CD39/CD73 and the imbalance of Th17 cells and regulatory T cells in allergic asthma

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Abstract. In the immune system, CD4⁺CD25⁺Foxp3⁺ regulatory T cells (Tregs) maintain self-tolerance and Th17 cells mediate inflammatory responses. CD39 is expressed on the surface of a subset of naturally occurring human Tregs that are important in constraining pathogenic Th17 cells. Additional studies have shown that Tregs differentiate into interleukin-17 (IL-17)⁺Foxp3⁺ T cells. Our previous study indicated an imbalance of Th17 and Tregs in allergic asthma; however, the underlying mechanisms remain poorly understood. Using quantitative PCR (qPCR), CD39 and CD73 mRNA levels in CD4⁺ T cells were investigated. Flow cytometry was used to analyze the proportion of IL-17⁺Foxp3⁺ T cells, and CD39 and CD73 expressed by CD4⁺ T cells and Tregs in the peripheral blood of the subjects. The results of the present study demonstrated an increased frequency of CD4⁺Foxp3⁺IL-17⁺ T cells in moderate to severe asthma. A deficiency in CD39 expressed on the surface of CD4⁺ T cells and Tregs was observed in asthma patients. The expression of CD39 and CD73 on the surface of CD4⁺ T cells and Tregs was negatively correlated with the number of Th17 cells. These results indicated that the plasticity of Tregs transforming to IL-17⁺Foxp3⁺CD4⁺ T cells, the reduced frequency of CD39⁺ Tregs and less effective suppression of IL-17 production by residual CD39⁺ Tregs leads to an imbalance of Th17 and Tregs in asthma. Therefore, enhanced CD39 activity is hypothesized to prevent the progression of asthma.

Introduction

CD4⁺CD25⁺Foxp3⁺ regulatory T cells (Tregs) have a broad immunosuppressive capacity and are central in regulating self-tolerance and homeostasis in the immune system (1). Th17 cells are important contributors to the inflammatory response (2). A number of studies have demonstrated that Th17 cells and Tregs exhibit unique profiles of cytokines and regulatory transcription factors (3-5). The differentiation and development of Th17 cells is dependent on the transcription factor, ROR-γt (6,7), while Tregs require TGF-β and the forkhead transcription factor, Foxp3 (8).

Although a number of T cell lineages exhibit distinct gene expression and regulation signatures, each subset retains substantial developmental plasticity (4). However, it is suggested that Th17 cells and Tregs exhibited greater developmental plasticity than Th1 and Th2 cells (9). A number of studies have shown that Tregs are able to differentiate into CD4⁺Foxp3⁺interleukin-17 (IL-17)⁺ T cells in the colitic microenvironment, colon carcinoma, psoriasis, allergic rhinitis and polyposis (10-12). However, the biological mechanisms of CD4⁺Foxp3⁺IL-17⁺ T cells remain poorly understood in allergic asthma. The mechanisms of Treg suppression on effector T cell (Teffs) remain unclear, but include cell-cell contact and the release of the soluble mediators, IL-10 and TGF-β. Surface molecules linked to Treg suppression, include the CCR4 and CCR8 chemokine receptors, CTLA-4, the CD103 integrin, the CD62L selectin and CD127 (13-17). CD39 is an ectonucleotidase that catalyzes ATP/ADP to form AMP, which is cleared by CD73 to form adenosine and CD39. CD73 expression was observed on the surface of CD4⁺ T cells, particularly in a subpopulation of Tregs (18,19). Extracellular ATP has multiple proinflammatory effects, including promoting the secretion of IL-17 and the maturation of dendritic cells, and inducing the apoptosis of Tregs (18-20); thus, its removal may result in anti-inflammatory effects. Adenosine, which functions via the A₂A adenosine receptor expressed on the surface of T cells, is critical in inhibiting the functions of activated Teffs (21). A previous study observed that CD39⁺ Treg cells from patients with multiple sclerosis (MS) suppressed pathogenic Th17 cells (20). However, the involvement of CD39⁺ Tregs in allergic asthma remains unclear.

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Table I. Primer sequences of CD39, CD73, Foxp3, ROR-γt and β-actin.

