



## Mycobacterium Vaccae Regulate $\gamma\delta$ T17 and $\gamma\delta$ Treg Cells in Mice Asthmatic Lung

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### ABSTRACT

**Background** Dysregulation of the balance between different T cell populations is believed to be an important basis for asthma.

**Objective** To observe the changes in  $\gamma\delta$ T subtypes in transgenic asthmatic mice after aerosol inhalation of *Mycobacterium vaccae*, and to further investigate the mechanism of *M. vaccae* in asthmatic mice and its relationship with  $\gamma\delta$ T cells.

**Methods** TCR- $\beta^{-/-}$  mice were exposed to atomized normal saline or *M. vaccae* for 5 days and the  $\gamma\delta$ T cells from the lung tissues were isolated. Changes in  $\gamma\delta$ T17 and  $\gamma\delta$ Treg populations were detected. Asthma was induced in BALB/c mice using ovalbumin, which was then transplanted with control or *M. vaccae*-primed  $\gamma\delta$ T cells. First we analyzed the content of  $\gamma\delta$ T cells that secrete IL-17 (IL-17  $\gamma\delta$ T cells) and Foxp3<sup>+</sup>  $\gamma\delta$ T cells in lung tissues and then measured the content of IL-17 in the bronchoalveolar lavage fluid (BALF) by ELISA.

**Results** Exposure to *M. vaccae* increased and decreased the relative proportions of Foxp3<sup>+</sup>  $\gamma\delta$ T cells and IL-17<sup>+</sup>  $\gamma\delta$ T cells, respectively, thereby decreasing airway reactivity and inflammation levels in asthmatic mice, and significantly decreasing IL-17 levels in BALF. Furthermore, mice treated with these primed T cells showed a decrease in IL-17<sup>+</sup>  $\gamma\delta$ T cells, and a concomitant increase in Foxp3<sup>+</sup>  $\gamma\delta$ T cells in their lung tissues. Furthermore, adoptive transfer of *M. vaccae*-primed  $\gamma\delta$ T cells decreased GATA3 and NICD and increased T-bet in lung.

**Conclusions** The *M. vaccae*-primed  $\gamma\delta$ T cells alleviated the symptoms of asthma by reversing Th2 polarization in the lungs and inhibiting the Notch/GATA3 pathway.

**Keywords:** Asthma, *Mycobacterium vaccae*,  $\gamma\delta$ T cells

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## INTRODUCTION

The regulatory imbalance between Th1/Th2 cells, such as the lack of Th1 cells and/or the appearance of overactive Th2 cells, is the most fundamental immunological mechanism in the pathogenesis of asthma (1, 2). Recent studies have shown that imbalances in other T cell populations, such as Th17/Treg cells are also involved in the development and progression of asthma (3, 4). In addition, the  $\gamma\delta$ T cell subsets are also involved in the pathogenesis of asthma, which can promote airway remodeling by increasing the production and releasing multiple inflammatory factors and cytokines, ultimately leading to the development of asthma (5, 6).  $\gamma\delta$ T cells release large amounts of Th2 cytokines such as IL-17, which recruits other granulocytes and exacerbates the inflammatory process (7, 8).

High levels of IL-17 are strongly associated with airway inflammation, reactivity, and airway remodeling in asthmatics (9-11). The  $\gamma\delta$ T17 cells are directly involved in the pathogenesis of ovalbumin (OVA)-induced asthma (12), which show an aberrant  $\gamma\delta$ T17/ $\gamma\delta$ Treg ratio (4). The  $\gamma\delta$ Tregs is an immunosuppressive  $\gamma\delta$ T cell subset of T cells manifested by  $\gamma\delta$ -T cell receptors (TCRs) and Foxp3 expression (13).

The T-bet/Stat-3 ratio is a decisive factor in Th1/Th2 polarization and stability (14), and an indicator of immune imbalance in asthmatic patients (15). A recent study showed that Shengjiang powder biased T cell polarization towards the Th2 phenotype in patients with sepsis patients by regulating T-bet and GATA3 (16). Furthermore, ectopic expression of T-bet and Foxp3 accompanied by GATA-3 and ROR $\gamma$ T knockdown inhibited the production of IL-17, IL-13, IL-5, and OVA-specific IgE in asthmatic mice, which inhibited eosinophil infiltration and alleviated airway inflammation (17). Consistent with this, Ginsenoside Rb alleviated allergic symptoms in OVA-induced asthmatic mice GATA3 and upregulated T-bet (18). Similarly, oral feeding

of *Lactobacillus bulgaricus* N45.10 reduced allergic asthma-triggered inflammation and airway remodeling in mice by upregulating T-bet and reducing GATA3 levels, which altered the proportion of Th1/Th2 cytokines (19). A recently concluded clinical trial reduced the severity of asthma by altering the T-bet/GATA3 ratio in the mononuclear cells of the airways involved in the inflammatory response (20). Finally, there is evidence that the Notch/Hes1 signaling pathway and IL-17 cytokines are closely associated with the development of  $\gamma\delta$ T cells (21). Th2 cell differentiation in asthma patients is mainly regulated through the Notch signaling (22, 23), while Treg cells can restore the balance between Th1 and Th2 cells through the Dll4/Notch signaling pathway and thus reverse the pathophysiological process of asthma (24). We have previously reviewed the role of the Notch signaling pathway in asthma and Th1/Th2 regulation (25).

