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Priya Rajavelu, ... , Thomas R. Korfhagen, Jeffrey A. Whitsett

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Research Article

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Airway epithelial SPDEF integrates goblet cell differentiation and pulmonary Th2 inflammation

Priya Rajavelu, Gang Chen, Yan Xu, Joseph A. Kitzmiller, Thomas R. Korfhagen, and Jeffrey A. Whitsett

Perinatal Institute, Divisions of Neonatology, Perinatal and Pulmonary Biology, Cincinnati Children's Hospital Medical Center, and the University of Cincinnati College of Medicine, Cincinnati, Ohio, USA.

Epithelial cells that line the conducting airways provide the initial barrier and innate immune responses to the abundant particles, microbes, and allergens that are inhaled throughout life. The transcription factors SPDEF and FOXA3 are both selectively expressed in epithelial cells lining the conducting airways, where they regulate goblet cell differentiation and mucus production. Moreover, these transcription factors are upregulated in chronic lung disorders, including asthma. Here, we show that expression of SPDEF or FOXA3 in airway epithelial cells in neonatal mice caused goblet cell differentiation, spontaneous eosinophilic inflammation, and airway hyperresponsiveness to methacholine. SPDEF expression promoted DC recruitment and activation in association with induction of *Il33*, *Csf2*, thymic stromal lymphopoietin (*Tslp*), and *Ccl20* transcripts. Increased *Il4*, *Il13*, *Ccl17*, and *Il25* expression was accompanied by recruitment of Th2 lymphocytes, group 2 innate lymphoid cells, and eosinophils to the lung. SPDEF was required for goblet cell differentiation and pulmonary Th2 inflammation in response to house dust mite (HDM) extract, as both were decreased in neonatal and adult *Spdef*^{-/-} mice compared with control animals. Together, our results indicate that SPDEF causes goblet cell differentiation and Th2 inflammation during postnatal development and is required for goblet cell metaplasia and normal Th2 inflammatory responses to HDM aeroallergen.

Introduction

The tracheobronchial tree is lined by a diversity of epithelial cells including basal, ciliated, serous, and goblet cells that make unique contributions to mucociliary clearance, innate immunity, and recruitment and activation of professional immune cells that mediate inflammatory responses to environmental exposures (1–5). The nature and extent of innate and acquired immune responses to respiratory pathogens and allergens are influenced genetically and developmentally and by exposure to specific pathogens and toxicants. There is increasing evidence that pulmonary immune responses are initiated early in development, in utero, and during infancy (6–10). Mechanisms by which early exposures of the lung to pathogens instruct the acquisition of innate and acquired immunity are of considerable interest. For example, wheezing following infection of infants to rhinovirus (RV) strongly predicts subsequent risks for development of asthma later in life (11, 12). The observations that the incidence of childhood asthma is lower in infants exposed to an abundance of bacterial and fungal microbes supports the concept that early environmental exposures influence the development of asthma (13). Mechanisms influencing the acquisition of Th1- and Th2-mediated immunity in infancy have important implications for

host defense and allergic responses that mediate subsequent responses to environmental stimuli throughout life (14). Recent studies in gnotobiotic mice demonstrate an age-dependent programming of invariant NK cells that influences subsequent allergic responses in the lung (14). Respiratory epithelial cells are increasingly recognized as important modulators of innate immune responses that govern inflammatory and immune responses to pathogens and toxicants via their synthesis of chemokines and cytokines, growth factors that regulate the migration and activation of the diverse immune cells to sites of injury (refs. 15–17, for review). Airway epithelial cells mount rapid and robust responses to a diversity of pathogen-associated molecular pattern molecules (PAMPs), damage-associated molecular pattern molecules (DAMPs) that influence both innate immune and repair processes that maintain pulmonary homeostasis. The airway epithelial barrier is maintained by a diversity of specific epithelial cell types that each contribute uniquely in response to airway injury. Basal and nonciliated progenitor cells serve as precursors from which ciliated and goblet cells are derived (2, 18). Airway goblet cells synthesize and secrete a variety of mucins and other proteins that play important roles in mucociliary clearance and host defense (19). While the activation of goblet cells and mucus secretion are normal responses to acute injury and infection, chronic goblet cell hyperplasia and metaplasia are features of common chronic pulmonary diseases, including asthma, cystic fibrosis (CF), and chronic obstructive pulmonary disease (COPD), wherein mucus hyperproduction contributes to the mortality and morbidity associated with these disorders (19–21).

Airway goblet cell differentiation is influenced by the transcription factors SAM-pointed domain-containing ETS-like factor (SPDEF) and forkhead ortholog A3 (FOXA3), which are

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selectively expressed in airway goblet cells and induced by toxicants, aeroallergens, viral infections, and cytokines. SPDEF is sufficient and required for goblet cell differentiation and mucus production in the mouse airway (22, 23). SPDEF and FOXA3 are induced following exposure to RV, aeroallergens, and Th2 cytokines, including IL-13, that enhance goblet cell differentiation and regulate a network of genes involved in the synthesis and packaging of airway mucins (22–25). SPDEF induces FOXA3 in airway epithelial cells, the latter also being sufficient to cause goblet cell metaplasia in the adult mouse lung (24). The present study was designed to test whether goblet cell differentiation induced by SPDEF or FOXA3 in the perinatal period influences the development of pulmonary innate immune responses. We demonstrated that expression of either SPDEF or FOXA3 in nonciliated airway epithelial cells of the developing mouse lung caused extensive goblet cell differentiation and Th2-mediated pulmonary inflammation. SPDEF was required for Th2 inflammatory responses to house dust mite (HDM) extract in neonatal and adult mice and was necessary for FOXA3-induced goblet cell differentiation and Th2 inflammation.

Results

SPDEF causes Th2-mediated inflammation and airway hyperresponsiveness. To assess the potential role of SPDEF in the regulation of goblet cell metaplasia and inflammation in the developing lung, *Spdef* mRNA was conditionally expressed in nonciliated respiratory epithelial cells under the control of the *Scgb1a1* promoter induced by administration of doxycycline to the dam from E16 to P15. On P15, SPDEF caused extensive goblet cell metaplasia in conducting airways, increasing *Foxa3*, *Muc5ac*, and *Muc5b* mRNAs and immunostaining (Figure 1A and Table 1). In contrast to the lack of inflammation following transient expression of SPDEF in adult mice (22, 23), SPDEF caused widespread inflammatory infiltrates consisting primarily of eosinophils in the neonatal mice (Figure 1, A and C). Airway hyperresponsiveness (AHR), determined by methacholine challenge tests, was augmented in association with increased α SMA (*Acta2*) mRNA (Figure 1B and Table 1). Consistent with the observed eosinophilic inflammation, mRNAs encoding cytokines and chemokines that mediate Th2 inflammation were significantly increased, including those mediating DC activation (*Il33*, *Il25*, *Csf2*); likewise, Th2 cytokines (*Il13*, *Ccl17*) and eosinophilic chemoattractants (*Ccl24* and *Ccl20*) were increased. SPDEF induced *Ear11* mRNA, which was associated with the eosinophilic inflammation noted histologically (Table 1).

Consistent with increased expression of Th2 cytokines and chemokines, the numbers of ST2⁺ and IL-17RB⁺ innate lymphoid cells (ILCs), CD3⁺ T lymphocytes, and IL-4⁺ T cells were increased by SPDEF, while IFN- γ CD3⁺ cell numbers were unaltered (Figure 1, D and E, and Supplemental Figure 1; supplemental material available online with this article; doi:10.1172/JCI79422DS1). Taken together, expression of SPDEF in nonciliated respiratory epithelial cells in the newborn period caused extensive goblet cell metaplasia, spontaneous Th2 pulmonary inflammation, and increased AHR, findings likely mediated by airway epithelial cell elaboration of the cytokines and chemokines that recruit and activate DCs, ILCs, and lymphocytes to mediate a Th2 inflammatory cascade in the neonatal-postnatal lung.