<table>
<thead>
<tr>
<th>Gene</th>
<th>Primer</th>
<th>Length (bp)</th>
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<tr>
<td>CD39</td>
<td>5'-CTG ATT CCT GGG AGC ACA T-3'</td>
<td>143</td>
</tr>
<tr>
<td></td>
<td>5'-GAC ATA GGT GGA GTG GGA GAG-3'</td>
<td></td>
</tr>
<tr>
<td>CD73</td>
<td>5'-TTC TAA ACA GCA GCA TCT CT-3'</td>
<td>214</td>
</tr>
<tr>
<td></td>
<td>5'-AAC ATT TCA TCC GTG TGT CT-3'</td>
<td></td>
</tr>
<tr>
<td>Foxp3</td>
<td>5'-ATG CGA CCC CCT TTC ACC TAC-3'</td>
<td>155</td>
</tr>
<tr>
<td></td>
<td>5'-TGG CGG ATG GCG TTC TTC-3'</td>
<td></td>
</tr>
<tr>
<td>ROR-γt</td>
<td>5'-GGC TCC CTG GAT GAA TAG AAT G-3'</td>
<td>190</td>
</tr>
<tr>
<td></td>
<td>5'-AGG CAG AGG CAG AAA ATG TAA AG-3'</td>
<td></td>
</tr>
<tr>
<td>β-actin</td>
<td>5'-AAG GTG ACA GCA GTC GGT T-3'</td>
<td>195</td>
</tr>
<tr>
<td></td>
<td>5'-TGT GTG GAC TTG GGA GAG G-3'</td>
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</tbody>
</table>

In the current study, the unrecognized functions of CD39 expressed by CD4+ T cells and Tregs in controlling Th17 cells were investigated. An understanding of these functions may demonstrate the uncontrolled function of pathogenic T cells in allergic asthma.

Subjects and methods

Subjects and sample preparation. Patients with allergic asthma from outpatient clinics at the Department of Pulmonary Medicine, Ruijin Hospital (Shanghai, China) were consecutively recruited into the study. Asthma severity was assessed based on the Global Initiative for Asthma (GINA) (22). All participants performed a forced expiratory volume in the first second (FEV1 %pred) test, asthma control questionnaire (ACQ) and allergen tests. Patients had not been treated with systemic glucocorticoids for one month prior to the study and had never been treated with other immunosuppressive agents or undergone desensitization therapy. Healthy donors, with normal pulmonary function and negative allergy tests, were selected as normal controls. Heparinized peripheral venous blood (8 ml) was collected from each participant. Written informed consent was obtained from all individuals and the study received ethical approval from the Research Ethics Board of Ruijin Hospital, Shanghai Jiao Tong University School of Medicine (Shanghai, China).

Flow cytometry and antibodies. Expression markers on the surface of T cells were determined by fluorescence-activated cell sorting analysis, following surface staining or intracellular staining with specific anti-human antibodies. Antibodies included CD4/CD25-fluorescein isothiocyanate (FITC)/allophycocyanin (APC), CD39-PE-cy7, CD73-Percp5.5, Foxp3-PE, IL-17A-FITC and CD4-PE-cy5 (eBiosciences, San Diego, CA, USA). For the analysis of Th17 cells and CD4+Foxp3+IL-17+ T cells, 1 ml blood was added to 1 ml Dulbecco's modified Eagle's medium (Gibco-BRL, Carlsbad, CA, USA) and the mixture was stimulated with 20 ng/ml phorbol 12-myristate-13-acetate and 1 µg/ml ionomycin in the presence of 2 mmol/ml monensin (eBiosciences). Following culture (4 h; 37°C; 5% CO2), red cell lysing solution was added to lyse red blood cells and cells were washed once in phosphate-buffered saline. Cells were incubated with CD4-PE-cy5 at 4°C for 30 min, fixed and permeabilized with Perm/Fix solution (eBiosciences) according to the manufacturer's instructions, and were stained with Foxp3-PE and IL-17A-FITC. For the analysis of CD39+ and CD73+ Tregs, cells were incubated with CD4/CD25-FITC/APC and CD39-PE-cy7 or CD73-Perp5.5 at 4°C for 30 min, fixed and permeabilized with Perm/Fix solution, and stained with Foxp3-PE.

