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RESEARCH ARTICLE

Effect of TGF-β1 on eosinophils to induce cysteinyl leukotriene E₄ production in aspirinexacerbated respiratory disease

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Abstract

Cysteinyl leukotriene (cysLT) overproduction and eosinophil activation are hallmarks of aspirin-exacerbated respiratory disease (AERD). However, pathogenic mechanisms of AERD remain to be clarified. Here, we aimed to find the significance of transforming growth factor beta 1 (TGF-β1) in association with cysteinyl leukotriene E₄ (LTE₄) production, leading to eosinophil degranulation. To evaluate levels of serum TGF-β1, first cohort enrolled AERD (n = 336), ATA (n = 442) patients and healthy control subjects (HCs, n = 253). In addition, second cohort recruited AERD (n = 34) and ATA (n = 25) patients to investigate a relation between levels of serum TGF-β1 and urinary LTE₄. The function of TGF-β1 in LTE₄ production was further demonstrated by ex vivo (human peripheral eosinophils) or in vivo (BALB/c mice) experiment. As a result, the levels of serum TGF-β1 were significantly higher in AERD patients than in ATA patients or HCs (P = .001; respectively). Moreover, levels of serum TGF- β 1 and urinary LTE₄ had a positive correlation (r= 0.273, P = .037). In the presence of TGF- β 1, leukotriene C₄ synthase (LTC₄S) expression was enhanced in peripheral eosinophils to produce LTE₄, which sequentially induced eosinophil degranulation via the p38 pathway. When mice were treated with TGF-\(\beta\)1, significantly induced eosinophilia with increased LTE₄ production in the lung tissues were noted. These findings suggest that higher levels of TGF-\(\beta\)1 in AERD patients may contribute to LTE₄ production via enhancing LTC₄S expression which induces eosinophil degranulation, accelerating airway inflammation.

Introduction

Aspirin-exacerbated respiratory disease (AERD) typically presents moderate-to-severe phenotypes of asthma, chronic rhinosinusitis (CRS) and/or nasal polyps with persistent eosinophilia in the upper and lower airway mucosa. In addition, cysteinyl leukotriene (cysLT) overproduction is a hallmark of AERD, and the increased level of cysLTs derived from mast cells and

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eosinophils is a characteristic feature of AERD, in which leukotriene C_4 synthase (LTC₄S) is a key enzyme for converting arachidonic acid to cysLTs [1]. To date, type 1 cysteinyl leukotriene receptor (cysLT₁R) antagonists and 5-lipoxygenase inhibitors have been used in the management of AERD [2]; however, there are unmet needs for its pathogenic mechanisms and therapeutic targets.

Eosinophilia in peripheral blood and upper/lower airway mucosa are commonly found in AERD patients [3]. Although both mast cells and eosinophils are critical for inducing airway inflammation in the pathogenesis of AERD, emerging evidence supports an important role of eosinophils in its pathogenesis [4,5]. AERD patients have shown that significantly elevated levels of eosinophil-derived granule proteins, such as eosinophil cationic protein (ECP) and eosinophil-derived neurotoxin (EDN), compared to aspirin-tolerant asthma (ATA) patients [6,7]. The direct effect of aspirin on eosinophil activation on releasing granule proteins has previously been demonstrated [8]. In addition, some studies have suggested that granule proteins play a crucial role in enhancing Th2 immune response among allergic diseases [9,10].

Transforming growth factor beta 1 (TGF- β 1) has also been suggested to contribute to immune responses and structural changes in the lungs of asthmatic patients [11]. Moreover, this mediator is strongly expressed in nasal mucosa in response to inflammation, but not in normal nasal mucosa [12]. So far, the role of TGF- β 1 in airway remodeling has mostly been highlighted; however, a few studies have shown that TGF- β 1 down-regulates cyclooxygenase (COX)-2 in airway epithelial cells and then reduces prostaglandin E2 production [13]. Furthermore, enhanced LTC4S expression in fibroblasts and monocytes in the presence of TGF- β 1 has been revealed [14,15], suggesting that TGF- β 1 may contribute to cysLT production in AERD patients. Considering that persistent eosinophilic inflammation is a key feature of AERD and that studies evaluating the function of TGF- β 1 in eosinophilic airway inflammation are still lacking, the present study focuses on the effect of TGF- β 1 on eosinophil activation in AERD pathogenesis.

We hypothesized that TGF- β 1 plays an important role in the severity of eosinophilic airway inflammation in AERD patients. This study compared the levels of serum TGF- β 1 between AERD and ATA patients, investigated the association between TGF- β 1 and LTE₄ in the clinical cohort, and evaluated the effect of TGF- β 1 on LTE₄ production *ex vivo* or *in vivo*.