Table 1. SPDEF induced mRNAs related to goblet cell differentiation and Th2 inflammation

RNA	Fold increase (mean \pm SEM)
	<i>Scgb1a1</i> -rtTA TRE2- <i>Spdef</i>
<i>Spdef</i>	283 \pm 49.8 ^B
<i>Foxa3</i>	3.1 \pm 0.5 ^A
<i>Muc5ac</i>	2.0 \pm 0.4 ^A
<i>Acta2</i>	2.2 \pm 0.5 ^A
<i>Tslp</i>	1.5 \pm 0.2
<i>Ccl20</i>	4.2 \pm 1.15 ^A
<i>Il33</i>	2.0 \pm 0.2 ^A
<i>Csf2</i>	1.6 \pm 0.1 ^A
<i>Il25</i>	2.1 \pm 0.3 ^A
<i>Il4</i>	1.2 \pm 0.1
<i>Il13</i>	12.1 \pm 1.2 ^A
<i>Ccl17</i>	2.4 \pm 0.5 ^A
<i>Ccl11</i>	3.1 \pm 1.3 ^A
<i>Ccl24</i>	4.2 \pm 1.1 ^A
<i>Ear11</i>	14.5 \pm 4.5 ^A
<i>Il5</i>	3.1 \pm 1.6

qRT-PCR was performed on whole lungs from *Spdef*-expressing and control mice to assess *Spdef*, *Foxa3*, *Muc5ac*, and *Acta2* mRNAs and those encoding Th2-associated chemokines and cytokines, as described in Figure 1. Data represent the mean \pm SEM. ^A*P* < 0.05 and ^B*P* < 0.01 by unpaired, 2-tailed Student's *t* test. *n* = 6–8 mice for each analysis.

FOXA3 causes goblet cell metaplasia and Th2 inflammation in neonatal mice. In adult mice and in human airway epithelial cells in vitro, FOXA3 induced SPDEF and caused goblet cell metaplasia (24). To test whether FOXA3 was sufficient to cause goblet cell metaplasia in neonatal mice, FOXA3 was conditionally expressed in nonciliated airway epithelial cells under the control of the same *Scgb1a1* promoter by placing the dams on doxycycline from E16.5 to P15. When FOXA3 was selectively expressed in Club cells, SPDEF was induced and SCBG1A1 (CCSP) staining decreased but was still present in the goblet cells induced by FOXA3, where it was coexpressed with MUC5B (Figure 2A and Supplemental Figure 2). In contrast, ciliated cells did not express MUC5B or SCBG1A1, consistent with the specificity of the *Scgb1a1* promoter in Club cells. FOXA3 caused extensive eosinophilic inflammation and increased AHR in response to methacholine, findings similar to those observed after expression of SPDEF (Figure 2, A–C). FOXA3 caused goblet cell differentiation associated with increased expression of SPDEF, MUC5AC, and MUC5B (Figure 2A and Supplemental Figure 2). As observed in SPDEF-expressing mice, pulmonary infiltrates consisted of increased numbers of eosinophils and ILCs, including ST2⁺, IL-17RB⁺, and ICOS⁺ lineage-negative cells; CD103⁺ DCs and IL-4⁺ CD3 lymphocytes were also increased (Figure 2, D–G). mRNAs, including *Tslp*, *Il33*, and *Csf2*, that encode factors mediating DC chemotaxis and activation were significantly increased by FOXA3. *Il13*, *Il4*, *Ccl17*, and *Ccl24* mRNAs were increased, consistent with the increased Th2-mediated inflammation induced by FOXA3 (Table 2). Taken together, FOXA3 expression in respiratory epithelial cells of postnatal mice induced SPDEF and caused goblet cell metaplasia

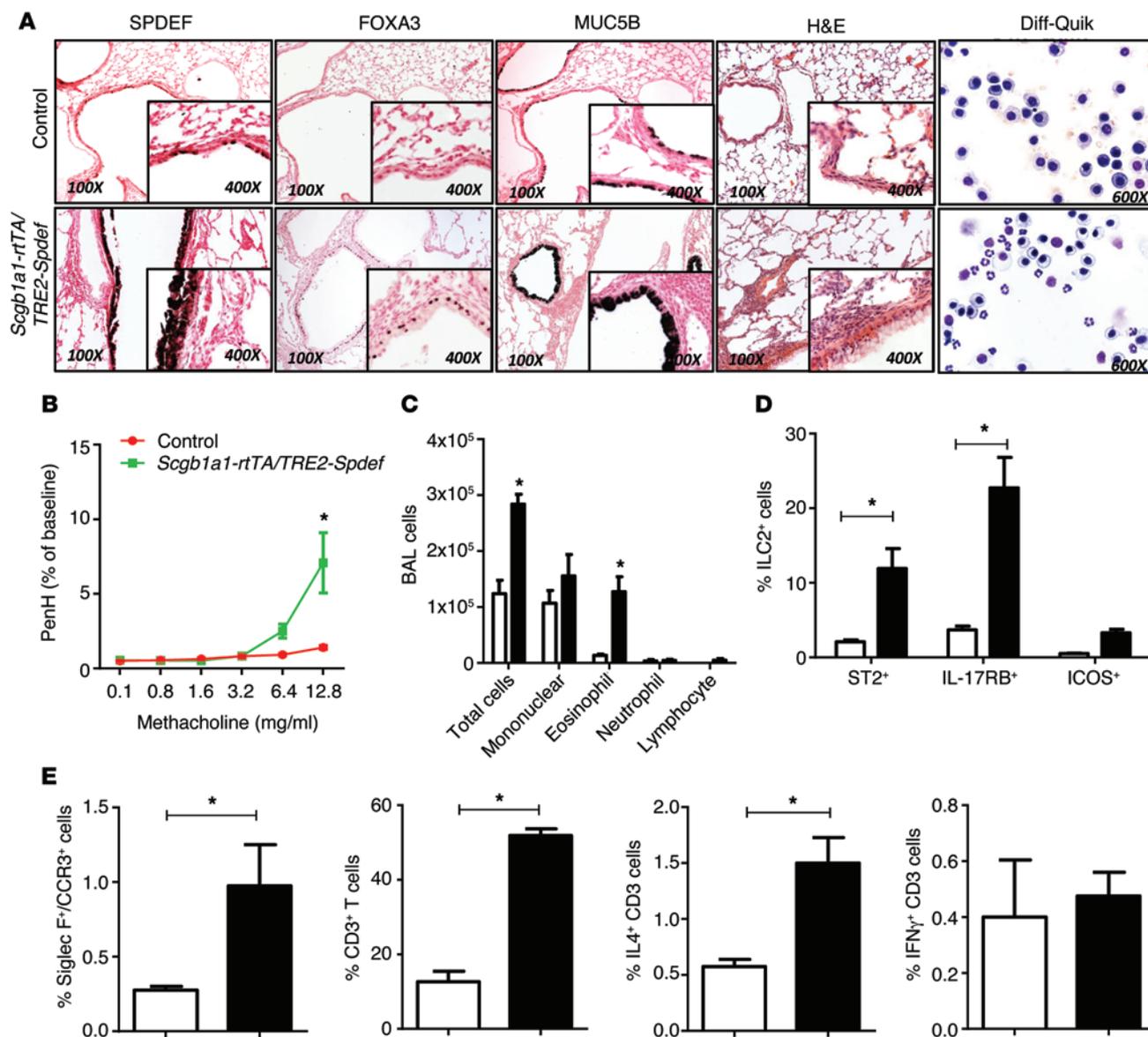


Figure 1. SPDEF causes goblet cell differentiation, pulmonary inflammation, and AHR. Dams of *Scgb1ab-rtTA TRE-Spdef* (black bars) and control littermate single-transgenic mice (white bars) were placed on doxycycline from E16.5 to P15. (A) At P15, lung histology of the pups was assessed by H&E ($n = 8$) and BALF ($n = 4$) staining with Diff-Quik. Immunostaining for SPDEF, FOXA3, and MUC5B demonstrated extensive goblet cell differentiation ($n = 8$). (A and C) Cells isolated from BALF were stained with Diff-Quik and showed eosinophilic infiltrates. Data represent the mean \pm SEM. * $P < 0.05$ compared with controls using an unpaired, 2-tailed Student's t test. (B) AHR is represented as PenH in response to methacholine. Data represent the mean \pm SEM of 6 mice per group. * $P < 0.05$ by 2-way ANOVA. (E) Flow cytometric analysis of lung cells obtained at P15 demonstrated increased numbers of SiglecF⁺CCR3⁺ in *Scgb1a1-rtTA Spdef* mice (black bar) compared with cell numbers detected in littermate control mice (white bar). (D and E) ST2⁺, IL-17RB⁺, and ICOS⁺ ILCs and total CD3⁺ and CD3⁺ IL-4-producing T cells were increased, and IFN- γ -producing CD3⁺ T cells were unaltered. Data represent the mean \pm SEM. * $P < 0.05$ compared with controls using an unpaired, 2-tailed Student's t test. $n = 4$ /group. Examples of FACS analyses are provided in Supplemental Figure 1.

and Th2-dominated pulmonary inflammation, both of which are essentially phenocopying effects of SPDEF.

