Inflammation in an Asthma Model

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The Epithelial Anion Transporter Pendrin Is Induced by Allergy and Rhinovirus Infection, Regulates Airway Surface Liquid, and Increases Airway Reactivity and Inflammation in an Asthma Model

Yasuhiro Nakagami,* Silvio Favoreto, Jr.,§ Guohua Zhen,* Sung-Woo Park,* Louis T. Nguyenv,* Douglas A. Kuperman,§ Gregory M. Dolganov,v Xiaozhu Huang,* Homer A. Boushey,† Pedro C. Avila,§ and David J. Erle2*†‡

Asthma exacerbations can be triggered by viral infections or allergens. The Th2 cytokines IL-13 and IL-4 are produced during allergic responses and cause increases in airway epithelial cell mucus and electrolyte and water secretion into the airway surface liquid (ASL). Since ASL dehydration can cause airway inflammation and obstruction, ion transporters could play a role in pathogenesis of asthma exacerbations. We previously reported that expression of the epithelial cell anion transporter pendrin is markedly increased in response to IL-13. Herein we show that pendrin plays a role in allergic airway disease and in regulation of ASL thickness. Pendrin-deficient mice had less allergen-induced airway hyperreactivity and inflammation than did control mice, although other aspects of the Th2 response were preserved. In cultures of IL-13-stimulated mouse tracheal epithelial cells, pendrin deficiency caused an increase in ASL thickness, suggesting that reductions in allergen-induced hyperreactivity and inflammation in pendrin-deficient mice result from improved ASL hydration. To determine whether pendrin might also play a role in virus-induced exacerbations of asthma, we measured pendrin mRNA expression in human subjects with naturally occurring common colds caused by rhinovirus and found a 4.9-fold increase in mean expression during colds. Studies of cultured human bronchial epithelial cells indicated that this increase could be explained by the combined effects of rhinovirus and IFN-γ, a Th1 cytokine induced during virus infection. We conclude that pendrin regulates ASL thickness and may be an important contributor to asthma exacerbations induced by viral infections or allergens.

Asthma is a common chronic disease of children and adults that is characterized by airway inflammation, hyperreactivity, mucus overproduction, and airway obstruction (1). Asthma exacerbations are commonly triggered by infections with respiratory viruses, especially rhinovirus, and can also be triggered by other environmental factors, including allergens (2–4). The airway epithelial cell is a prominent participant in asthma exacerbations. Epithelial cell mucus production is increased in asthma (5), and plugging of the airways with mucus is a prominent and sometimes fatal feature of severe asthma exacerbations (6). Rhinovirus infects airway epithelial cells directly, inducing changes in gene expression (7), and also induces leukocyte production of IFN-γ (8), which has effects on epithelial and other cells in the airway (9). Airway epithelial cells also play central roles in the response to allergens. In this setting, Th2 cells and other cells produce IL-13 and IL-4, cytokines that act directly on epithelial cells to induce mucus production and airway hyperreactivity (10, 11). A recent clinical trial found that inhalation of an inhibitor of IL-13 and IL-4 reduced the late asthmatic response to allergen challenge (12), suggesting that these mechanisms discovered in mouse models are relevant to humans with asthma.

Changes in the airway surface liquid (ASL)3 that covers the epithelium can cause airway disease. ASL is comprised of a mucus layer and a periciliary layer (13). Ciliary beating and coughing propel ASL toward the upper airways, thereby removing mucus and inhaled pathogens and particles (14). ASL thickness is regulated by active ion transport across epithelial cells (15). In cystic fibrosis, mutations in the cystic fibrosis transmembrane conductance regulator gene (CFTR) result in decreased chloride secretion, ASL dehydration, impaired mucus clearance, and airway inflammation and obstruction (16). Similar pathology was seen in mice with ASL dehydration resulting from transgenic overexpression of the epithelial sodium channel (ENaC) (17). Changes in ion transport have also been implicated in asthma pathogenesis. IL-13 or IL-4 stimulation alters chloride conductance and converts human

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3 Abbreviations used in this paper: ASL, airway surface liquid; BAL, bronchoalveolar lavage; CFTR, cystic fibrosis transmembrane conductance regulator; CLCA1, chloride channel calcium activated 1; Ct, threshold cycle; ENaC, epithelial sodium channel; MTEC, mouse tracheal epithelial cells; NHBE, normal human bronchial epithelial; RV16, rhinovirus serotype 16; Smc1a, sodium channel, nonvoltage-gated, type 1, α; Slc26a4, solute carrier family 26 member 4.

