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ORIGINAL ARTICLE

The histone deacetylase inhibitor trichostatin A suppresses murine innate allergic inflammation by blocking group 2 innate lymphoid cell (ILC2) activation

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ABSTRACT

Background Group 2 innate lymphoid cells (ILC2) are an important source of the type 2 cytokines interleukin (IL)-5 and IL-13 that are critical to the allergic airway phenotype. Previous studies reported that histone deacetylase (HDAC) inhibition by trichostatin A (TSA) downregulated adaptive allergic immune responses; however, the effect of HDAC inhibition on the early innate allergic immune response is unknown. Therefore, we investigated the effect of TSA on innate airway inflammation mediated by ILC2 activation.

Methods BALB/c mice were challenged intranasally with *Alternaria* extract, exogenous recombinant mouse IL-33 (rmIL-33) or the respective vehicles for four consecutive days following TSA or vehicle treatment. Bronchoalveolar lavage (BAL) fluids and lungs were harvested 24 h after the last challenge.

Results We found that TSA treatment significantly decreased the number of ILC2 expressing IL-5 and IL-13 in the lungs challenged with *Alternaria* extract or rmIL-33 compared with vehicle treatment ($p<0.05$). TSA treatment significantly decreased protein expression of IL-5, IL-13, CCL11 and CCL24 in the lung homogenates from *Alternaria* extract-challenged mice or rmIL-33-challenged mice compared with vehicle treatment ($p<0.05$). Further, TSA treatment significantly decreased the number of perivascular eosinophils and mucus production in the large airways that are critical components of the asthma phenotype ($p<0.05$). TSA did not change early IL-33 release in the BAL fluids; however, TSA decreased lung IL-33 expression from epithelial cells 24 h after last *Alternaria* extract challenge compared with vehicle treatment ($p<0.05$).

Conclusions These results reveal that TSA reduces allergen-induced ILC2 activation and the early innate immune responses to an inhaled protease-containing aeroallergen.

INTRODUCTION

Asthma is a chronic lung disorder that is characterised by inflammation and swelling in the airway that results in mild-to-severe breathing difficulty during an exacerbation.¹ The mechanisms that initiate allergic immune responses in asthma development are incompletely understood. The T-helper (Th) 2 cytokines, IL-4, IL-5, IL-9 and IL-13 are

Key messages

What is the key question?

► Does histone deacetylase (HDAC) inhibition suppress the early immune response to an inhaled aeroallergen by blocking type 2 immune responses mediated by group 2 innate lymphoid cells (ILC2)?

What is the bottom line?

► Trichostatin A downregulated the number of ILC2 expressing IL-5 and IL-13 following *Alternaria* extract challenge with consequent attenuation of lung eosinophilia and mucus hypersecretion.

Why read on?

► This is the first report that an HDAC inhibitor suppressed innate immune-mediated allergic inflammation, identifying a novel target for therapeutics in allergen-induced lung diseases.

critical in the pathogenesis of allergic asthma as these cytokines regulate the Th2-type response including B cell class switching to IgE production, eosinophilia, mucus production, and bronchoconstriction.^{2–3} The main source of these cytokines was long considered to be Th2 cells produced in the setting of adaptive immune responses. Recently, alternative ways to induce Th2 cytokine production have been discovered in both mouse and human studies.^{4–6} Group 2 innate lymphoid cells (ILC2) are resident in the lung and are also an important source of IL-5 and IL-13. ILC2 are lineage negative (Lin[−]) cells that do not have cell surface expression of T cell, B cell, macrophage, dendritic cell (DC), neutrophil or natural killer (NK) cell markers, yet these cells have surface expression of ST2 (IL-33 receptor), CD25 (IL-2R α) and CD127 (IL-7R α).⁵ ILC2 are activated by epithelial-derived cytokines such as IL-33, TSLP, IL-25 or their combination, which results in ILC2 production of IL-5 and IL-13. In vivo mouse studies revealed that proteases or aeroallergens containing protease activity, such as *Alternaria alternata* induced IL-33 release from

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type II epithelial cells.⁷ In addition, exogenous IL-33 administration induced Th2-type airway inflammation mediated by ILC2, but not CD4⁺ T cells.⁸

Epigenetic regulation of gene expression has been reported as a novel approach to treat allergy and other immune-mediated diseases.^{9–10} One epigenetic mechanism for regulating gene expression is the acetylation and deacetylation of histone by histone acetyltransferases (HATs) and histone deacetylases (HDACs), respectively. The acetylation of histones by HATs increases DNA accessibility and promotes gene expression, whereas the removal of acetyl groups by HDACs has the opposite effect.¹¹ There are 11 isoforms of HDACs and there are also many non-histone protein substrates of HDACs in addition to histone proteins.¹² Compounds that have the ability to inhibit multiple HDAC isoforms are called pan-HDAC inhibitors (HDACi). Currently, there are three pan-HDACi drugs, vorinostat, romidepsin, and belinostat, that have been approved by the US Food and Drug Administration (FDA) for cancer treatment.¹³ Trichostatin A (TSA) is not approved by FDA, but is a pan-HDACi that inhibits enzyme activity of the HDAC 1, 2, 3, 4, 6 and 10 isoforms.