Materials and methods

Ethics

Two cohorts of adult asthmatic patients were assessed in this study approved by the Institutional Review Board of Ajou University Hospital (AJIRB-GEN-SMP-13-108; AJIRB-BMR-SUR-15-498). All patients provided written informed consent to participate in this study by signing the consent form.

Patient cohorts, clinical parameters and serum cytokine levels

The clinical significance of TGF- β 1 level in AERD was evaluated in which AERD (n = 336), ATA (n = 442) patients and healthy control subjects (HCs; n = 253) were recruited. Among AERD and ATA patients enrolled in the first cohort, we enrolled AERD (n = 34) and ATA (n = 25) patients who had wanted to participate voluntarily in the second cohort study to investigate the role of TGF- β 1 in association with LTE₄ metabolite levels/eosinophil activation markers. AERD were diagnosed according to clinical features previously described [16]. The diagnosis of AERD was based on a positive response to lysine-aspirin bronchoprovocation test (L-ASA BPT). The presence of CRS and nasal polyps was confirmed using paranasal sinus

X-rays, CT scans and/or rhinoscopy as well as clinical symptoms. The degree of airway obstruction was evaluated using spirometry. The degree of airway hyperresponsiveness was examined by methacholine bronchial challenge test. Atopy status was defined as previously described [17]. The levels of serum IgE were quantified using UniCAP® system (Thermo-Fisher Scientific, Waltham, MA, USA). The levels of serum TGF- β 1 from every study subject were measured using ELISA (R&D systems, Minneapolis, MN, USA). Sputum collection and neutrophils/eosinophils counting were performed as previously described [18], which were always the same between the first and second cohorts. To determine TGF- β 1-low/high groups, the cutoff value (48.1 ng/mL) was set at mean plus 2 standard deviations of the test values. In the second cohort, the urine and serum of each patient were simultaneously collected in the morning time during the enrollment period. The urinary LTE4 metabolite levels were measured using ultra-high-performance liquid chromatography system as previously described [19]. In addition, the levels of serum EDN were measured using the ELISA kit (SKIMS-BIO, Seoul, Korea).

Stimulation of peripheral eosinophils from asthmatic patients

Peripheral eosinophils were isolated from asthmatic patients as previously described [20]. To stimulate eosinophils, the cells (1×10^6) were seeded on a 24-well plate and maintained in RPMI-1640 medium (Sigma-Aldrich, St. Louis, MO, USA) supplemented with 2% fetal bovine serum (FBS; ThermoFisher Scientific). Then, the cells were treated with human recombinant 10 ng/mL IL-5 (Sigma-Aldrich) and 5 ng/mL TGF- β 1 (R&D systems). To investigate the effect of cysLTs on eosinophil degranulation, the cells were treated with LTE₄ (Cayman Chemical, Ann Arbor, MI, USA) for 4 hours in the presence of 10 ng/mL IL-5 (Sigma-Aldrich). The function of montelukast (Sigma-Aldrich; 0.1 and 1 μ M) against LTE₄ was also investigated. To confirm eosinophil degranulation, eosinophils were seeded on Poly L-lysine-coated slides (Polysciences, Warrington, PA, USA). Then the cells were incubated overnight with antieosinophil peroxidase antibody (Cell Signaling, Minneapolis, MN, USA), followed by Alexa fluor 488 donkey anti-rabbit (ThermoFisher Scientific) for 1 hour. 4',6-diamidino-2-phenylindole (Sigma-Aldrich), and was observed using a Zeiss LSM710 confocal microscope (Carl Zeiss AG, Oberkochen, Germany).

Interactions between eosinophils and airway epithelial cells

A549 cells (American Type Culture Collection, Manassas, VA, USA) were used to investigate the role of airway epithelial cells in eosinophilic airway inflammation. The cells (5×10^5) were seeded on a 24-well plate in RPMI with 10% FBS. Then RPMI with 2% was used when A549 cells were treated with peripheral eosinophils (1×10^6) from asthmatic patients for 24 hours. In addition, A549 cells were treated with EDN (Athens Research & Technology, Athens, GA, USA; 1, 10 and 100 ng/mL) for 24 hours to demonstrate the effect of granule proteins on airway epithelial cell stimulation. To collect supernatant, culture medium was centrifuged at 12,000 rpm for 20 min at 4°C.