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bronchial epithelial cells from an absorptive to a secretory phenotype, which may help to “flush” particulates and mucus from the airway lumen (18). In a recent genome-wide analysis, CLCA1 (calcium-activated chloride channel 1) was the most highly up-regulated gene in bronchial epithelial cells of asthmatics compared with healthy controls (19). Despite its name, CLCA1 is not a channel but CLCA1 does affect chloride conductance by an unknown mechanism (20). Increased expression of CLCA1 may be important in asthma pathogenesis since overexpression of its mouse ortholog (known as Gob-5 or C1ca3) was reported to cause airway inflammation and airway hyperreactivity (21), although the precise contributions of CLCA1 and other CLCA family members are still controversial (22–24). Other molecules involved in the transport of ions, including the Na\(^+-K\^-Cl\^-\) cotransporter NKCC1 (25), may also play a role, but the contributions of these molecules to asthma pathogenesis is unknown.

We previously reported that mRNA encoding the anion transporter pendrin (solute carrier family 26 member 4, Slc26a4) is increased in acute allergic airway disease (26), suggesting that pendrin could contribute to allergic airway disease and might play a role in asthma. Pendrin is a 780–aa anion transporter that is expressed in the inner ear, the kidney, and the thyroid (27). Pendrin can exchange chloride for other anions, including bicarbonate, oxalate, and hydroxyl ions (28). In Pendred syndrome, a common form of hereditary hearing loss caused by mutations in pendrin-deficient (Slc26a4\(^-/-\)) mice on a 129SvEv genetic background (29), impaired airway function by affecting ASL hydration. ASL thickness is important for mucociliary function, airway hyperreactivity, and inflammation (33). However, the contributions of pendrin to allergic airway disease have not yet been directly addressed.

We hypothesized that allergen-induced increases in pendrin contribute to altered airway function by affecting ASL hydration. To address this issue, we took advantage of the availability of pendrin-deficient (Slc26a4\(^-/-\)) mice (30). Although these mice are deaf, thyroid function is unimpaired and kidney function is normal except during sodium restriction (30, 34). We used Slc26a4\(^-/-\) mice to study the role of pendrin in the in vivo response to allergen and Slc26a4\(^-/-\) mouse tracheal epithelial cells (MTEC) to investigate how pendrin affects ASL thickness. To determine whether pendrin is also induced by viral infection, the most common precipitant of asthma exacerbations, we also analyzed the expression of SLC26A4 in nasal epithelium from human subjects with naturally occurring rhinovirus infections and in cultured NHBE cells infected with rhinovirus. The results of these studies indicate that pendrin is induced by stimuli that provoke asthma exacerbations and contributes to airway inflammation and hyperreactivity.

Materials and Methods

**Mice**

Pendrin-deficient (Slc26a4\(^-/-\)) mice on a 129SvEv genetic background were generously provided by E. Green (30). Mice were bred and maintained under pathogen-free conditions. Slc26a4\(^+/-\) and Slc26a4\(^+/-\) littermates were used as controls for the in vivo allergen challenge model. All experiments were performed according to the guidelines of the University of California, San Francisco (UCSF) Committee on Animal Research.

**Sensitization and challenge with OVA**

Age- and sex-matched 6–8-wk-old mice were sensitized by i.p. injection of OVA (Sigma-Aldrich) on days 0, 7, and 14 as reported previously (26). The sensitizing emulsion consisted of 50 μg OVA and 10 mg of aluminum potassium sulfate in 200 μl of saline. On days 21, 22, and 23, the sensitized mice were lightly anesthetized by isoflurane inhalation and challenged with 100 μg OVA in 30 μl of saline administered intranasally. Control mice were treated in the same way, except that OVA was omitted during both the sensitization and challenge phases.

