$ T cells increased exponentially in lung tissues (Fig. 2b). These results suggest that pulmonary $\gamma\delta$ T cells produce mainly IL-17A at the fibrosis phase after bleomycin exposure. To confirm the role of IL-17A in bleomycin-induced pulmonary fibrosis, experiments were performed in IL-17A-deficient (IL-17A $^{-/-}$) mice. The collagen accumulation was reduced on day 21 after bleomycin exposure in IL-17A $^{-/-}$ mice (Fig. 2c). Furthermore, the amounts of collagen in lung tissues and BALF on 21 days after bleomycin exposure was significantly lower in IL-17A $^{-/-}$ mice than in WT mice (Fig. 2d). The above findings suggest that IL-17A is an exacerbation factor in the development of pulmonary fibrosis. Conversely, pulmonary $\gamma\delta$ T cells play an inhibitory role in the development of pulmonary fibrosis despite the production of inflammatory cytokine such as IL-17A.

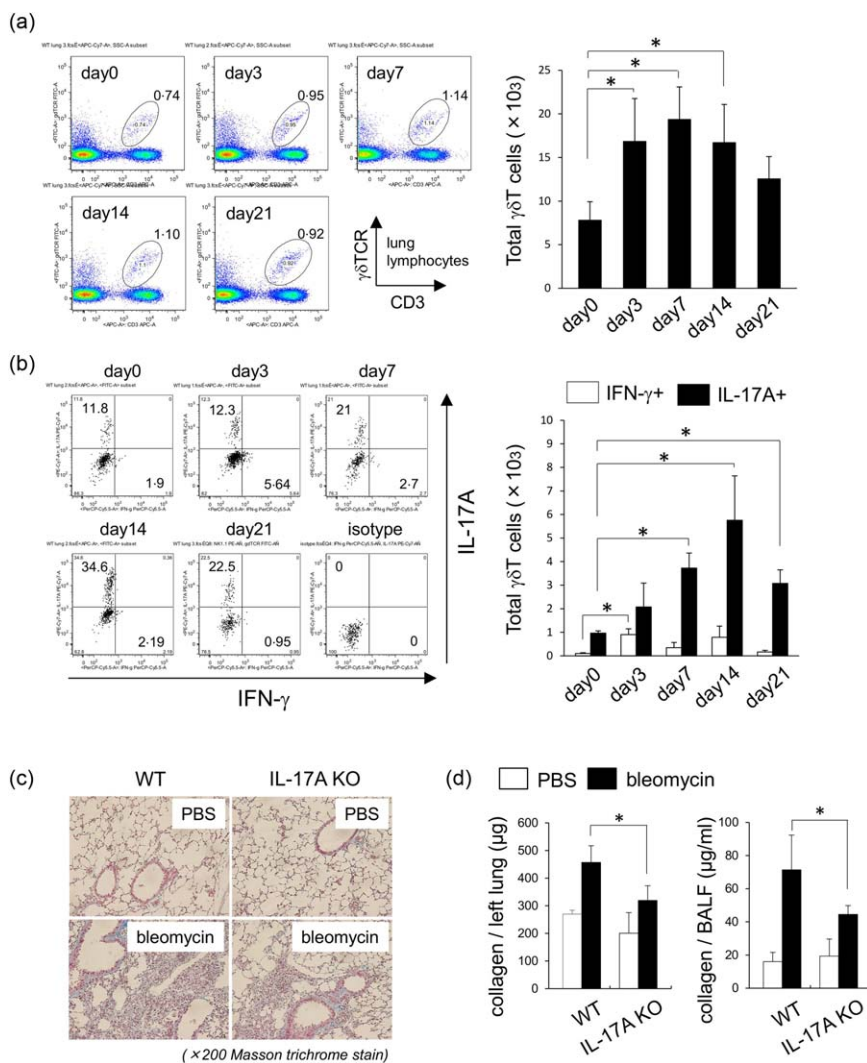


Fig. 2. Phenotypes of pulmonary $\gamma\delta$ T cells in bleomycin-induced fibrosis.

(a) Pulmonary lymphocytes were harvested from wild-type (WT) mice ($n = 3$) on days 0, 3, 7, 14 and 21 after intratracheal instillation of bleomycin. Cells were stained for CD3 ϵ , $\gamma\delta$ T cell receptor (TCR) δ and analysed by flow cytometry. Data are representative of at least three independent experiments. Data are mean \pm standard deviation (s.d.). $*P < 0.05$.

(b) Pulmonary lymphocytes were harvested from WT mice ($n = 3$) on days 0, 3, 7, 14 and 21 after bleomycin exposure and stimulated by phorbol myristate acetate (PMA)/ionomycin for 6 h. Cells were stained for CD3 ϵ , $\gamma\delta$ TCR, interferon (IFN)- γ , interleukin (IL)-17A and analysed by flow cytometry. CD3 ϵ ⁺ $\gamma\delta$ TCR⁺ cells were gated. Data are representative of at least two independent experiments. Data are mean \pm s.d. $*P < 0.05$.

(c) Lung tissues were removed from WT ($n = 3$) and interleukin (IL)-17A^{-/-} mice ($n = 4$) on day 21 after bleomycin exposure. Paraffin sections were stained with Masson's trichrome. Original magnification $\times 200$. (d) Lung tissues and bronchoalveolar lavage fluid (BALF) were obtained from WT ($n = 3$) and IL-17A^{-/-} mice ($n = 4$) on day 21 after bleomycin exposure. Collagen production was determined by sircol assay. Data are representative of at least three independent experiments. Data are mean \pm s.d. $*P < 0.05$.

$\gamma\delta$ T cells induce Th1 cells but reduce Th17 cells

The number of total cells in BALF was significantly lower in TCR δ ^{-/-} mice compared with WT mice at 7 days after bleomycin exposure (Fig. 3a). The numbers of dendritic cells and neutrophils in BALF were significantly lower in TCR δ ^{-/-} mice than WT mice at days 3 and 7. These results suggest that $\gamma\delta$ T cells induce infiltration of dendritic cells and neutrophils into the lung, resulting in severe inflammation.

