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IL-33 and Thymic Stromal Lymphopoietin Mediate Immune Pathology in Response to Chronic Airborne Allergen Exposure

Koji Iijima,* Takao Kobayashi,* Kenichiro Hara,* Gail M. Kephart,* Steven F. Ziegler,† Andrew N. McKenzie,‡ and Hirohito Kita*

Humans are frequently exposed to various airborne allergens in the atmospheric environment. These allergens may trigger a complex network of immune responses in the airways, resulting in asthma and other chronic airway diseases. In this study, we investigated the immunological mechanisms involved in the pathological changes induced by chronic exposure to multiple airborne allergens. Naive mice were exposed intranasally to a combination of common airborne allergens, including the house dust mite, Alternaria, and Aspergillus, for up to 8 wk. These allergens acted synergistically and induced robust eosinophilic airway inflammation, specific IgE Ab production, type 2 cytokine response, and airway hyperresponsiveness in 4 wk, followed by airway remodeling in 8 wk. Increased lung infiltration of T cells, B cells, and type 2 innate lymphoid cells was observed. CD4+ T cells and type 2 innate lymphoid cells contributed to the sources of IL-5 and IL-13, suggesting involvement of both innate and adaptive immunity in this model. The lung levels of IL-33 increased quickly within several hours after allergen exposure and continued to rise throughout the chronic phase of inflammation. Mice deficient in IL-33R (Il1r1−/−) and thymic stromal lymphopoietin receptor (Tslpr−/−) showed significant reduction in airway inflammation, IgE Ab levels, and airway hyperresponsiveness. In contrast, mice deficient in IL-25R or IL-1R showed minimal differences as compared with wild-type animals. Thus, chronic exposure to natural airborne allergens triggers a network of innate and adaptive type 2 immune responses and airway pathology, and IL-33 and thymic stromal lymphopoietin most likely play key roles in this process. *The Journal of Immunology, 2014, 193: 000–000.

Exposure and sensitization to allergens are risk factors for developing asthma and allergic airway diseases (1). Allergens play a key role in triggering and exacerbating asthma and allergy symptoms (2). In particular, exposure to airborne allergens derived from animals, arthropods, and molds is considered an important risk factor for asthma (3–5). Simultaneous exposure to several allergens is also common (6, 7). Indeed, exposure to multiple allergens and increased levels of total allergens in the home are significantly associated with developing asthma (8). In addition, certain allergens—such as Alternaria, the house dust mite (HDM), mouse, and cockroach—are detected together at high levels in home environments (8). Most patients with allergic asthma are sensitized to multiple allergens (9). Therefore, the conventional asthma models in mice that rely on OVA sensitization and challenge, or exposure to a single allergen, may not capture the impact and complexity of chronic exposure to multiple airborne allergens in humans.

Recent studies suggest that the biological properties of allergen molecules most likely play important roles in developing type 2 immunity to inhaled allergens. For example, the response to HDM is mediated by recognition of the major HDM allergen Der p 2 and perhaps endotoxin contained in fecal pellets through TLR4 (10, 11). Alternatively, many Th2-inducing allergens have enzymatic activities, such as proteases from HDM and fungi (12–14) and phospholipases from bee venom (15), which may trigger protease-activated receptors or other as-yet-unidentified receptor(s). Allergen proteases may also increase the passage of allergens across the epithelial barrier (12, 13). Thus, several immune receptors and pathways are most likely triggered by exposure to natural allergens. It would be critical to identify key fundamental factor(s) that plays a central role in these complex immune responses to allergens.

Using a mouse model, we sought to investigate the mechanisms involved in pathological processes induced by chronic exposure to airborne allergens. To accomplish our goal of mimicking natural allergen exposure in humans, we simultaneously exposed animals for a prolonged period to several allergens that are relevant to human asthma. Previously, combined sensitization of mice to HDM, ragweed, and Aspergillus broke tolerance and established resistant asthma in humans (5, 14). Chronic intranasal exposure of naive animals to a combination of these allergens triggered robust type 2 immune responses, which involve both innate and adaptive immunity; the allergens worked synergistically to induce

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*Division of Allergic Diseases, Department of Internal Medicine, Mayo Clinic, Rochester, MN 55905; †Immunology Program, Benaroya Research Institute, Seattle, WA 98101; and ‡Medical Research Council Laboratory of Molecular Biology, Cambridge CB2 0QH, United Kingdom

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Address correspondence and reprint requests to Prof. Hirohito Kita, Allergic Diseases Research Laboratory, Guggenheim Building, Room 401A, Mayo Clinic Rochester, 200 First Street SW, Rochester, MN 55905. E-mail address: kita.hirohito@mayo.edu

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Abbreviations used in this article: AHR, airway hyperresponsiveness; BAL, bronchoalveolar lavage; B.M., basement membrane; HDM, house dust mite; ILC2, type 2 innate lymphoid cell; OAAH, OV A, Alternaria, Aspergillus, and HDM; PAS, periodic acid–Schiff; TSLP, thymic stromal lymphopoietin.

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a strong response. Importantly, among various immunological factors, IL-33 appeared to be particularly important in mediating the process because it is significantly increased during both the early and chronic phases of exposure and because blockade of the pathway attenuated airway pathology. Thymic stromal lymphopoietin (TSLP) also contributed significantly, whereas IL-25 and IL-1 appeared to play minimal roles.