Several studies have indicated that TSA enhances inflammatory gene expression mediated by nuclear factor-kappa B (NF-κB).^{14–16} In contrast, many studies indicate that HDACi suppressed expression of proinflammatory genes.⁹ For instance, TSA repressed lipopolysaccharide (LPS)-induced protein expression of TNF-α, IL-6, IL-1β¹⁷ and IL-12p40¹⁸ in bone marrow derived macrophages. Further, TSA suppressed LPS-induced or poly I:C-induced mRNA expression of proinflammatory cytokines and chemokines in human microglial cells.¹⁹ In vivo experiments, TSA administration during antigen sensitisation or challenge suppressed ovalbumin (OVA)-induced airway inflammation and hyperresponsiveness in a mouse model dependent upon adaptive immune responses^{20–21} and ameliorated airway constriction in an *Aspergillus fumigatus* antigen-sensitised and challenged allergic mouse model.²² However, no studies have investigated the effect of HDACi on the innate allergic inflammatory response that is mediated by ILC2.

We hypothesised that HDAC inhibition negatively regulates innate allergic immune responses through suppression of ILC2 function. To test this hypothesis, we investigated the effects of HDAC inhibition by TSA on *Alternaria* extract-induced innate allergic inflammation model in mice. In this study, we found that TSA treatment significantly decreased the number of ILC2-expressing IL-5 and IL-13 in the lungs of mice challenged with *Alternaria* extract compared with vehicle treatment. In addition, TSA treatment decreased protein expression of IL-5, IL-13, CCL11 (eotaxin) and CCL24 (eotaxin-2) in the bronchoalveolar lavage (BAL) fluid and lung homogenates compared with vehicle treatment. Moreover, TSA treatment significantly decreased the number of perivascular eosinophils and mucus production in the large airways that are critical components of the allergic asthma phenotype. Furthermore, TSA treatment significantly decreased the number of ILC2-expressing IL-5 and IL-13 in the lungs and the protein expression of IL-5, IL-13, CCL11 and CCL24 in the lung homogenates of mice challenged with exogenous recombinant mouse IL-33 (rmIL-33) compared with vehicle treatment.

METHODS

Mice

Eight-week to 10-week-old female wild-type (WT) BALB/c mice were obtained from Charles River Laboratories. IL-33^{Citrine/+} reporter mice were generated by crossbreeding WT BALB/c

mice and IL-33^{Citrine/Citrine} mice.²³ Animal experiments were approved by the Institutional Animal Care and Use Committee at Vanderbilt University and were conducted according to the guidelines for the Care and Use of Laboratory Animals prepared by the Institute of Laboratory Animal Resources, National Research Council.

HDAC inhibitor treatment and *Alternaria* extract challenge or recombinant mouse IL-33 challenge

TSA (2.0 mg/kg in 50 μL dimethyl sulfoxide (DMSO); Selleckchem) or 50 μL DMSO as vehicle was administered subcutaneously to mice anaesthetised with isoflurane daily for 2 days prior to either *Alternaria* extract challenge (GREER), rmIL-33 challenge (Peprotech) or the respective vehicle challenge (figures 1A, 4A and 6A) and continued until the day prior to harvest. From day 0 to day 3, either 5 μg (protein amount) of *Alternaria* extract in 100 μL of phosphate buffered saline (PBS), 0.5 μg of rmIL-33 in 100 μL of 0.1% BSA/PBS or 100 μL of respective vehicles was used as an intranasal challenge to mice anaesthetised with ketamine/xylazine 1 h after TSA or its vehicle treatment. In an additional experiment, TSA or its vehicle was administered concomitantly with *Alternaria* extract challenge or PBS challenge (figure 8A). One hour or 24 h after the last challenge of *Alternaria* extract, rmIL-33 or the respective vehicles, the mice were sacrificed and BAL fluid and whole lungs were harvested for cytokine and chemokine measurements.

Flow cytometry

Lungs were harvested 24 h after the last challenge of *Alternaria* extract, rmIL-33 or the respective vehicles. The lungs were minced and digested with collagenase IV (1 mg/mL, Sigma-Aldrich) and DNase I (50 IU/mL, Sigma-Aldrich) in RPMI1640 with 5% fetal bovine serum (FBS) for 1 h at 37°C, and a single cell suspension was obtained by grinding and passing the digested lung through a 70 μm cell strainer. The cells were stimulated with phorbol 12-myristate 13-acetate (PMA) (10 ng/mL), ionomycin (1 μM) and BD GolgiStop (BD Biosciences) at the manufacturer's recommended concentration for 6 h. Then, both cell surface and intracellular cytokine staining were performed to enumerate IL-5⁺ and IL-13⁺ ILC2 and CD4T cells. ILC2 were identified as Lin[−] CD45⁺ CD25⁺ CD127⁺ cells. CD4⁺ T cells were identified as CD3⁺ CD4⁺ cells. To determine the IL-33-expressing cells (citrine reporter positive), cell surface staining was performed to distinguish the type of white blood cells and epithelial cells. CD4⁺ T cells were identified as CD3⁺ CD4⁺ cells. B cells were identified as CD3[−] CD19⁺ cells. Macrophages were identified as F4/80⁺ CD11c^{high} cells. DCs were identified as F4/80[−] CD11c⁺ cells. Eosinophils were identified as siglec F⁺ Gr-1^{int} cells. Granulocytes were identified as Gr-1^{high} cells. Epithelial cells were identified as CD45[−] EpCAM⁺ cells. All samples were measured on a BD LSR II Flow Cytometer and analysed using FlowJo Software.