Polymerase chain reaction

Total RNA was isolated from human peripheral eosinophils using $TRIzol^{\circledR}$ (ThermoFisher Scientific), according to the manufacturer's instructions. Then, 1 μ g of total RNA was synthesized to the single-stranded cDNA using primers (LTC₄S, Forward: 5'-AGGTGGGCTGGTTCC TATCTA-3' and Reverse: 5'-CCCATGGCTATCCTACCATTT-3'; GAPDH, Forward: 5'-GCAAA GTCAAGGCTGAGAAC-3' and Reverse: 5'-ATGGTGGTGAAGACGCCAGT-3'). The PCR

products were separated by electrophoresis using a 1% ethidium bromide-stained agarose gel and visualized by ultraviolet transillumination.

Western blot analysis

To separate proteins (total protein concentration of cell lysate; $50 \mu g$), 10% sodium dodecyl sulphate-polyacrylamide gel electrophoresis was used. Then the gels were transferred to PVDF membrane (BIO-RAD, Hercules, CA, USA). The antibodies used were as follows: TGF- β 1 receptor (TGFR1; Abcam, Cambridge, United Kingdom; 1:1,000; 45 kDa), TGF- β 2 receptor (TGFR2; Abcam; 1:1,000; 75 kDa), LTC₄S (Sigma-Aldrich; 1:500; 40 kDa), p38 (Cell Signaling Technology; 1:1000; 38 kDa), phospho-p38 (Cell Signaling Technology; 1:500; 38 kDa), and actin (Santa Cruz, Dallas, TX, USA; 1:1,000; 42 kDa).

In vivo mouse model

All experimental protocols were approved by the Institutional Animal Care and Use Committee of Ajou University (IACUC-2017-0067). Female 6-week-old BALB/c wild-type mice (Jackson Laboratory, Bar Harbor, ME, USA) were maintained under specific pathogen-free conditions. To demonstrate the effect of TGF- β 1 on LTE₄ production, mice (n = 6 mice per group) were intranasally injected with 0.1 µg of mouse recombinant TGF- β 1 (R&D systems) for 5 days. Eosinophil numbers in bronchoalveolar lavage fluid (BALF) were determined by Diff-quick staining (Dade Behring, Dudingen, Switzerland). Moreover, LTE₄ (MyBioSource, San Diego, CA, USA) and EDN (LifeSpan BioSciences, Seatle, WA, USA) in BALF were measured using ELISA kits.

Statistical analysis

All statistical analyses were performed using IBM SPSS software, version 26.0 (IBM Corp., Armonk, NY, USA). *P* values < .05 was considered statistically significant. GraphPad Prism 8.0 software (GraphPad Inc., San Diego, CA, USA) was used to create graphs.

Results

Higher levels of serum TGF-\(\beta\)1 in AERD patients

Demographic data from the study subjects of the first cohort are described in Table 1. The presence of nasal polyps and decrease in FEV₁ (%) after lysine-aspirin bronchoprovocation test were significantly higher in AERD patients than in ATA patients (P = .001 and P = .001, respectively). In addition, lower baseline FEV₁ (%) and PC₂₀ methacholine values were noted in the AERD patients compared to the ATA patients (P = .026 and P = .001, respectively), whereas total IgE, total eosinophil count and sputum eosinophil/neutrophils (%) were not significantly different between the 2 groups. However, the levels of serum TGF-β1 were significantly higher in the AERD patients than in the ATA patients (P = .001). When asthmatic patients were divided into the TGF-β1-low and -high subgroups (the cutoff value, 48.1 ng/ mL), the TGF-β1-high subgroup showed lower baseline FEV₁ (%) than the TGF-β1-low subgroup within the AERD group (P = .034), while no differences were found within the ATA group (Table 2). In this study, we enrolled the second cohort to verify the reproducibility of clinical data. As in the result of the first cohort, the levels of serum TGF-\(\beta\)1 were also significantly higher in the AERD group than in the ATA group (P = .026; Table 3). In addition, the levels of urinary LTE₄ were significantly higher in the AERD group than those in the ATA group (P = .001; Table 3). These findings indicate that higher levels of TGF-β1 may have an important role in AERD pathogenesis.