Materials and Methods

Reagents

Extracts of natural allergens, including *Alternaria alternata*, *Aspergillus fumigatus*, and *Dermatophagoides pteronyssinus*, were purchased from Greer Laboratories (Lenoir, NC). These extracts contain undetectable levels of endotoxin (<1 ng/mg extract, <0.15 µg/dose). Endotoxin-free OVA was prepared in our laboratory, as previously described (17).

**Mice**

BALB/c, C57BL/6, Rag1−/−, and Il1r1−/− mice were purchased from The Jackson Laboratory (Bar Harbor, ME). ST2−/− (Il1r1−/−) (18), Il17rb−/− (19), Tslpr−/− (20), and Il13−/−GFP mice (19) were generated on the BALB/c background and have been bred in the animal facility at the Mayo Clinic (Rochester, MN). Il13−/−GFP mice (21) were a gift of K. Takatsu (University of Toyama, Toyama, Japan). Six- to 12-week-old female mice were used in this study. All experiments in this study were approved by the Mayo Clinic Institutional Animal Care and Use Committee.

*Acute allergen exposure model*

Mice were exposed intranasally to a mixture of allergen extracts (10 µg each) in 50 µl PBS, *Alternaria, Aspergillus*, and *Dermatophagoides* (OAAH) in 50 µl PBS, three times per week, for up to 8 wk under isoflurane inhalation anesthesia. The levels of protein in the 10 µg doses of *Alternaria, Aspergillus*, and *Dermatophagoides* extracts, and HDM extracts were 2.0, 3.5, and 3.2 µg, respectively. Each allergen extract plus OVA or a mixture of allergen extracts without OVA were also used for certain experiments. Alternatively, each allergen alone (10 µg/dose or 30 µg/dose in 50 µl PBS) was used for some experiments. Twenty-four hours after the final exposure, airway hyperresponsiveness (AHR) to inhaled methacholine was analyzed by whole-body plethysmography, as previously described (Buxco Electronics, Sharon, CT) (22). Mice were nebulized with increasing doses of methacholine (1.6–50 mg/ml), and AHR was represented as a percentage of baseline enhanced pause. AHR in anesthetized mice was also analyzed in certain experiments by Flexivent forced oscillation technique (Scireq, Montreal, Canada) (23). Thereafter, 100 µl blood was collected by retro-orbital bleeding in 10% heparinized PBS. Red blood cells were then killed, and bronchoalveolar lavage (BAL) was performed by intratracheal instillation of 1.0 ml HBSS. In some experiments, the lungs were also collected, homogenized in 0.5 ml PBS, and centrifuged at 10,000 rpm for 10 min; the protein concentrations in the supernatant were quantified with the bicinchoninic acid (BCA) Protein Assay kit (Thermo Fisher, Rockford, IL). Total leukocyte counts in BAL fluids were determined using a hemocytometer after staining with Randox’s stain. For cell differentiation, cytopsin preparations from BAL fluids were determined using a hemocytometer after staining with Wright-Giemsa. Cytokine levels in the supernatants of BAL fluids and lung homogenates were measured using ELISA. We also collected the spleen for analyses of cytokine production in vitro.

*Acute allergen exposure model*

Naive mice were exposed once intranasally to a mixture of allergen extracts (10 µg each of OAAH in 50 µl PBS). One, 3, 6, 12, or 24 h after the exposure, mice were killed, and BAL samples and the lungs were collected. Cytokine levels in the supernatants of BAL fluids and lung homogenates were measured using ELISA. ELISA

The levels of IL-1α, IL-1β, IL-4, IL-5, IL-13, IL-17E (IL-25), IL-33, IFN-γ, and TSLP in the supernatants of lung homogenates and BAL fluids were measured using ELISA kits (R&D Systems, Minneapolis, MN) following the procedure recommended by the manufacturer. The levels of OVA-specific IgE or IgG1 in the plasma were measured using sandwich ELISA, as previously described (17). To measure the levels of allergen-specific IgG1 Abs in the plasma, ELISA plates were coated with recombinant Alt a 1, natural Asp r 1 (Asp f 1 homolog), or recombinant Der p 1 (all from Indoor Biotechnologies, Charlottesville, VA) overnight at 4°C (15 µg/ml in 0.1 M carbonate buffer [pH 9.5]). The plates were blocked with PBS containing 1% BSA for 2 h; Plasma samples, which were diluted 1:20 in PBS containing 1% BSA and 0.05% Tween 20, were added to the plates and incubated for 4 h. Thereafter, the plates were incubated with HRP-conjugated rat anti-mouse IgG1 (BD Biosciences, San Jose, CA) for 1 h. After washing, tetramethylbenzidine substrate (Thermo Scientific, Rockford, IL) was added, and the reaction was stopped with 2 M H2SO4. The absorbance at 450 nm was read in a microplate autoreader (Thermomax; Molecular Devices).

**Splenocyte cytokine production in vitro**

Splenocytes were suspended in RPMI 1640 medium (Invitrogen, Carlsbad, CA) supplemented with 10% FBS, 100 U/ml penicillin, and 100 µg/ml streptomycin. Cells were cultured at 4 × 10⁵ cells/well in 96-well round-bottom tissue culture plates in the presence or absence of 100 µg/ml OVA Ag. Five days later, the concentration of IL-13 in the cell-free supernatants was measured using ELISA.