*Mycobacterium* infection can inhibit airway inflammation and hyperresponsiveness in asthma (26). Inactivated *M. phlei* in BCG inhibits the Th2 subset and relieves the symptoms of asthma (27). Studies show that the inhalation of inactivated *M. phlei* aerosol can restore the Th1/Th2 balance in the murine asthma model and reduce the inflammatory lesions in the lungs (28-30). At the molecular level, inactivated *M. phlei* restored the T-bet/GATA-3 balance and inhibited Th2 cytokine secretion (31). Inactivated *M. phlei* reduced the immune response mediated by IL-17<sup>+</sup>  $\gamma\delta$ T cells, thus attenuating the pathophysiological changes in asthmatic mice (32). *Mycobacterium vaccae* and inactivated *Mycobacterium phlei* have similar functions, because they are both of the genus *Mycobacterium*, and have the same immunomodulatory effect as other T cells to regulate the immune response of the body, aerosol bacteria can stimulate the body's immune response through mucosal immunity, cellular immune response regulates airway inflammation and airway hyperresponsiveness. A recent

study showed that *M. vaccae* atomization can reduce the level of hyperresponsiveness and inflammatory factors in the airways and thus achieve the prevention of allergic bronchial asthma (33). To further elucidate the immunomodulatory mechanism of asthma, we exposed TCR- $\beta^{-/-}$  mice to atomized *M. vaccae* and transplanted the primed  $\gamma\delta$ T cells into OVA-induced asthmatic mice, and we wanted to observe the pathogenesis of asthma by analyzing changes in various immunological and molecular indices, and also research the functions of inactivated *M. vaccae* in asthmatic mice. *Mycobacterium* has a regulatory effect on  $\gamma\delta$ T. This article focuses on its regulatory effect on T-bet/GATA3 pathway and  $\gamma\delta$ T17/ $\gamma\delta$ Treg.

## MATERIALS AND METHODS

Male TCR- $\beta^{-/-}$  mice (Mice with the  $\alpha\beta$ T gene removed and  $\gamma\delta$ T cells present in mice) (Beijing Baiaosaitu Company) and male BALB/c mice (Hunan Slake Jingda Experimental Animal Co). All animals (4-6 weeks, weighing 18-22 g) grew in a specific pathogen-free (SPF) environment and survived according to the feeding requirements of the Animal Welfare Act. Ethical approval for this study was obtained from the Animal Ethics Committee of Guangxi Medical University (IACUC: 201711033.)

The TCR- $\beta^{-/-}$  mice were randomized into the control and *M. vaccae*-infected groups (n=20 each). Therefore, the mice were exposed to *M. vaccae* (22.5  $\mu$ g *M. vaccae* in 10 ml normal saline) or the same volume of normal saline for 20 min per day for 5 consecutive days. The lung tissue was collected to isolate the  $\gamma\delta$ T cells. The isolated  $\gamma\delta$ T cells were stored in PBS solution and placed on ice at 4°C to detect the activity of  $\gamma\delta$ T cells to ensure that the  $\gamma\delta$ T cells were active. The removed  $\gamma\delta$ T cells were immediately transfused into BALB/c mice (Not TCR- $\beta^{-/-}$ ). The BALB/c mice were randomly divided into the normal control, asthma model, and