Cytokine measurements

Quantikine ELISA or DuoSet ELISA kits from R&D were used to measure the concentration of cytokines and chemokines in BAL fluids and lung homogenates according to manufacturer's instructions. Values below the detection limit were assigned a value at half of the lower limit of detection to allow for statistical analysis.

Cell differential counting

Cytospin preparations of BAL fluid cells were performed for each sample. The slides were stained with a Richard-Allan

Figure 1 (A) Mice were challenged with *Alternaria* extract intranasally for four consecutive days. Trichostatin A (TSA) or its vehicle was administered subcutaneously on day -2 and -1, and then every 1 h before *Alternaria* extract challenge on day 0-3. (B) The number of IL-5-expressing and IL-13-expressing group 2 innate lymphoid cell (ILC2) in the lung from the mice challenged with PBS or *Alternaria* extract following vehicle or TSA treatment. (C) Representative plots of IL-5 and IL-13 intracellular staining in ILC2 in the lungs of mice challenged with PBS or *Alternaria* extract following vehicle or TSA treatment. The results are shown as mean±SEM of four mice in PBS-challenged groups and six mice in *Alternaria* extract-challenged groups. * $p < 0.05$.

Scientific Three-Step Stain Kit (Thermo Scientific) to determine cell differentials. Differential counts were based on counting 200 cells per slide.

Histological analyses of lung sections

Twenty-four hours after the last challenge of *Alternaria* extract or PBS, mice were sacrificed and the lung tissue was fixed in 10% formalin solution, paraffin-embedded, cut in 6 μ m sections, mounted and stained with anti-major basic protein (MBP)

antibody generously provided by Dr. James J. Lee (Mayo Clinic, Scottsdale, Arizona, USA) to evaluate eosinophils.²⁴ Slides were examined by a pathologist blinded to experimental groups to score eosinophil infiltration using a 0-4 scoring system: 0, no eosinophils; 1, a few circulating eosinophils and rarely one or two eosinophils are seen in perivascular spaces; 2, increased circulating eosinophils and eosinophils are prominent in perivascular spaces of large to medium vessels; 3, circulating eosinophils are prominent and eosinophilic perivascular cuffs are observed

Figure 2 Bronchoalveolar lavage (BAL) fluids and lungs were harvested 24 h after the last *Alternaria* extract challenge to measure the protein expression of IL-5, IL-13, CCL11, CCL17, CCL22 and CCL24. The results are shown as mean±SEM of four mice in PBS-challenged groups and six mice in *Alternaria* extract-challenged groups. *p<0.05.

in large, medium and small vessels; 4, increased circulating eosinophils, prominent perivascular cuffing by eosinophils in all vessels and interstitial infiltrates of eosinophils.

Paraffin-embedded lung tissue slices from mice sacrificed 48 h after the last challenge of *Alternaria* extract or PBS were stained with periodic acid–Schiff's (PAS) solution to visualise mucus and mucous-producing cells. Slides were scored by a pathologist

blinded to the experimental groups. All airways in the section were scored (15–40 airways) according to a five-point scale: 0, no PAS-positive cells; 1, <5% PAS-positive cells; 2, 5–10% PAS-positive cells; 3, 10–25% PAS-positive cells and 4, >25% PAS-positive cells.

Statistical analysis

All data were reported as mean±SEM. The data were compared by one-way analysis of variance (ANOVA) with Bonferroni-multiple pairs comparisons test using GraphPad Prism 5 (GraphPad Software). p Values <0.05 (p<0.05) were considered significant between two groups.

RESULTS

TSA treatment suppresses the number of mouse lung ILC2-producing IL-5 and IL-13 in response to airway *Alternaria* extract challenge

To establish the effect of TSA on cytokine production by ILC2, we enumerated the number of IL-5-expressing and IL-13-expressing lung ILC2 following either four consecutive days of PBS or *Alternaria* extract challenge in TSA-treated or vehicle-treated mice. The gating strategy for ILC2 is shown in online supplementary figure S1. ILC2 was identified as Lin[−]CD45⁺CD25⁺CD127⁺ cell. *Alternaria* extract challenge significantly increased the number of lung ILC2 that expressed either IL-5 (p<0.05) or IL-13 (p<0.05) compared with PBS-challenged mice (figure 1B). There was a statistically significant decrease in the number of IL-5-expressing and IL-13-expressing ILC2 (p<0.05) in the *Alternaria* extract-challenged mice treated with TSA compared with the *Alternaria* extract-challenged mice treated with vehicle for TSA (figure 1B). Figure 1C shows the representative flow cytometry dot plots of IL-5 and IL-13 intracellular staining in the lung ILC2 from mice challenged with PBS or *Alternaria* extract following TSA or vehicle treatment.