**Lung immunohistochemistry**

To identify lymphocytes in mouse lungs, lung tissue was embedded in OCT compound, frozen, and sectioned. Sections were fixed in ice-cold acetone and incubated with a dual endogenous enzyme block (Dako, Carpinteria, CA) and a blocking reagent (Background Sniper for CD3 and B220, Rodent Block M for CD4 and CD8; Biocare Medical, Concord, CA). The sections were then stained with rat anti-mouse CD11c (145-2C11; BioLegend, San Diego, CA), CD4 (GK1.5; eBioscience, San Diego, CA), CD8α (YTS 168AG; Abcam, Cambridge, MA), B220 (RA3-6B2; eBioscience), or the appropriate isotype-matched rat IgG control (eBioscience). Staining was visualized using a rat-on-mouse HRP-polymer kit and a diaminobenzidine detection system (Biocare Medical). Sections were counterstained with hematoxylin QS (Vector Laboratories, Burlingame, CA).

**Flow cytometry**

Lungs were minced using a gentleMACS Dissociator (Miltenyi Biotec, Auburn, CA) and digested with Liberase Research Grade (Roche, Mannheim, Germany) in RPMI 1640 medium in the presence of DNase I solution (STEMCELL Technologies, Vancouver, Canada) for 1 h at 37°C. After digestion, single lung cells were hemolyzed with ammonium-chloride-potassium buffer, and washed with PBS containing 0.1% sodium azide and 1% BSA. To quantitate the numbers of lymphocytes, single lung cells were stained with FITC-CD3ε (145-2C11), PE-B220 (RA3-6B2, PerCP complex (PerCP)-CD4 (RM4-5), and allophycocyanin-CD8 (53-6-7; BD Biosciences) for 30 min at 4°C. After washing in PBS containing 0.1% sodium azide and 1% BSA buffer, the expression levels of CD3, CD4, CD8, and B220 were detected by FACs (BD FACSCalibur; BD Biosciences). To examine the expression levels of cytokines by type 2 innate lymphoid cells (ILC2s) or CD4+ T cells, single lung cells from Il13−/−GFP mice or Il5−/−/venus mice were stained with a PE-conjugated lineage mixture CD3 (145-2C11), CD41 (mC5-3), CD163/2 (24G2), B220 (RA3-6B2), allophycocyanin-CD25 (PC61), PerCP-Cy5.5-CD44 (IM7; BD Biosciences) or PE-CD4, and PerCP-CD3. Lung ILC2s and CD4+ T cells were identified as Lin CD25−CD44high cells and CD3+CD4+ cells, respectively, as previously described (24).

**Quantitation of airway remodeling**

For histological analyses, lung tissues were fixed in 10% formalin and embedded in paraffin. Six-millimeter sections were stained with periodic acid–Schiff (PAS) or Masson’s trichrome. Computer analysis was performed using image-analysis software (KS400 Image Analysis System; Carl Zeiss Microscopy, Thornwood, NY), the threshold for the lumen was also used to analyze the stained area. The threshold for the total tissue was also recorded in randomly selected airway sections (×10) per tissue section and normalized as unit per mm B.M.

**Statistical analysis**

Data are expressed as the mean ± SEM for the mice indicated. The statistical significance of the differences between the various treatment groups was assessed with Student t test by using PRISM software and InStat software (both from GraphPad Software, La Jolla, CA; p < 0.05 was considered significant).
Results

Chronic exposure to natural airborne allergens induces robust type 2 immune responses followed by airway remodeling

To investigate the immunological mechanisms involved in airway pathology induced by chronic exposure to multiple airborne allergens, we intranasally administered a combination of allergen extracts three times per week for up to 8 wk to naive BALB/c mice (Fig. 1A); BAL and lung specimens were analyzed 24 h after the last exposure. No systemic immunization, such as i.p. or s.c. injection of allergens, was used throughout the procedure. For allergens, we used HDM, Alternaria, and Aspergillus. HDM and Alternaria are commonly found in homes, and exposure to them increases the risk of developing asthma (3, 5, 8). Aspergillus is implicated in severe treatment-resistant asthma (5, 14). Although the cockroach is also implicated in human asthma (4), we did not include it in this study because cockroach extracts are highly contaminated with endotoxin (i.e., 7000 EU/mg). A mixture of HDM, Alternaria, and Aspergillus extracts (10 μg/dose each) was...

FIGURE 1. Chronic airway exposure of naive mice to natural airborne allergens induces type 2 immune responses, eosinophilic inflammation, and airway remodeling. (A) Schematic representation of the exposure protocol. Naive BALB/c mice were exposed intranasally three times/wk to a combination of OVA and extracts of OAAH for up to 8 wk. (B) BAL fluids were analyzed for the number of inflammatory cells. (C) Plasma specimens were analyzed for the levels of anti-OVA Abs. (D) Lung homogenates were analyzed for cytokine levels. (B–D) Results are the mean ± SEM (n = 5–6 in each group). *p < 0.05, **p < 0.01, compared with mice exposed to PBS. n.d., Not determined. (E) Representative photomicrographs of lung specimens. Scale bar, 100 μm. (F) The magnitude of airway remodeling was analyzed by computer-assisted image analysis. Results are the mean ± SEM (n = 10 in each group). *p < 0.05, **p < 0.01, compared with mice exposed to PBS.
spiked with a small amount of endotoxin-free OVA (10 μg/dose), which allows us to monitor the development of Ag-specific adaptive immunity; thus, the allergen mix was named OAAH (i.e., short for OVA, Alternaria, Aspergillus, and HDM).