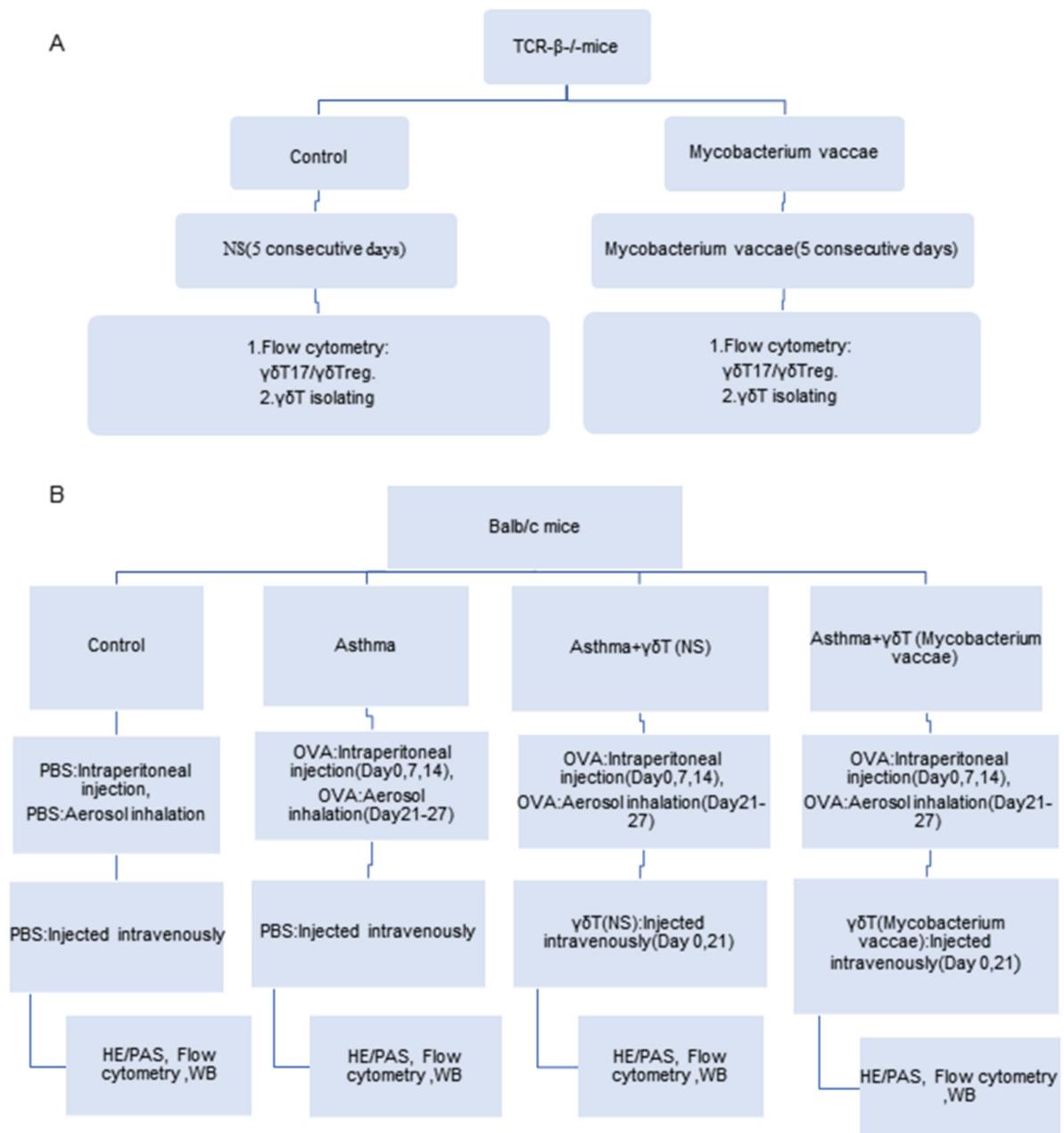
the  $\gamma\delta$ T cell control groups (TCR- $\beta^{-/-}$  mice were exposed to 10ml normal saline for 5 consecutive days, 20 min daily, and  $\gamma\delta$ T cells were isolated from lung tissue, injected to the BALB/c mice of OVA-induced asthma) and sensitized  $\gamma\delta$ T cell adoptive transfer groups (TCR- $\beta^{-/-}$  mice were exposed to *M. vaccae* (22.5  $\mu$ g *M. vaccae* in 10 ml normal saline) for 5 consecutive days for 20 min every day, and lung tissues were isolated from  $\gamma\delta$ T cells, injected to the BALB/c mice of OVA-induced asthma) 38 (n=8 each). To construct the asthma model, we injected 25 mg OVA (Sigma-Aldrich, USA) intraperitoneally on days 0, 7, and 14, respectively. From day 22, OVA at a concentration of 2% (mg/mL) was released in a closed chamber for 7 consecutive days. The mice in adoptive transfer groups T cells were injected intravenously with  $10^6$   $\gamma\delta$ T cells extracted from the *M. vaccae* or saline-treated TCR- $\beta^{-/-}$  mice before the first intraperitoneal injection and on the first day of atomization. Lung tissues were harvested for testing (Figure 1). Therefore, lung tissues, bronchio-alveolar lavage fluid (BALF), and lung tissue were harvested for various tests.

### Airway Reactivity

The mice in each experimental group were placed on specific test devices to detect specific airway resistance (sRaw) after stimulation with 6.25 mg/ml, 12.5 mg/ml, 25 mg/ml, and 50 mg/ml of methacholine and PBS, respectively.

### Enzyme-linked Immunosorbent Assay (ELISA)

After 40 minutes of airway reactivity test, the animals were anesthetized with 10% (300 mg/kg) chloral hydrate intraperitoneal injection, no symptoms of peritoneal inflammation in mice were observed, when mouse movement decreased, and suffering was considered humane endpoints that required immediate intervention and sudden death of cervical dislocation. No symptoms of peritoneal inflammation in mice were seen. The animals were intubated and 500  $\mu$ M of



**Figure 1.** The flow chart of TCR- $\beta^{-/-}$  mice. The mice were randomly divided into 2 groups and intervene shown above. In the *Mycobacterium vaccae* group, mice were Aerosol bacteria with *Mycobacterium vaccae* over 5 consecutive days, then, flow cytometry was performed to detect  $\gamma\delta T17/\gamma\delta Treg$  and  $\gamma\delta T$  isolating. The control group was intervened with NS. Figure 1-1. The flow chart of OVA sensitization model of the asthmatic BALB/C mice. The mice were randomly divided into 4 groups and treated shown above. In the asthmatic group, mice were sensitized with 25 $\mu$ g of OVA emulsified in 1mg of aluminum hydroxide in a total volume of 200 $\mu$ l on days 0, 7, and 14 by intraperitoneal administration. Beginning on the 21st day, the mice were challenged with 2% OVA for 30 min per day by an ultrasonic nebulizer in a closed chamber for 7d to establish models. The group C was intervened with  $\gamma\delta T$  cell (NS) for injected intravenously on day 0, 21. The group D was intervened with  $\gamma\delta T$  cell (*Mycobacterium vaccae*) for injected intravenously on day 0, 21. The control group was treated with PBS instead of OVA at the sensitization and challenge stages.

precooled PBS was slowly injected into the trachea before BALF was removed and stored in liquid nitrogen. IL-17 levels were measured

using an ELISA kit (Multi-Sciences: Wuhan Magee Biotechnology Co., Ltd.) according to the instructions.

### Histopathology

4% paraformaldehyde was used to fix lung tissue, followed by paraffin embedding and sectioning. The 4  $\mu$ m-thick sections were stained with hematoxylin-eosin (HE) and periodic acid-Schiff (PAS) according to standard protocols. The extent of pulmonary inflammation was observed under a light microscope and classified according to the semiquantitative scoring method (34). The sections were sequentially placed in xylene and gradient alcohol, followed by HE and immunofluorescence staining. To perform immunofluorescence staining, the corresponding primary antibodies were added: polyclonal goat anti-mouse T-bet antibody (1:1000; ab91109, Abcam), GATA3 antibody (1:1500; ab106625, Abcam), incubated overnight in a wet box, and the next day horseradish peroxidase-labeled secondary antibody (1: 800; ab205719, Abcam) was added, incubated for 2 h at room temperature. After being washed, they were observed under a fluorescent microscope and photographed.