We also examined the roles of IFN- γ ⁺CD4⁺ T and IL-17A⁺CD4⁺ T cells in the development of pulmonary fibrosis. After exposure to bleomycin, the proportion of IFN- γ ⁺CD4⁺ T cells in lung tissues at day 3 was significantly lower in TCR δ ^{-/-} mice than WT mice (Fig. 3b). In contrast, the proportion of IL-17A⁺CD4⁺ T cells at days 3, 7, 14 and 21 days after bleomycin exposure was significantly higher in TCR δ ^{-/-} mice. The proportion of pulmonary regulatory T cells (T_{reg}) [forkhead box protein 3 (Foxp3⁺)CD4⁺ T] cells was not significantly different

between WT and TCR δ ^{-/-} mice (Fig. 3c). These findings suggest that $\gamma\delta$ T cells seem to induce IFN- γ ⁺CD4⁺ T cells and suppress IL-17A⁺CD4⁺ T cells in mice with bleomycin-induced IP.

$\gamma\delta$ T cells attenuate bleomycin-induced pulmonary fibrosis via the suppression of Th17 cells

To confirm the role of $\gamma\delta$ T cells in bleomycin-induced pulmonary fibrosis, we transferred $\gamma\delta$ T cells expanded by IL-2, IL-7 and IL-15 into bleomycin-treated TCR δ ^{-/-} mice. As shown in Fig. 4a, $\gamma\delta$ T cells reduced collagen accumulation and fibrosis fraction in TCR δ ^{-/-} mice. The amounts of collagen in BALF was not decreased significantly, but tended to be lower in TCR δ ^{-/-} mice infused with $\gamma\delta$ T cells compared with control mice (Fig. 4b). In $\gamma\delta$ T cells infused TCR δ ^{-/-} mice, pulmonary IL-17A⁺CD4⁺ T cells were significantly lower compared with control mice, whereas pulmonary IFN- γ ⁺CD4⁺ T cells were not (Fig. 4c). After exposure of TCR δ ^{-/-} mice to bleomycin, pulmonary CD4⁺

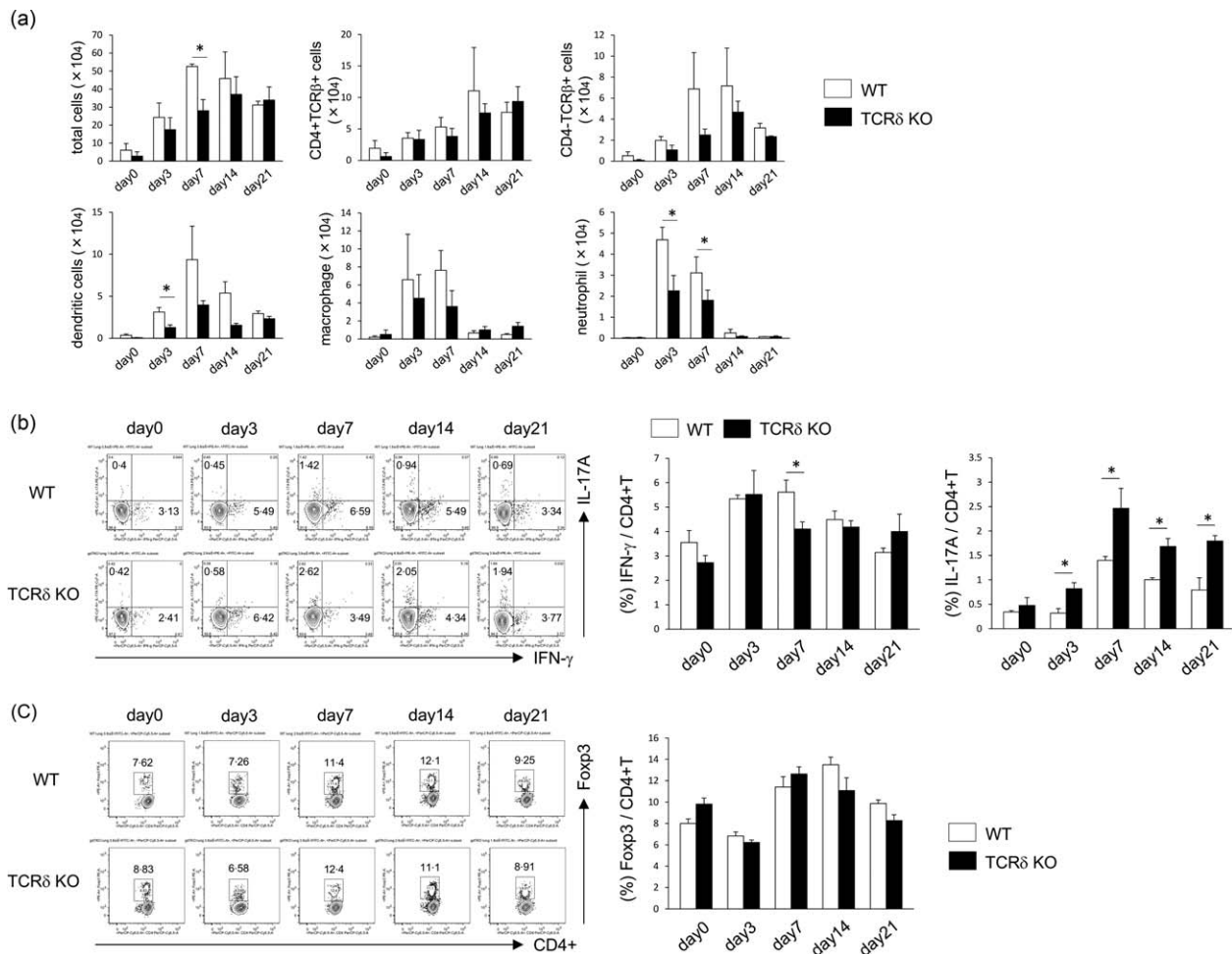


Fig. 3. Inflammatory and cellular changes in lung tissues of mice exposed to bleomycin. (a) wild-type (WT) ($n = 3$) and T cell receptor (TCR) $\delta^{-/-}$ mice ($n = 4$) were infused intratracheally with bleomycin. At days 0, 3, 7, 14 and 21, the number of total, CD4⁺TCR β ⁺, CD4⁻TCR β ⁺, dendritic cells (CD11c⁺), macrophages (CD11c⁻CD11b⁺Gr-1⁻) and neutrophils (CD11c⁻CD11b⁺Gr-1⁺) was analysed by flow cytometry (FCM). Data are representative of at least two independent experiments. Data are mean \pm standard deviation (s.d.). * $P < 0.05$. (b) Pulmonary lymphocytes were harvested from WT ($n = 3$) and TCR $\delta^{-/-}$ mice ($n = 4$) on days 0, 3, 7, 14 and 21 after bleomycin exposure and stimulated by phorbol myristate acetate (PMA)/ionomycin for 6 h. Cells were stained for TCR β , CD4, interferon (IFN)- γ , interleukin (IL)-17A and analysed by flow cytometry. CD4⁺TCR β ⁺ cells were gated. Data are representative of at least three independent experiments. Data are mean \pm s.d. * $P < 0.05$. (c) Pulmonary lymphocytes were harvested from WT ($n = 3$) and TCR $\delta^{-/-}$ mice ($n = 4$) on days 0, 3, 7, 14 and 21 after bleomycin exposure. Cells were stained for TCR β , CD4, forkhead box protein 3 (Foxp3) and analysed by flow cytometry. CD4⁺TCR β ⁺ cells were gated. Data are representative of at least two independent experiments. Data are mean \pm s.d. * $P < 0.05$.