Sputum Neu (%)

TGF-β1 (ng/mL)

NA

.001

NA

.021

| Variables | AERD (n = 336) | ATA (n = 442) | HCs (n = 253) | P value | | |
|-------------------------------|-------------------|-------------------|-----------------|-------------|--------------|-------------|
| | | | | AERD vs.ATA | AERD vs. HCs | ATA vs. HCs |
| Age (y) | 42.2 ± 13.9/336 | 44.7 ± 14.4/442 | 31.6 ± 10.6/253 | .018 | .001 | .001 |
| Female sex (%) | 64.9/336 | 61.9/442 | 54.7/253 | .385 | .012 | .065 |
| Atopy (%) | 52.1/330 | 48.8/412 | 28.0/200 | .367 | .001 | .001 |
| Nasal polyp (%) | 42.1/271 | 18.2/148 | NA | .001 | NA | NA |
| Severe asthma (%) | 22.5/329 | 16.8/440 | NA | .048 | NA | NA |
| Baseline FEV ₁ (%) | 85.3 ± 20.2/316 | 88.8 ± 19.6/335 | NA | .026 | NA | NA |
| Fall of FEV ₁ (%) | 16.1 ± 5.7/189 | 6.5 ± 3.8/185 | NA | .001 | NA | NA |
| PC ₂₀ (mg/mL) | 3.2 ± 4.6/239 | 5.3 ± 5.7/246 | NA | .001 | NA | NA |
| Total IgE (kU/L) | 358.4 ± 550.1/328 | 365.9 ± 590.0/410 | 74.0 ± 111.6/66 | .857 | .001 | .001 |
| TEC (/μL) | 413.9 ± 412.2/310 | 384.0 ± 368.1/344 | NA | .327 | NA | NA |
| Sputum Eos (%) | 23.9 ± 35.2/217 | 21.6 ± 32.5/216 | NA | .470 | NA | NA |
| | | | | | | |

Table 1. Demographic data from the study subjects enrolled in the first cohort of adult asthmatic patients.

P values were obtained by Pearson's Chi-square test for categorical variables (sex, atopy, nasal polyp, severe asthma, baseline FEV₁, Fall of FEV₁, sputum Eos and Neu) and Student's t test for continuous variables (age, PC₂₀, total IgE, TEC, TGF-β1).

NA

 $22.5 \pm 11.3/175$

AERD, aspirin-exacerbated respiratory disease; ATA, aspirin-tolerant asthma; HCs, healthy control subjects; FEV_1 , forced expiratory volume in 1 s; PC_{20} , the provocative concentration of methacholine required to cause a 20% fall in FEV_1 ; Fall of FEV_1 ; decrease in FEV_1 after the inhalation of lysin aspirin; IgE, immunoglobulin E; TEC, total eosinophil count; Eos, eosinophils; Neu, neutrophils; NA, not available.

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57.5 ± 34.6/166

 $33.1 \pm 14.2/191$

Function of TGF-β1 in LTC₄S expression and LTE₄ production

The levels of serum TGF- β 1 and urinary LTE₄ showed a significantly positive correlation (r = 0.273, P = .037; Fig 1). When peripheral eosinophils from asthmatic patients were treated with TGF- β 1, expression of LTC₄S in the cells was markedly upregulated (Fig 2A). In addition, levels of LTC₄S in the cells were enhanced by TGF- β 1 treatement (Fig 2B), but dexamethasone did not effectively reduce levels of LTC₄S (Fig 2C). In the presence of TGF- β 1, high levels of LTC₄S with increased in peripheral eosinophils from the AERD patients was noted compared to those of the ATA patients (Fig 2D). Moreover, the TGF- β 1 receptor (TGFR1) within

.561

.001

Table 2. Characteristics of asthmatic patients with high (≥48.1 ng/mL) and low TGF-β1 (<48.1 ng/mL) levels in the first cohort.

59.6 ± 33.2/192

 $28.4 \pm 15.7/304$

| Variables | AERD | | P value | ATA | | P value |
|-------------------------------|------------------|-------------------|---------|------------------|-------------------|---------|
| | High (n = 23) | Low (n = 168) | | High (n = 27) | Low (n = 277) | |
| Age (y) | 40.0 ± 13.8/23 | 43.9 ± 13.2/168 | .192 | 41.5 ± 11.9/27 | 45.8 ± 14.7/277 | .149 |
| Female sex (%) | 56.5/23 | 70.8/168 | .164 | 59.3/27 | 60.3/277 | .917 |
| Atopy (%) | 43.5/23 | 46.7/165 | .774 | 54.2/24 | 50.0/256 | .696 |
| Nasal polyps (%) | 57.1/21 | 47.5/118 | .413 | 42.9/7 | 14.0/57 | .056 |
| Severe asthma (%) | 30.4/23 | 29.3/167 | .914 | 25.9/27 | 19.3/274 | .414 |
| Baseline FEV ₁ (%) | 75.4 ± 19.2/21 | 85.5 ± 20.2/153 | .034 | 82.5 ± 23.6/16 | 89.2 ± 20.0/207 | .207 |
| PC ₂₀ (mg/mL) | 2.1 ± 2.7/16 | 3.5 ± 4.8/115 | .278 | $7.2 \pm 7.0/23$ | 6.2 ± 6.2/255 | .577 |
| Total IgE (kU/L) | 303.5 ± 285.5/23 | 363.4 ± 619.2/164 | .649 | 553.8 ± 910.6/22 | 371.7 ± 599.0/255 | .193 |
| TEC (/μL) | 352.3 ± 282.4/23 | 425.9 ± 410.2/150 | .408 | 496.7 ± 370.4/13 | 368.1 ± 352.5/210 | .204 |
| Sputum Eos (%) | 26.6 ± 39.5/19 | 21.1 ± 34.1/122 | .520 | 24.5 ± 30.6/10 | 19.0 ± 32.1/141 | .602 |
| Sputum Neu (%) | 58.0 ± 39.2/13 | 54.4 ± 34.2/95 | .727 | 62.3 ± 29.8/9 | 62.7 ± 33.8/118 | .973 |