SPDEF is required for HDM-induced pulmonary inflammation and AHR. Since SPDEF and FOXA3 were sufficient to induce Th2-mediated pulmonary inflammation in the postnatal period, we tested whether the absence of SPDEF influenced goblet cell metaplasia and pulmonary inflammation following exposure to HDM extract in both neonatal and adult mice. Adult *Spdef*^{+/+} and *Spdef*^{-/-} littermates were treated i.n. daily for 3 days with 100 μ g HDM extract. AHR, goblet cell differentiation, eosinophilic infil-

tration, and Th2 inflammatory markers were markedly suppressed in the *Spdef*^{-/-} mice (Figure 3, A-C, and Table 3). In control mice, HDM caused Th2-associated lung inflammation, goblet cell differentiation, and increased AHR. Induction of *Foxa3*, *Muc5b*, and *Muc5ac* mRNAs and those associated with goblet cell differentiation seen after HDM exposure were inhibited in the *Spdef*^{-/-} mice (Table 3). Likewise, expression of mRNAs encoding the DC-related chemoattractants *Il33* and *Ccl20* and Th2 lymphocyte-associated *Ccl17*, *Il4*, *Il13* were suppressed, as was eosinophilic infiltration (Table 3). Consistent with the latter findings, the

Table 2. FOXA3 induced mRNAs related to goblet cell differentiation and Th2 inflammation

mRNA	Fold increase (mean ± SEM)
	<i>Scgb1a1-rtTA Foxa3</i>
<i>Foxa3</i>	59.54 ± 21.1 ^A
<i>Spdef</i>	1.61 ± 0.3
<i>Muc5ac</i>	4.40 ± 1.1 ^A
<i>Muc5b</i>	2.50 ± 0.5 ^A
<i>Tslp</i>	1.45 ± 0.2 ^A
<i>Ccl20</i>	2.00 ± 0.4
<i>Il33</i>	3.47 ± 0.6 ^A
<i>Csf2</i>	2.13 ± 0.3 ^A
<i>Il4</i>	3.52 ± 0.5 ^A
<i>Il13</i>	8.71 ± 1.5 ^A
<i>Ifnγ</i>	0.95 ± 1.5
<i>Ccl17</i>	3.56 ± 0.8 ^A
<i>Ccl11</i>	0.92 ± 0.1
<i>Ccl24</i>	2.49 ± 0.4 ^A

qRT-PCR was performed on whole lungs from *Foxa3*-expressing and control mice, as described in Figure 2. qRT-PCR showed that FOXA3 increased *Muc5b*, *Muc5ac*, *Tslp*, *Il33*, *Csf2*, *Il4*, *Il13*, *Ccl17*, and *Ccl24* mRNAs. Data represent the mean ± SEM. ^A*P* < 0.05 compared with controls using an unpaired, 2-tailed Student's *t* test. *n* = 8–10 mice/group.

induction of *Ear11* and *Ccl24* mRNAs after HDM treatment was inhibited in the *Spdef*^{-/-} mice. *Il17* mRNA and IL-17⁺CD3⁺ cells were decreased, and recruitment of ST2⁺ ILCs and T cells expressing IL-4, IL-13, or IL-17 were reduced in mice lacking SPDEF following HDM exposure (Figure 3, D and E, and Table 3). To assess whether effector T cells were deficient in *Spdef*^{-/-} mice, anti-IgD serum was administered to directly induce T cell activation independently of pulmonary sensitization (26). Anti-IgD directly induced IL-4 expression and caused lymph node and spleen enlargement, demonstrating intact effector T cell responses in *Spdef*^{-/-} mice. Consistent with intact T cell effector functions, serum IgE and IgG1 levels were similar in *Spdef*^{-/-} and *Spdef*^{+/+} mice (Supplemental Figure 3). Taken together, SPDEF is required for HDM-induced goblet cell differentiation, Th2 inflammation, and pulmonary AHR responses in adult mice. To test whether SPDEF is required for inflammatory responses to HDM in the developing lung, neonatal *Spdef*^{+/+} and *Spdef*^{-/-} mice were treated with 50 μg HDM extract. Consistent with our findings in adult mice, SPDEF was required for normal Th2 responses, mucous metaplasia, eosinophilic inflammation, and AHR in the neonatal mice exposed to HDM (Figure 4 and Table 4).

SPDEF is required for FOXA3-induced pulmonary inflammation. Since expression of either SPDEF or FOXA3 was sufficient to induce Th2 inflammatory responses in neonatal mice, we tested whether SPDEF was required for the effects of FOXA3. Th2 inflammatory mediators, goblet cell differentiation, and AHR caused by expression of FOXA3 were significantly inhibited in neonatal *Spdef*^{-/-} as well as in *Spdef*^{+/+} mice (Supplemental Figure 4). Increased expression of *Muc5ac*, *Muc5b*, *Tslp*, *Il4*, and *Il13* mRNAs was decreased in FOXA3-expressing *Spdef*^{-/-} and *Spdef*^{+/+} mice compared with that observed in *Spdef*^{+/+} littermates (Supple-

mental Figure 4D). Methacholine challenge tests demonstrated that the effects of FOXA3 on AHR were inhibited in *Spdef*^{-/-} and *Spdef*^{+/+} mice (Supplemental Figure 4, A and B). Thus, SPDEF plays a central role in both HDM- and FOXA3-induced goblet cell metaplasia and pulmonary inflammation.

FOXA3 enhances pulmonary responses to HDM. While FOXA3 was not required for the induction of Th2-mediated inflammation by aeroallergen in adult mice (24, 25), the present findings in neonatal mice demonstrated that FOXA3 was sufficient to induce Th2-mediated inflammation. Since the effects of SPDEF and FOXA3 in pulmonary inflammation were similar, we assessed whether FOXA3 further enhanced inflammatory responses during aeroallergen challenge. *Scgb1a1-rtTA OTet7-Foxa3* conditional mice were exposed to HDM extract (50 μg) delivered i.n. AHR to HDM extract was increased in FOXA3-expressing mice (Supplemental Figure 5). Likewise, *Spdef* and other goblet cell-associated *Muc5ac* and *Muc5b* mRNA levels were further increased in the FOXA3-expressing mice after exposure to HDM. mRNAs indicating Th2-related DC and lymphocytic activity were increased by expression of FOXA3 after HDM exposure (Supplemental Figure 5C). Consistent with these findings, *Spdef* haploinsufficiency inhibited the enhanced AHR response to HDM in *Foxa3*-expressing transgenic mice (Supplemental Figure 4B).