T cells were inclined towards IL-17A⁺CD4⁺ T cells (Fig. 4d). The transfer of $\gamma\delta$ T cells in TCR $\delta^{-/-}$ mice improved the ratio of IL-17⁺CD4⁺ T cells to IFN- γ ⁺CD4⁺ T cells (Fig. 4d).

To examine whether IL-17A plays essential roles in the development of pulmonary fibrosis in TCR $\delta^{-/-}$ mice, IL-17A^{-/-}TCR $\delta^{-/-}$ mice were treated with bleomycin. At day 21, collagen accumulation in lung tissues and collagen production in BALF were significantly lower in IL-17A^{-/-}TCR $\delta^{-/-}$ mice than TCR $\delta^{-/-}$ mice (Fig. 4e,f). However, pulmonary collagen accumulation and production were not significantly different between IL-17A^{-/-} and IL-17A^{-/-}TCR $\delta^{-/-}$ mice. These findings support the notion

that $\gamma\delta$ T cells suppressed pulmonary fibrosis through the suppression of IL-17⁺CD4⁺ T cells.

$\gamma\delta$ T cells suppress Th17 cell differentiation *in vitro* via IFN- γ production

To examine whether $\gamma\delta$ T cells affect Th17 cell differentiation, CD4⁺ T cells were cultured under several conditions *in vitro*. CD4⁺ cells isolated from TCR $\delta^{-/-}$ mice showed normal differentiation of Th17 cells compared with WT mice (Fig. 5a). To investigate the effects of IFN- γ in Th17 cell differentiation, we used IFN- γ neutralization antibodies. The number of IL-17A⁺CD4⁺ T cells was significantly