P values were obtained by Pearson's Chi-square test for categorical variables and Student's t test for continuous variables.

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Table 3. Demographic data of the study subjects enrolled in the second cohort.

| Variables | AERD (n = 34) | ATA (n = 25) | P value |
|-------------------------------------|----------------------|----------------------|---------|
| Age (y) | $44.5 \pm 10.3/34$ | 49.2 ± 19.1/25 | .266 |
| Female sex (%) | 70.6/34 | 76.0/25 | .770 |
| Atopy (%) | 32.4/34 | 40.0/25 | .544 |
| Nasal polyp (%) | 64.0/25 | 18.2/11 | .014 |
| Severe asthma (%) | 52.9/34 | 32.0/25 | .109 |
| Baseline FEV ₁ (%) | $86.6 \pm 20.3/30$ | 94.5 ± 15.3/15 | .195 |
| Fall of FEV ₁ (%) | 17.8 ± 4.5/21 | 5.4 ± 2.1/9 | .001 |
| PC ₂₀ (mg/mL) | $3.2 \pm 4.3/25$ | $4.8 \pm 5.6/16$ | .308 |
| Total IgE (kU/L) | $232.7 \pm 242.9/32$ | $280.2 \pm 312.8/20$ | .542 |
| TEC (/μL) | $493.3 \pm 292.9/30$ | 428.9 ± 280.5/16 | .475 |
| Sputum Eos (%) | $30.8 \pm 41.6/25$ | $20.8 \pm 32.3/15$ | .429 |
| Sputum Neu (%) | $36.6 \pm 37.0/19$ | 59.8 ± 34.0/14 | .411 |
| TGF-β1 (ng/mL) | $36.9 \pm 15.2/34$ | 27.7 ± 15.3/25 | .026 |
| LTE ₄ (ng/mL creatinine) | $0.4 \pm 0.3/34$ | $0.1 \pm 0.2/25$ | .001 |

P values were obtained by Pearson's Chi-square test for categorical variables and Student's t test for continuous variables.

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eosinophils was highly expressed in peripheral eosinophils from the AERD patients than those from the ATA patients, while TGFR2 was not (S1 Fig). These results imply that eosinophils from AERD patients may be more sensitive to TGF- β 1 in association with LTE₄ production because of highly expressed TGFR1 on the surface of eosinophils.

Induction of eosinophil degranulation by LTE₄ treatment

As TGF- β 1 markedly enhanced LTE₄ production by eosinophils, sequential effects of LTE₄ on peripheral eosinophils were further investigated. In this study, we found significantly elevated levels of serum EDN in the AERD group compared to the ATA group (P=.036; Fig 3A). In addition, the levels of urinary LTE₄ were positively correlated with serum EDN (r=0.314, P=.035; Fig 3B). When peripheral eosinophils from asthmatic patients were treated with LTE₄, phosphorylation of p38 was significantly elevated in the cells; however, montelukast (cysLT receptor 1 antagonist) could inhibit phosphorylation of signaling molecules (Fig 4A). In addition, LTE₄ enhanced levels of EDN released from the eosinophils (Fig 4B). The eosinophils observed using confocal microscopy also showed granule proteins released by LTE₄ stimulation (Fig 4C), indicating that LTE₄ is important for inducing eosinophil degranulation through the p38 pathway.