To determine the contribution of CD4⁺ T cells to the IL-5 and IL-13 expression 24 h after four consecutive days of *Alternaria* extract challenge, intracellular cytokine staining was performed by flow cytometry (see online supplementary figure S2). The number of CD3⁺CD4⁺IL-5⁺ and CD3⁺CD4⁺IL-13⁺ cells was significantly increased in *Alternaria* extract-challenged groups compared with PBS-challenged groups (p<0.05), while no change was detected in the number of CD3⁺CD4⁺IL-5⁺ and CD3⁺CD4⁺IL-13⁺ cells between TSA-treated group and vehicle-treated group in *Alternaria* extract-challenged mice. Online supplementary figure S2B shows the representative flow cytometry dot plots of IL-5 and IL-13 intracellular staining in the lung CD3⁺CD4⁺ T cells from mice challenged with PBS or *Alternaria* extract following TSA or vehicle treatment. In the *Alternaria* extract-challenged groups, there was a statistically significant 12-fold increase in the number of IL-5-expressing ILC2 (p<0.05) (figure 1B) compared with IL-5-expressing CD3⁺CD4⁺ cells (see online supplementary figure S2A), and a 2.5-fold increase in the number of IL-13-expressing ILC2 (p<0.05) (figure 1B) compared with IL-13-expressing CD3⁺CD4⁺ cells (see online supplementary figure S2A). These results indicate that ILC2, but not CD3⁺CD4⁺ T cells are the major source of IL-5 and IL-13 in response to this four consecutive days airway *Alternaria* extract-challenge protocol.

TSA treatment suppresses *Alternaria* extract-induced cytokine and chemokine expression in the lung and airway

We found that *Alternaria* extract challenge increased the number of IL-5-expressing and IL-13-expressing ILC2, and TSA

Figure 3 (A) Cell differentials from BAL fluids harvested 24 h after the last *Alternaria* extract challenge. (B) Representative sections and (C) eosinophil score as determined by anti-major basic protein (MBP) antibody staining 24 h after the last *Alternaria* extract challenge. (D) Representative sections and (E) mucus score as determined by periodic acid–Schiff's (PAS) staining 48 h after the last *Alternaria* extract challenge. The results are shown as mean±SEM of four mice in PBS-challenged groups and six mice in *Alternaria* extract-challenged groups. * $p < 0.05$.

treatment suppressed the *Alternaria* extract-induced cytokine expressing ILC2 compared with vehicle treatment. Next, we measured IL-5 and IL-13 protein in BAL fluid and lung homogenates. *Alternaria* extract challenge significantly increased the protein expression of IL-5 ($p < 0.05$) and IL-13 ($p < 0.05$) in both BAL fluids and lung homogenates compared with PBS-challenged groups. We found a statistically significant decrease in *Alternaria* extract-induced protein expression of IL-5 ($p < 0.05$) and IL-13 ($p < 0.05$) in both the BAL fluids and lung homogenates from TSA-treated mice compared with vehicle-treated mice (figure 2). Further, there was a statistically significant decrease in *Alternaria* extract-induced protein

expression of eosinophil-recruiting chemokines, CCL11 ($p < 0.05$) and CCL24 ($p < 0.05$), in both the BAL fluids and lung homogenates from TSA-treated mice compared with vehicle-treated mice. Meanwhile, there was no statistically significant difference in protein expression of Th2-recruiting chemokines, CCL17 (TARC) and CCL22 (MDC), between TSA-treated and vehicle-treated groups after *Alternaria* extract challenge (figure 2). Further, we measured proinflammatory cytokine and chemokine expression in BAL fluid and lung homogenates. *Alternaria* extract challenge significantly increased the protein expression of the neutrophil-recruiting chemokine KC (CXCL1) ($p < 0.05$) in both the BAL fluids and lung

Figure 4 (A) Mice were challenged with *Alternaria* extract intranasally for four consecutive days. Liraglutide or its vehicle was administered subcutaneously on day -2 and -1, and then every 4 h before and after *Alternaria* extract challenge on day 0–3. The BAL fluids and lungs were harvested 1 and 24 h, respectively, after last *Alternaria* extract challenge. The protein level of IL-33 (B and C) and IL-1 α (D and E) in the BAL fluids and lung homogenates was measured by ELISA. The results are shown as mean $\pm$ SEM of four mice in PBS-challenged groups and six mice in *Alternaria* extract-challenged groups. * p <0.05.

homogenates, TNF- α (p <0.05) in the BAL fluids and granulocyte macrophage colony stimulating factor (GM-CSF) (p <0.05) in the lung homogenates (see online supplementary figure S3). TSA treatment significantly decreased *Alternaria* extract-induced KC (p <0.05) and GM-CSF (p <0.05) in the lung homogenates, and TNF- α (p <0.05) in the BAL fluids compared with vehicle treatment (see online supplementary figure S3). These results suggest that TSA treatment suppresses *Alternaria* extract-induced innate allergic immune and inflammatory responses.