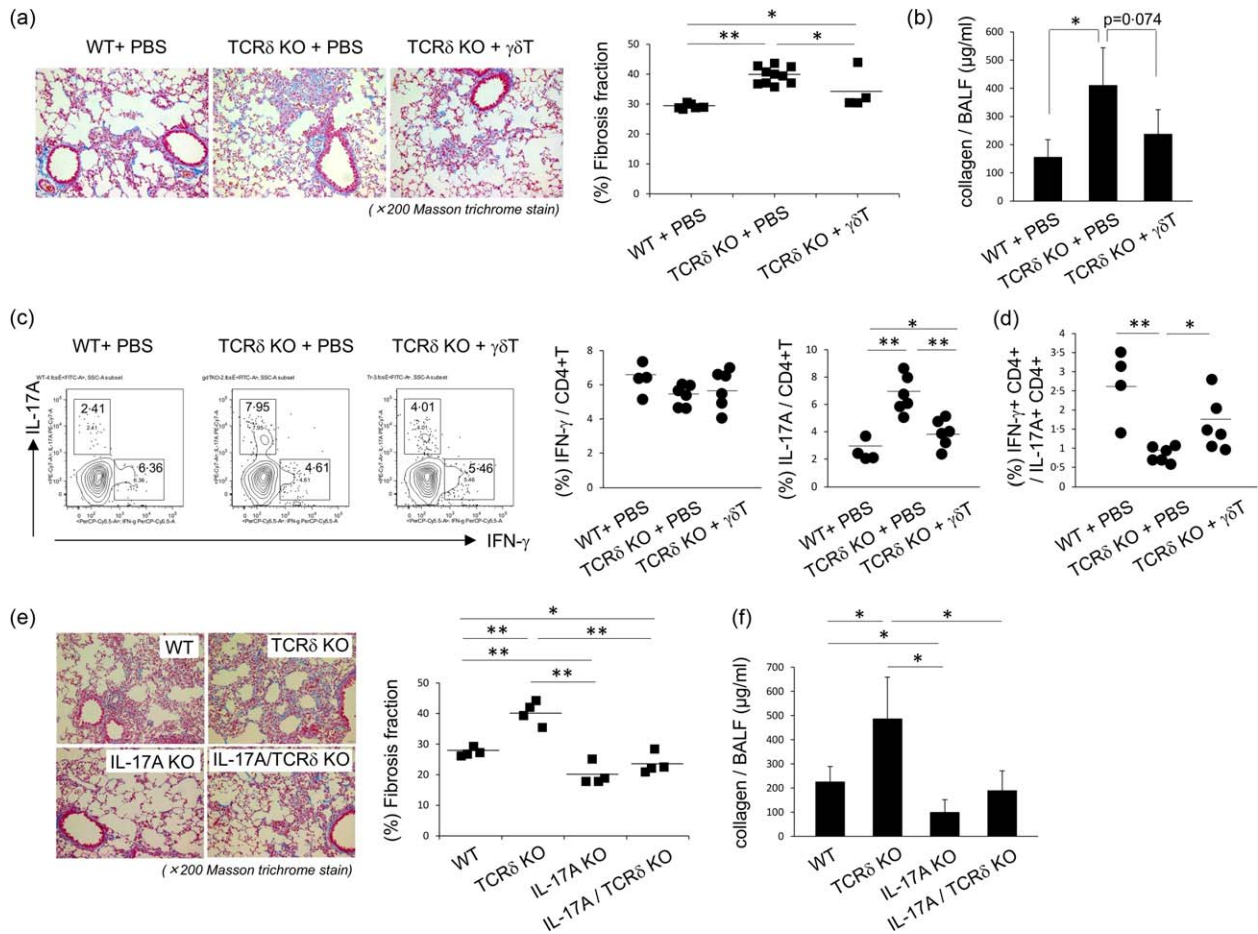


Fig. 4. Improvement of pulmonary fibrosis by $\gamma\delta$ T cells. (a) Wild-type (WT) mice were harvested and pulmonary $\gamma\delta$ T cells were purified, as described in Materials and methods. Purified pulmonary $\gamma\delta$ T cells were expanded by interleukin (IL)-2, IL-7 and IL-15 for 12 days. Then, the expanded $\gamma\delta$ T cells were transferred into T cell receptor (TCR) $\delta^{-/-}$ mice (1×10^5 /mice). Six WT mice received phosphate-buffered saline (PBS), 10 TCR $\delta^{-/-}$ mice received PBS and four TCR $\delta^{-/-}$ mice received $\gamma\delta$ T cells. The lung tissues were removed on day 21 after bleomycin exposure. Paraffin sections were stained with Masson's trichrome. Original magnification $\times 200$. Fibrosis fraction was measured by quantitative image analysis, as described in Materials and methods. Data are representative of at least two independent experiments. Data are mean \pm standard deviation (s.d.). * $P < 0.05$, ** $P < 0.01$. (b) Bronchoalveolar lavage fluid (BALF) was obtained on day 17 after bleomycin exposure and collagen production was determined. Data are representative of at least two independent experiments. Data are mean \pm s.d. * $P < 0.05$. (c) At 18 days after bleomycin exposure, pulmonary lymphocytes were harvested and then stimulated with phorbol myristate acetate (PMA)/ionomycin for 6 h. Cells were stained for CD3 ϵ , CD4, interferon (IFN)- γ , IL-17A and analysed by flow cytometry. CD4 $^+$ CD3 ϵ^+ cells were gated. Data are representative of at least two independent experiments. Data are mean \pm s.d. * $P < 0.05$, ** $P < 0.01$. (d) At 18 days after bleomycin exposure, pulmonary lymphocytes were harvested and then stimulated with PMA/ionomycin for 6 h. Cells were stained by CD3 ϵ , CD4, IFN- γ , IL-17A and analysed by flow cytometry (FCM). Data are the proportion of IFN- γ^+ CD4 $^+$ T/IL-17A $^+$ CD4 $^+$ T cells in each mouse. Data are representative of at least two independent experiments. Data are mean \pm s.d. * $P < 0.05$, ** $P < 0.01$. (e) WT ($n = 4$), TCR $\delta^{-/-}$ ($n = 4$), IL-17A $^{-/-}$ ($n = 4$) and IL-17A/TCR $\delta^{-/-}$ ($n = 4$) mice were treated with bleomycin. After 21 days, paraffin sections were stained with Masson's trichrome. Original magnification $\times 200$. Fibrosis fraction was measured by quantitative image analysis. Data are representative of at least two independent experiments. Data are mean \pm s.d. * $P < 0.05$, ** $P < 0.01$. (f) WT ($n = 4$), TCR $\delta^{-/-}$ ($n = 4$), IL-17A $^{-/-}$ ($n = 4$) and IL-17A/TCR $\delta^{-/-}$ ($n = 4$) mice were treated with bleomycin. BALF was obtained on day 21 after bleomycin exposure and collagen production was determined. Data are representative of at least two independent experiments. Data are mean \pm s.d. * $p < 0.05$.

higher in the presence of anti-IFN- γ mAb (Fig. 5b). These results suggest that IFN- γ suppresses Th17 cell differentiation *in vitro*. To elucidate whether $\gamma\delta$ T cells play a direct role in Th17 cell differentiation *in vitro*, we used pulmonary $\gamma\delta$ T cells derived from WT and IFN- $\gamma^{-/-}$ mice. After co-culture with CD4 $^+$ T cells and $\gamma\delta$ T cells derived from

WT mice, the proportion of IL-17A $^+$ CD4 $^+$ T cells was significantly lower than $\gamma\delta$ T cells derived from IFN- $\gamma^{-/-}$ mice (Fig. 5c). To confirm further the direct effects of IFN- γ from $\gamma\delta$ T cells, we used CD4 $^+$ T cells from IFN- $\gamma^{-/-}$ mice. As shown in Fig. 5d, $\gamma\delta$ T cells derived from WT mice suppressed IL-17 $^+$ CD4 $^+$ T cells, whereas $\gamma\delta$ T cells derived