Effect of granule proteins in airway epithelial cells

When airway epithelial cells (A549 cells) were co-cultured with peripheral blood eosinophils with/without LTE₄, significantly elevated levels of TGF- β 1 in culture supernatant were noted; however, the effect of eosinophils in airway epithelial cells was partially attenuated by montelu-kast treatment (S2 Fig). In particular, EDN could enhance TGF- β 1 production from airway epithelial cells. Although dexamethasone tends to inhibit the effect of granule proteins in airway epithelial cells, it could not fully attenuate TGF- β 1 production from the cells (S3 Fig), suggesting limited action of corticosteroids against eosinophil granule proteins.

Enhanced LTE₄ and EDN production by TGF-β1 treatment in vivo

To demonstrate the effect of TGF- β 1 on cysLT production, mice were intranasally injected with TGF- β 1 with or without montelukast for every 5 days (Fig 5A). As a result, the total cell

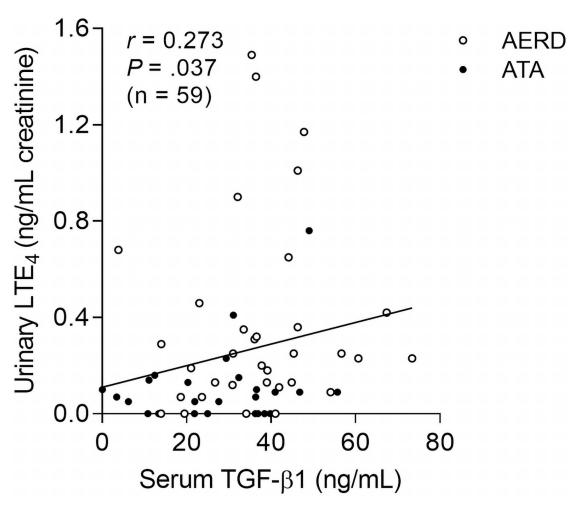


Fig 1. Association between levels of serum TGF- β 1 and urinary LTE₄ in the study subjects. Data are represented as Spearman correlation coefficient r(P value).

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and eosinophil but not macrophage number in BALF was markedly elevated in mice treated with TGF- β 1. Although montelukast did not fully reduce total cell count, the number of eosinophils was significantly decreased (Fig 5B). In addition, production of LTE₄ and EDN was markedly elevated when mice were treated with TGF- β 1, but montelukast could attenuated these mediators (Fig 5C and 5D). These findings show that TGF- β 1 may contribute to eosinophilic airway inflammation through induction of LTE₄ production.

Discussion

This is the first study to demonstrate the pathophysiological function of TGF- β 1 in AERD in the 2 clinical cohorts of adult asthmatic patients. It was found that the levels of serum TGF- β 1 were higher in the AERD patients than in the ATA patients. AERD patients with higher TGF- β 1 levels had lower FEV₁ (%) and PC₂₀ methacholine values, suggesting that TGF- β 1 may be involved in the lung. In addition, *ex vivo* and *in vivo* studies confirmed an association between TGF- β 1 and cysLT overproduction in AERD pathogenesis. Furthermore, increased LTE₄ could induce eosinophil degranulation, which further stimulates airway epithelial cells to

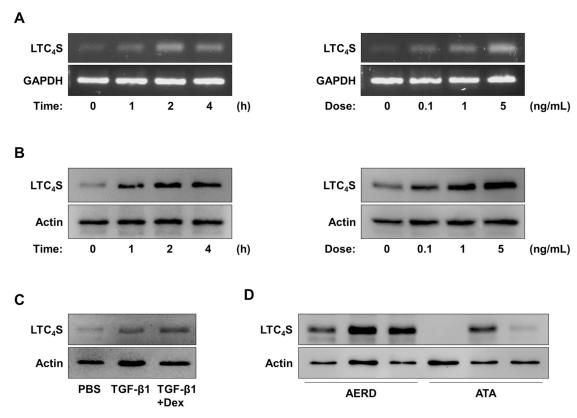


Fig 2. Effect of TGF- β 1 on LTC₄S expression in human peripheral eosinophils. Effect of TGF- β 1 on (A) LTC₄S expression and (B) LTC₄S levels in peripheral eosinophils in a time- or dose-dependent manner (samples from 3 asthmatic patients were pooled). (C) Function of dexamethasone against TGF- β 1 treatment (samples from 3 asthmatic patients were pooled). (D) Comparison of LTC₄S levels between ARED and ATA patients (n = 3 asthmatic patients per group).