TSA treatment suppresses the number of perivascular eosinophils and mucus production in response to airway *Alternaria* extract challenge

Based on our finding that TSA treatment decreased protein expression of IL-5, CCL11, and CCL24 in both BAL fluid and lung homogenates following *Alternaria* extract challenge for four consecutive days, we hypothesised that the reduction of IL-5, CCL11, and CCL24 by TSA treatment decreases eosinophil accumulation in the lung following *Alternaria* extract challenge. To test this hypothesis, we measured BAL cell counts and differentials in *Alternaria* extract-challenged or PBS-challenged mice treated with TSA or vehicle. *Alternaria* extract challenge significantly increased the number of total BAL cells, eosinophils, lymphocytes and neutrophils compared with PBS-challenged groups (p <0.05). There was a statistically significant decrease in the number of total BAL cells, eosinophils, lymphocytes and neutrophils in the *Alternaria* extract-challenged mice treated with TSA compared with the *Alternaria* extract-challenged mice treated with vehicle (p <0.05) (figure 3A). However, there were no differences in the number of macrophages between the TSA-treated and vehicle-treated groups after *Alternaria* extract challenge. Next, we assessed perivascular eosinophils by immunohistopathology. Representative immunohistopathology sections stained with anti-MBP antibody are shown in

figure 3B. We found that there were very few eosinophils around alveolar spaces and no perivascular eosinophils following intranasal PBS challenge, while there was a significant increase in perivascular eosinophils (p <0.05) in the mice challenged with *Alternaria* extract and challenged with vehicle. However, TSA treatment resulted in a significant decrease of perivascular eosinophils (p <0.05) in the lung challenged with *Alternaria* extract (figure 3C).

We also found that TSA treatment decreased lung protein expression of IL-13, a critical inducer of airway mucus production,²⁵ in both BAL fluid and lung homogenates following *Alternaria* extract challenge (figure 2). Therefore, we tested the hypothesis that TSA treatment suppresses mucus production during innate immune responses to *Alternaria* extract. Representative histopathology sections stained with PAS solution are shown in figure 3D. There was no mucus or mucous-producing cells in the lungs of mice challenged with PBS. *Alternaria* extract challenge for four consecutive days significantly increased production and secretion of mucus (p <0.05) from large airway epithelial cells 48 h after the last *Alternaria* extract challenge. However, TSA treatment significantly decreased the production and secretion of mucus (p <0.05) compared with mice challenged with *Alternaria* extract treated with vehicle (figure 3E). Taken together, these results indicate that TSA treatment suppresses acute eosinophilic lung inflammation and airway mucus secretion in innate allergic responses to *Alternaria* extract challenge.

TSA treatment suppresses *Alternaria* extract-induced IL-33 in the lung homogenate, but not in BAL fluid

In this study, we found that TSA treatment suppressed the *Alternaria* extract-induced cytokine-expressing ILC2 compared with vehicle treatment. ILC2 are capable of producing large amounts of IL-5 and IL-13 following stimulation with

IL-33, a combination of IL-33 and IL-25, TSLP, IL-2 or IL-7.⁵

The full-length IL-33 that has biological activity is released in the extracellular space after cell damage such as necrosis.²⁶ Many airborne allergens contain proteases that induce rapid IL-33 release in BAL fluids after airway administration.^{7, 26} First, we measured IL-33 protein level in BAL fluid and lung homogenates in the time course experiment after the last *Alternaria* extract challenge in the four consecutive days challenge protocol. The IL-33 protein level in BAL fluid peaked at 1 h after last *Alternaria* extract challenge; thereafter, the IL-33 protein level was decreased in a time-dependent manner (see online supplementary figure S4). In contrast, the IL-33 protein level in the lung homogenates was increased in a time-dependent manner and peaked at 24 h. Therefore, we assayed IL-33 protein level in BAL fluid and lung homogenates 1 and 24 h, respectively, after the last *Alternaria* extract challenge to determine the effect of TSA treatment (figure 4A). *Alternaria* extract challenge significantly increased the protein level of IL-33 ($p<0.05$) in BAL fluid compared with PBS-challenged groups 1 h after the last challenge. However, the protein level of IL-33 in BAL fluid was not changed between TSA-treated and vehicle-treated groups with *Alternaria* extract challenge (figure 4B). On the other hand, we found a statistically significant decrease in *Alternaria* extract-induced protein expression of IL-33 ($p<0.05$) in lung homogenates from TSA-treated mice compared with vehicle-treated mice (figure 4C). These results suggest that TSA treatment suppresses the expression of IL-33 in the lung but does not suppress the IL-33 release in the airway.

It has been reported that IL-33 expression is controlled by IL-1 α .²⁷ We therefore measured IL-1 α protein level in the lung homogenate 24 h after the last *Alternaria* extract challenge. *Alternaria* extract challenge significantly increased the protein level of IL-1 α ($p<0.05$) in the lung homogenates compared with PBS challenge 1 h after the last challenge. We found a statistically significant decrease in *Alternaria* extract-induced protein expression of IL-1 α ($p<0.05$) in lung homogenates from TSA-treated mice compared with vehicle-treated mice (figure 4D). This result suggests that intracellular IL-1 α expression may be associated with IL-33 expression in the lung.

TSA treatment suppresses IL-33 expression in the lung epithelial cells

We found that TSA treatment suppressed protein expression of IL-33 in lung homogenate 24 h after the last *Alternaria* extract challenge. To investigate the localisation of IL-33 expression and the effect of TSA on per cell IL-33 expression level, we used IL-33^{Citrine/+} reporter mice. There was expression of citrine in the lung from naïve IL-33^{Citrine/+} reporter mice, but no citrine expression was observed in the lung from naïve WT mice (see online supplementary figure S5A). We found that CD45⁺ EpCAM⁺ cells expressed citrine fluorescence in PBS-challenged mouse lungs by flow cytometry (figure 5A,B). There was a significant increase in the mean fluorescent intensity (MFI), percentage and number of the lung CD45⁺ EpCAM⁺ citrine⁺ cells in the *Alternaria*-challenged mice treated with vehicle compared with PBS-challenged mice treated with vehicle ($p<0.05$). However, TSA treatment resulted in a decrease of the MFI, percentage and number in the lung CD45⁺ EpCAM⁺ citrine⁺ cells compared with *Alternaria* extract-challenged mice that were treated with vehicle ($p<0.05$) (figure 5B,C). In contrast, citrine fluorescence was not detected in T cells, B cells, DCs, macrophages, eosinophils or the other granulocytes (see online supplementary figure S5B). These results indicate that

TSA treatment reduced the number of lung epithelial cells expressing IL-33 following *Alternaria* extract challenge and decreased the amount of IL-33 expressed by individual cells in response to this same challenge.