Anti-TSLP ameliorates the inflammatory effects of FOXA3. Thymic stromal lymphopoietin (TSLP) is expressed by respiratory epithelial cells and is a known mediator of Th2-dependent inflammation in multiple organs, including skin, intestine, and lung (27, 28). We interrogated previous FOXA3-associated ChIP data from our recent studies (24) and identified strong binding of FOXA3 to the promoter region of the human *TSLP* gene (Supplemental Figure 6). To test the role of TSLP in the induction of FOXA3-generated pulmonary inflammation in neonatal mice, TSLP activity was inhibited by injection of these mice with anti-TSLP Ab on P3 during the induction of FOXA3 (29, 30). Anti-TSLP substantially inhibited pulmonary inflammation and MUC5B staining (Supplemental Figure 7A). TSLP Ab also inhibited *Spdef*, *Muc5ac*, *Csf2*, *Ccl11*, *Ear11*, *Il4*, *Tslp*, *Il5*, *Ccl24*, *Il13*, and *Ccl17*, indicating that TSLP is an important mediator of FOXA3-induced inflammation (Supplemental Figure 7B).

Discussion

Pulmonary innate and acquired immunity is rapidly acquired following birth and exposure of pulmonary tissues to commensal and pathogenic microbes, allergens, and toxicants that recruit BM-derived immune cells to the lung. The present findings demonstrate that the selective expression of SPDEF or FOXA3, both transcription factors normally restricted to conducting airways and submucosal gland epithelial cells, was sufficient to induce airway goblet cell differentiation in the neonatal-postnatal period. Either transcription factor was sufficient to cause the spontaneous pulmonary Th2 and eosinophilic inflammation associated with enhanced recruitment and activation of ST2⁺ group 2 ILCs, DCs, and Th2-expressing lymphocytes, mediated in part by the expression of cytokines and chemokines by respiratory epithelial cells. Remarkably, SPDEF was required for Th2 responses to HDM extract in both neonatal and adult mice and for the inflammatory effects of FOXA3. The present findings demonstrate a critical role for SPDEF and goblet cells in the acquisition of pulmonary Th2 immune responses.

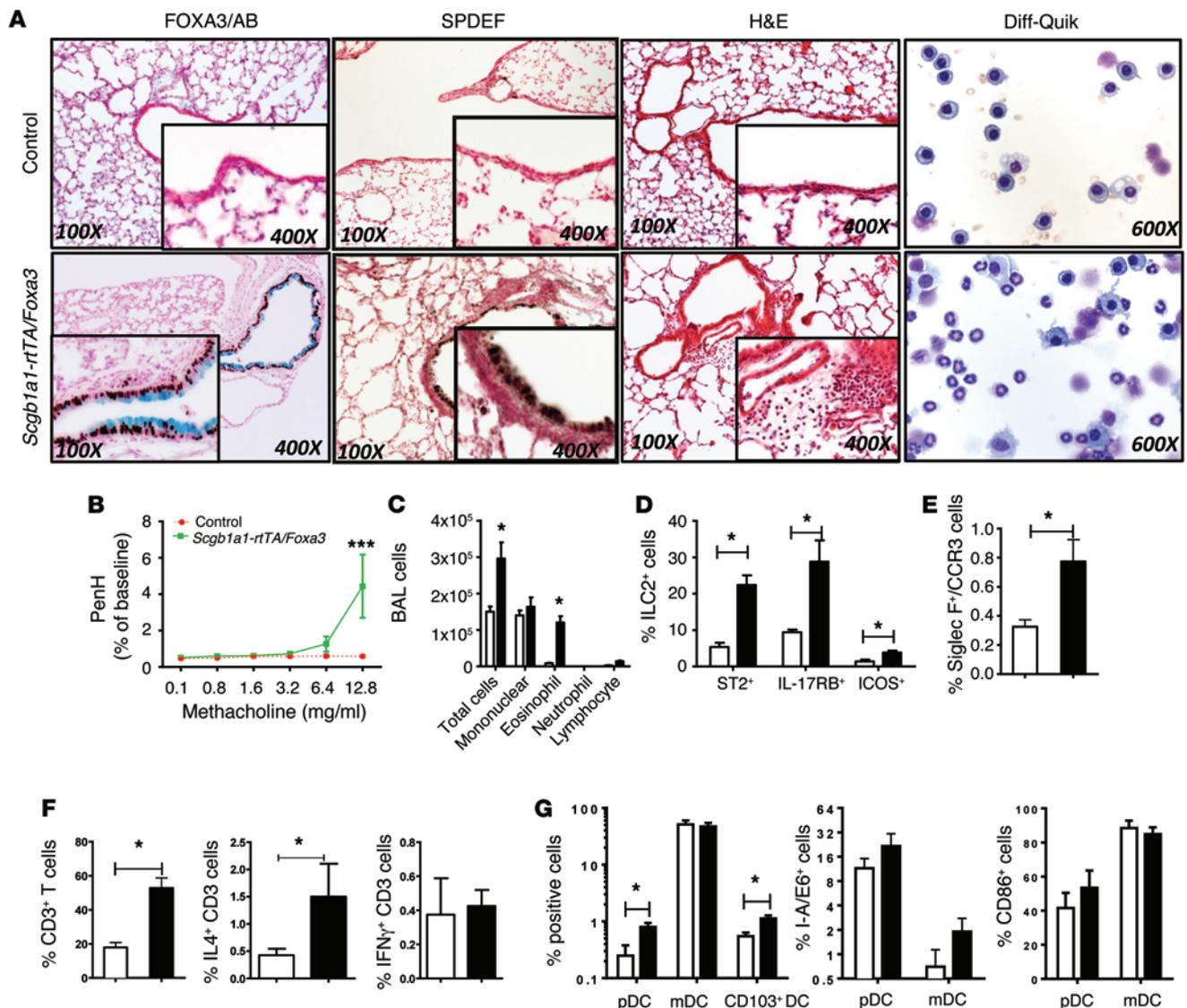


Figure 2. Conditional expression of FOXA3 in airway epithelial cells causes pulmonary inflammation and goblet cell metaplasia. (A–C) Control littermates (white bars) and *Scgb1ab-rtTA OTet7-Foxa3* dams (black bars) were placed on doxycycline from E16.5 to P15. Lung tissue was obtained from the pups at P15. *Scgb1ab-rtTA OTet7-Foxa3* pups (black bars) developed spontaneous eosinophilic inflammation, goblet cell metaplasia, and AHR. Diff-Quik staining of BALF ($n = 4$) demonstrated eosinophilic infiltration. (E) Flow cytometric analysis of lung cells obtained at P15 demonstrated increased numbers of SiglecF⁺CCR3⁺, (D–F) ST2⁺, IL-17RB⁺, and ICOS⁺ ILCs and CD3⁺ IL-4-producing T cells. IFN- γ -producing CD3⁺ T cells were unaltered. (G) pDCs (CD11b⁻CD11c⁻B220⁺), CD103⁺ mDCs (CD11b⁺CD11c⁻B220⁺) were increased and mDCs were unchanged in lung from *Scgb1a1-rtTA OTet7-Foxa3* pups. Activation markers (I-A/E and CD86) on pDCs and mDCs are shown in G. Data represent the mean \pm SEM. * $P < 0.05$ compared with controls using an unpaired, 2-tailed Student's t test. $n = 4$ /group.