### Flow Cytometry

The lung cell precipitate was resuspended in serum-supplemented RPMI 1640 medium and diluted to a density of 10<sup>9</sup>/ml, and incubated with 25ng/ml PMA, 1ng/ml ionomycin, and monensin at room temperature for 4 hours. The cells were harvested, washed with PBS, and stained with PerCP-Cy5-5 anti-CD3 and APC anti- $\gamma\delta$ T antibodies at 4°C for 30 minutes, fixed and permeabilized at 4°C for 20 minutes, stained with PE anti-IL-17 or PE anti-Foxp3 antibody at 4°C in the dark for 30 min, resuspended in 200  $\mu$ l PBS, and analyzed by flow cytometry with FlowJo 7.6 software.

### Western Blotting

The prepared cold RIPA lysate containing PMSF was added to the lung tissue, followed by homogenization. The precipitate was discarded after centrifugation and the supernatant was taken for determination of

the protein concentration while boiling with the addition of 5XLoading buffer. A 10% separation gel was used for SDS-PAGE, followed by the transfer of protein bands to PVDF membranes. 5% nonfat milk was used to block and primary antibodies (anti-GAPDH (1:5000), anti-NICD (1:2500), anti-GATA3 (1:5000), and anti-T-bet (1:5000) were added after completion, and incubated at 4°C overnight, TBST was washed 3 times and then incubated for 2 hours at room temperature by adding the corresponding labeled secondary antibody (1:1000). Licor Odyssey software was used for the analysis of positive bands.

### Statistical Analysis

SPSS software (version 22.0) was used to perform a statistical analysis of the data. The data were expressed as mean $\pm$ standard deviation (mean $\pm$ SD). All experiments were carried out independently at least three times. The unpaired t-test was used for the comparison between the two data groups. One-way analysis of variance (ANOVA) was used to analyze the differences between three and more groups, and the Bonferroni test was used for pairwise comparisons between the groups. Pearson's correlation was used to measure the correlation between the samples. P value<0.05 was considered statistically significant.