TSA treatment suppresses the number of mouse lung ILC2-producing IL-5 and IL-13 in response to airway rmIL-33 challenge

In this study, there was no difference of IL-33 protein level in the BAL fluid between TSA-treated and vehicle-treated mice 1 h after last *Alternaria* extract challenge (figure 4B); meanwhile, TSA treatment suppressed the *Alternaria* extract-induced IL-5-expressing and IL-13-expressing ILC2 compared with vehicle treatment (figure 1B). These data suggest that TSA treatment decreases ILC2 activation by inhibiting IL-33-induced activation of ILC2, rather than by inhibiting *Alternaria* extract-induced IL-33 release by epithelial cells. Therefore, we tested the hypothesis that TSA treatment suppresses IL-33-mediated ILC2 activation in vivo. To determine the effect of TSA on IL-33-mediated ILC2 activation, we enumerated the number of IL-5-expressing and IL-13-expressing lung ILC2 following either four consecutive days of rmIL-33 challenge or the vehicle for rmIL-33 in TSA-treated or vehicle-treated mice (figure 6A). rmIL-33 challenge significantly increased the number of lung ILC2 that expressed either IL-5 or IL-13 compared with vehicle-challenged mice ($p<0.05$) (figure 6B). There was a statistically significant decrease in the number of IL-5-expressing and IL-13-expressing ILC2 in the rmIL-33-challenged mice treated with TSA compared with the rmIL-33-challenged mice treated with the vehicle for TSA ($p<0.05$) (figure 6B). Figure 6C shows the representative flow cytometry dot plots of IL-5 and IL-13 intracellular staining in the lung ILC2 from mice challenged with rmIL-33 or its vehicle following TSA treatment or TSA-vehicle treatment. This result indicates that TSA treatment inhibits IL-33-induced activation of ILC2.

TSA treatment suppresses rmIL-33-induced allergic pulmonary inflammation

We assessed cell differentials and cytokine and chemokine expression level in the BAL fluid and lung homogenates 24 h after rmIL-33 challenge for four consecutive days (figure 6A). IL-1 α protein expression was not changed by exogenous rmIL-33 challenge in lung homogenate (figure 7A). This result is consistent with a previous report that IL-33 neutralisation by soluble ST2 administration had no effect in house dust mite extract-induced lung IL-1 α protein expression in mouse model.²⁷ On the other hand, protein expression of IL-5 and IL-13 in both BAL fluids and lung homogenates was significantly increased in rmIL-33-challenged mice compared with vehicle-challenged mice ($p<0.05$); and TSA treatment significantly decreased the rmIL-33-induced IL-5 and IL-13 protein expression compared with TSA-vehicle treatment ($p<0.05$) (figure 7B). CCL11 and CCL24 were also increased in lung homogenates from rmIL-33-challenged mice compared with its vehicle-challenged mice ($p<0.05$); and TSA treatment significantly decreased the rmIL-33-induced CCL11 and CCL24 expression in the lung homogenates compared with treatment with the vehicle for TSA ($p<0.05$) (figure 7B). rmIL-33 challenge significantly induced CCL17 and CCL22 expression in the lung homogenates ($p<0.05$), but TSA treatment did not decrease these chemokines compared with treatment with the vehicle for TSA (figure 7B). Further, there was a statistically significant decrease in the number of total BAL cells and eosinophils in the rmIL-33-challenged mice treated with TSA compared with the

Figure 5 (A) Representative histograms of citrine (IL-33) expression on CD45⁺ EpCAM⁺ cells in the lungs from WT mice (filled grey area) or IL-33^{Citrine/+} reporter mice challenged with vehicle–PBS (solid black line), trichostatin A (TSA)–PBS (dotted green line), vehicle–*Alternaria* extract (solid blue line) and TSA–*Alternaria* extract (dotted orange line). (C) The MFI, the percentage and the number of Citrine⁺ CD45⁺ EpCAM⁺ cells in the lung of mice challenged with PBS or *Alternaria* extract following vehicle or TSA treatment. The results are shown as mean±SEM of six mice in PBS-challenged groups and seven mice in *Alternaria* extract-challenged groups. *p<0.05.

rmIL-33-challenged mice treated with vehicle (p<0.05) (figure 7C). These results suggest that TSA treatment suppresses IL-33-induced Th2 cytokine and eosinophil chemotactic chemokine expression, and subsequent eosinophil accumulation in the airway.