**Assessment of airway reactivity, inflammation, OVA-specific IgE, and mucus**

Airway reactivity was measured as previously reported (35). Briefly, on day 24, mice were anesthetized, intratracheally intubated, mechanically ventilated, and paralyzed, and airway resistance was measured at baseline and following i.v. administration of increasing doses of acetylcholine (0.03, 0.1, 0.3, 1, and 3 μg/g body weight). Bronchoalveolar lavage (BAL) fluid leukocytes and serum OVA-specific IgE levels were measured as previously described (26). Paraffin-embedded 5-μm lung sections were stained with H&E and with periodic acid-Schiff for evaluation of mucus production. The severity of peribronchial inflammation was scored by a blinded observer using the following features: 0, normal; 1, few cells; 2, a ring of inflammatory cells 1 cell layer deep; 3, a ring of inflammatory cells 2–4 cells deep; 4, a ring of inflammatory cells of >4 cells deep (36). Epithelial cell mucous content was determined using stereology as previously reported (11).

**Analysis of mouse gene expression by quantitative RT-PCR**

Total RNA from mouse lungs was isolated using TRIzol (Invitrogen) and purified using the RNaseasy kit (Qiagen). First-strand cDNA synthesis was performed using the SuperScript First-Strand cDNA Synthesis System (Invitrogen). Total RNA from MTEC cells was isolated using the RNeasy kit (Qiagen), and first-strand cDNA synthesis was performed using the QuantiTect Reverse Transcription Kit (Qiagen). Mouse Slc26a4 primers were 5’-CATCTCGAAGACAGCTCAACTC-3’ and 5’-CATCTTCGCTCCTCATCAT-3’. Mouse sodium channel, nonvoltage-gated, type I (Scnn1a) primers were 5’-GGAGGGTTCTGTAACCTCAACATC-3’ and 5’-TGGAGACAGCAGCATCCGGCT-3’. Mouse Scnn1b primers were 5’-CTCGAGTCATCGGCGAACGC-3’ and 5’-CCAGTGGCCGAGGAGAG-3’. Mouse CLCA1 primers were 5’-GCATTCATCTCTGCCTCCAT-3’ and 5’-GGACATTCCTTGTGATG-3’. Mouse Clca3 primers were 5’-GCCC TCTAGGCAATGTGAGG-3’ and 5’-CTCTGGTGTGGGCCATCT-3’. Mouse β-actin (Actb) primers were 5’-TGTGATTACACTGGAGCA-3’ and 5’-GGGGTGTGGAGGACTTTAAAA-3’. Mouse Gapd primers were 5’-GGGTTCTGCTTCTCACCCA-3’ and 5’-TGCCGTCTGGTACCT-3’. Sequences of other primers and probes used herein have been previously reported (31). SYBR Green real-time PCR was performed using the 7900HT Fast Real-Time PCR System (Applied Biosystems). The threshold cycle (Ct) of each transcript was normalized to the average Ct for Actb and Gapd. Fold differences were determined by the 2^(-ΔΔCt) method (37).

**Culture of MTEC**

MTEC were isolated from age-matched 6–14-wk-old pendrin-deficient mice and wild-type littermate controls and cultured as previously described (38). Cells were considered to be confluent when transepithelial resistance was >1000 Ω cm². After reaching confluence, MTEC were cultured at air-liquid interface for 14–17 days in the absence or presence of 20 ng/ml murine IL-13 (PeproTech) as indicated in the text.

**Measurement of ASL thickness**

ASL thickness was measured using previously reported methods (39). Briefly, bextradin-conjugated BCECF (2’,7’-bis(2-carboxethyl)-5(6)-carboxyfluorescein) (Invitrogen) was suspended in FC-72 (3M) at a concentration of 10 mg/ml, and 20–50 μl of the suspension was deposited onto the ASL 3 mm before measurement. ASL thickness was measured with a Nikon EZ-C1 laser scanning confocal system and a Nikon Eclipse FN1 upright microscope. Fluorescent images were obtained using a ×10 objective (Plan Fluor, numerical aperture 0.30, working distance 15.8 mm). ASL
thickness was determined from the fluorescence peak width (width at half-maximal fluorescence) along the z-axis. Standards made using known volumes of BCECF-containing solution sandwiched between coverglasses were used for calibration. ASL was exposed to a fully humidified 5% CO₂ atmosphere, and air and a stage were maintained at 37°C using a Tempontol heating system (PeCon).