Quantitative PCR (qPCR). CD4+ T cells were obtained from peripheral blood using a human CD4+ T cell enrichment cocktail (Stem Cell Technologies Inc., Vancouver, BC, Canada) by Ficoll-Hypaque density centrifugation. Total RNA in the CD4+ T cells was isolated with TRizol reagent (Invitrogen Life Technologies, Carlsbad, CA, USA) and was reverse transcribed into cDNA (Promega Corporation, Madison, WI, USA). Primers were designed by Invitrogen Life Technologies and synthesized by BioTNT (Shanghai, China), according to the manufacturer's instructions. For amplification, the SYBR-Green I qPCR kit was used (BioTNT). Each reaction was run in triplicate on the ABI villa7 real time PCR system (Invitrogen Life Technologies, Carlsbad, CA, USA) and was normalized to housekeeping gene β-actin transcripts. Specific primers used are listed in Table I.

Statistical analysis. GraphPad Prism 5 software (GraphPad Software, Inc., La Jolla, CA, USA) was used for statistical analysis. Homogeneity of variance in the three groups was tested first. If each group showed homogeneity, analysis was performed using one way analysis of variance followed by Student-Newman-Keuls test and the data are presented as the mean ± SD. When heteroscedasticity was present in each group, data were analyzed using the Mann-Whitney test and are presented as medians (interquartile range). Pearson's correlation was used to analyze the relevance. P<0.05 was considered to indicate a statistically significant difference.

Results

General characteristics of subjects. According to the GINA guidelines, allergic asthmatics may be divided into two
subgroups, intermittent to mild and moderate to severe. No significant differences were identified in terms of age and gender between the patients and normal control groups. FEV1 (%pred) in patients with moderate to severe asthma was significantly lower than those with intermittent to mild asthma, but the scores for ACQ were of the opposite trend (Tables II and III; note that the number of patients differs between Tables II and III as the patients were recalled for the analysis in table III and not all were available).

Tregs differentiate into CD4+Foxp3+IL-17+ T cells in the microenvironment of allergic asthma. The relative frequency of CD4+Foxp3+IL-17+ T cells over total Tregs was 4.69% (2.90-8.01) in healthy controls. However, the frequency in patients with intermittent to mild asthma and moderate to severe asthma was 6.10 (4.63-8.70) and 16.80% (9.13-32.65), respectively. CD4+Foxp3+IL-17+ T cells were detected in the peripheral blood of patients with allergic asthma. Higher levels were observed in patients with moderate to severe asthma compared with those with intermittent to mild asthma (P<0.01) and healthy controls (P<0.01); however, no significant difference was identified between intermittent to mild asthma and healthy controls (Fig. 1).

Decreased CD39 and CD73 on the surface of CD4+ T cells in allergic asthma contributes to the imbalance of Th17 Tregs. Results demonstrated that the proportions of Th17 cells (23) and CD4+Foxp3+IL-17+ T cells were increased in asthma patients. Therefore, increased Th17 cells and CD4+Foxp3+IL-17+ T cells were hypothesized to be correlated with decreased CD4+CD39+ and CD4+CD73+ T cells. Thus, the mRNA and protein levels of CD39 and CD73 in CD4+ T cells were investigated. The correlation between the levels CD39 and CD73 expressed by CD4+ T cells and Th17 cells, CD4+Foxp3+IL-17+ T cells and disease severity were analyzed.

CD39 mRNA in CD4+ T cells was significantly lower in patients with moderate to severe asthma compared with those with intermittent to mild asthma [0.15 (0.10-0.19)x10^{-3} vs. 0.27 (0.19-0.39)x10^{-3}; P<0.01] and healthy controls [0.15 (0.10-0.19)x10^{-3} vs. 0.55 (0.34-0.70)x10^{-3}; P<0.001] and there was also a significant difference between intermittent to mild asthma and healthy controls (P<0.01; Fig. 2); however, CD73 mRNA in CD4+ T cells was significantly lower in patients with intermittent to mild asthma compared with healthy controls (Fig. 3). The relative frequency of CD4+CD39+ T cells in the peripheral blood was significantly higher in healthy controls compared with patients with intermittent to mild asthma (10.77±3.94 vs. 4.43±4.13%; P<0.001) and with moderate to severe asthma (10.77±3.94 vs. 4.43±4.13%; P<0.001); however, there was no significant difference identified between the two subgroups of asthma (Fig. 4). No significant differences in the levels of CD4+CD73+ T cell levels were identified among these three groups (Fig. 4).