TSA treatment concomitant with *Alternaria* extract challenge suppresses allergic pulmonary inflammation

To determine the effect of TSA treatment as a potential therapeutic procedure, we administered TSA or its vehicle concomitantly with *Alternaria* extract challenge (figure 8A). We found a

Figure 6 (A) Mice were challenged with rmlL-33 or its vehicle intranasally for four consecutive days. Trichostatin A (TSA) or its vehicle was administered subcutaneously on day -2 and -1, and then every 1 h before rmlL-33 challenge on day 0–3. (B) The number of IL-5-expressing and IL-13-expressing ILC2 in the lung from the mice challenged with rmlL-33 or its vehicle following TSA or TSA-vehicle treatment. (C) Representative plots of IL-5 and IL-13 intracellular staining in ILC2 in the lungs of mice challenged with rmlL-33 or its vehicle following TSA or TSA-vehicle treatment. The results are shown as mean ± SEM of four mice in PBS-challenged groups and six mice in *Alternaria* extract-challenged groups. * $p < 0.05$.

statistically significant decrease in *Alternaria* extract-induced protein expression of IL-33 ($p < 0.05$) in lung homogenates from TSA-treated mice compared with vehicle-treated mice, but there was no change in *Alternaria* extract-induced IL-1 α expression between TSA-treated mice and vehicle-treated mice (figure 8B). Further, we found a statistically significant decrease in *Alternaria* extract-induced protein expression of IL-5, IL-13, CCL11 and CCL24 in both the BAL fluids and lung homogenates from TSA-treated mice compared with vehicle-treated mice ($p < 0.05$) (figure 8C). Meanwhile, there was no statistically significant difference in protein expression of CCL17 and CCL22, between TSA-treated and vehicle-treated groups with *Alternaria* extract challenge (figure 8C). In addition, there was a statistically significant decrease in the number of total BAL cells and eosinophils in the *Alternaria* extract-challenged mice treated with TSA compared with the *Alternaria*

extract-challenged mice treated with vehicle ($p < 0.05$) (figure 8D). These results suggest that concomitant TSA treatment also suppresses *Alternaria* extract-induced innate eosinophilic airway inflammation.

DISCUSSION

In this study, we report for the first time that an HDACi significantly decreases the early innate immune responses to aeroallergen challenge mediated by ILC2 activation. TSA treatment significantly decreased the number of lung ILC2-expressing IL-5 and IL-13 after four consecutive days *Alternaria* extract challenge. In addition, TSA decreased IL-5, IL-13, CCL11 and CCL24 protein expression, the number of eosinophils, lymphocytes and neutrophils in the BAL fluid, and perivascular eosinophil accumulation induced by *Alternaria* extract challenge. Further, mucus score in the airways 48 h after the last *Alternaria*

Figure 7 (A and B) The BAL fluids and lungs were harvested 24 h after last rmIL-33 challenge to measure the protein expression of IL-1 α , IL-5, IL-13, CCL11, CCL17, CCL22 and CCL24. (C) Cell differentials from BAL fluids harvested 24 h after the last rmIL-33 challenge. The results are shown as mean $\pm$ SEM of four mice in PBS-challenged groups and 11 mice in rmIL-33-challenged groups. * p <0.05.

Figure 8 (A) Mice were challenged with *Alternaria* extract and trichostatin A (TSA) concomitantly for four consecutive days. (B and C) The BAL fluids and lungs were harvested 24 h after last *Alternaria* extract challenge to measure the protein expression of IL-1 α , IL-5, IL-13, IL-33, CCL11, CCL17, CCL22 and CCL24. (D) Cell differentials from BAL fluids harvested 24 h after the last *Alternaria* extract challenge. The results are shown as mean $\pm$ SEM of four mice in PBS-challenged groups and six mice in *Alternaria* extract-challenged groups. * p <0.05.

extract challenge was downregulated by TSA treatment. Although TSA treatment did not decrease *Alternaria* extract-induced IL-33 release in the BAL fluids 1 h after the last *Alternaria* extract challenge, *Alternaria* extract-induced IL-33 and IL-1 α protein level in the lung homogenates 24 h after the

last *Alternaria* extract challenge was significantly decreased by TSA treatment. IL-33 expression was localised in lung epithelial cells. Furthermore, TSA treatment significantly decreased the number of lung ILC2-expressing IL-5 and IL-13, the protein expression of IL-5, IL-13, CCL11 and CCL24 in the lung

homogenates and the number of eosinophils in the BAL fluids after four consecutive days of exogenous rmIL-33 challenge.

Epigenetic regulation plays an important role in immune responses.²⁸ However, the mechanisms by which HDAC inhibition regulates immune responses are still unclear.

Since histone acetylation regulates DNA-templated processes by influencing local chromatin structure,¹¹ transcriptional activation is generally correlated with histone acetylation and repression is associated with histone deacetylation.^{29–30} However, histone acetylation levels differ in cell types or regions of tissue.³¹ Since previous studies have shown that effects of HDACi were dependent on the type of diseases and the type of HDAC inhibition, the therapeutic effects of HDACi treatment for inflammatory or allergic diseases remain controversial. Several studies reported that TSA enhances inflammatory gene expression mediated by NF- κ B^{14–16}; meanwhile, we found that TSA treatment decreased *Alternaria* extract-induced proinflammatory cytokines and chemokines such as KC, TNF- α and GM-CSF (see online supplementary figure S3), as well as type 2 cytokine and chemokine such as IL-5, IL-13 and eotaxin (figure 2). Further, many studies have reported that HDACi have strong anti-inflammatory effects as well as antitumor effects. For instance, TSA attenuated OVA sensitisation and challenge-induced allergic airway inflammation by reducing IgE levels, mucus and proinflammatory cell infiltration scores, and airway hyperresponsiveness in mouse models of adaptive immunity-mediated OVA-induced inflammation.^{20–21} In addition, HDACi treatment ameliorated collagen-induced rheumatoid arthritis³² and DNFB-induced allergic contact dermatitis³³ in a rodent model. Our finding and these animal studies suggest the possibility of novel therapeutic strategies for chronic allergic diseases using HDACi treatment.

