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# Role of prostaglandin  $D<sub>2</sub>/CRTH2$  pathway on asthma exacerbation induced by Aspergillus fumigatus

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## **Summary**

Aspergillus fumigatus is often associated in asthmatic patients with the exacerbation of asthma symptoms. The pathomechanism of this phenomenon has not been fully understood. Here, we evaluated the immunological mechanisms and the role of the prostaglandin  $D_2$ / Chemoattractant Receptor-Homologous Molecule Expressed on Th2 Cells (CRTH2) pathway in the development of Aspergillus-associated asthma exacerbation. We studied the effects of A. fumigatus on airway inflammation and bronchial hyper-responsiveness in a rat model of chronic asthma. Inhalation delivery of A. fumigatus conidia increased the airway eosinophilia and bronchial hyper-responsiveness in ovalbumin-sensitized, challenged rats. These changes were associated with prostaglandin  $D_2$  synthesis and CRTH2 expression in the lungs. Direct inflammation occurred in ovalbumin-sensitized, challenged animals, whereas pre-treatment with an antagonist against CRTH2 nearly completely eliminated the A. fumigatus-induced worsening of airway eosinophilia and bronchial hyper-responsiveness. Our data demonstrate that production of prostaglandin  $D_2$  followed by eosinophil recruitment into the airways via a CRTH2 receptor are the major pathogenic factors responsible for the A. fumigatus-induced enhancement of airway inflammation and responsiveness.

Keywords: airway responsiveness; Aspergillus fumigatus; asthma; CRTH2; eosinophil; prostaglandin  $D_2$ .

## Introduction

Asthma is an inflammatory disease of the airways that may be worsened by numerous extrinsic factors. The most common trigger is continuous exposure to allergens of which fungal agents are important factors. There is overwhelming evidence for the presence of fungal sensitization in patients with asthma. There is also a strong association between fungal exposure and severity of asthma. Epidemiological studies have associated mould exposure with the development, persistence and severity of asthma. However, the mechanism of these changes has not been established. Several lines of evidence suggest that eosinophils play an important role in patients with asthma during fungal infection-induced exacerbation.<sup>1–9</sup> However, little is known regarding the mechanisms of recruitment and activation of eosinophils in asthmatic airways and enhancement of airway responsiveness during fungal infection.

Infection with fungus immediately activates the innate immune system via several pathways.<sup>10</sup> Aspergillus fumigatus, the most common species colonizing the lungs of patients with chronic lung disease, is recognized by membrane or cytosolic receptors, such as cognate C-type lectin and Toll-like receptors. $11-13$  In asthmatic patients, the changed milieu in the airways facilitates the adhesion of A. fumigatus spores, which allows the colonization and eventual release of spores.<sup>14</sup> This process activates signalling pathways linked to the production of inflammatory cytokines and chemokines, as well as the production of reactive oxygen. The activation of these pattern-recognition receptors by A. fumigatus triggers the synthesis of interleukins<sup>15,16</sup> and other cytokines/chemokines in airway epithelial cells, dendritic cells and other immune cells.<sup>17</sup> However, little is known about its impact on the axis of lipid mediators and their receptors.

Recent studies have shown that prostaglandin  $D_2$ (PGD2), a major cyclooxygenase 3 metabolite synthesized in activated mast cells and macrophages, acting as a potent chemoattractant of eosinophils, is an important mediator involved in eosinophilic airway inflammation.<sup>18–23</sup> The bioactivity of  $PGD<sub>2</sub>$  is mediated by two G protein-coupled receptors, DP (DP1) and Chemoattractant Receptor-Homologous Molecule Expressed on Th2 Cells (CRTH2) (DP2), and both in vitro and in vivo eosinophil trafficking is mediated mostly by CRTH2, which is preferentially expressed on eosinophils, basophils and T helper type 2 lymphocytes.<sup>20,21,24–26</sup> Reports recently showed that CRTH2 agonists administered into the trachea led to translocation of eosinophils from the bloodstream into the airway, $27-29$  and that CRTH2-deficient mice exhibited decreased infiltration of eosinophils and other inflammatory cells during chronic allergic skin inflammation.<sup>30</sup>

As reported herein, we developed rat models mimicking the enhanced asthmatic responses induced by A. fumigatus. In these models, we used the inhalation delivery of A. fumigatus conidia to simulate fungal exposure and showed that it worsened allergen-induced eosinophilic airway inflammation and airway hyperresponsiveness (AHR). In these experiments, we also establish that  $PGD<sub>2</sub>$ , acting via the CRTH2 receptor, is the essential modulator of eosinophilic airway inflammation and bronchial hyper-responsiveness induced by A. fumigatus spore inhalation.