Respiratory epithelial cells are increasingly recognized as active mediators of innate host defense and inflammation, playing multiple roles in mucociliary clearance, barrier function, and recruitment and activation of professional immune cells, the latter being mediated by the elaboration of diverse cytokines and chemokines (15–17). In contrast to the Th2 inflammatory responses seen in the present study in neonatal mice after expression of SPDEF or FOXA3, the expression of these transcription factors in adult mice caused reversible goblet cell differentiation from nonciliated airway epithelial cells (Club cells) and increased mucus production, but did not cause acute inflammation (22–24). Consistent with those studies, subsets of goblet cells induced by expression of FOXA3 coexpressed CCSP and MUC5B, albeit at

lower levels for CCSP (Supplemental Figure 2). The present findings support the concept that SPDEF and FOXA3, perhaps in part related to goblet cell differentiation, influence the recruitment and activation of professional immune cells during the adaptation to exposures to aeroallergens and microbes after birth. SPDEF and FOXA3, transcription factors expressed selectively in nonciliated airway epithelial cells, activate a number of genes regulating mucus metaplasia, including *MUC5AC*, *MUC5B*, *AGR2*, and other proteins involved in the synthesis, glycosylation, and packaging of mucus (23, 31). While a rare subset of airway epithelial cells in normal mouse lung, goblet cells are relatively abundant in conducting airways of the neonatal mouse, and their numbers increase in the first weeks of life (32). Acute goblet cell metaplasia is associ-

Table 3. SPDEF was required for Th2 responses to HDM extract in adult mice

mRNA	Fold change (mean ± SEM)	
	WT + HDM	<i>Spdef</i> ^{-/-} + HDM
<i>Foxa3</i>	0.8 ± 0.2	0.1 ± 0.1 ^A
<i>Acta2</i>	1.7 ± 0.032	1.2 ± 0.03 ^B
<i>Muc5ac</i>	3.1 ± 1.0	1.6 ± 0.1
<i>Muc5b</i>	0.9 ± 0.8	0.4 ± 0.2
<i>Tslp</i>	1.2 ± 0.1	1.05 ± 0.2
<i>Ccl20</i>	2.4 ± 0.5	0.7 ± 0.1 ^A
<i>Il33</i>	12.1 ± 1.0	3.9 ± 0.9 ^A
<i>Csf2</i>	2.9 ± 0.07	2.4 ± 0.2
<i>Il25</i>	1.1 ± 0.1	1.3 ± 0.5
<i>Il4</i>	39.5 ± 7.8	14.1 ± 8.4 ^A
<i>Il13</i>	227 ± 54	67 ± 29.5 ^A
<i>Ccl17</i>	2.7 ± 0.2	1.2 ± 0.1 ^A
<i>Il17a</i>	11.9 ± 5.0	2.7 ± 0.2 ^A
<i>Ear11</i>	2489 ± 464	346 ± 225 ^A
<i>Ccl11</i>	9.1 ± 1.7	5.2 ± 1.8
<i>Ccl24</i>	27.1 ± 0.1	3.2 ± 1.0 ^A

qRT-PCR was performed on whole lungs from *Spdef*^{-/-} and *Spdef*^{+/+} mice exposed to HDM extract, as described in Figure 3. WT saline-treated mice represent the baseline for comparison of *Spdef*^{-/-} and *Spdef*^{+/+} mice. After exposure to HDM extract, mRNAs regulating DC chemotaxis and activation (*Il33* and *Ccl20*), Th2- and Th17 lymphocyte-associated mRNAs (*Il4*, *Il13*, *Ccl17*, and *Il17a*), and eosinophil-associated mRNAs (*Ear11* and *Ccl24*) were decreased in lungs from adult *Spdef*^{-/-} mice compared with those from *Spdef*^{+/+} mice, as assessed by qRT-PCR. ^A*P* < 0.05 and ^B*P* < 0.01 by 2-way ANOVA, followed by Bonferroni's post test. *n* = 6/group.

ated with bacterial and viral infections and exposure to allergens and toxicants (17, 20, 21, 33, 34). Chronic goblet cell metaplasia is associated with increased SPDEF and FOXA3 expression in the setting of asthma, COPD, CF, and other chronic inflammatory diseases of the lung (24). We speculate that early infectious or other inflammatory exposure inducing goblet cell differentiation via SPDEF may serve to enhance Th2 polarization of pulmonary innate immunity to influence subsequent inflammatory responses to environmental challenges.

While mechanisms by which SPDEF induces Th2 inflammatory responses in neonatal mice are likely complex, expression of cytokines and chemokines, known to be products of epithelial cells, e.g., IL-33, TSLP, IL-25, and granulocyte-macrophage CSF (GM-CSF), were spontaneously induced by expression of SPDEF and FOXA3 in neonatal mice. Professional immune cells are actively recruited to the lung following birth by exposure to the environment and release of chemokines by pulmonary cells; for example, monocyte and macrophage recruitment is mediated by the expression of GM-CSF by pulmonary parenchymal cells (35, 36). In the present studies, SPDEF and FOXA3 induced chemokines and cytokines known to attract and activate group 2 ILCs and DCs, recruiting and sensitizing lymphocytes that together elaborate Th2-associated cytokines such as IL-4, IL-5, and IL-13 (37, 38). Since IL-13 and IL-4 regulate SPDEF in airway epithelial cells through IL-4 receptor signaling and STAT6 (22), their expres-

sion by Th2 lymphocytes and ILCs may serve to further enhance SPDEF and FOXA3 expression. Since a number of cytokines and chemokines were induced by SPDEF and FOXA3, the mediators of pulmonary Th2 inflammation are likely complex. TSLP, a Th2 chemokine expressed by epithelial cells that are expressed, for example, in skin, gastrointestinal tract, and lung in response to injury and allergens (38–40), was induced by either SPDEF or FOXA3; anti-TSLP Ab substantially inhibited FOXA3-induced Th2-mediated inflammation. However, anti-TSLP decreased expression of SPDEF, perhaps influencing expression of Th2 cytokines (e.g., IL-4 and IL-13) known to induce SPDEF. Surprisingly, anti-TSLP treatment resulted in increased expression of FOXA3 (Supplemental Figure 7), perhaps indicating lack of a counterregulatory response, although it is unclear in the present experiments whether the increase in *Foxa3* mRNA resulted from the endogenous *Foxa3* gene or from changes in the activity of the *Scgb1a1* transgene expressing the *Foxa3* cDNA. Recent in vivo and in vitro studies showed that FOXA3 induced TSLP expression in human airway epithelial cells in vitro (24); ChIP of FOXA3-associated transcriptional targets identified direct binding of FOXA3 to the promoter region of the human *TSLP* gene, supporting the concept that TSLP represents an important signaling molecule produced by the airway epithelial cells that regulate Th2 inflammatory responses to FOXA3 and SPDEF. FOXA3 binding was also detected in the first intron and in the 3' region of the *CCL17* gene (Supplemental Figure 6).

While FOXA3 was not required for goblet cell differentiation and inflammation following antigen sensitization in mice (24, 25), we demonstrated that SPDEF was required for normal Th2 inflammatory responses to HDM allergen in both postnatal and adult mice. Both SPDEF and FOXA3 have antiinflammatory effects on airway epithelial cells, probably functioning by distinct mechanisms. SPDEF bound MyD88 and TRIF, inhibiting NF-κB and interferon responses (41). In contrast, FOXA3 inhibited IFN responses to RV, thereby impairing RV clearance, which was mediated in part by the inhibitory effects of FOXA3 on IRF3 and IFN production (24). Taken together with its ability to enhance expression of Th2-polarizing cytokines and chemokines, we speculate that inhibition of Th1 and IFN responses by SPDEF and FOXA3 may favor Th2 responses to inflammatory stimuli. HDM activates NF-κB through TLR4, the latter being required for normal Th2 responses in the lung (33, 34). While SPDEF and FOXA3 regulate the expression of a number of modulators of Th2 inflammatory responses to HDM, the precise drivers of the spontaneous Th2 inflammatory responses caused by the expression of SPDEF and FOXA3 are unclear. Increased expression of both DC and ILC chemoattractants and activation of Th2 lymphocytes induced by SPDEF or FOXA3 likely serve to enhance Th2-type inflammation in the present models. Alternatively, goblet cells induced by SPDEF may have a direct role in enhancing innate immune responses. Recent findings that goblet cells provide conduits for transepithelial delivery of tolerogenic antigens from the intestinal lumen to CD103⁺ DCs provide a precedent for a role for goblet cells in communicating with DCs to influence innate immune responses in epithelia (42).