Previous toxicity and dose-ranging studies in vivo in rats showed that TSA was well tolerated in single doses of up to 5 mg/kg administered by subcutaneous injection.³⁴ In our mouse model of four consecutive days of *Alternaria* extract challenge, an anti-inflammatory effect without toxicity was observed at doses of both 0.5 and 2.0 mg/kg TSA; however, there was no statistically significant difference in protein expression of IL-5 and IL-13 between 0.5 mg/kg TSA-treated and vehicle-treated groups after *Alternaria* extract challenge (data not shown). 2.0 mg/kg TSA treatment significantly decreased the lung protein expression of IL-5 and IL-13 compared with vehicle treatment (figure 2). Therefore, we investigated the effect of TSA at the dose of 2.0 mg/kg in subsequent experiments.

In this study, we focused on ILC2, a cell type that produces high levels of Th2 cytokines such as IL-5 and IL-13 in response to IL-33. IL-33 is a nuclear protein constitutively expressed in epithelial cells, and is released as an endogenous alarm signal, known as an alarmin, in response to tissue damage.^{26–35} This nuclear cytokine has been reported as a pathogenic factor in several inflammatory diseases such as asthma, arthritis, inflammatory bowel disease and multiple sclerosis.^{36–37} Recent studies revealed that IL-33 is a critical cytokine to expand ILC2 proliferation and activation.^{4–5} We found that IL-33 protein release in the BAL fluids 1 h after 4th *Alternaria* extract challenge was not different between TSA and its vehicle treatment. On the other hand, TSA treatment significantly decreased the number of ILC2-expressing IL-5 and IL-13 24 h after four consecutive days of *Alternaria* extract challenge or rmIL-33 challenge. These results suggest that TSA inhibits IL-33-induced ILC2 activation but does not attenuate early IL-33 release by *Alternaria* extract challenge. These are critical findings because it reveals a

suppressive effect by HDACi on IL-33-induced ILC2 activation and its subsequent attenuation of innate allergic immune responses.

We further found that IL-1 α expression in the lung homogenates was increased by *Alternaria* extract challenge, but not exogenous rmIL-33 challenge, and TSA significantly decreased the *Alternaria* extract-induced IL-1 α expression (figures 4D and 7A). IL-1 α is also known as an endogenous alarmin to induce inflammation and initiates an autocrine feedback for the expression of other cytokines, including GM-CSF and IL-33.²⁷ We found that TSA suppressed lung IL-33 expression in epithelial cells after four consecutive days *Alternaria* extract challenge (figure 4C). Taken together, IL-1 α expression level may be associated with intracellular IL-33 expression in lung epithelial cells.

Currently, there are no HDACi approved for the treatment of allergic diseases. However, HDACi have been investigated in rodent models of allergic inflammation. A recent study reported that a selective HDAC6 inhibitor (tubastatin) showed anti-inflammatory and antirheumatic effects in rodent models.³⁸ Further, the oral administration of another selective HDAC6 inhibitor compound, ACY-1215, attenuated TNCB-induced skin inflammation, including contact hypersensitivity.³⁹ In vitro studies suggest possible mechanisms by which HDACi suppress the activity of inflammatory cells or epithelial cells. HDAC6, a member of the cytoplasmic class IIb HDACs, is a unique enzyme that deacetylates non-histone proteins, such as α -tubulin and heat shock protein 90 (Hsp90). HDAC6 is a key protein involved in aggresome formation that may act as an adaptor protein linking polyubiquitinated proteins to the dynein motor complex for transport. Inhibition of HDAC6 blocks aggresome formation.⁴⁰ In addition, inhibition of HDAC6 results in hyperacetylation of Hsp90; consequently, inactivation of Hsp90 chaperone activity leads facilitating the structural maturation and complex assembly of client proteins including steroid hormone receptors and selected kinases.⁴¹ Since HDAC6 enzymatic activity seems to be required for the regulation of cell morphology, HDAC6 activity in lymphocytes may regulate cell migration and immune synapse formation.⁴²

In this study, we showed that HDAC inhibition by TSA reduces innate allergic airway inflammation by inhibiting ILC2 activation using a mouse model of four consecutive days of *Alternaria* extract challenge or rmIL-33 challenge. Further, the inhibition of ILC2 activation attenuated eosinophilia in the lung and mucus hypersecretion from the airway epithelial cells. These results suggest that histone acetylation is involved in the control of development of allergic airway inflammation mediated by ILC2 activation stimulated with IL-33. HDACi drugs require further experimental analysis before their use in human clinical trials, but our results suggest a promising therapeutic effect for the treatment of allergic lung disorders.

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