## Materials and methods

#### Animals

Specific pathogen-free 5- to 6-week-old, male Wistar rats weighing between 140 and 180 g were supplied by the Shanghai Laboratory Animal Centre (Shanghai, China). The animals were maintained in a 12/12 hr light/dark cycle at a room temperature of 23° and relative humidity 40%; they were provided with standard laboratory rat chow and water ad libitum. The experimental protocol was reviewed and approved by the Shanghai Jiaotong University School of Medicine Ethical Committee for Laboratory Animals.

## Protocols for allergen exposure and A. fumigatus inhalation

A total of 40 male Wistar rats were randomly divided into five groups ( $n = 8$ /group): control group (NS); A. fumigatus spore-exposed group  $(NS + AF)$ ; ovalbumin  $(OVA)$ -sensitized and OVA-challenged group (OVA); OVA-sensitized, OVA-challenged and A. fumigatus spore-exposed group (OVA + AF); and CRTH2 antagonist OC00459 treated group  $(OVA + AF + Treat)$ . Saline  $(0.1 \text{ ml})$  was intraperitoneally administered to a control group (NS) and an AF group ( $NS + AF$ ) on days 0 and 7. The other groups were sensitized and challenged with OVA. Briefly, rats were sensitized by an intraperitoneal injection of 1 mg OVA (Albumin Chicken egg, Grade V; Sigma Chemical Co., St. Louis, MO) with 100 mg aluminium hydroxide (Sigma

Chemical) in 1 ml saline on days 0 and 7. Rats were challenged via the airways with aerosolized 2% OVA for 30 min, 3 days per week, from days 14 to 37 by an ultrasonic nebulizer (PARI-BOY N037; PARI, Starnberg, Germany). Five days after the last challenge (day 42), rats were given via the airways aerosolized A. fumigatus spore suspension 5 days per week from days 42 to 67 (Fig. 1). Nonsensitized and non-challenged rats receiving aerosolized saline were treated in the same fashion. From days 42 to 67, a single oral dose of 5 mg/kg OC000459 by gavage in 10% DMSO/saline solution was administered to an antagonisttreated group  $(OVA + AF + Treat)$ . Airway hyper-responsiveness was assessed 72 h after the last inhalation, and tissues and cells were obtained for further assays.

#### Measurement of AHR

Airway hyper-responsiveness was assessed by a change in airway function after the challenge with aerosolized methacholine (Sigma-Aldrich, St Louis, MO) using barometric plethysmography (Buxco Electronics Inc., Troy, NY) as previously described.31,32 Briefly, pressure differences were measured between the main chamber of the plethysmograph, which contained conscious, spontaneously breathing animals, and the reference chamber (box pressure signal). Rats were challenged with aerosolized saline for baseline measurements or with methacholine (3125–50 mg/ml) for 3 min, and readings were taken and averaged for 3 min after each nebulization. Data were expressed using the dimensionless parameter, enhanced pause (Penh).

#### Sample collection

Rats were anaesthetized with an intraperitoneal injection of 1% sodium pentobarbital. Approximately 10 ml of blood was collected by heart bleed. The serum was isolated and stored at  $-80^\circ$  until assay.

The trachea was cannulated and the left lungs were lavaged via the tracheal cannula with 3.5 ml PBS (pH 72) three times. Lavage fluid was recovered by gentle aspiration with a syringe. The collected bronchoalveolar lavage fluid was then centrifuged at  $401$  g for 7 min at 4°. The supernatant from the first wash was collected and stored at  $-80^{\circ}$  for cytokine determination.

The right lower lobe was harvested, inflated with 10% neutral buffered formalin, fixed in formalin for 16 hr, and paraffin-embedded for histological and immunohistochemical analyses. The other lobe was stored in liquid nitrogen for further examination.