Chronic and multiple exposures to OAAH induced a robust increase in BAL eosinophils in 4 wk; eosinophils made up ~70% of total BAL cells (Fig. 1B). Plasma concentrations of OVA-specific IgE and IgG1 Abs continued to rise for at least 8 wk (Fig. 1C). Increased lung levels of type 2 cytokines, including IL-4, IL-5, and IL-13, were also observed in 4 wk (Fig. 1D). The IL-5 and IL-13 levels appeared to decline in 8 wk. Notably, a marked increase in the lung levels of IL-33 was observed (please note the y-axis scale in Fig. 1D) in 4 wk, and the levels remain elevated for at least 8 wk. In contrast, TSLP was undetectable, and the IL-25 levels were not affected by allergen exposure. Lung pathology specimens showed apparent evidence of structural changes in 8 wk, including increased PAS-positive airway epithelial cells and connective tissues stained with Masson’s trichrome; these changes were minimal or not apparent in 4 wk (Fig. 1E, 1F). Together, these findings suggest that chronic exposure to multiple allergens induces robust eosinophilic inflammation and IL-5 and IL-13 production, followed by airway remodeling. Continued increases in specific IgE and IgG1 Abs in peripheral blood and IL-33 in the lungs for 8 wk were also noted.

Allergens act synergistically to induce robust responses

We performed subsequent mechanistic studies by focusing on the 4-wk time point, which represents the peak for airway eosinophilia and cytokine levels (Fig. 1B, 1D). First, we examined the contribution of each allergen at the dose used in this study (i.e., 10 μg/dose). Exposure to OVA alone, OVA plus Aspergillus, or OVA plus HDM produced minimal airway inflammation (Fig. 2A), whereas exposure to OVA plus Alternaria induced modest airway eosinophilia. Importantly, the three allergens when combined induced marked increases in BAL lymphocytes and eosinophils, ~10-fold and 3-fold higher, respectively, compared with OVA plus Alternaria. The production of specific Abs, in particular of the IgE isotype, was also enhanced dramatically by exposure to a combination of three allergens (Fig. 2B).

**FIGURE 2.** Allergens work synergistically to induce robust airway inflammation and Ab production. (A) Naive mice were exposed intranasally to PBS, OVA alone, OVA plus each allergen (10 μg/dose), or OVA plus a mixture of all allergens (Alternaria, Aspergillus, HDM; 10 μg/dose each), as indicated, three times/wk, for 4 wk. BAL fluids were analyzed for the number of inflammatory cells. (B) Plasma was analyzed for anti-OVA Ab levels; ALT, Alternaria; ASP, Aspergillus. (C) Naive mice were exposed to each allergen (10 μg/dose) or a combination of all allergens (Alternaria, Aspergillus, HDM; 10 μg/dose each) for 4 wk. BAL fluids were analyzed for the number of inflammatory cells. (D) The levels of IL-13 in BAL fluids were analyzed. (E) The levels of IgG1 Abs to Alt a 1, Asp r 1 (Asp f 1 homolog), and Der p 1 in the plasma were analyzed. Results are the mean ± SEM [n = 3 in (A) and (B), and n = 5 in (C)–(E)]. *p < 0.05, **p < 0.01, compared with mice exposed to PBS, or between groups, as indicated by the horizontal lines.
Second, to examine whether the OVA that was used to monitor adaptive immunity was necessary to induce type 2 responses to allergens, we compared the effects of the OAAH allergen mix and one without OVA. Removal of OVA from OAAH did not affect the magnitude and characteristics of airway inflammation (Supplemental Fig. 1A). The plasma levels of total IgE and BAL levels of IL-13 were not affected by omitting OVA from OAAH, and OVA-specific IgE was undetectable in animals exposed to OAAH without OVA (Supplemental Fig. 1B).

Third, to examine the effects of allergens alone without OVA, mice were exposed to each allergen (10 μg/dose) or a mixture of all three allergens (10 μg/dose each) for 4 wk. Exposure to Aspergillus or HDM produced minimal airway inflammation (Fig. 2C), whereas exposure to Alternaria induced modest airway eosinophilia. A mixture of the three allergens induced marked increases in BAL lymphocytes and eosinophils and BAL levels of IL-13 (Fig. 2D, 2E). Mice also developed IgG1 Abs to Alt a 1, Asp r 1, and Der p 1 in plasma when they were exposed to a mixture of the three allergens (Fig. 2E); exposure to Alternaria alone induced a significant increase in anti-Alt a 1 Ab.

Finally, the effects of each allergen were examined at a higher dose (i.e., 30 μg/dose). At this dose, not only Alternaria but also HDM induced airway eosinophilia and specific IgG1 Abs (Supplemental Fig. 2). Altogether, these findings suggest allergens act synergistically during chronic exposure even if the effects of each allergen may not be apparent in a relatively low dose.