Correlation analysis indicated that CD39 mRNA expression was positively correlated with Foxp3 mRNA expression (r=0.484; P<0.001) and negatively correlated with ROR-γt (r=-0.272, P=0.051). In addition, the frequency of Th17 cells was negatively correlated with the relative frequencies of CD4+CD39+ T cells and CD4+CD73+ T cells (r=-0.348, P<0.05).

### Table II. Characteristics of subjects for analyzing the expression of CD39 and CD73 protein in CD4+ T cells and Treg cells.

<table>
<thead>
<tr>
<th>Characteristics</th>
<th>Normal control (n=20)</th>
<th>Intermittent to mild asthma (n=23)</th>
<th>Moderate to severe asthma (n=15)</th>
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<tr>
<td>Age (years)</td>
<td>35.45±11.9</td>
<td>39.52±12.28</td>
<td>43.60±15.07</td>
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<tr>
<td>Gender (male/female)</td>
<td>6/14</td>
<td>11/12</td>
<td>4/11</td>
</tr>
<tr>
<td>FEV1 (% pred)</td>
<td>94.14±5.77</td>
<td>79.16±12.60a</td>
<td>66.51±7.04ab</td>
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<tr>
<td>ACQ</td>
<td>-</td>
<td>12.11±1.73</td>
<td>18.42±1.38b</td>
</tr>
</tbody>
</table>

*p<0.01, vs. normal control and *p<0.01, vs. mild asthma. FEV1, forced expiratory volume in 1 sec; ACQ, asthma control questionnaire.

### Table III. Characteristics of subjects for analyzing the expression of CD39, CD73, Foxp3 and ROR-γt mRNA in CD4+ T cells.

<table>
<thead>
<tr>
<th>Characteristics</th>
<th>Normal control (n=23)</th>
<th>Intermittent to mild asthma (n=17)</th>
<th>Moderate to severe asthma (n=12)</th>
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<tr>
<td>Age (years)</td>
<td>37.26±11.88</td>
<td>40.24±11.92</td>
<td>44.33±14.29</td>
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<td>Gender (male/female)</td>
<td>10/13</td>
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<tr>
<td>FEV1 (% pred)</td>
<td>94.03±5.39</td>
<td>81.24±11.35a</td>
<td>64.34±6.03ab</td>
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<tr>
<td>ACQ</td>
<td>-</td>
<td>11.52±2.29</td>
<td>18.07±1.44b</td>
</tr>
</tbody>
</table>

*p<0.01 vs. normal control and *p<0.01, vs. mild asthma. FEV1, forced expiratory volume in 1 sec; ACQ, asthma control questionnaire.

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CD39+ and CD73+ Treg cells are significantly decreased in the peripheral blood of patients with allergic asthma. Impaired function of Tregs was responsible for airway inflammation in allergic asthma. Preliminary studies indicated that the functional deficiency of Tregs results from a decrease in the number and capacity of cells secreting IL-10 and TGF-β, as well as a number of other factors. Tregs have been observed to coexpress CD39 and CD73 enzymes, which catalyze ATP and ADP into adenosine where adenosine binds to the A_{2A} receptor expressed by Teffs to suppress inflammatory responses (18). CD39+ Treg cells have been reported to be important in constraining pathogenic Th17 cells in MS (21). Further studies are required to determine the mechanisms of the imbalance of Th17 and Tregs in allergic asthma.

The proportions of CD39+ and CD73+ Treg cells over total Tregs in the peripheral blood were significantly higher in healthy controls compared with patients with intermittent to mild asthma [85.18 (79.04-88.12) vs. 36.02 (27.47-66.91)%; P<0.001 and 34.78 (26.30-54.23) vs. 10.50 (7.69-13.98)%; P<0.001, respectively; Figs. 5 and 6]. The relative frequency of Th17 cells demonstrated a similar trend in the opposite direction. The frequency of Th17 cells was observed to be negatively correlated with the relative frequencies of CD39+ and CD73+ Treg cells (r=-0.377, P<0.05 and r=-0.428, P<0.05, respectively; Table IV). The relative frequency of CD4+Foxp3+IL-17+ T cells was not shown to be correlated with the relative frequencies of CD39+ and CD73+ Tregs (Table IV).
Discussion

Our previous study demonstrated an increased proportion of Th17 cells and a decreased frequency of Tregs in asthma (23). Correlation analysis indicated that the frequency of peripheral blood Th17 cells was negatively correlated with the percentage of Tregs (23). Thus, the immunosuppressive capacity of Tregs was decreased in the inhibition of inflammatory responses mediated by Th17 cells and there was an imbalance of Th17 cells and Tregs in asthma. However, the underlying mechanisms remain unclear.