Collection of human nasal samples during common colds

The clinical study was approved by the UCSF Committee of Human Research, and all subjects signed a written informed consent. Asthmatic and nonasthmatic control subjects were recruited within 3 days of onset of common colds through advertisements on the internet and campus flyers and by contacting subjects from previous studies. Subjects were characterized by spirometry, allergy skin testing, and methacholine bronchial reactivity as we previously described (40). Subjects had three visits at 1–3 d after cold symptom onset. In visit 1, nasal lavage was used to identify rhinovirus and other viruses as previously described (41). Subjects recorded nasal and chest symptoms during the week after visit 1 and the week before visit 3. Nasal cold symptoms were scored by adding together daily scores for each of eight symptoms (nasal discharge, nasal congestion, sneezing, throat discomfort, tiredness, cough, fever/chills, and headache); each scored from 0 to 3 during 1 wk (yielding a possible weekly range of 0–168). The final nasal cold symptom score was calculated as the difference between score for the week after visit 1 (cold) and the score for the week before visit 3 (baseline). Chest symptoms were the change of scores for five symptoms (shortness of breath, chest tightness, wheezing, cough, and sputum production; each scored 0–10; possible weekly range, 0–350). At each visit, three nasal lavages with 10 ml of normal saline solution each were performed before collecting nasal mucosal samples using Rhino-Probe curettes (Arlington Scientific). Mucosal samples were collected from the inferomedial aspect of the inferior turbinate on one side at visit 1, from the other side at visit 2, and from either side at visit 3. Nasal mucosal samples were immediately placed in RNasy kit solution with β-mercaptoethanol, frozen at bedside in dry ice, and stored at −80°C. Nasal mucosal samples had >95% epithelial cells.

Evaluation of SLC26A4 mRNA in human nasal mucosal samples

After RNA isolation and cDNA synthesis, SLC26A4 mRNA was quantified using a validated two-step real-time PCR method (25). Primer and probe sequences have been previously reported (26).

In vitro infection of NHBE cells with rhinovirus

NHBE cells from normal individuals (Lanza, passages 1–4) were seeded onto 6.5-mm Transwell inserts (BD Biosciences) at 5 × 10⁴ cells/insert and grown to 100% confluence and then cultured at air-liquid interface for 2 wk to promote epithelial differentiation. Rhinovirus serotype 16 (RV16) was grown in HeLa cells, purified by sucrose gradient, and quantified by a PFU assay using standard virological methods. NHBE cells were infected on the apical side with RV16 (multiplicity of infection of 5) for 6 h at 35°C with mild agitation at 2 rpm, after which virus was washed off with PBS and the cells were cultured for 24 h. Some cells were pretreated with 10 ng/ml human IFN-γ (R&D Systems) for 24 h before RV16 infection. Control cells received a sham inoculum. SLC26A4 mRNA expression was measured using the same primer and probe sequences used for analysis of the nasal mucosal samples, and changes in gene expression were calculated using the 2⁻ΔΔCt method.

Statistical analyses

With the exception of the pendrin measurements from nasal epithelial cells from human subjects, data are presented as means ± SEM and significance testing was done using Student’s t test, Tukey’s test, or Tukey-Kramer’s
test after ANOVA, or Kruskal-Wallis’s test. 
p values < 0.05 were considered significant. For comparing characteristics of asthmatic and nonasthmatic subjects, we used the Mann-Whitney’s rank test for continuous variables or the Fisher exact test for categorical variables. Levels of SLC26A4 mRNA among the three visits were analyzed using generalized estimating equations (xtgee command of Stata). Pairwise comparisons between two visits were then analyzed using Wilcoxon’s signed-rank test. KaleidaGraph (ver. 3.6, Synergy Software) and Stata (Ver. 8, StataCorp) software packages were used for statistical analyses.