Adaptive immunity is necessary to drive chronic type 2 immunity to allergens

Recent studies suggest that acute airway exposure to certain natural or model allergens, such as Alternaria (24, 25) and the cysteine protease papain (26), activates type 2 innate immunity, and type 2 ILC2s are sufficient to mediate airway inflammation even in the absence of adaptive immunity. Therefore, to examine the role of adaptive immunity in the chronic allergen exposure model, we used BALB/c Rag1−/− mice, which are deficient in mature T cells and B cells. Whereas wild-type mice showed increases in BAL eosinophils, lymphocytes, and IL-13, as well as in plasma anti-OVA IgE, these responses were abolished in Rag1−/− mice (Fig. 3A, 3B).

Immunohistochemical analyses revealed occasional CD3+ T cells and B220+ B cells scattered in the lung parenchyma in wild-type mice exposed to PBS (Fig. 3C). When wild-type mice were exposed to the allergens for 4 wk, CD3+ cells and B220+ cells increased in number and accumulated in the perivascular and peribronchial regions. B220+ cells formed distinct clusters, whereas CD3+ cells were distributed rather diffusely. As estimated by flow cytometry, the total numbers of CD3+ cells and B220+ cells in the lungs increased by 5–6-fold in OAAH-exposed animals as compared with PBS-exposed animals (Fig. 3D); increases in both the CD4+ and CD8+ subsets of T cells were observed.

To verify the involvement of adaptive immunity, we cultured splenocytes from PBS- or OAAH-exposed animals with OVA Ag in vitro. Splenocytes from OAAH-exposed animals, but not those from PBS-exposed animals, produced IL-13 when stimulated with OVA in vitro (Fig. 3E). Altogether, unlike acute models, intact adaptive immunity is necessary for airway inflammation in the chronic exposure model.

Both CD4+ T cells and ILC2s are sources of type 2 cytokines

We then investigated whether innate immunity is involved. Because ILC2s are recognized as an important source of type 2 cytokines during the acute and innate phase of the immune response (27, 28), we quantitated ILC2s in the lungs of this chronic exposure model. Lung ILC2s were identified as lineage-negative (Lin−) and CD25 and CD44 double-positive cells (Fig. 4A), as described previously (24). ILC2s were present in the lungs of naive (data not shown) as well as PBS-exposed animals, and the number increased by >2-fold when animals were exposed to OAAH for 4 wk (Fig. 4B).

To examine the functions of these ILC2s during chronic inflammation, we used cytokine reporter mice, including the IL-5 reporter IL-5venus mice and IL-13 reporter IL-13eGFP mice. In PBS-exposed mice, a small fraction of CD44high cells within the Lin− population expressed IL-5venus or IL-13eGFP (Fig. 4C); no other cell populations within the Lin− population expressed these cytokines. Because the Lin− CD44high cell population in the lungs consists exclusively of ILC2s (24) (Fig. 4A), these findings suggest that a small fraction of ILC2s expresses IL-5 or IL-13 when mice are exposed to PBS. Importantly, when mice were exposed to OAAH, the proportion of IL-5− or IL-13−producing ILC2s increased by ∼3–7-fold, and ∼15–20% of total ILC2s showed ongoing production of these cytokines (Fig. 4C, 4D).

To examine the functions of T cells, similar analyses were performed by gating on CD3+ T cells. Unlike ILC2s, IL-5venus or IL-13eGFP signals were undetectable or minimal within the CD3+ cell population in PBS-exposed animals. When mice were exposed to OAAH, the prevalence of IL-5− or IL-13−positive cells increased dramatically (Fig. 4E). These cytokine-positive CD3+ T cells expressed CD4 (Fig. 4E), and no CD4− cells were positive for IL-5 or IL-13. Approximately 1.4–2.0% of total CD4+ T cells expressed cytokines when mice were exposed to OAAH (Fig. 4F). On average, 12.3 × 103 and 7.7 × 103 ILC2s/lung were positive for IL-5venus and IL-13eGFP, respectively, and 55.5 × 103 and 58.3 × 103 CD4+ T cells/lung were positive for IL-5venus and IL-13eGFP, respectively, after OAAH exposure (n = 3). Altogether, both ILC2s and CD4+ T cells most likely contribute to the increased IL-5 and IL-13 production in mice after chronic airway exposure to airborne allergens; CD4+ T cells appear to outnumber ILC2s.

Innate cytokines, including IL-33, increase rapidly after allergen exposure in naive mice

The molecular and cellular mechanisms that initiate and maintain type 2 immunity remain topics of active investigation. Evidence suggests that epithelial cells make important contributions by producing several cytokines that facilitate the development of type 2 immunity, including IL-25, IL-33, and TSLP (29–31). IL-1 family cytokines, such as IL-1α and IL-1β, have been considered prototypic innate cytokines with diverse immunological activities (32), and IL-1α has been shown to induce IL-33 release from airway epithelial cells (33). Therefore, to further investigate the immunological mechanisms of chronic type 2 immune responses to airborne allergens, we focused on the roles for these cytokines.

We investigated whether these innate cytokines are produced when mice are exposed acutely to the combination allergen OAAH. Naive BALB/c mice were exposed to PBS or OAAH only once, and the kinetic changes of these cytokines in the lungs were analyzed. Substantial amounts of IL-33 (600 pg/mg tissue) were detected in the lungs of naive nontreated mice, suggesting that this cytokine is constitutively produced (Fig. 5A). When mice were exposed to OAAH, the lung levels of IL-33, IL-1α, and IL-1β increased quickly within one to several hours and peaked at 6 h, and the levels declined to the baseline values by 12 or 24 h. The increase in TSLP was also observed, whereas the levels were considerably lower than IL-33, IL-1α, or IL-1β. The kinetic changes in these four cytokines were similar. In contrast, IL-25 levels did not change significantly for up to 24 h.