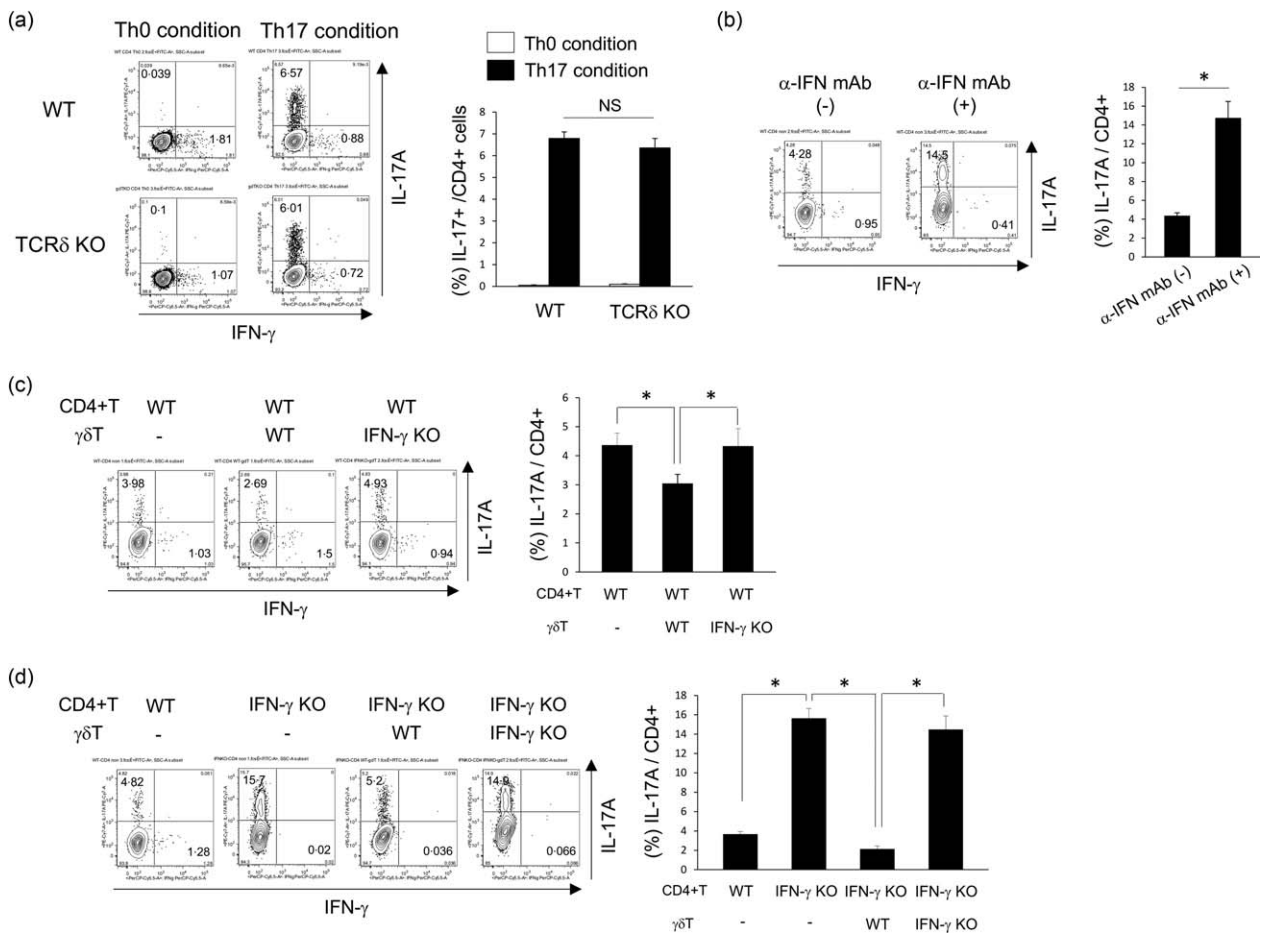


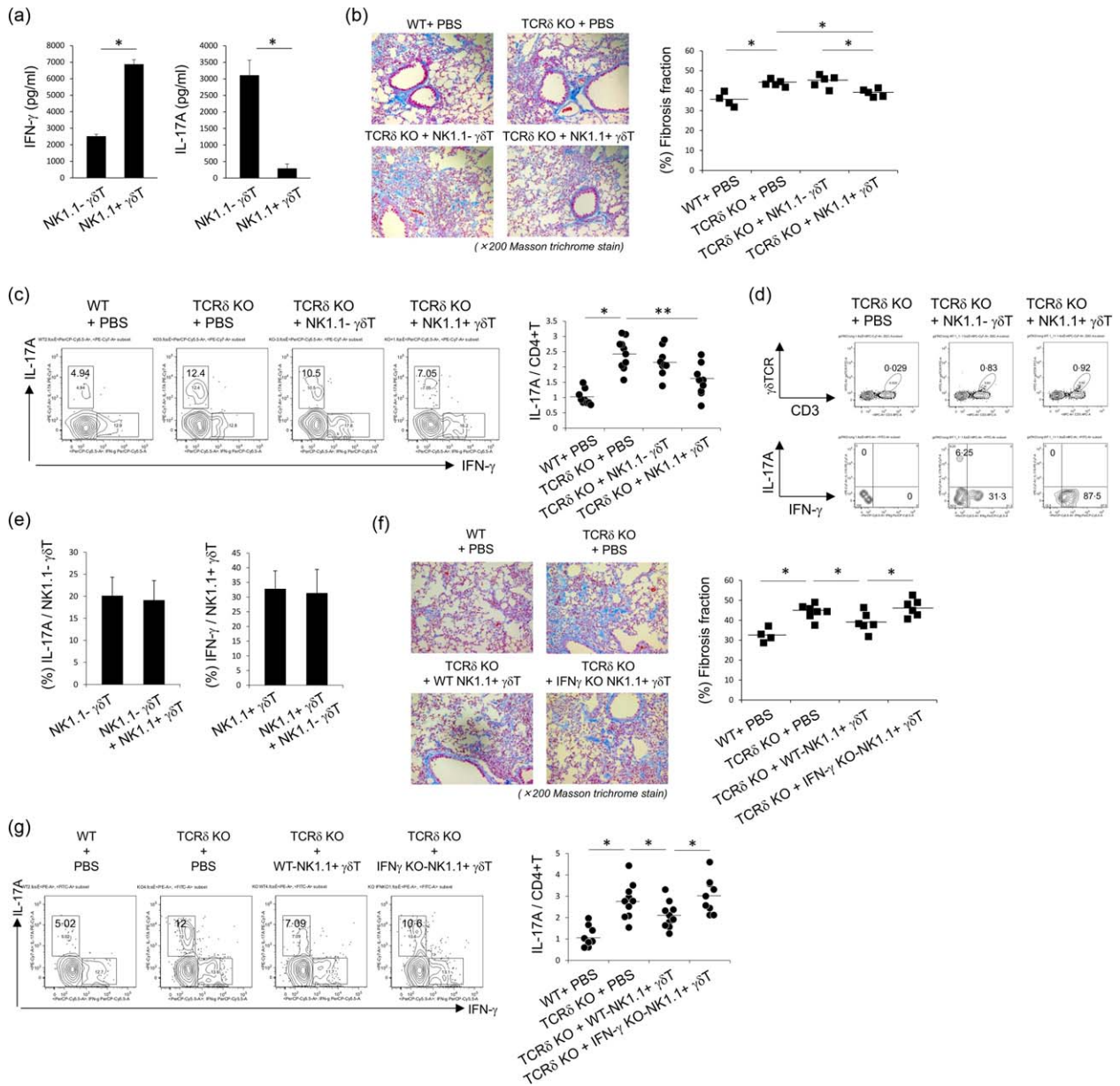
Fig. 5. Effects of interferon (IFN)- γ ⁺ $\gamma\delta$ T cells in T helper 17 (Th17) cell differentiation *in vitro*. (a) Splenic CD4⁺ T cells were isolated from wild-type (WT) and T cell receptor (TCR) δ ^{-/-} mice and cultured in Th0 conditions [anti-CD3 monoclonal antibody (mAb), anti-CD28 mAb, anti-interferon (IFN)- γ mAb and anti-interleukin (IL)-4 mAb], Th17 conditions (anti-CD3 mAb, anti-CD28 mAb, transforming growth factor (TGF)- β , IL-6, anti-IFN- γ mAb and anti-IL-4 mAb) for 96 h, as described in Materials and methods. Cells were stained for CD3 ϵ , CD4, IFN- γ , IL-17A and analysed by flow cytometry. CD4⁺CD3 ϵ ⁺ cells were gated. Data are representative of at least two independent experiments. Data are mean \pm standard deviation (s.d.). * P < 0.05. (b) Splenic CD4⁺ T cells from WT mice were cultured in Th17 conditions (anti-CD3 mAb, anti-CD28 mAb, TGF- β , IL-6 and anti-IL-4 mAb) with/without anti-IFN- γ mAb for 96 h, as described in Materials and methods. Cells were stained for CD3 ϵ , CD4, IFN- γ , IL-17A and analysed by flow cytometry. CD4⁺CD3 ϵ ⁺ cells were gated. Data are representative of at least two independent experiments. Data are mean \pm s.d. * P < 0.05. (c,d) Pulmonary $\gamma\delta$ T cells were harvested from WT and IFN- γ ^{-/-} mice and purified, as described in Materials and methods. The purified pulmonary $\gamma\delta$ T cells were expanded by IL-2, IL-7 and IL-15 for 12 days. Splenic CD4⁺ T cells from WT and IFN- γ ^{-/-} mice were co-cultured with the expanded $\gamma\delta$ T cells derived from WT and IFN- γ ^{-/-} mice under Th17 conditions (anti-CD3 mAb, anti-CD28 mAb, TGF- β , IL-6 and anti-IL-4 mAb) for 96 h, as described in Materials and methods. Cells were stained for CD3 ϵ , CD4, IFN- γ , IL-17A and analysed by flow cytometry. CD4⁺CD3 ϵ ⁺ cells were gated. Data are representative of at least three independent experiments. Data are mean \pm s.d. * P < 0.05.