## Quantification of eosinophils in the bronchoalveolar lavage

After supernatant removal, cells were resuspended in 05 ml PBS (pH 74). Total cell numbers were counted on



Figure 1. Experimental protocols. Rats were divided into five groups: non-sensitized control group (NS); Aspergillus fumigatus spore-exposed group (NS + AF); ovalbumin (OVA) -sensitized and OVA-challenged group (OVA); OVA-sensitized, OVA-challenged and A. fumigatus sporeexposed group (OVA + AF); and CRTH2 antagonist OC00459-treated group (OVA + AF + Treat). In the OVA-sensitized and OVA-challenged receiving saline (OVA) group, rats were sensitized by two intraperitoneal injections of OVA/aluminium hydroxide and then received three consecutive days of an aerosolized OVA challenge from days 14 to 37 for 4 weeks. To evaluate the effect of A. fumigatus, OVA-sensitized and OVAchallenged mice received 5 consecutive days of an aerosol inhalation of  $1 \times 10^6$  colony-forming unit/ml spore suspension from days 42 to 67 for 4 weeks. The OVA + AF + Treat group received OC000459 at a dose of 5 mg/kg daily. Non-sensitized and non-challenged mice receiving saline or A. fumigatus represented the NS and NS + AF groups, respectively. Values are means $\pm$ SEM. \*P < 0.05.

a haemocytometer and  $1 \times 10^3$  to 5  $\times 10^3$  cells were spun onto glass microscope slides (cytospin 3; Shandon Scientific, Runcorn, UK). The cell slides were air dried for 24– 36 hr, fixed and stained with a Wright–Giemsa stain set (Jiancheng Bioengineering Institute, Nanjing, China). Differential cell counts of at least 300 cells in duplicate slides were counted in a blinded fashion according to morphological criteria. The number of eosinophils recovered was calculated and expressed as absolute cell numbers.

#### Detection of PGD<sub>2</sub>

Concentration of  $PGD<sub>2</sub>$  in serum and bronchoalveolar lavage fluid supernatants was measured by the Prostaglandin D<sub>2</sub>-MOX EIA Kit (Detection Limit 80% B/B0 31 pg/ml; Cayman Chemical, Ann Arbor, MI). This assay is based on the competition between sample  $PGD<sub>2</sub>$ and a PGD<sub>2</sub>-acetylcholinesterase conjugate (PGD<sub>2</sub> tracer) for a limited number of  $PGD<sub>2</sub>$  monoclonal antibody binding sites. Sample extraction procedures for protein removal and eicosanoid stabilization were performed according to the provider's instructions, and the  $PGD<sub>2</sub>$ concentration was determined following the manufac-

turer's instructions. The assay has a detection range between 250 and 2 pg/ml, and intra- and inter-assay coefficient of variation was < 10%.

## Immunohistochemical analysis of lung CRTH2 expression

Sections of lung specimens underwent immunoperoxidase staining using antibodies directed against CRTH2 (1 : 400) (PAB13243; Abnova, Taipei City, Taiwan). The sections were deparaffinized in xylene and rehydrated in methanol. Endogenous peroxidases were blocked by 5%  $H_2O_2$  treatment. For better antigen retrieval of CRTH2, the samples were boiled in a microwave oven for 2–3 min in citrate buffer. Samples were then washed with PBS. Samples were then incubated with the primary antibody at room temperature for 1 hr. After washing, the revelation was performed with the use of appropriate secondary antibody, the Readyto-go SABC-AP anti-rabbit IgG kit (Boster, Wuhan, Hubei, China) according to the supplier's recommendations. Immunoreactivity was visualized by a treatment with diaminobenzidine (Sigma-Aldrich), and the slides were counterstained with Mayer's haematoxylin.

#### Measurement of total CRTH2 protein expression

Preparation of protein extracts. For preparation of protein extracts, tissue was crushed in the frozen state in a cryotube by shaking with a sterile steel ball. The crushing was resuspended in cold RIPA lysis buffer (Tris–HCl pH 76 25 mM; NaCl 150 mM; Nonidet-P40 1%; sodium deoxycholate 1%; sodium dodecyl sulphate 01%; Pierce, Rockford, IL) supplemented with protease inhibitors (Complete, Roche, Basel, Switzerland) and subsequently swirled for 10 min on ice. The extracts were then centrifuged at 14 000  $g$  for 15 min at 4°. The supernatants were analysed for protein content by the Bio-Rad protein assay based on the Bradford method (Bio-Rad, Hercules, CA).