When supernatants of BAL fluids were analyzed, IL-33 was detectable within 1 h after exposure to OAAH. The IL-33 levels...
declined quickly, and IL-33 became undetectable 3 h after the exposure. No TSLP was detectable in BAL fluid supernatants at any time points. These findings suggest that expression of IL-1 family cytokines and TSLP increases transiently when naïve mice are exposed to allergens.

IL-33 and to a lesser degree TSLP play pivotal roles in chronic inflammation

To examine the roles of these innate cytokines, we investigated mice that are deficient in their receptors, including ST2<sup>−/−</sup> (i.e., IL-33R deficient), TSLPR<sup>−/−</sup> (TSLP receptor deficient), Il17rb<sup>−/−</sup> (i.e., IL-25R deficient), and Il1r1<sup>−/−</sup> (i.e., deficient in the receptors for IL-1α and IL-1β) mice. Wild-type mice or receptor knockout mice were exposed to PBS or OAAH for 4 wk. PBS-exposed wild-type mice or knockout mice did not show any apparent signs of airway inflammation: BAL eosinophils, <0.1 × 10⁵ cells; BAL lymphocytes, <0.1 × 10⁵ cells; and IL-4, IL-5, and IL-13 levels in lung homogenate, <5.0 pg/mg tissue. When exposed to OAAH, ST2<sup>−/−</sup> mice showed 70–80% reduction in BAL eosinophil and lymphocyte numbers (Fig. 6A, 6B) and in plasma levels of total IgE and OVA-specific IgE (Fig. 6C, 6D), compared with wild-type mice exposed to OAAH. TSLPR<sup>−/−</sup> mice also showed ~50% reduction in BAL eosinophils as well as in plasma IgE Ab levels. Although Il17rb<sup>−/−</sup> mice showed partial reduction in BAL eosinophil and lymphocyte numbers, IgE Abs were not affected. In contrast, Il1r1<sup>−/−</sup> mice did not show significant changes in any of these immunological parameters; the number of BAL eosinophils rather increased in these Il1r1<sup>−/−</sup> mice (p < 0.05).

Fig. 7A–C shows the levels of lung cytokines. The levels of IL-4, IL-5, and IL-13 were significantly decreased in ST2<sup>−/−</sup> mice; IL-5 levels appeared to be affected the most. Partial reduction in IL-5 was also observed in TSLPR<sup>−/−</sup> and Il17rb<sup>−/−</sup> mice, and no apparent changes were observed in Il1r1<sup>−/−</sup> mice. We finally...
examined the roles of these cytokines in airway reactivity to inhaled methacholine. When naive wild-type mice were exposed to OAAH for 4 wk, they developed enhanced reactivity to inhaled methacholine as compared with PBS-exposed wild-type mice (Fig. 7D). Analysis of airway reactivity with a forced oscillation technique also verified this observation (Fig. 7E). Although we evaluated a full-range dose-response curve for inhaled methacholine (Supplemental Fig. 3), the data at 6 mg/ml methacholine are summarized in Fig. 7F for clarity. Wild-type BALB/c mice or C57BL/6 mice exposed to OAAH showed significant increases in airway reactivity to methacholine, compared with those exposed to PBS. The increase in reactivity was significantly attenuated in \textit{ST2}\textsuperscript{−/−} and TSLPR\textsuperscript{−/−} mice (\(p<0.05\)), but not in \textit{Il17rb}\textsuperscript{−/−} or \textit{Il1rl}\textsuperscript{−/−} mice. Together, these findings suggest that IL-33 and, to a lesser degree, TSLP play pivotal roles in airway inflammation, IgE Ab production, and AHR when mice are exposed to multiple airborne allergens for a prolonged period. The role of IL-25 is most likely minimal, and IL-1\(\alpha/\beta\) may play no role or rather act as an inhibitor of airway eosinophilia in this model.

**FIGURE 4.** ILC2s and CD4\(^{+}\) T cells are most likely the source of IL-5 and IL-13. (A) Naive wild-type mice were exposed intranasally to PBS or allergens (OAAH) for 4 wk. Lung single-cell suspensions were gated for the lineage-negative (Lin\(^{-}\)) cell population, and ILC2s were identified as CD25\(^{+}\)CD44\(^{\text{high}}\) cells. (B) The number of ILC2s in the lung specimens was quantitated. (C) IL-5\textsuperscript{venus} mice (upper panels) or IL-13\textsuperscript{eGFP} mice (lower panels) were exposed to PBS or allergens (OAAH) for 4 wk. Lung single-cell suspensions were gated for the Lin\(^{-}\) cell population, and the expression levels of IL-5\textsuperscript{venus} or IL-13\textsuperscript{eGFP} in the CD44\(^{\text{high}}\) population were analyzed by flow cytometry. (D) The proportions of CD44\(^{\text{high}}\)IL-5\textsuperscript{venus} cells or CD44\(^{\text{high}}\)IL-13\textsuperscript{eGFP} cells among the ILC2 population were quantitated. (E) Lung single-cell suspensions were gated for the CD3\(^{+}\) population, and the expression levels of IL-5\textsuperscript{venus} or IL-13\textsuperscript{eGFP} in the CD4\(^{+}\) cell population were analyzed by flow cytometry. (F) The proportions of CD44\(^{\text{high}}\)IL-5\textsuperscript{venus} cells or CD44\(^{\text{high}}\)IL-13\textsuperscript{eGFP} cells among the CD4\(^{+}\) T cell population were quantitated. Results are the mean \(\pm\) SEM \(n=4\) in (B); \(n=3\) in (D) and (F) and are representative of two independent experiments. *\(p<0.05\), **\(p<0.01\), compared with mice exposed to PBS.
Discussion