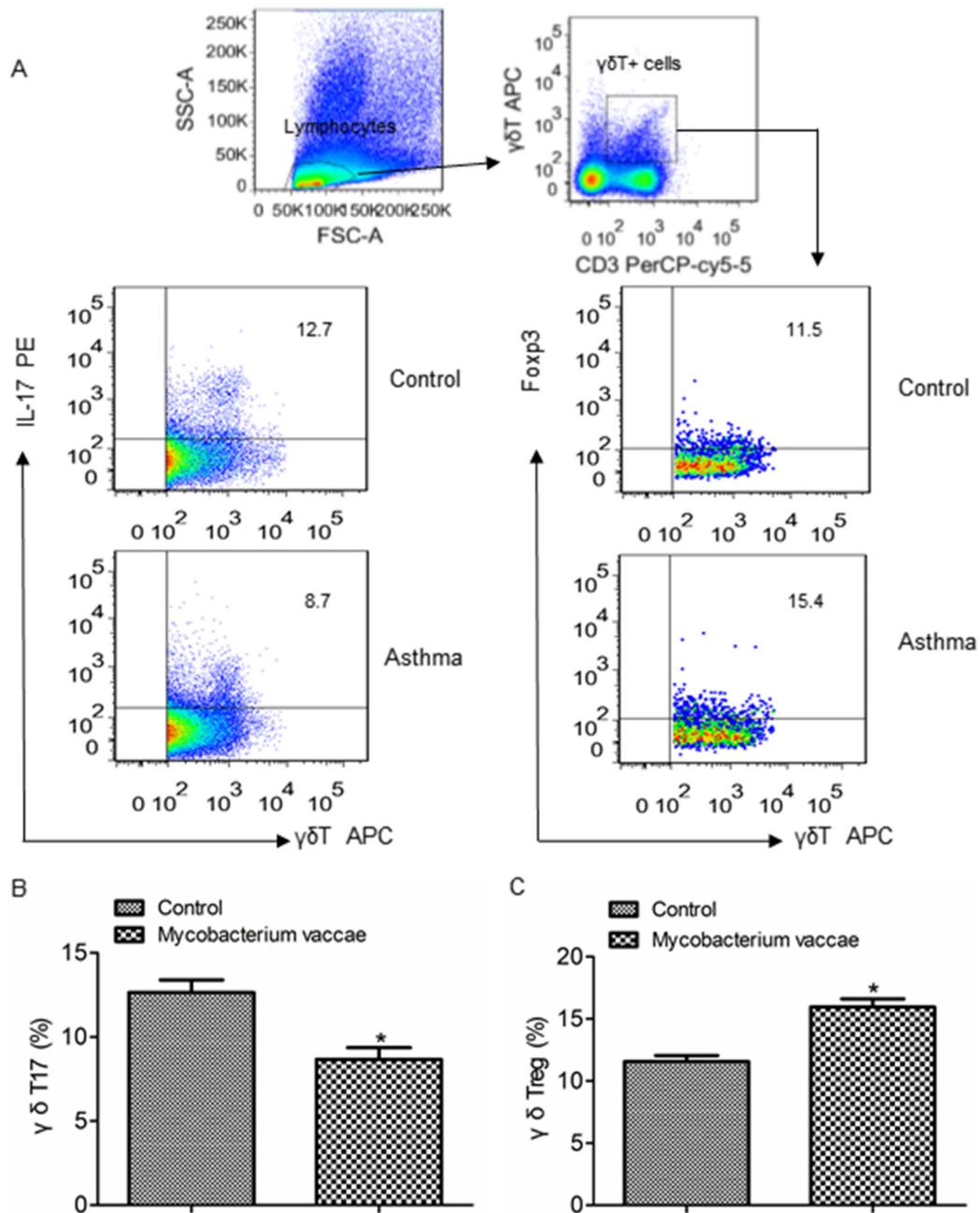
## RESULTS

### Percentage of $\gamma\delta$ Treg and $\gamma\delta$ T17 Cells

The TCR- $\beta^{-/-}$  mice exposed to *M. vaccae* had fewer  $\gamma\delta$ T17 and more  $\gamma\delta$ Treg cells in their lung tissue compared with the control group (Figure 2), indicating that the bacteria can skew the  $\gamma\delta$ T17/ $\gamma\delta$ Treg ratio.

### Airway Reactivity

As shown in Figure 3-A, the asthmatic mice had significantly higher airway responsiveness after MCH stimulation compared with the control animals (P<0.05), which was markedly decreased by *M.*



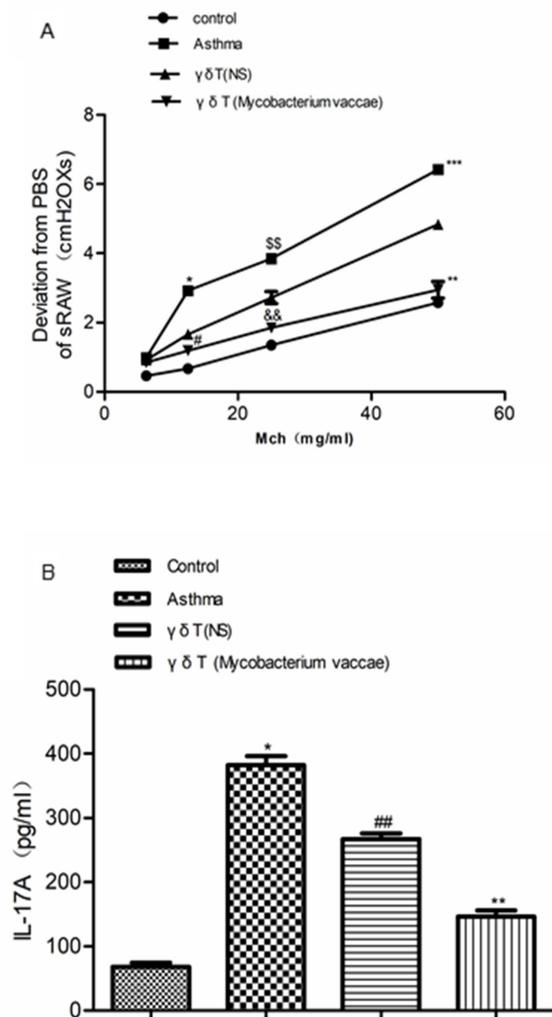
**Figure 2.** Flow cytometry plots showing relative percentages of lung tissues IL-17+ cell percentages and Treg+ $\gamma\delta$ T cell percentages in the control and infected mice. n = 20, \*p<0.05 vs control.

*vaccae*-primed  $\gamma\delta$ T cells (P<0.05). On the contrary, non-primed control  $\gamma\delta$ T cells did not affect the response to MCH (P>0.05). *M. vaccae*-sensitized  $\gamma\delta$ T cells alleviated airway reactivity and inflammation in OVA-exposed mice.

### Lung Histopathology

The control mice had regular bronchial morphology with normal alveolar interval and

little infiltration of inflammatory cells around the bronchus and blood vessels, no epithelial hyperplasia or thickening of the bronchial wall was observed. Asthmatic stimulation increased the infiltration of inflammatory cells, which narrowed the bronchial lumen and thickened its wall. On the contrary, adoptive transfer of naive T cells (NS) or *M. vaccae*-primed  $\gamma\delta$ T cells mitigated the above patho-inflammatory changes, and the



**Figure 3.** SRaw growth curve showing the airway responsiveness of OVA-stimulated mice after stimulation with 12.5mg/ml, 25mg/ml and 50mg/ml MCH, n=8, \*P<0.05, #P<0.05, <sup>SS</sup>P<0.01, <sup>\*\*\*</sup>P<0.001, vs control, #P<0.05, <sup>&&</sup>P<0.01, <sup>\*\*</sup>P<0.01 vs asthma (Figure 3-A). Effect of  $\gamma\delta$ T cells on BALF IL-17 level in asthmatic mice. Results of ELISA indicating the amount of IL-17A released in BALF in (a) control, (b) asthma, (c) non-primed  $\gamma\delta$ T cells groups and (d) *M. vaccae*-primed  $\gamma\delta$ T cells. N=8, \*P<0.05 vs control, \*\* P<0.01 vs asthma, <sup>##</sup> P>0.05 vs asthma (Figure 3-B).

primed cells resulted in a more obvious effect. Furthermore, PAS staining showed excessive mucus exudation and goblet cell hyperplasia in asthmatic mice compared with only a few intraluminal bronchial goblet cells and a lack of mucus leakage in the healthy controls. T cells primed with nave and *M. vaccae*-primed  $\gamma\delta$ T cells decreased the amount of mucous in the bronchi and perivascular space, and the

change was more obvious with the primed cells (Figure 4, Table 1).