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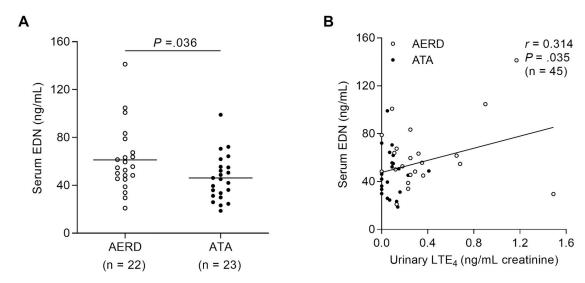


Fig 3. Relation between levels of urinary LTE₄ and serum EDN. (A) Levels of serum EDN in the study subjects. Data are presented as mean. P values were obtained by Student's t test. (B) A correlation between the levels of serum EDN and urinary LTE₄. Data are represented as Spearman correlation coefficient t (P value).

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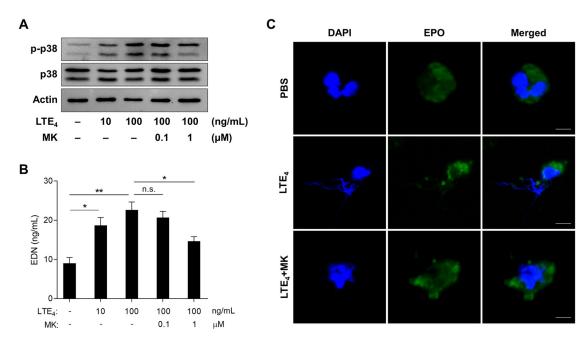


Fig 4. Function of LTE₄ in eosinophil degranulation. (A) Phosphorylation of p38 in peripheral eosinophils (samples from 3 asthmatic patients were pooled). (B) Levels of EDN released from the cells. Data are presented as mean \pm SD, n = 5. *P < .05 and **P < .01 were obtained by the Mann-Whitney test. n.s., not significant. (C) Images of eosinophils observed using confocal microscopy. Scale bar, 5 μ m. DAPI, 4',6-diamidino-2-phenylindole (blue); EPO, eosinophil peroxidase (green); MK, montelukast.

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produce TGF- β 1, resulting in the formation of the vicious circle. These provide a new insight into AERD pathogenesis via the TGF- β 1-LTE₄-eosinophil axis.

In the present study, significantly elevated levels of serum TGF-β1 were noted in the AERD patients compared to the ATA patients having a positive correlation with the levels of urinary

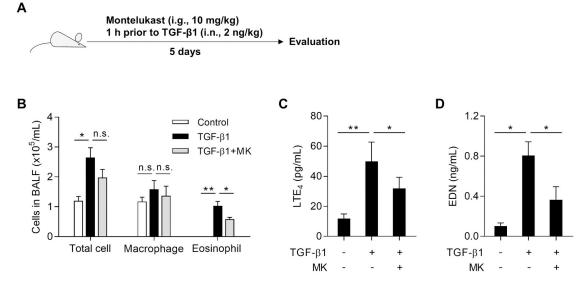


Fig 5. Roles of TGF-β1 in the lipoxygenase pathway to produce LTE₄ in vivo. (A) Experimental schedule. (B) Differential cell count. Levels of (C) LTE₄ and (D) EDN in bronchoalveolar lavage fluid. Data are presented as mean \pm SD, n = 6. *P < .05 and **P < .01 were obtained by the Mann-Whitney test. n.s., not significant. i.n., intranasal injection; i.g., intragastric administration; MK, montelukast.

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LTE₄. Previously, TGF- β 1 polymorphisms have been suggested as a risk factor for AERD development, and TGF- β 1 was associated with the prevalence of CRS in AERD patients, but not in ATA patients [21]. Nevertheless, the function of TGF- β 1 in the pathogenesis of AERD has not been fully understood. Therefore, we aimed to find the functional effect of TGF- β 1 in eosinophils, especially LTE₄ production and eosinophil activation. Although the role of TGF- β 1 in the arachidonic acid pathway was not well studied, a previous study revealed that TGF- β 1 contributed to changes in LTC₄S expression and cysLT₁R in astrocytes [22]. Our *ex vivo* study demonstrated significantly enhanced LTC₄S expression and LTE₄ production in response to TGF- β 1 in peripheral eosinophils from asthmatic patients. These suggest that TGF- β 1 may be an essential factor for LTE₄ production.