from IFN- γ ^{-/-} mice did not. Taken together, the results showed that $\gamma\delta$ T cells seem to suppress Th17 cell differentiation via IFN- γ production.

IFN- γ ⁺ $\gamma\delta$ T cells attenuate bleomycin-induced pulmonary fibrosis

To examine the effects of IFN- γ from $\gamma\delta$ T cells in the progression of pulmonary fibrosis, $\gamma\delta$ T cells were separated into two subsets based on the expression of NK1.1. As shown in Fig. 6a, NK1.1⁺ $\gamma\delta$ T cells showed large amounts of IFN- γ production compared with NK1.1⁻ $\gamma\delta$ T cells. In TCR δ ^{-/-} mice, NK1.1⁺ $\gamma\delta$ T cells reduced collagen accumu-

lation and fibrosis fraction in the lungs, whereas NK1.1⁻ $\gamma\delta$ T cells did not (Fig. 6b). The transfer of NK1.1⁺ $\gamma\delta$ T cells in TCR δ ^{-/-} mice reduced pulmonary IL-17A⁺CD4⁺ T cells (Fig. 6c). After NK1.1⁺ $\gamma\delta$ T cells transferred into TCR δ ^{-/-} mice, pulmonary $\gamma\delta$ T cells showed large amounts of IFN- γ production *in vivo* (Fig. 6d). To examine the interactions for cytokine production in $\gamma\delta$ T cell subsets, NK1.1⁻ $\gamma\delta$ T cells and NK1.1⁺ $\gamma\delta$ T cells were co-cultured *in vitro*. IFN- γ production from NK1.1⁺ $\gamma\delta$ T cells and IL-17A production from NK1.1⁻ $\gamma\delta$ T cells were not significantly different in the presence or absence of other $\gamma\delta$ T cell subsets (Fig. 6e).



Furthermore, to confirm the role of IFN- γ ⁺ $\gamma\delta$ T cells in bleomycin-induced pulmonary fibrosis, NK1.1⁺ $\gamma\delta$ T cells from WT and IFN- γ ^{-/-} mice were transferred into bleomycin-treated TCR δ ^{-/-} mice. The results showed that NK1.1⁺ $\gamma\delta$ T cells from WT mice reduced collagen accumulation and fibrosis fraction in the lungs, whereas NK1.1⁺ $\gamma\delta$ T cells from IFN- γ ^{-/-} mice did not (Fig. 6f). In TCR δ ^{-/-} mice receiving NK1.1⁺ $\gamma\delta$ T cells from IFN- γ ^{-/-} mice, pulmonary IL-17A⁺ CD4⁺ T cells were not reduced (Fig. 6g).

Discussion

In this study, we demonstrated that $\gamma\delta$ T cells play a regulatory role in the development of bleomycin-induced pulmonary fibrosis. In TCR δ ^{-/-} mice, the severity of pulmonary

fibrosis correlated with accumulation of IL-17A⁺ CD4⁺ T cells in lung tissues. Infusion of $\gamma\delta$ T cells in TCR δ ^{-/-} mice attenuated pulmonary fibrosis and decreased pulmonary IL-17A⁺ CD4⁺ T cells significantly compared with control mice. After exposure to bleomycin, pulmonary $\gamma\delta$ T cells secreted large amounts of IFN- γ in WT mice. The results also showed suppression of Th17 cell differentiation in the presence of IFN- γ ⁺ $\gamma\delta$ T cells *in vitro*. Collectively, these results suggest that IFN- γ ⁺ $\gamma\delta$ T cells seem to suppress IL-17⁺ CD4⁺ T cells responses in bleomycin-induced pulmonary fibrosis (summarized schematically in Fig. 7).

The presence of $\gamma\delta$ T cells is critical for a controlled immune response in the lung. In various lung diseases, $\gamma\delta$ T cells play an effective or suppressive role by secreting various cytokines such as IFN- γ , IL-17A, IL-22 and IL-4