### IL-17 Levels

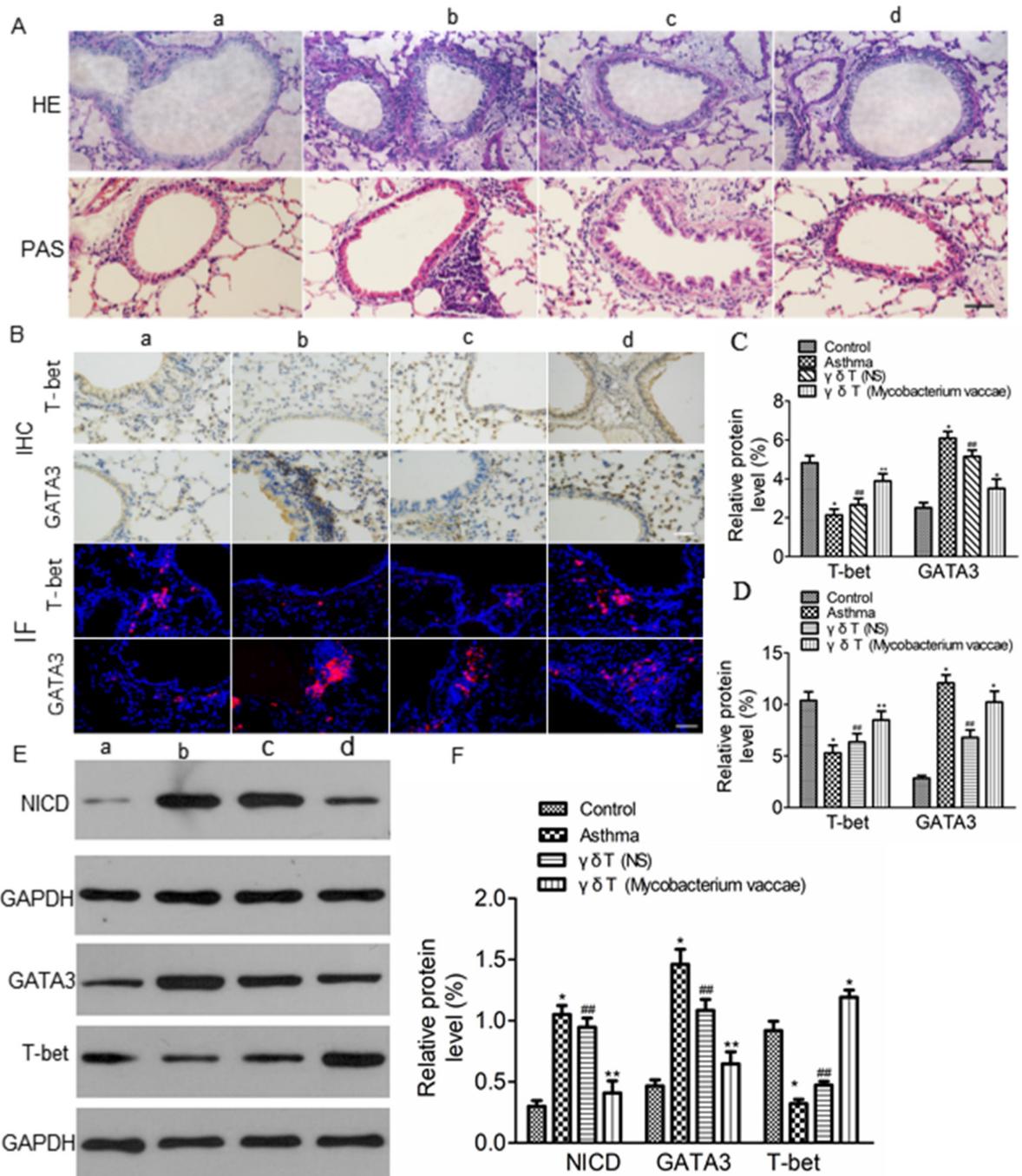
Consistent with histopathological findings, IL-17A levels were significantly higher in BALF of asthmatic mice compared with the normal controls and were restored to almost baseline levels by adoptive transfer of *M. vaccae*-primed  $\gamma\delta$ T cells (Figure 4, P<0.05). On the contrary, non-primed  $\gamma\delta$ T cells had no ameliorative effect on BALF IL-17 levels (P<0.05, (Figure 3).

### *M. Vaccae*-primed $\gamma\delta$ T Cells Reversed Th2 Polarization in Asthmatic Mice

Compared with the control mice, the in situ expression of T-bet was significantly lower and that of GATA-3 and NICD was significantly higher in the lung tissues of asthmatic mice (P<0.05 for both). Consistent with the findings so far, T cells treated with *M. vaccae*-primed  $\gamma\delta$ T cells reversed the levels of both in these mice (P<0.05), whereas the control  $\gamma\delta$ T cells had no significant effect (P>0.05; Figure 4).