Results

Allergen-induced airway hyperreactivity and inflammation are attenuated in pendrin-deficient mice

Pendrin-deficient mice and control mice were sensitized and then challenged by intranasal administration of OVA to produce an allergic response in the lung and airway. Respiratory system resistance was measured in sedated and mechanically ventilated mice after i.v. administration of acetylcholine (Fig. 1a). OVA challenge increased airway reactivity in control mice, as expected. Airway hyperreactivity was also observed in OVA-challenged pendrin-deficient mice, but the degree of hyperreactivity was significantly less than in OVA-challenged control mice. There were no differences in airway reactivity to acetylcholine between saline-challenged control and saline-challenged pendrin-deficient mice. BAL fluid was collected to assess the effects of OVA sensitization and challenge on inflammatory cell recruitment (Fig. 1b). Cells retrieved from saline-challenged mice were mostly macrophages. There were large increases in macrophages, eosinophils, lymphocytes, and neutrophils in OVA-challenged control mice. Increases were also observed in OVA-challenged pendrin-deficient mice, but the numbers of total cells, eosinophils, and neutrophils were significantly less than in OVA-challenged control mice. Histological analysis revealed large numbers of inflammatory cells around the conducting airways in OVA-challenged control mice, but inflammation was significantly reduced in OVA-challenged pendrin-deficient mice (Fig. 1, c and d).

Allergen-induced IgE production, mucus metaplasia, and Th2 cytokine mRNA production are unaffected by pendrin deficiency

We measured OVA-specific IgE in serum to determine whether reduced airway hyperreactivity in pendrin-deficient mice might be explained by a generalized impairment in the allergic response. OVA challenge led to production of similar amounts of OVA-specific IgE in control and pendrin-deficient mice (Fig. 2a), indicating that this aspect of the allergic response was preserved. We also found that there was no difference in allergen-induced mucus metaplasia between control and pendrin-deficient mice (Fig. 2b). Since mucus metaplasia depends on Th2 cytokines, especially IL-13, this result indicates that Th2 responses were intact in pendrin-deficient mice. Analysis of expression of mRNAs encoding the Th2 cytokines IL-4, IL-5, and IL-13 and the Th1 cytokine IFN-γ in lungs from OVA-challenged mice revealed no significant differences between control and pendrin-deficient mice (Fig. 2c).

Allergen challenge induces changes in pendrin and ion channel mRNA expression in vivo

Allergen challenge increased the pendrin transcript SLC26A4 (Fig. 3a), as we reported previously (26). We also investigated the expression of five other mRNAs encoding proteins involved in epithelial cell ion transport. Allergen challenge significantly reduced transcripts encoding all three ENaC subunits (SCNN1A, SCNN1B, and SCNN1G) and CFTR (Fig. 3b). Allergen challenge also led to a large increase in expression of Clca3 mRNA, as previously reported (21). There were no significant differences in expression of the three ENaC transcripts, CFTR, or Clca3 between OVA-challenged control and OVA-challenged pendrin-deficient mice.

IL-13 stimulation of MTEC induces changes in pendrin and ion channel mRNA expression in vitro

To identify effects of pendrin on ASL, we studied MTEC cultures from control and pendrin-deficient mice. Cells were cultured at
air-liquid interface, a method that allows cells to differentiate, polarize, and form an ASL layer. We first investigated how IL-13 stimulation of MTEC affected expression of transcripts encoding proteins involved in ion transport. IL-13 markedly up-regulated SLC26A4 mRNA in MTEC cultures from control and pendrin-deficient MTEC (Fig. 4a). We previously reported similar effects of IL-13 on the human orthologs of these mRNAs (SLC26A4 and CLCA1) in NHBE cells (31). Although Scnn1a expression was not affected by IL-13, Scnn1b and Scnn1g were both substantially decreased. Decreases in these 2 ENaC subunit mRNAs were substantially more marked than the decreases measured after in vivo allergen challenge (Fig. 4b). It is possible that the dose or duration of IL-13 stimulation used in the MTEC experiments accounts for this difference. Alternatively, since the whole lung samples analyzed in the allergen challenge model contain large numbers of ENaC-expressing alveolar epithelial cells (42), it seems likely the effects of IL-13 on Scnn1b and Scnn1g expression in airway epithelial cells were underestimated by the analysis of whole lung RNA. We did not identify any significant differences in expression of ENaC mRNAs, Cfr, or Clca3 between IL-13-stimulated control MTEC and IL-13-stimulated pendrin-deficient MTEC (Fig. 4b). Mucin 5 subtype a and c (Muc5ac) expression was similar in control and pendrin-deficient MTECs (data not shown).