Animal models of disease are critical in research to investigate mechanisms and to identify and validate targets for therapeutic interventions. Whereas mouse models have been invaluable in asthma research, their limitations have also been identified and discussed (34, 35). In this study, by attempting to mimic natural environmental exposure, three different airborne allergens that are relevant to human asthma were combined and administered to the airways of naive mice for a prolonged period, up to 8 wk. Although each allergen itself provoked minimal responses at the dose used in this study (i.e., 10 μg/dose), they synergistically induced robust eosinophilic airway inflammation and IgE Ab production in 4 wk. Airway remodeling continued to occur at least for 8 wk. The results from this new model extend previous observations and suggest that IL-33 is involved in mediating chronic type 2 immune responses to natural airborne allergens, including airway inflammation, production of IgE Ab, and AHR. In contrast, although IL-1α and IL-1β showed apparent increases during the acute phase (Fig. 5), mice deficient in IL-1R were not protected from developing chronic airway inflammation. A previous report also suggested that IL-1α is necessary for the induction of IL-33 release from airway epithelial cells when mice are exposed to HDM extract (33). However, the results from this study indicate that IL-33 can be involved in type 2 immunity independent of the IL-1 pathway. Another new series of studies focusing on the 8-wk time point or even longer is necessary to investigate the roles of IL-33 and potentially other cytokines in airway remodeling induced by chronic allergen exposure.

The role of TSLP in regulating type 2 immunity in the lungs has remained a topic of some controversy. Previously, when mice were sensitized by i.p. injection of the OVA Ag, TSLP was found to play a key role in the lung Th2 response (36, 37). Furthermore, transgenic overexpression of TSLP in the airway also induced asthma-like pathological changes (36). However, more recently, by exposing naive mice intranasally to HDM extract for 10 d, Chu et al. (38) demonstrated that Tslpr−/− mice showed comparable levels of IgE Ab and airway eosinophilia as did wild-type mice. The results from our study suggest that TSLP is involved in airway eosinophilia and IgE Ab production when animals are exposed to natural allergens for a prolonged period, although the contribution of TSLP may not be as robust as that of IL-33 (Figs. 6, 7). The mechanism and ability of TSLP to induce type 2 immunity are most likely different from that of IL-33. For example, overexpression of TSLP alone in the airways caused a relatively weak innate immune response (39). Both overexpression of TSLP and airway administration of the OVA Ag were required to induce a disease phenotype (39), suggesting that TSLP conditions the lung milieu toward the optimal development of adaptive immunity against innocuous Ags. In contrast, IL-33 appears to be involved in both the innate and adaptive phases of type 2 immunity by acting on a variety of cell types, including, but not limited to ILC2s, dendritic cells, CD4+ T cells, mast cells, and eosinophils (40). Therefore, potential differences in the observations regarding the role of TSLP in this study and those by Chu et al. (38) can be explained by the variance in the models, in particular the length of allergen exposure (e.g., 4 wk versus 10 d) and possibly the types of allergens used.

![FIGURE 5](http://www.jimmunol.org/) The lung levels of IL-33, TSLP, and IL-1 increase quickly following exposure to allergens. Naive BALB/c mice were exposed intranasally to PBS or allergens (OAAH) only once. (A) The kinetics changes of cytokines in lung homogenates were analyzed. (B) Changes in the kinetics of cytokine production in BAL fluid supernatants were analyzed. Results are the mean ± SEM (n = 3 in each group). *p < 0.05, **p < 0.01, compared with mice exposed to PBS.
IgE (C) (eosinophils were exposed intranasally to allergens (OAAH) for 4 wk. The number of type mice (BALB/c or C57BL/6) or mice deficient in cytokine receptors and IgE Ab production in response to chronic allergen exposure. Naive wild-