Fig. 6. Inhibitory effects of interferon (IFN)- γ ⁺ $\gamma\delta$ T cells in the progression of pulmonary fibrosis. (a) Pulmonary natural killer (NK)1.1⁻ $\gamma\delta$ T and NK1.1⁺ $\gamma\delta$ T cells were purified from wild-type (WT) mice and expanded by interleukin (IL)-2, IL-7 and IL-15 for 12 days as described in Materials and methods. The expanded $\gamma\delta$ T cells were stimulated by anti-CD3 and anti-CD28 monoclonal antibodies (mAbs) for 72 h. IFN- γ and IL-17A in the culture supernatant were measured by enzyme-linked immunosorbent assay (ELISA). Data are representative of at least three independent experiments. Data are mean \pm standard deviation (s.d.). * $P < 0.05$. (b) The expanded NK1.1⁻ $\gamma\delta$ T and NK1.1⁺ $\gamma\delta$ T cells from WT mice were transferred into T cell receptor (TCR) δ ^{-/-} mice (1×10^5 cells/mice). Four WT mice received phosphate-buffered saline (PBS), five TCR δ ^{-/-} mice received PBS, five TCR δ ^{-/-} mice received NK1.1⁻ $\gamma\delta$ T cells and five TCR δ ^{-/-} mice received NK1.1⁺ $\gamma\delta$ T cells. The lung tissues were removed on day 21 after bleomycin exposure. Paraffin sections were stained with Masson's trichrome. Original magnification was $\times 200$. Fibrosis fraction was measured by quantitative image analysis, as described in Materials and methods. Data are representative of at least two independent experiments. Data are mean \pm s.d. * $P < 0.05$. (c) At 18 days after bleomycin exposure, pulmonary lymphocytes were harvested and then stimulated with phorbol myristate acetate (PMA)/ionomycin for 6 h. Cells were stained for CD3 ϵ , CD4, IFN- γ , IL-17A and analysed by flow cytometry. CD4⁺CD3 ϵ ⁺ cells were gated. Data are shown as a ratio of IL-17A⁺ cells in CD4⁺ T cells compared with those in control mice. The value of control mice was shown by mean of two independent experiments as 1.0. Graph represents values of two independent mice, using the average of the control mice. Data are mean \pm s.d. * $P < 0.05$, ** $P < 0.01$. (d) The expanded NK1.1⁻ $\gamma\delta$ T and NK1.1⁺ $\gamma\delta$ T cells from WT mice were transferred into TCR δ ^{-/-} mice (2×10^6 cells/mice). At 3 days after bleomycin exposure, pulmonary lymphocytes were harvested and then stimulated with PMA/ionomycin for 6 hrs. Cells were stained for CD3 ϵ , TCR δ , IFN- γ , IL-17A and analysed by flow cytometry. Data are representative of at least two independent experiments. (e) The expanded NK1.1⁻ $\gamma\delta$ T (5×10^5 cells/ml) and NK1.1⁺ $\gamma\delta$ T cells (5×10^5 cells/ml) were co-cultured with anti-CD3 mAb and anti-CD28 mAb for 96 h. IFN- γ and IL-17A from NK1.1⁻ or NK1.1⁺ fractions were analysed by flow cytometry. Data are representative of at least two independent experiments. Data are mean \pm s.d. (f) Pulmonary NK1.1⁺ $\gamma\delta$ T cells were purified from WT and IFN- γ ^{-/-} mice and expanded by IL-2, IL-7 and IL-15 for 12 days, as described in Materials and methods. Then, the expanded $\gamma\delta$ T cells were transferred into TCR δ ^{-/-} mice (1×10^5 cells/mice). Four WT mice received PBS, seven TCR δ ^{-/-} mice received PBS, six TCR δ ^{-/-} mice received NK1.1⁺ $\gamma\delta$ T cells from WT mice and six TCR δ ^{-/-} mice received NK1.1⁺ $\gamma\delta$ T cells from IFN- γ ^{-/-} mice. The lung tissues were removed on day 21 after bleomycin exposure. Paraffin sections were stained with Masson's trichrome. Original magnification was $\times 200$. Fibrosis fraction was measured by quantitative image analysis, as described in Materials and methods. Data are representative of at least two independent experiments. Data are mean \pm s.d. * $P < 0.05$. (g) At 18 days after bleomycin exposure, pulmonary lymphocytes were harvested and then stimulated with PMA/ionomycin for 6 h. Cells were stained for CD3 ϵ , CD4, IFN- γ , IL-17A and analysed by flow cytometry. CD4⁺CD3 ϵ ⁺ cells were gated. Data are shown as a ratio of IL-17A⁺ cells in CD4⁺ T cells, compared with those in control mice. The value of control mice was shown by mean of two independent experiments as 1.0. Graph represents each values of two independent mice, using the average of the control mice. Data are mean \pm s.d. * $P < 0.05$.

[5,6,10]. In the pathogenesis of bleomycin-induced pulmonary fibrosis, it was reported that IL-17A⁺ $\gamma\delta$ T and CXCL10⁺ $\gamma\delta$ T cells involved as the inflammatory or inhibitory agents, respectively [5,11,21]. Conversely, we focused previously on IFN- γ -producing $\gamma\delta$ T cells and reported that these cells might participate in the pathogenesis of pulmonary fibrosis in systemic sclerosis (SSc) patients [24]. In the present study, we showed an accumulation of IFN- γ ⁺ $\gamma\delta$ T cells and IL-17A⁺ $\gamma\delta$ T cells in lung tissues after bleomycin exposure. It is reported that IFN- γ has anti-fibrotic properties while IL-17A has profibrotic effects, both *in vivo* and *in*

vitro [13–21]. Based on these reports, $\gamma\delta$ T cells seem to have opposite effects in the progression of bleomycin-induced pulmonary fibrosis. Conversely, we showed that pulmonary fibrosis was exacerbated in TCR δ ^{-/-} mice, suggesting that $\gamma\delta$ T cells might function as a regulatory T cells against bleomycin-induced pulmonary fibrosis. We speculate that $\gamma\delta$ T cells suppress bleomycin-induced pulmonary fibrosis through three potential mechanisms: uncontrollability of Th17 cells, over-expression of TGF- β and reduction of T_{regs}. These scenarios are discussed below.

First, does uncontrollability of IL-17A-producing CD4⁺ T cells play a role? IL-17A had been implicated in the regulation of bleomycin-induced pulmonary fibrosis via collagen and TGF- β production from fibroblasts [21]. In this study, we found a higher proportion of pulmonary IL-17A⁺ CD4⁺ T cells in TCR δ ^{-/-} mice compared with WT mice. Furthermore, using IL-17A^{-/-}TCR δ ^{-/-} mice, we found that IL-17A played a pivotal role in the development of pulmonary fibrosis in TCR δ ^{-/-} mice. In humans and mice, IL-17A⁺CD4⁺ $\alpha\beta$ TCR⁺ (Th17) cells are the major source of IL-17A [26]. Previous studies showed that Th17 cells exacerbated pulmonary fibrosis in bleomycin-induced IP mice [20,21]. Blocking the recruitment of CD3⁺ or CD4⁺ T cells resulted in fewer fibrotic lesions [27,28]. These findings supported that bleomycin-induced pulmonary fibrosis was dependent upon T cells and the absence of $\alpha\beta$ T cells might attenuate fibrosis. In TCR δ ^{-/-} mice treated with bleomycin,

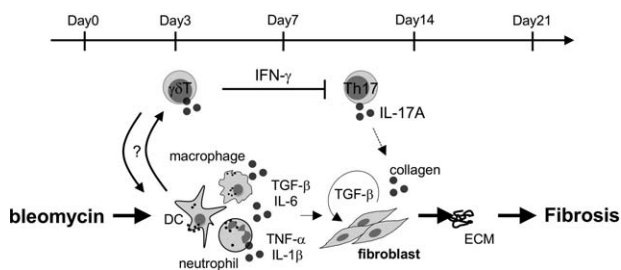


Fig. 7. Schematic diagram of the role of $\gamma\delta$ T cells in pulmonary fibrosis. Schematic diagram illustrating the role of $\gamma\delta$ T cells in the suppression of pulmonary fibrosis. After bleomycin exposure, $\gamma\delta$ T cells expanded or accumulated into lung tissues. These cells reduced pulmonary interleukin (IL)-17A⁺ CD4⁺ T cells and played regulatory roles in the pathogenesis of pulmonary fibrosis.