### Expression of IL-17 + $\gamma\delta$ T and Foxp3 + $\gamma\delta$ T

Asthmatic stimulation significantly increased the number of IL-17<sup>+</sup>  $\gamma\delta$ T cells in lung tissues compared with the control mice (23±3.1 vs 10±3.7; P<0.05), which was decreased by *M. vaccae*-primed  $\gamma\delta$ T cells (13±2.8; P<0.05). The Foxp3<sup>+</sup>  $\gamma\delta$ T cell count was lower in asthmatic mice compared with the healthy mice (2.4±0.6 vs 5.4±0.5; P<0.05), and restored after adoptive transfer of *M. vaccae*-primed  $\gamma\delta$ T cells (4.5±1.4 vs 2.4±0.6; P<0.05) but not the control  $\gamma\delta$ T cells (2.7±0.6 vs 2.4±0.6; P>0.05). Then, a positive correlation was established between the relative levels of NICD protein and the percentage of IL-17<sup>+</sup>  $\gamma\delta$ T cells (r=0.56, P<0.001). This strongly indicates that the  $\gamma\delta$ T17/ $\gamma\delta$ Treg ratio alters upon asthma induction to a more Th2 phenotype, and *M. vaccae*-primed  $\gamma\delta$ T cells restore this balance and reverse Th2 polarization. Furthermore, we also observed



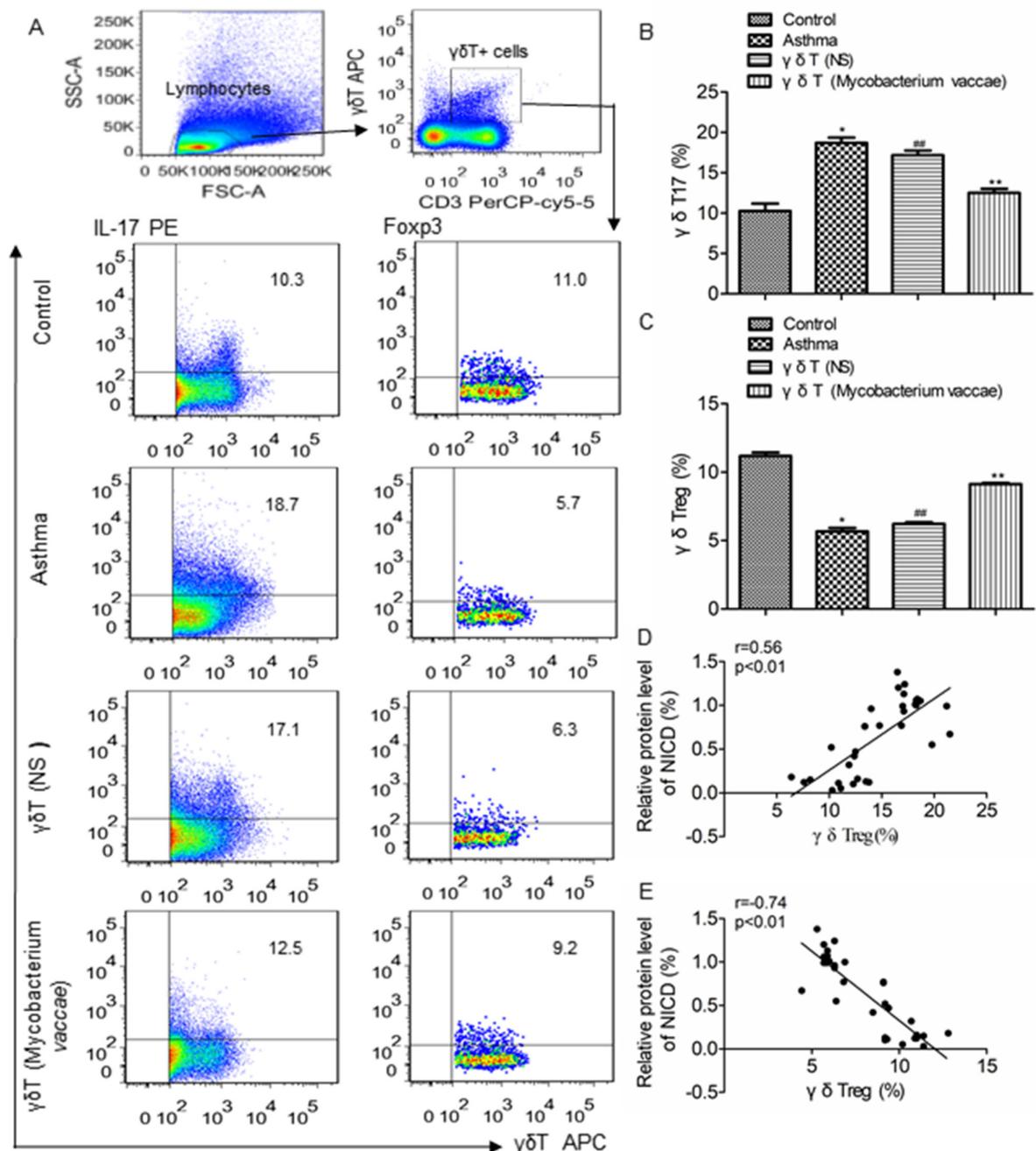
**Figure 4.** Effect of  $\gamma\delta T$  cells on the lung tissues of asthmatic mice. Representative images of HE-stained lung tissues showing the bronchio-alveolar architecture of the different animal groups. Original magnification – 400x, scale bar - 25 $\mu$ m). Representative images of PAS-stained tissues showing goblet cells and mucus exudation in the airway epithelium of different groups. Original magnification – 200x, scale bar - 50 $\mu$ m), n=8 (Figure 4-A). Representative IHC images (Figure 4-B,C) and IF images (Figure 4-B,D) showing the in situ expression intensity of T-bet and GATA3 and WB images (E,F) showing T-bet, GATA3 and NICD protein levels in the lung tissues of the different groups. \*P<0.05 vs control, \*\*P<0.01 vs asthma, \*\*P>0.05 vs asthma. Original magnification – 200x, scale bar - 50 $\mu$ m), n=8.

a significant negative correlation between the relative levels of NICD protein and the percentage of Foxp3<sup>+</sup>  $\gamma\delta T$  cells (r=-0.74, P<0.01). Taken together, *M. vaccae* alleviates

the symptoms of OVA-induced asthma by lowering the balance towards the  $\gamma\delta T$ reg cells through the inhibition of the Notch pathway. (Figure 5).