Pendrin deficiency results in increased ASL thickness

We measured ASL thickness of control and pendrin-deficient MTEC cultures stimulated with or without IL-13 using a confocal microscope. Typical z-axis fluorescence traces obtained from cultures of unstimulated and IL-13-stimulated control MTEC cells are shown in Fig. 5a. In the absence of IL-13, there was no significant difference in ASL thickness between control and pendrin-deficient MTEC cultures (Fig. 5b). IL-13 stimulation increased ASL thickness in both control and pendrin-deficient MTEC cultures. ASL in IL-13-stimulated pendrin-deficient MTEC cultures was significantly thicker than that in IL-13-stimulated control MTEC cultures, indicating that the effect of IL-13 on ASL was exaggerated in pendrin-deficient MTEC cultures.

SLC26A4 expression in nasal epithelial cells is increased during common colds

Nasal mucosa samples were collected from asthmatic and nonasthmatic human subjects at three time points after onset of cold symptoms. Samples from 22 subjects with confirmed rhinovirus infection were analyzed for this study. The characteristics of the subjects are shown in Table I. There were no obvious differences between nonasthmatic and asthmatic subjects or between nonatopic and atopic subjects, but there were relatively small numbers of subjects in these subgroups. When data from all 22 subjects were analyzed together (Fig. 6a), we found that SLC26A4 mRNA expression was significantly increased during acute colds (at visit 1, 1–3 d after onset of symptoms and at visit 2, 5–7 d after onset of symptoms) compared with a time long after resolution of symptoms (visit 3, ~42 d after onset of symptoms). Mean SLC26A4 expression was 4.9-fold higher at visit 1 and 2.2-fold higher at visit 2 compared with visit 3.

The combination of RV16 and IFN-γ increases SLC26A4 mRNA in NHBE cells

To determine whether rhinovirus directly increases SLC26A4 mRNA expression in human airway epithelial cells, we infected NHBE cells with RV16 in vitro. Under the conditions we used, RV16 infection alone had no effect on SLC26A4 mRNA expression (Fig. 6b). We considered the possibility that cytokines produced by other cells during rhinovirus infection contribute to SLC26A4 mRNA induction in vivo. Rhinovirus stimulates production of IFN-γ by mononuclear cells (8, 43–45), and we found that

Table I. Subject characteristics

<table>
<thead>
<tr>
<th>Characteristics</th>
<th>Nonasthmatics (n = 7)</th>
<th>Asthmatics (n = 15)</th>
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<tr>
<td>Age (years)</td>
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<td>33.7 ± 2.5</td>
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<td>Female-male</td>
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<td>13.2</td>
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<td>(liters) (%)</td>
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<td>94.2 ± 2.8</td>
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<td>Chest cold symptom score</td>
<td>44.3 ± 16.3</td>
<td>66.9 ± 14.6</td>
<td>0.37</td>
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</table>

a FEV1 indicates forced expiratory volume in 1 s.

b PC20 indicates provocative dose of methacholine causing a 20% fall in FEV1.
had a small but nonsignificant effect on model, we previously showed that and of human subjects with rhinovirus infection implicate pendrin The results of our studies of a mouse allergic airway disease model Discussion many genes known to be induced by IFN-γ were markedly upregulated in nasal epithelial cells during common colds (S. Favoreto, H. Boushey, and P. Avila, unpublished data). IFN-γ alone had a small but nonsignificant effect on SLC26A4 mRNA expression. However, the combination of RV16 infection and IFN-γ stimulation caused a larger and statistically significant effect on SLC26A4 expression in NHBE cells.