Western blotting. Samples were separated on a 10% SDS– PAGE and transferred to a polyvinylidene difluoride membrane (Roche). The anti-CRTH2 rabbit polyclonal antibody (PAB13243; Abnova) was used at a 1 : 1000 dilution. The secondary anti-rabbit antibody coupled with horseradish peroxidase (HSA0003) at a 1 : 3000 dilution was detected by chemiluminescence with the ECL system (Beyotime, Haimen City, Jiangsu, China), using glyceraldehyde 3-phosphate dehydrogenase (GAPDH) as loading control.

## Reverse transcription-PCR analysis of CRTH2 mRNA

Total RNA was isolated using the TRIzol® reagent (TaKa-Ra, Yasu, Shiga, Japan) according to the manufacturer's instructions. To remove contaminating DNA, isolated RNA was treated with RNase-free DNase I (Takara Bio Inc). The RNA was quantified using the RiboGreen RNA quantification kit (Molecular Probes, Eugene, OR), dissolved in diethylpyrocarbonate-treated H<sub>2</sub>O and stored at  $-80^\circ$  until use. One microgram of total RNA was reverse-transcribed using PrimeScript™ Master Mix (Takara Bio Inc.) as detailed in the manufacturer's guidelines. One-fiftieth of the reverse transcriptase reaction was analysed by real-time quantitative PCR. The following primers were used: GAPDH, sense 5′-CAC CCg CgA gTA CAA CCT TC-3′ and antisense 5′-CCC ATA CCC ACC ATC ACA CC-3′; CRTH2, sense 5′-gCT TCC AAA CCA CAg CAA CT-3′ and antisense 5′-CCA CCA CAA ACA ggA TgA gTC-3′.

Quantitative PCR analysis was performed in a total volume of 20  $\mu$ l containing template DNA, 0.05 nm of sense and antisense primers, 10 µl SYBR®Premix Ex Taq (Tli RNaseH Plus; Takara Bio Inc.). After incubation at 95° for 30 seconds (initial denaturation), the mixtures were subjected to 40 amplification cycles (5 seconds at 95° for denaturation and 30 seconds for annealing and extension at 60°). Incorporation of SYBR® Green dye into PCR products was monitored in real time using a  $CFX96^{TM}$ Real-Time PCR Detection System (Applied Biosystems, Foster City, CA) allowing determination of the threshold cycle (CT) at which exponential amplification of PCR products begins. After PCR, dissociation curves were generated with one peak, indicating the specificity of the amplification. A threshold cycle (CT value) was obtained from each amplification curve using the software provided by the manufacturer (Applied Biosystems).

Relative mRNA expression in chondrocytes was determined using the  $\Delta\Delta CT$  method, as detailed in the manufacturer's guidelines (Applied Biosystems). A  $\Delta CT$  value was first calculated by subtracting the CT value for the housekeeping gene GAPDH from the CT value for each sample. A  $\Delta\Delta$ CT value was then calculated by subtracting the  $\Delta$ CT value of the control from the  $\Delta$ CT value of each treatment. Fold changes compared with the control were then determined by raising 2 to the  $-\Delta\Delta CT$  power. Each PCR generated only the expected specific amplicon as shown by the melting-temperature profiles of the final product and by gel electrophoresis of test PCRs. Each PCR was performed in triplicate on two separate occasions for each independent experiment.

#### Statistical analysis

Statistical analysis was performed by one-way analysis of variance and Student–Newman–Keuls test.  $P$  value <0.05 was considered significant.

#### Results

## A. fumigatus increases airway responsiveness in OVAsensitized rats only

Analysis of the  $OVA + AF$  and  $OVA + NS$  groups showed significant differences in Penh (Fig. 2a). The inhalation of A. fumigatus significantly increased Penh in  $OVA + AF$  group rats relative to  $OVA + NS$  in an methacholine dose-dependent manner (Fig. 2a). Analysis of  $NS + AF$  and  $OVA + AF$  groups showed significant differences in Penh between these groups (Fig. 2a). To determine if AF without exposure to OVA has an effect on airway responsiveness, we compared Penh in the  $NS + AF$  group with Penh in the  $NS + NS$  group. No significant difference in Penh was observed in the  $NS + AF$  group when compared with the  $NS + NS$  group (Fig. 2a). Significant reduction in Penh was observed in the  $OVA + AF + Treat$  group when compared with the  $OVA + AF$  group (Fig. 2a).