**Table 1. Airway inflammation scores and mucus scores**

Group	Inflammation scores					Mucus scores				
	0	1	2	3	4	0	1	2	3	4
a	8	0	0	0	0	7	1	0	0	0
b	0	0	0	3	5	0	0	0	3	5
c	0	0	2	3	3	0	0	2	2	4
d	0	4	2	2	0	0	4	3	1	0



**Figure 5.** Flow cytometry plots showing the percentage of IL-17<sup>+</sup> γδT cells and Foxp3<sup>+</sup> γδT cells in the lung tissues of different groups. Foxp3<sup>+</sup> γδT cells in the lung tissues of different groups (Figure 5-A,B,C). Correlation between NICD protein levels in lung tissue and the percentage of IL-17<sup>+</sup> γδT cells ( $r=0.56$ ,  $P<0.001$ ) is shown (Figure 5-D). Correlation between NICD protein levels in lung tissue and the percentage of Foxp3<sup>+</sup> γδT cells is also shown ( $r=-0.74$ ,  $P<0.01$ ) (Figure 5-E). \* $P<0.05$  vs control, \*\* $P<0.01$  vs asthma, \*\* $P>0.05$  vs asthma. N=8.

## DISCUSSION

The pathogenesis of asthma is still not fully understood, and current studies suggest that immune T cells are closely associated with it. Among them, the Th1/Th2 imbalance of  $\gamma\delta$  T cells is widely accepted to be involved in the development and progression of asthma (34). Not only in asthma disease but also other allergic and autoimmune diseases, IL-17<sup>+</sup>  $\gamma\delta$ T and Treg subsets are involved in varying degrees (35). The  $\gamma\delta$ T17 cells, in particular, drive airway remodeling and hyperresponsiveness in asthma by secreting inflammatory cytokines and activating neutrophils. GATA3 is the key Th2 transcription factor (36) that promotes the synthesis of Th2 cytokines (37). On the other hand, the T-bet triggers Th1 differentiation by upregulating IFN- $\gamma$ . The Th1/Th2 imbalance in asthma is the result of an aberrant T-bet/GATA3 ratio, which causes hypersecretion of Th2-type cytokines and triggers airway inflammation.

We primed the  $\gamma\delta$ T cells in TCR- $\beta^{-/-}$  mice with *M. vaccae* and transplanted these cells into an OVA-induced asthma model. T cells treated with *M. vaccae*-primed  $\gamma\delta$ T cells not only alleviated airway hyperresponsiveness and inflammatory damage in lung tissues of asthmatic mice but also decreased IL-17 levels in BALF. Furthermore, the mice treated with these primed  $\gamma\delta$ T cells also showed a higher proportion of Foxp3<sup>+</sup> relative to IL-17<sup>+</sup> T cells in their lung tissues, which corresponded to a higher ratio of T-bet / GATA3. In addition, T cells treated with *M. vaccae*-primed  $\gamma\delta$ T cells inhibited the Notch pathway in inflamed lungs, as indicated by the decreased levels of NICD. These results are consistent with the antiasthmatic effects of *Mycobacteria* and its culture supernatants, as well as those of the BCG vaccine, in mice models (38, 39). We hypothesize that the improvement effect of primed T cells was directly due to the cells injected themselves, which is supported by the involvement of  $\gamma\delta$ T17/ $\gamma\delta$ Treg cells in the

occurrence and development of asthma (40), or indirectly via inhibition of the Notch/GATA3 pathway and Th2 polarization. The specific regulatory mechanism will have to be verified in subsequent studies.

*M. vaccae* and *M. phlei* have an immunomodulatory effect on their hosts. Aerosol bacteria can stimulate mucosal immunity, which triggers airway inflammation and hyperresponsiveness. A previous meta-analysis showed that *M. vaccae* immunotherapy can effectively treat pulmonary TB (41). Another study showed that *M. vaccae* administered before chronic subordinate colony housing (CSC) strongly promoted active stress coping (42). These experimental data indicate the clinical application of *M. vaccae*. The imbalance of  $\gamma\delta$ T17/ $\gamma\delta$ Tregs is correlated with the pathogenesis of OVA-induced asthma (4). Inhalation of inactivated *M. phlei* can correct the Th1/Th2  $\gamma\delta$ T cell balance in asthma (43), thereby reducing lung inflammatory lesions. Consistent with this, aerosol bacteria from inactivated *M. phlei* reduced airway inflammation in the asthmatic mice, upregulated T-bet and GATA-3 mRNA levels, and inhibited Th2 cytokine secretion (31). In this study, aerosolized *M. vaccae* decreased the proportion of  $\gamma\delta$ T17 cells and increased that of  $\gamma\delta$ Tregs, which reduced AHR, inflammation, and IL-17 levels in BALF. The mechanism was first the direct effect of injected  $\gamma\delta$ T cells on asthma. Second, the effect of infused Treg on the infused  $\gamma\delta$ Treg in the mouse body can inhibit the Notch pathway, regulate Th1/Th2 differentiation, and then regulate the proportion of T-bet / GATA3. Our findings provide the theoretical basis for new therapeutic approaches against asthma.

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**Conflict of Interest:** None declared.

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