Discussion

The results of our studies of a mouse allergic airway disease model and of human subjects with rhinovirus infection implicate pendrin in airway dysfunction during asthma exacerbations. In the allergic model, we previously showed that Slc26a4 mRNA expression was markedly increased in response to Th2 cytokines (26). Herein we show that pendrin makes a significant contribution to allergen-induced airway hyperreactivity and airway inflammation in this model. Both airway hyperreactivity and inflammation have been shown to be influenced by ASL hydration in other models (17), and we found that pendrin expression affects ASL thickness in MTEC cultures. Furthermore, we demonstrated that SLC26A4 mRNA was up-regulated in human subjects during naturally occurring colds caused by rhinovirus, suggesting that changes in pendrin expression may be important not only for allergen-induced exacerbations but also for viral infections, the most common triggers of asthma exacerbations.

Allergen-induced airway hyperreactivity and inflammation were reduced in pendrin-deficient mice. These effects cannot be explained by a generalized defect in the allergic response, since production of OVA-specific IgE, mucus, and Th2 cytokine mRNAs was unaffected by pendrin deficiency. Instead, it seems likely that pendrin deficiency affects hyperreactivity and inflammation due to its effects on ASL. A causal association between ASL dehydration and airway inflammation and obstruction has been established in cystic fibrosis (16) and in transgenic mice overexpressing ENaC (17). In allergic airway disease, IL-13 stimulation causes increased production of mucus (11) along with increased secretion of ions and water (18). We found that pendrin-deficient MTEC cultures were better hydrated (had thicker ASL) than control cells following IL-13 stimulation, and this may facilitate mucus clearance and improve airway function. Mechanistic relationships between ASL hydration, airway inflammation, and airway obstruction and hyperreactivity are complex and incompletely understood despite extensive analysis of various asthma and cystic fibrosis models. It is clear that airway epithelial cells can affect inflammation by producing cytokines that affect maturation and recruitment of eosinophils and other leukocytes (46). We examined expression of mRNAs encoding cytokines that are induced in the allergic model and are known to play a role in eosinophil recruitment, including IL-5 (Fig. 2c) and the chemokines CCL2 (MCP-1), CCL11 (eotaxin-1), and CCL24 (eotaxin-2) (data not shown), but found no differences in pendrin-deficient mice. We cannot formally exclude the possibility that other pendrin-expressing cells in the lung are involved in allergic airway disease, but we are not aware of reports of pendrin expression in cells other than epithelial cells and we detected minimal if any Slc26a4 mRNA in BAL fluid inflammatory cells (data not shown). Our findings are consistent with the hypothesis that pendrin’s effects on inflammation and airway reactivity are due to direct effects on ASL, although other explanations for these effects cannot be excluded.

Nakao and colleagues have very recently described the results of other, complementary approaches to studying the role of pendrin in airway disease (33). These investigators found that forced expression of pendrin using a Sendai virus vector led to increased expression of mucus. We found no effect of pendrin deficiency on levels of mucus in response to allergen challenge using a highly quantitative stereology-based approach, indicating that pendrin expression is not required for allergen-induced mucus production in this model. Similar results were reported in studies of another allergen-induced epithelial cell gene, Clca3 (Gob-5): overexpression led to increased mucus production (21, 24), but gene deletion had no effect on mucus production in response to allergen challenge (22, 24). This may be explained by the existence of redundant pathways leading to mucus production (24). Nakao and colleagues also found that forced expression of pendrin caused airway inflammation and hyperreactivity (33). Our findings of decreased inflammation and hyperreactivity in allergen-challenged pendrin-deficient mice provide additional evidence for the involvement of pendrin in these processes and show that pathological responses to allergen are diminished when pendrin is absent.