uncontrolled Th17 cells might lead to exacerbation of pulmonary fibrosis. Infusion of $\gamma\delta$ T cells into TCR $\delta^{-/-}$ mice treated with bleomycin reduced the percentage of lung fibrosis and collagen accumulation and lowered the percentage of pulmonary IL-17A⁺CD4⁺ T cells compared with control mice. These findings suggest that $\gamma\delta$ T cells seem to suppress the activity of pulmonary Th17 cells and attenuate the development of fibrosis. However, several studies reported that $\gamma\delta$ T cells induced the generation and activation of Th17 cells [29,30]. Do *et al.* [31] also reported that CCR6⁺ $\gamma\delta$ T cells play a crucial role in enhancing *in-vivo* Th17 cell differentiation and exacerbation of T cell-mediated colitis. Conversely, there is information on whether $\gamma\delta$ T cells suppress Th17 cell differentiation. To explore the inhibitory mechanisms of $\gamma\delta$ T cells, we evaluated Th17 cell differentiation *in vitro*. The results showed that $\gamma\delta$ T cells derived from WT mice suppressed Th17 cell differentiation *in vitro*, whereas $\gamma\delta$ T cells derived from IFN- $\gamma^{-/-}$ mice did not. Thus, $\gamma\delta$ T cells might suppress Th17 cell differentiation depending on IFN- γ production. IFN- γ is known as one of the Th1 cytokines and could suppress Th17 cell differentiation both *in vivo* and *in vitro* [32,33].

The current study shows that $\gamma\delta$ T cells can be classified into two major subsets based on distinct cytokine profiles, IFN- γ -producing $\gamma\delta$ T cells and IL-17A-producing $\gamma\delta$ T cells. The expression of NK1.1 and CD27 *versus* Scart-2 and CCR6 segregate with the commitment of $\gamma\delta$ T cells to produce IFN- γ and IL-17A, respectively [12,34–36]. In the present study, we also indicated that NK1.1⁺ $\gamma\delta$ T cells showed large amounts of IFN- γ production and were attenuated by bleomycin-induced pulmonary fibrosis through the suppression of pulmonary Th17 cell activities. However, NK1.1⁺ $\gamma\delta$ T cells derived from IFN- $\gamma^{-/-}$ mice did not attenuate the progression of lung fibrosis. Following exposure to bleomycin, we showed an increase in pulmonary IFN- γ^{+} $\gamma\delta$ T cells in WT mice. Previously, Haas *et al.* [35] and our group [8] showed that $\gamma\delta$ T cells can produce large amounts of IFN- γ in response to various stimuli, such as TCR, IL-12 and IL-18. Bleomycin-induced pulmonary fibrosis is characterized by over-expression of IL-12 and IL-18 in lung tissues [37,38]. Therefore, IFN- γ seems to be induced by pulmonary $\gamma\delta$ T cells through IL-12 or IL-18 and it seems to suppress the pulmonary Th17 cells response in bleomycin-induced pulmonary fibrosis.

Secondly, does excess production of fibrotic factors play a role in exacerbation of pulmonary fibrosis in TCR $\delta^{-/-}$ mice? TGF- β is known as one of major profibrotic factors *in vivo* and *in vitro* [39]. After bleomycin exposure, the expression of pulmonary TGF- β mRNA and collagen production increased in TCR $\delta^{-/-}$ mice compared with WT mice. Previous reports showed that TGF- β induced fibroblast proliferation and collagen production in bleomycin-induced pulmonary fibrosis [40]. Conversely, inhibition of TGF- β signalling by TGF- β neutralizing antibodies or TGF- β receptor blockers attenuated bleomycin-induced

pulmonary fibrosis [41,42]. Furthermore, Wilson *et al.* [21] reported that the profibrotic activity of TGF- β in the development of pulmonary fibrosis may be attributed to the induction of IL-17A from T cells. In TCR $\delta^{-/-}$ mice treated with bleomycin, the increase in pulmonary IL-17A⁺ CD4⁺ T cells could be one possible reason for the excessive production of TGF- β . Consequently, over-expression of TGF- β might induce collagen synthesis and cause exacerbation of pulmonary fibrosis in TCR $\delta^{-/-}$ mice.

After bleomycin exposure, myeloid cells infiltrated into the BALF by several chemokines and exacerbated pulmonary fibrosis [43]. It was reported that $\gamma\delta$ T cells also produced chemokines such as CCL3, CCL5 and CXCL10 in several stimuli [11,44–46]. In the present study, we showed that $\gamma\delta$ T cells might induce the infiltration of dendritic cells and neutrophils into the BALF. However, the transfer of $\gamma\delta$ T cells in TCR $\delta^{-/-}$ mice attenuated bleomycin-induced pulmonary fibrosis. Thus, these results demonstrated that severe cell infiltrations in BALF did not cause the enhanced severity of fibrosis observed in TCR $\delta^{-/-}$ mice.

Thirdly, is insufficiency of regulatory T cells relative to Th17 cells important? In the present study, we showed that the proportion of pulmonary FoxP3⁺ CD4⁺ T cells was not significantly different between WT and TCR $\delta^{-/-}$ mice after bleomycin exposure. FoxP3⁺CD4⁺ T cells are T_{regs} and can suppress Th17 cell responses [47]. In peripheral tissues, the expression of FoxP3 in CD4⁺ T cells was induced in the presence of TGF- β [48]. However, pulmonary FoxP3⁺CD4⁺ T cells were not increased in TCR $\delta^{-/-}$ mice, even under the condition of TGF- β over-expression. Our findings demonstrated that FoxP3⁺CD4⁺ T cells did not play a role in the increase of Th17 cells in TCR $\delta^{-/-}$ mice. Thus, the regulatory role of $\gamma\delta$ T cells in the development of pulmonary fibrosis seem to be involved directly in the suppression of IL-17A production by CD4⁺ T cells.

In conclusion, our results demonstrated that $\gamma\delta$ T cells attenuated bleomycin-induced pulmonary fibrosis. The regulatory role of $\gamma\delta$ T cells in pulmonary fibrosis seems to be mediated through the suppression of IL-17A⁺CD4⁺ T cell activity via IFN- γ production. These findings should enhance our understanding of the pathogenesis of pulmonary fibrosis and that targeting $\gamma\delta$ T cells is a potentially useful therapeutic strategy in the management of pulmonary fibrosis.

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Disclosure

None declared.

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