FIGURE 6. IL-33 and TSLP play critical roles in airway inflammation

Although the expression profile of IL-33 has been well characterized in normal tissues (41, 42), relatively little is known regarding its expression during inflammation. One of the major findings in this study is the dynamic change in lung IL-33 levels in response to chronic allergen exposure. IL-33 was the most abundant cytokine examined in lung tissues from naive nontreated animals (Fig. 5). IL-33 levels then increased quickly, similarly to other IL-1 family cytokines, after acute allergen exposure. In contrast, IL-33 secretion into the airway lumen was detectable only at the earliest time point of examination (i.e., 1 h), suggesting increased total lung levels of IL-33 do not necessarily predict secretion of the protein to the extracellular space. Interestingly, whereas IL-33 levels in the lungs decreased temporarily to base-

line in 24 h, they were augmented dramatically by 20-fold during the chronic phase of allergen exposure, and remained elevated for at least 8 wk (Fig. 1D). IL-33 can be produced by airway tissue cells, such as epithelial cells, fibroblasts, vascular endothelial cells, and smooth muscle cells, as well as by immune cells such as dendritic cells and macrophages (43–45). Studies using genetic tools, such as IL-33 reporter mice (46, 47), combined with immunological techniques, will be necessary to map the source(s) of IL-33 precisely during each of the acute and chronic phases of allergen exposure. Questions also remain as to the mechanism of enhanced IL-33 production and release. The direct effects of allergens on airway epithelial cells (48, 49), the influence of the cytokine milieu (50), and potential autocrine mechanisms of the IL-33/ST2 pathway (46) may need to be considered.

Unlike conventional OVA-mediated asthma models in mice (35), type 2 airway inflammation induced by chronic exposures to natural airborne allergens is most likely mediated by complex and perhaps intertwining networks of innate and adaptive immunity. ILC2s have been known to provide a critical early source of type 2 cytokines, such as IL-5 and IL-13, when animals are exposed acutely to allergens (24–26). In contrast, the roles of ILC2s in chronic inflammation have not been well understood. In this study, when exposed to allergens for a prolonged period (i.e., 4 wk), both CD4+ T cells and ILC2s were identified as the potential source of IL-5 and IL-13 (Fig. 4). The results are consistent with previous analyses of the source of IL-13 in OVA-induced type 2 responses, indicating that CD4+ T cells and ILC2s are the sources of IL-13 (51). Nonetheless, T cells were essential for robust eosinophilic inflammation and type 2 cytokine production (Fig. 3A, 3B), consistent with previous findings (52). Although further studies will be necessary to identify key element(s) that regulates the network of innate and adaptive immunity, potential crosstalk between CD4+ T cells and ILC2s may also need to be considered. For example, T cell–derived cytokines, such as IL-2, sustain the numbers and activities of ILC2s (24). ILC2s may also regulate the differentiation and longevity of CD4+ T cells. For example, IL-13 derived from lung ILC2s most likely enhances development of Th2-type Ag-specific CD4+ T cells by optimizing Ag presentation by dendritic cells (53). A new mouse model to remove ILC2s specifically and to leave the other immune cell compartments intact, such as those using the Cre-Lox system (54), will most likely dissect the importance of ILC2s in the chronic phase of type 2 airway inflammation in vivo.

A potential weakness and at the same time a strength of this study is the complexity of its model. Each allergen itself, such as HDM and Alternaria, is reported to be sufficient to trigger several immune pathways when administered to the airways. For example, cysteine proteases, Der p 2, LPS, and chitin in the HDM extract could activate protease-activated receptors, TLR4, and dectin, as well as other as yet unidentified receptors; proteases also increase epithelial tight junction permeability (12). Protease-like activities in Alternaria extract can trigger ATP release, resulting in the activation of purinergic receptors on airway epithelia cells (48). A more recent study suggests that, among various natural allergens, serine protease activity from Alternaria uniquely induces IL-33 secretion in mice (55). Conceivably, a combination of these allergens can trigger multiple receptors that may work synergistically or even in some instances antagonistically to regulate airway immunity. Therefore, our model is unlikely to be optimal to dissect immune mechanisms in detail, such as the identification of specific receptor(s) to recognize allergens. Nonetheless, we could also consider this model relevant to a natural condition in which human airways are exposed constantly to a number of environmental
factors, including multiple allergens as well as other environmental stresses. Considering its complexity, the evidence obtained in this study in support of the central importance of IL-33 and perhaps TSLP may be highly valuable. Further studies on the molecular and cellular mechanisms that are involved in IL-33 and TSLP production, in particular those in chronically inflamed airway tissues, will help us better understand the immunological mechanisms of asthma and develop novel therapeutic strategies for the disease.

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FIGURE 7. IL-33 and TSLP play significant roles in type 2 cytokine production and the development of AHR in response to chronic allergen exposure. (A–C) Naïve wild-type mice or mice deficient in cytokine receptors were exposed to allergens, as in Fig. 6. The lung levels of IL-4 (A), IL-5 (B), or IL-13 (C) were analyzed. Results are the mean ± SEM (n = 5 in each group) and are representative of two independent experiments. *p < 0.05, **p < 0.01, compared with wild-type mice. (D–F) Naïve wild-type mice or mice deficient in cytokine receptors were exposed intranasally to PBS or allergens (OAAH) for 4 wk. Airway reactivity to inhaled methacholine was analyzed, as described in Materials and Methods. (D) The dose response to methacholine in wild-type BALB/c mice as examined by whole-body plethysmography is presented. Results are the mean ± SEM (n = 20 in each group, a pool of 4 experiments). *p < 0.05, compared with mice exposed to PBS. (E) The dose response to methacholine in wild-type BALB/c mice as examined by a forced oscillation technique is presented. Results are the mean ± SEM (n = 5 in each group). *p < 0.05, compared with mice exposed to PBS. (F) Airway reactivity to 6 mg/ml methacholine is presented. Results are the mean ± SEM (n = 5 in each group). *p < 0.05, between the groups indicated by horizontal lines.

Disclosures
The authors have no financial conflicts of interest.

References


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