We found that pendrin deficiency affected ASL in a manner consistent with what is known about pendrin function outside the lung. Heterologous expression of pendrin in Xenopus oocytes (47), SF9 insect cells (47), HEK-293 kidney cells (48), and Fischer rat thyroid cells (32) has been shown to increase transport of chloride and other ions. Studies of pendrin-deficient mice established that pendrin contributes to absorption of chloride and water in the ear and the kidney. In the inner ear, pendrin is expressed at high levels and deficiency of pendrin results in impaired chloride and water uptake from endolymph (30). In the kidney, pendrin has no detectable effects on renal function or fluid or electrolyte balance under normal conditions (49). However, mineralocorticoid administration results in a substantial increase in pendrin expression by cortical collecting duct cells (50). Mineralocorticoids caused hypertension and weight gain in wild-type mice but not in pendrin-deficient mice, indicating that mineralocorticoid-induced pendrin expression leads to increased chloride and water absorption in the kidney (50). Similarly, we found that the effect of pendrin on ASL thickness in MTEC cultures was only apparent after stimulation with IL-13, which substantially increased pendrin expression. IL-13 increased ASL thickness in both control and pendrin-deficient MTEC cultures, likely due to multiple effects on expression of ion transporters including reduced expression of ENaC. The increase in ASL was significantly larger in pendrin-deficient MTEC cultures, which is consistent with the hypothesis that pendrin, which

![Figure 6](http://www.jimmunol.org/)

**FIGURE 6.** SLC26A4 is increased in nasal epithelial cells during common colds. a. Boxplots representing SLC26A4 expression during common colds (1–3 d and 5–7 d after onset) and at baseline (≈42 days after onset). Normalized gene copy number was determined by quantitative RT-PCR. Data are from 22 subjects, including 15 subjects with asthma. **p < 0.01 vs visit 3. b. Effects of RV16 and IFN-γ on SLC26A4 expression in NHBE cells. Cells were infected with RV16 and/or stimulated with 10 ng/ml IFN-γ for 24 h as indicated. Results are means and error bars represent SEM (n = 3 wells/condition). **p < 0.05 vs Control; †p < 0.05 vs RV16 alone.**
localizes to the apical membrane of airway epithelial cells (33), increases total chloride uptake and water absorption in these cells. However, other direct or indirect effects of pendrin on absorption or secretion of ions and water might also account for our findings. In any case, our findings demonstrate that pendrin expression has a significant effect on ASL thickness after IL-13 stimulation.

We found that naturally occurring colds caused by rhinovirus, a very common trigger of exacerbations, were associated with increased expression of SLC26A4. Our in vitro studies of NHBE cells indicate that the combination of direct effects of rhinovirus on NHBE cells and indirect effects mediated by IFN-γ, a cytokine produced by leukocytes during rhinovirus infections, could account for this increase. Additionally, the present study confirms that allergen, another trigger of asthma exacerbations, increases Slc26a4 mRNA expression in mice. We have also reported that lung expression of Slc26a4 is increased in mouse models of bacterial infection (51), where the recently reported (32) ability of pendrin to secrete thiocyanate, an anion with a role in antimicrobial defense, may be important. These results suggest that increases in pendrin expression could contribute to inflammation and airway dysfunction in viral infections, thought to be the predominant cause of asthma exacerbations, as well as in bacterial infections and allergen-induced exacerbations. We previously reported that bronchial epithelial expression of SLC26A4 was not increased in human subjects with stable, mild to moderate asthma compared with healthy, nonasthmatic controls (26). This suggests that pendrin plays a more important role in airway dysfunction during asthma exacerbations than during stable asthma.

Inhibition of pendrin could be an effective strategy for treatment of asthma exacerbations. General evidence in support of strategies designed to augment ASL comes from a recent demonstration that inhalation of hypertonic saline had beneficial effects in cystic fibrosis (52). Pendrin is one of several molecules that regulate ASL and represent possible therapeutic targets in asthma and other airway diseases. For treatment of cystic fibrosis and chronic bronchitis, considerable efforts have been made to develop therapies designed to increase ASL by inhibiting ENaC (53) or stimulating chloride channels or other effects of this drug (23). Pendrin pre-dominant cause of asthma exacerbations, as well as in bacterial infection of airway disease traits despite redundancy of calcium-activated chloride channel family members.

Disclosures

The authors have no financial conflicts of interest.

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