SPDEF and FOXA3 were sufficient to enhance AHR assessed by “enhanced pause” (Penh); however, it is unclear whether

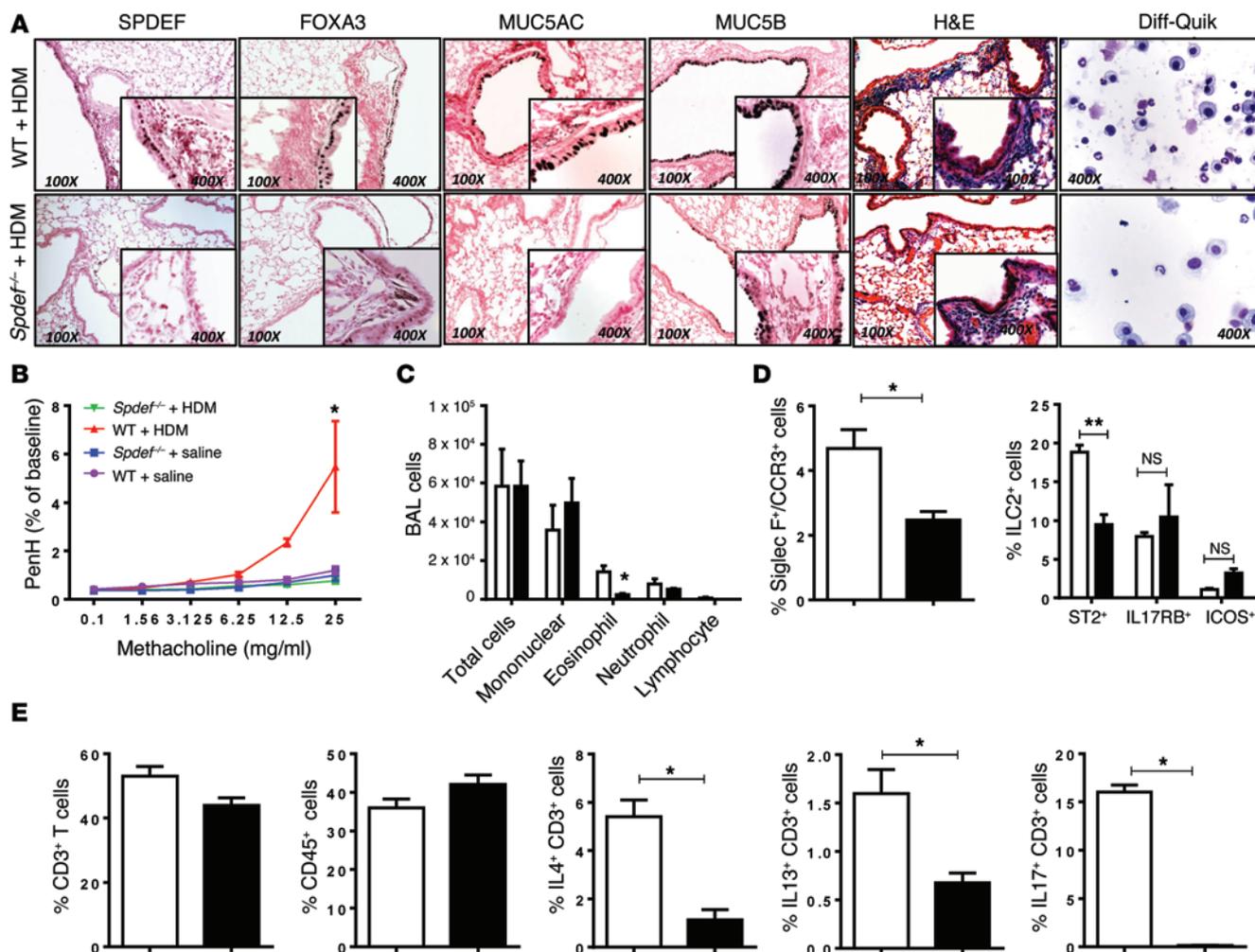


Figure 3. SPDEF is required for Th2 inflammatory responses, AHR, and recruitment of ST2⁺ ILCs and IL-4⁺, IL-13⁺, and IL-17⁺ T cells in response to HDM extract in adult mice. HDM extract (100 μ g) was administered i.n. daily for 3 days to adult control and *Spdef*^{-/-} mice, which were sacrificed 2 days later. (A) Decreased mucous metaplasia and inflammatory responses to HDM were seen in *Spdef*^{-/-} mice. (A and C) BALF cells ($n = 4$) were stained with Diff-Quik, which showed eosinophilic and neutrophilic infiltrates, and differential cell counts were quantified. Goblet cell differentiation (A) and the increase in *Acta2* mRNA (Table 3) caused by HDM challenge were modestly decreased in *Spdef*^{-/-} mice. (B) AHR was decreased in *Spdef*^{-/-} mice. Data represent the mean \pm SEM. Significance was analyzed using 2-way ANOVA with Bonferroni's post tests. $n = 6$ /group. (D and E) Flow analysis of lung cells from *Spdef*^{-/-} (white bars) and *Spdef*^{-/-} mice (black bars) after HDM challenge. CD45⁺ and CD3⁺ cell numbers were not altered, and ST2⁺ ILC numbers were decreased. (E) Numbers of IL-4⁺, IL-13⁺, and IL-17⁺ T cells were decreased in *Spdef*^{-/-} mice. Gating strategies are depicted in Supplemental Figure 1. Data represent the mean \pm SEM. * $P < 0.05$ compared with controls using an unpaired, 2-tailed Student's *t* test. $n = 4$ /group.

these findings represent their effects on goblet cell metaplasia or on bronchial smooth muscle. *Acta2* mRNA was increased by expression of *Spdef* and was decreased in both neonatal and adult *Spdef*^{-/-} mice after HDM extract exposure. Since *Acta2* mRNA is expressed in both vascular and bronchial smooth muscle cells, it is unclear whether these findings are directly related to changes in AHR. Given that increased IL-4 receptor signaling is known to enhance AHR (43, 44), the present findings may be related to increased Th2 cytokine production on airway smooth muscle precursors, to direct effects of IL-4 receptor signaling on airway smooth muscle activity, or to the effects of this signaling on goblet cell differentiation. SPDEF is selectively expressed in airway goblet cells but is also detected in breast, prostate, and intestinal epithelial cells (45, 46). In our past and present studies, SPDEF staining in pulmonary tissues was confined to goblet

cells (22–24). While lack of HDM-induced AHR and Th2 inflammation in *Spdef*^{-/-} mice is likely mediated by the cell-autonomous effects of SPDEF in airway epithelial cells, it remains possible that nonepithelial cells were involved in the present study. Direct activation of Th2 lymphocytes was observed in *Spdef*^{-/-} mice injected with anti-IgD. These findings and the effects of epithelium-specific expression of both FOXA3 and SPDEF on Th2 inflammatory responses in the present studies support a primary role for lung epithelial SPDEF. The ability of SPDEF to induce FOXA3 likely further enhances the Th2 polarization of inflammatory responses and goblet cell differentiation. In human cells, SPDEF and FOXA3 activate each other, with ChIP-Seq and RNA analyses supporting the concept that FOXA3 binds to and activates the *SPDEF* promoter to enhance its transcription (24). In those studies, FOXA3-binding sites also were identified in the

Table 4. SPDEF was required for Th2-associated mRNA responses to HDM extract in postnatal mice

mRNA	Fold increase in WT + HDM vs. <i>Spdef</i> ^{-/-} + HDM (mean ± SEM)
<i>Foxa3</i>	4.9 ± 1.1 ^B
<i>Acta2</i>	1.9 ± 0.3 ^B
<i>Muc5ac</i>	3.2 ± 0.9 ^B
<i>Muc5b</i>	2.4 ± 0.5 ^B
<i>Tslp</i>	5.3 ± 1.8 ^B
<i>Ccl20</i>	2.1 ± 0.5 ^A
<i>Il33</i>	2.4 ± 0.8 ^A
<i>Csf2</i>	1.6 ± 0.5
<i>Il25</i>	2.1 ± 0.3
<i>Il4</i>	3.5 ± 1.1 ^A
<i>Il13</i>	1.6 ± 0.3
<i>Il17a</i>	5.2 ± 1.2 ^A
<i>Il5</i>	3.2 ± 0.9
<i>Ear11</i>	2.7 ± 0.9
<i>Ccl11</i>	2.8 ± 0.9

qRT-PCR was performed on whole lung from young *Spdef*^{-/-} and control mice exposed to HDM extract, as described in Figure 4. mRNAs regulating DC chemotaxis and activation (*Tslp*, *Il33*, and *Ccl20*), Th2- and Th17 lymphocyte-associated mRNAs (*Il4* and *Il17a*), and eosinophil-associated mRNAs (*Il5*) were significantly decreased in postnatal *Spdef*^{-/-} mice, as assessed by qRT-PCR. ^A*P* < 0.05 and ^B*P* < 0.01 by unpaired, 2-tailed Student's *t* test. *n* = 6 per group.

promoter regions of the genes known to play roles in goblet cell differentiation, including human *SPDEF*, *MUC5B*, and *AGR2* genes, supporting a direct role for FOXA3 in the regulation of some aspects of goblet cell metaplasia. Actions of FOXA3 on gene expression in human airway epithelial cells in vitro indicated both SPDEF-dependent and -independent roles for FOXA3 in human airway cells in vitro (24). In the present studies, inflammatory responses to HDM were exacerbated by expression of FOXA3 (Supplemental Figure 5).