It has been shown that Tregs may be converted to Th17 cells (10,12). In the current study, Tregs in patients with asthma were observed to exhibit the capacity to produce IL-17 (CD4+Foxp3+IL-17+ T cells). CD4+Foxp3+IL-17+ T cells were hypothesized to represent Tregs in the middle stages of transformation into Th17 cells. Furthermore,
CD4⁺Foxp3⁺IL-17⁺ T cell levels were associated with disease severity. Thus, the imbalance of Th17 cells and Tregs in asthma was hypothesized to be due to the increased capacity of Tregs to transform into Th17 cells. A balance of Foxp3 and ROR-γt has been reported in Tregs, and Foxp3 is known to inhibit the expression of ROR-γt (4). In addition, CD39 promotes the expression of Foxp3, which amplifies and stabilizes the expression of CD39 (24). In the present study, CD39 mRNA was observed to be negatively correlated with ROR-γt mRNA and positively correlated with Foxp3 mRNA in CD4⁺ T cells. This observation indicates that decreased Foxp3 and CD39 and increased ROR-γt in asthma may be a mechanism of plasticity of Tregs transforming to Th17 cells in asthma.

The correlation between the imbalance of Th17 and Tregs in asthma and the expression of CD39 and CD73 by CD4⁺ T cells, particularly Tregs, was then investigated. To the best of our knowledge, this study demonstrated for the first time that a subset of human Tregs expressed CD39 and CD73 in asthma and the relative frequencies of CD4⁺CD39⁺ T cells, CD39⁺ Tregs and CD73⁺ Tregs in the peripheral blood were significantly lower in patients with intermittent to mild asthma compared with healthy controls. These results suggested that CD39 and CD73 may be involved in the occurrence and progression of allergic asthma. CD39 and CD73 expression by Tregs was not identified to be significantly different between asthma subgroups, but there was an increasing trend in the expression observed in patients with moderate to severe asthma. Tregs from individuals with asthma were hypothesized to constitute a deficiency in the mechanism of immunosuppression by increasing CD39 and CD73 expression, although Tregs do not inhibit the progression of asthma.

A number of studies have indicated that human Foxp3⁺ Tregs, while capable of suppressing proliferation and IFN-γ production, do not suppress the IL-17 production of Teffs (25,26). Mechanisms of suppression of Foxp3 Tregs include cell-cell contact, the release of soluble mediators (IL-10 and TGF-β) and the consumption of IL-2. In the present study, another mechanism, the negative correlation between the decreased relative frequency of CD39⁺ Tregs and increased Th17 cells, was introduced. Thus, the results indicated that CD39⁺ Tregs may inhibit the production of IL-17, which is not consistent with certain studies, which have indicated that total Foxp3⁺ Tregs do not suppress IL-17 production by T cells.

The relative frequencies of CD39⁺ and CD73⁺ Tregs were shown to be reduced in intermittent to mild asthma and there was an increasing trend in moderate to severe asthma. However, the increased relative frequency of CD39⁺ and CD73⁺ Tregs did not inhibit the progression of asthma. These observations indicated that CD39⁺ and CD73⁺ Treg populations in moderate to severe asthma are less effective at suppressing Th17 responses than the same cells from healthy controls. The current study indicated that the upregulation of Th17 cells in asthma may be due to a decrease in the relative frequency and impaired function of CD39⁺ and CD73⁺ Tregs in asthma.

In the present study, the relative frequencies of CD4⁺Foxp3⁺IL-17⁺ T cells, Th17 cells and CD39 and CD73 expression by CD4⁺ T cells and Tregs was determined and their correlation was analyzed. Future studies are likely to investigate the suppressive function of CD39⁺ Tregs and their biological characteristics in asthma.

In conclusion, the plasticity of Tregs transforming to IL-17⁺Foxp3⁺CD39⁺ T cells, the reduced frequency CD39⁺ Tregs and the efficacy of suppression of IL-17 production by residual CD39⁺ Tregs, may lead to the imbalance of Th17 and Tregs in allergic asthma. The observations indicate that enhancing CD39 activity may be beneficial in preventing the progression of asthma.

References