## A. fumigatus increases lung eosinophilia in OVAsensitized rats

Significant enhancement of the eosinophil ratio in bronchoalveolar lavage fluid was generally seen in OVA-sensitized, OVA-challenged rats (NS  $11.34 \pm 2.71\%$  versus



Figure 2. Inhalation of Aspergillus fumigatus increased bronchial hyper-responsiveness (a) and recruited eosinophils into the airways (b) of ovalbumin (OVA) -sensitized and OVA-challenged rats. Treatment with OC000459 prevents the development of airway hyper-responsiveness (AHR) (a) and eosinophil accumulation in bronchoalveolar lavage (BAL) fluid (b). Enhanced pause (Penh) values for increasing concentrations of inhaled methacholine and the ratio of eosinophils in BAL fluid were measured in the five groups: non-sensitized control group (NS); Aspergillus fumigatus spore-exposed group (NS + AF); ovalbumin (OVA) -sensitized and OVA-challenged group (OVA); OVA-sensitized, OVA-challenged and A. fumigatus spore-exposed group (OVA + AF); and CRTH2 antagonist OC00459-treated group (OVA + AF + Treat)., 72 hr after the last inhalation. Eo, eosinophil. Values are means  $\pm$  SEM. \*P < 0.05.

OVA 24.93  $\pm$  8.81%, P < 0.05). Inhalation of A. fumigatus in NS-sensitized, challenged rats did not increase the ratio of eosinophils in bronchoalveolar lavage fluid (NS + AF 13.34  $\pm$  2.05%, P = 0.425 versus NS 11.34  $\pm$  2.71%) but produced a significant increase in the ratio of eosinophils in bronchoalveolar lavage fluid of OVA-sensitized and OVA-challenged rats (OVA + AF 31.24  $\pm$  3.81% versus OVA 24.93  $\pm$  8.81%,  $P < 0.05$ ; Fig. 2b). In bronchoalveolar lavage fluid from the treated group, the ratio of eosinophils was significantly lower compared with the  $OVA + AF$  group  $(OVA + AF + Treat 6.38 \pm 1.68\%$  versus OVA + AF 31.24  $\pm$  3.81%  $P < 0.05$ ; Fig. 2b).

## Role of  $PGD<sub>2</sub>/CRTH2$  receptor in A. fumigatusinduced exacerbation of eosinophilic airway inflammation in OVA-sensitized rats

In the present study we observed that A. fumigatus increased the concentration of  $PGD<sub>2</sub>$  in the bronchoalveolar lavage fluid of OVA-exposed animals significantly, though not in bronchoalveolar lavage fluid of NS-exposed animals (Fig. 3). To determine the mechanisms involved in  $PGD<sub>2</sub>$ -mediated eosinophil chemotaxis we first examined the expression of PGD<sub>2</sub> receptor CRTH2. Expression of CRTH2 was identified in lung tissue by immunohistochemical analysis (Fig. 4). Microscopic examination of the lung immunostained slides showed higher expression for CRTH2 in samples from OVA-sensitized rats compared with NS group (Fig. 4). CRTH2 expression was lower in the OVA group than in the OVA +  $AF$  group (Fig. 4) and lower in the treated rats than in the



Figure 3. Prostaglandin  $D_2$  (PGD<sub>2</sub>) levels measured by using ELISA in bronchoalveolar lavage (BAL) fluid samples. Administration of Aspergillus fumigatus alters  $PGD<sub>2</sub>$  levels in BAL fluid. Levels of  $PGD<sub>2</sub>$ in BAL fluid were measured in the five groups: non-sensitized control group (NS); Aspergillus fumigatus spore-exposed group (NS + AF); ovalbumin (OVA) -sensitized and OVA-challenged group (OVA); OVA-sensitized, OVA-challenged and A. fumigatus spore-exposed group (OVA + AF); and CRTH2 antagonist OC00459-treated group (OVA + AF + Treat), 72 hr after the last challenge. Values are means  $\pm$  SEM.  $*P < 0.05$ .

 $OVA + AF$  group (Fig. 4). These results indicate the involvement of CRTH2 in the pathophysiology of eosinophilic inflammation in asthma.