While goblet cell differentiation and mucus production are the noisome accompaniments of acute pulmonary infection or injury, enhancement of Th2 inflammation mediated by expression of FOXA3 and SPDEF may influence innate immune responses in chronic conditions associated with mucous metaplasia and may therefore play a role in the sensitization of patients with chronic pulmonary diseases that may be linked to susceptibility to infection and inflammation, as seen in asthma, COPD, and CF. Likewise, persistent mucus hyperproduction, goblet cell metaplasia, and inflammation following acute infection or injury may influence subsequent inflammatory and immune responses to both commensal and pathogenic microbes and aeroallergens.

In summary, we demonstrate that SPDEF and FOXA3 induced goblet cell differentiation and Th2-mediated inflammatory responses after birth, an active time of recruitment and activation of innate and acquired immunity. Exposure to both commensal and pathogenic microbes and antigens influences the acquisition of immune responses after birth that are likely to have long-term

effects on the patterning of subsequent immune and inflammatory responses of the lung. Both SPDEF and FOXA3 are induced by aeroallergens and viral exposures that cause goblet cell metaplasia. Induction of goblet cell metaplasia via SPDEF and FOXA3 in airway epithelial cells may play a role in the establishment of Th2 immune polarization, influencing subsequent response patterns to environmental exposures. SPDEF and, by inference, goblet cells are required for normal pulmonary responses to HDM extract.

Methods

Mouse models. *Scgb1a1-rtTA* (line 2) mice were either bred with *OTet7-TRE2-Spdef* mice or *OTet7-Foxa3-IRES-EGFP* mice, and dams were placed on doxycycline from E16.5 to P15 (22, 24). *Spdef*^{-/-} mice were obtained from the laboratory of Hans Clevers (Netherlands Institute of Developmental Biology, Utrecht, The Netherlands) and backcrossed onto BALB/c mice until an F9 generation was obtained (46). Similarly, the *Spdef*^{+/+} WT mice were also backcrossed onto BALB/c mice to obtain an F9 generation. In some experiments, *Scgb1a1-rtTA Foxa3-IRES* mice were bred with *Spdef*^{-/-} mice to obtain *Scgb1a1-rtTA Foxa3-IRES Spdef*^{-/-} mice, which were subsequently bred again with *Spdef*^{-/-} mice to obtain *Scgb1a1-rtTA Foxa3-IRES Spdef*^{-/-} and *Scgb1a1-rtTA Foxa3-IRES Spdef*^{-/-} mice. During this breeding, the dams were placed on doxycycline from E16.5 to P15. The transgenes were identified by PCR using the primer sets described in Supplemental Table 1.

Bronchoalveolar lavage fluid collection and cell counting. Mice were euthanized, the lungs were lavaged twice with 0.2 to 0.5 ml ice-cold PBS with 0.1 mM EDTA to collect bronchoalveolar lavage fluid (BALF) cells, and cytopins of BALF cells (30,000–50,000 cells) were prepared and stained with a Diff-Quik Stain Kit (Polysciences Inc.). Differential counts were determined in 300 to 400 cells.

AHR. AHR was measured in unrestrained mice using whole-body plethysmograph (Buxco Respiratory Products, Data Sciences International). Mice were exposed to increasing concentrations of methacholine (0.1–25 mg/ml). AHR is represented as Penh in response to methacholine. Penh values measured during this period were averaged and expressed as absolute Penh values.

Histology, immunohistochemistry, and immunofluorescence confocal microscopy. Paraffin-embedded lung tissues from P15 mice were sectioned and subjected to antigen retrieval using citrate buffer. IHC analyses were performed using goat anti-FOXA3 (1:100, SC-5361; Santa Cruz Biotechnology Inc.); guinea pig anti-SPDEF (1:3,000; generated in the Whitsett Laboratory); mouse mAb against MUC5AC (1:500, ab3649; Abcam); and rabbit anti-MUC5B (1:100, H-300; Santa Cruz Biotechnology Inc.). Immunofluorescence confocal microscopic analyses of MUC5B, acetylated tubulin (1:100, T7451; Sigma-Aldrich), and CCSP (guinea pig anti-CCSP-G210, generated in the Whitsett Laboratory at 1:15,000) were performed on 7-μm-thick sections of paraffin-embedded tissue and imaged with a Nikon A1Rsi Confocal Microscope.

Quantitative RT-PCR. Total RNA was prepared from whole-lung tissues from P15 mice collected in TRIzol (Invitrogen) according to the manufacturer's protocol, and cDNA was generated by reverse transcription using 500 ng to 1 μg total RNA (iScript cDNA Synthesis Kit; Bio-Rad Laboratories). The cDNAs were amplified in a StepOne-Plus Real-time PCR System (Applied Biosystems, Life Technologies) with the TaqMan probes (Applied Biosystems, Life Technologies) described in Supplemental Table 2. The Ct values of the target genes