Here, we found highly expressed TGFR1, but not TGFR2 in peripheral eosinophils from AERD patients than from ATA patients. Previously, TGF- β 1 has been shown to enhance the expression of TGFR1, and activation of Smad and MAPK/ERK in fibroblasts [23]. These findings are one of the plausible mechanisms explaining how eosinophils could produce more LTE₄ in association with an increased level of TGF- β 1 in AERD patients. In addition, previous studies have shown that AERD patients present moderate to severe phenotypes with lower levels of FEV₁ (%) and PC₂₀ methacholine compared to ATA patients [24,25]. Here, we also showed that AERD patients with higher levels of serum TGF- β 1 with lower levels of FEV₁ (%), suggesting that TGF- β 1 may contribute to presenting more severe phenotypes with lung dysfunction. As conventional anti-inflammatory medications have limited effects in the TGF- β 1-mediated inflammatory pathway [2], a new therapeutic strategy to suppress the pathway noted in the present study is required in long-term management of AERD.

Overproduction of cysLTs is the key finding in the pathogenesis and progression of AERD pathogenesis. In particular, LTE₄ is involved in persistent eosinophilia via enhancement of eosinophil recruitment to the airway mucosa and bronchoconstriction [26]. In this study, LTE₄ could stimulate eosinophils to secrete EDN through p38 phosphorylation, similar to eotaxin (a potent stimulator of eosinophil chemotaxis) which binds to a CC chemokine receptor and induces eosinophil degranulation through activation of the ERK–p38 pathway [27]. However, LTE₄ certainly activates eosinophils via cysLT₁R rather than other receptors as montelukast reduced levels of granule proteins released from eosinophils. A previous paper has also been shown that LTE₄ is able to release granule proteins through binding to cysLT₁R (major) and other receptors (minor) [28]. These implicate that increased levels of LTE₄ may be responsible for enhancing eosinophil degranulation as well as eosinophil activation/recruitment, exacerbating type 2 airway inflammation in AERD patients where leukotriene receptor antagonists have partially suppressive effects.

Increased eosinophil number and activation markers in blood, sputum and tissues are common characteristics of bronchial asthma. The eosinophils communicate with several cell types involved in the pathogenesis of asthma; however, eosinophil-epithelial cell interactions have been extensively highlighted to play an important role in the processes of chronic airway inflammation as airway epithelial cells are a regulator of both innate and adaptive immune responses to host defence [29,30]. Following stimulation by multiple factors, airway epithelial cells produce large quantities of cytokines, chemokines and growth factors, such as TGF- β 1, enhancing type 2 immune response [31,32]. Although the mechanism of TGF- β 1 production from airway epithelial cells has not been fully elucidated, eosinophil granule proteins, such as ECP and EDN, are a possible factor contributing to the stimulation of the airway [33–35]. The function of EDN in enhancing airway remodeling in patients with eosinophilic CRS has been shown [36]. Furthermore, a recent study in adult asthmatic cohorts demonstrated that higher serum EDN was associated with severe asthma with asthma exacerbation [37]. In addition, a novel effect of EDN in airway epithelial cells on releasing TGF- β 1 was noted in our *in vitro*

study, where steroid may have a limited action. Taken together, these findings provide a possible mechanism of how activated eosinophils to induce TGF- $\beta1$ production contributing to chronic progressive type 2 airway inflammation in AERD pathogenesis.

This study has some limitations. First, the effect of TGF- $\beta1$ in multiple cells, such as mast cells, neutrophils or platelets, has not been determined. Secondly, further clinical trials in AERD patients according to the results of serum TGF- $\beta1$ levels and eosinophil activation status are needed to validate our findings.

In conclusion, TGF- β 1 has a novel function contributing to cysLT overproduction through induction of LTC₄S expression in eosinophils of AERD patients. Moreover, increased LTE₄ induces eosinophil degranulation via the p38 pathway which further stimulates airway epithelial cells, suggesting that TGF- β 1 plays a key role in enhancing eosinophilic airway inflammation, leading to poor clinical outcomes of AERD patients.

Supporting information

S1 Fig. Expression of TGF- β 1 and TGF- β 2 receptors in human peripheral eosinophils. (n = 3 asthmatic patients per group) TGFR1, TGF- β 1 receptor; TGFR2, TGF- β 2 receptor. (PDF)

S2 Fig. Effect of eosinophils on airway epithelial cells by secreting granule proteins. Levels of TGF- β 1 released from A549 cells when co-cultured with peripheral eosinophils with/without LTE4 or montelukast (MK). The data are presented as means \pm SD, n = 5. *P < .05 and **P < .01 were obtained by the Mann-Whitney test. n.s., not significant. (PDF)

S3 Fig. Function of dexamethasone (Dex) against eosinophil granule proteins to suppress TGF- β 1 production from airway epithelial cells. The data are presented as means \pm SD, n = 5. *P < .05 was obtained by the Mann-Whitney test. n.s., not significant. EDN, eosinophilderived neurotoxin. (PDF)

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