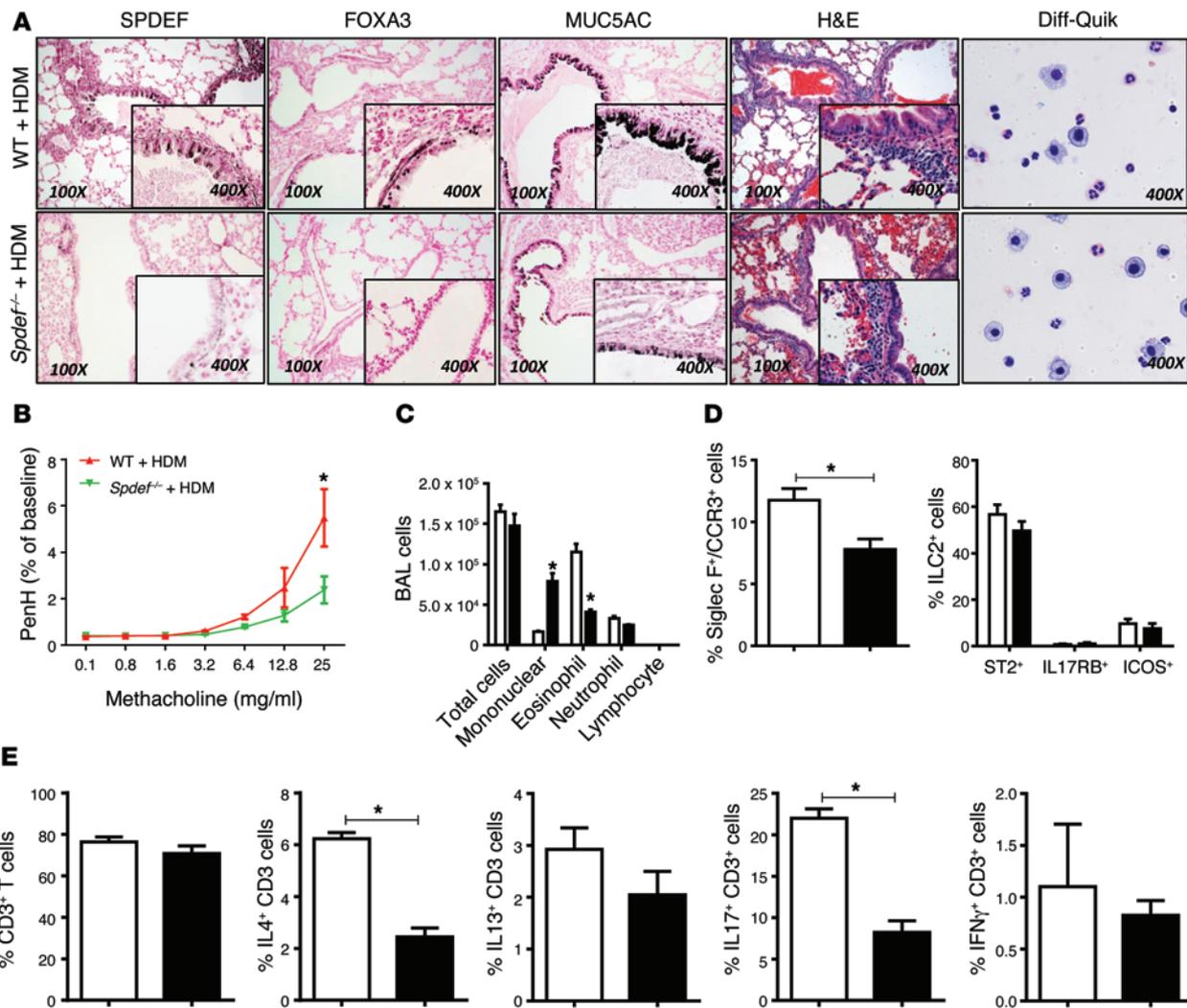


Figure 4. SPDEF is required for AHR, Th2, and TH17 inflammatory responses to HDM in postnatal mice. HDM (50 μ g) was administered i.n. daily for 3 days to PND15 control and *Spdef*^{-/-} mice, and the mice were sacrificed 2 days later. (A) Decreased mucous metaplasia and inflammatory responses to HDM were observed in *Spdef*^{-/-} mice. (A and C) BALF cells ($n = 4$) were stained with Diff-Quik, demonstrating eosinophilic and neutrophilic infiltrates. Goblet cell differentiation (A) and increased *Acta2* mRNA caused by HDM challenge (Table 4) were modestly decreased in *Spdef*^{-/-} mice. (B) *Spdef*^{-/-} mice were less responsive to HDM (green) than were controls (red line). Data represent the mean \pm SEM, using 2-way Anova with Bonferroni's post tests. $n = 6$ /group. (D and E) Flow analysis of lung cells from *Spdef*^{+/+} (white bars) and *Spdef*^{-/-} mice (black bars) after HDM challenge. Siglec-F⁺CCR3⁺ eosinophils were decreased, and ST2⁺, IL-17RB⁺, ICOS⁺ ILCs and CD3⁺ cells were unaltered. CD3⁺ IL-4⁺ and CD3⁺ IL-17⁺ T cells were decreased in *Spdef*^{-/-} mice, and CD3⁺ IFN- γ ⁺ T cells were unaltered. Gating strategies are depicted in Supplemental Figure 1. Data represent the mean \pm SEM. * $P < 0.05$ compared with controls using an unpaired, 2-tailed Student's t test. $n = 4$ /group.

were normalized to that of the endogenous housekeeping gene 18S rRNA to obtain the Δ Ct. The average Δ Ct of the target genes in transgenic lung tissues was subtracted from the average Δ Ct of the target genes in control lung tissues to obtain the $\Delta\Delta$ Ct. The change in gene expression relative to that in control lungs was calculated as $2^{-\Delta\Delta Ct}$ and represented graphically (47).

Flow cytometric analysis of lung cells. Whole-lung cell suspensions were prepared as previously described (48) and used for flow cytometric analysis. Briefly, lung cell suspensions were prepared by enzymatically digesting the lung tissue using 5,000 caseinolytic units of dispase (Discovery Labware Inc., Corning). Single-cell suspensions were then stained with the fluorochrome-labeled Abs described in Supplemental Table 3. For detection of eosinophils in lung cell suspensions, SiglecF and CCR3 were used. For analysis of ILCs, a lineage cocktail

with IL7R α , FLT3, ST2, IL-17RB, and ICOS was used. For analysis of myeloid DCs (mDCs) and plasmacytoid DCs (pDCs), CD45, CD11c, CD11b, B220, and CD103 were used along with CD86 and I-A/E as activation markers. CD45 and CD3 were used for analysis of T cells. For intracellular cytokine detection, whole-lung cell suspensions were cultured in the presence of brefeldin A (BioLegend), followed by surface staining, then permeabilization with Cytofix/Cytoperm solution (BD Biosciences), and staining for IL-4 and IFN- γ . The stained lung cell samples were acquired on a BD FACSCanto III flow cytometer using FACS Diva software.

Anti-TSLP Ab treatment. The pups obtained after breeding *Scgb1a1-rtTA* (line 2) and *OTet7-Foxa3-IRES-EGFP* mice were injected i.p. on P3 with either anti-TSLP Ab (50 μ g) (clone eBio28F12; eBioscience) or control Ab (50 μ g) (clone eBR2a; eBioscience). On P8, the

pups were euthanized and lung specimens collected for total RNA and pathology. Dams were placed on doxycycline from E16.5 to P8.

Pulmonary allergen challenge. Mice underwent 3 consecutive i.n. challenges with HDM allergen (*Dermatophagoides pteronyssinus*) or HDM extract (GREER) (50 µg for neonatal mice; 100 µg for adult mice), or with saline alone. After 48 hours of allergen exposure, mice were euthanized and lung tissues collected. The timing of HDM challenges and anti-TSLP treatment is provided in the figure legends.

Total IgE, HDM-specific IgE, and IgG1 ELISA. Total IgE and HDM-specific IgE and IgG1 were measured in serum by ELISA. Serum was diluted 1:100 for IgE and 1:10,000 for IgG1. Total IgE levels were measured using the appropriate kit (BD Biosciences). For measuring HDM-specific IgG1 and IgE, plates were coated with 5 µg/well HDM in carbonate coating buffer (pH 9.6) overnight at 4°C. The following day, plates were blocked with 10% FCS in PBS for 1 hour and incubated with samples for 2 hours at room temperature. HRP anti-mouse RaM IgG1 (1:1,000) (BD Biosciences) or biotinylated anti-mouse IgE (2 µg/ml, clone R35-118; BD Biosciences) was used to detect IgG1 and IgE Abs, respectively, and the reaction was developed with a tetramethylbenzidine substrate reagent set (1:1; BD Biosciences).

In some experiments, the adult *Spdef*^{+/+} and *Spdef*^{-/-} mice were injected i.p. with 200 µl donkey anti-mouse IgD serum, a gift of F. Finkelman (Division of Immunobiology, Cincinnati Children's Hospital Medical Center, Cincinnati, Ohio, USA). On day 7, serum was collected and ELISA for IgE and IgG1 was performed.

Statistics. Values are expressed as the mean ± SEM. All statistical analysis was performed using GraphPad Prism 6 (GraphPad Software). Data were analyzed using a 2-tailed, unpaired Student's *t* test or 2-way ANOVA, and Bonferroni's correction was used for multiple comparisons. *P* values of 0.05 or less were considered statistically significant.

Study approval. Animals were housed in pathogen-free conditions according to protocols approved by the IACUC of the Cincinnati Children's Hospital Research Foundation.

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Address correspondence to: Jeffrey A. Whitsett, Cincinnati Children's Hospital Medical Center, Division of Pulmonary Biology, 3333 Burnet Avenue, Cincinnati, Ohio 45229-3039, USA. Phone: 513.803.2790; E-mail: jeff.whitsett@cchmc.org.

Gang Chen's present address is: CF Center of University of North Carolina at Chapel Hill, Chapel Hill, North Carolina, USA.

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