We also investigated the effect of A. fumigatus on the expression of CRTH2 by quantitative PCR (Fig. 5). In



Figure 4. Immunohistochemical staining of CRTH2 in lung tissue from the five groups; non-sensitized, non-challenged saline inhaled rats (NS) (a), non-sensitized, non-challenged and Aspergillus fumigatus spores inhaled rats  $(NS + AF)$  (b), OVA-sensitized, OVA-challenged and salineinhaled rats (OVA) (c), OVA-sensitized, OVA-challenged and A. fumigatus spores inhaled rats (OVA + AF) (d), and OVA-sensitized, OVA-challenged and A. fumigatus spores inhaled rats receiving OC000459 (OVA + AF + Treat) (e). (f) PBS negative control.

comparison with the NS group, CRTH2 mRNA levels were significantly elevated in all OVA-sensitized and challenged groups (Fig. 5). CRTH2 mRNA levels were significantly decreased in  $OVA + AF + Treat$  group versus the  $OVA + AF$  group, and the lowest level was found in the  $NS + AF$  group (Fig. 5).

These results were confirmed by Western blot analysis using specific anti-CRTH2 antibodies (Fig. 6). Higher expression was observed in OVA-sensitized with respect to NS rats. Inhalation of A. fumigatus increased the total CRTH2 protein contents. As seen in Fig. 6, after treatment the expression of CRTH2 protein significantly decreased. Equal loading of protein in tissue homogenates was also determined by Western blot using an anti-GAP-DH antibody. All blots were repeated three times and one representative image is shown.

These results indicate that infiltrated cells express functional CRTH2 and suggest that  $PGD<sub>2</sub>$  might modulate eosinophil chemotaxis through activation of CRTH2, which may be mediated by an increase of transcription at the CRTH2 locus.

#### **Discussion**

Eosinophilia is a hallmark in allergic asthma, and is blamed for the chronicity of the disease. Exacerbations of asthma are known to be associated with eosinophilic inflammation in children and adults.<sup>33-35</sup> For example, asthmatics with refractory eosinophilic inflammation have significantly higher exacerbation rates, $33$  whereas eosinophil-targeted therapies such as anti-interleukin-5 significantly reduce asthma exacerbation rates. $33-35$  In murine models of allergen-induced AHR, depletion of eosinophils protects mice from developing mucus accumulation, AHR and/or airway remodelling, suggesting that eosinophilia plays a role in the pathogenesis of asthma.<sup>36,37</sup>



Figure 5. qRT-PCR of CRTH2 mRNA transcripts in lung tissue. Results were expressed as the number of CRTH2 gene copies normalized to the number of glyceraldehyde-3-phosphate dehydrogenase gene copies (housekeeping gene). Values are means  $\pm$  SEM.  $*P < 0.05$ . #The difference between groups tended to be different  $(P = 0.05)$  but did not reach significance.

The immunomodulatory effect of A. fumigatus and its components have been well documented.<sup>38-41</sup> Exposure to A. fumigatus extracts leads to eosinopoiesis and the subsequent increase in peripheral blood eosinophils, reiterating that fungal antigens are responsible for the recruitment of eosinophils into the affected tissues.<sup>42</sup> In our previous studies we showed that prolonged exposure of OVA-sensitized and OVA-challenged rats to A. fumigatus spores resulted in enhanced levels of serum IgE, pulmonary eosinophils and bronchial hyper-responsiveness.14,43 Although some authors reported accumulation of eosinophils following A. fumigatus inhalation,  $44-47$  in this study only a few eosinophils were observed after A. fumigatus exposure alone. However, when A. fumigatus inhalation was combined with OVA allergen challenge of sensitized mice, the number of eosinophils was also significantly enhanced. Similarly, studies in humans have also shown significant blood eosinophilia, as a result of infection with A. fumigatus.<sup>48-50</sup> Therefore, the current interest in airway eosinophilic inflammation is to discover the molecular events resulting in eosinophil recruitment and hence to develop an effective mode of therapy.

The accumulation of eosinophils in the lung is regulated by chemokines such as eotaxin, as well as a number of different cytokines such as interleukin-5, granulocyte– macrophage colony-stimulating factor and granulocyte colony-stimulating factor.<sup>51–53</sup> Mast-cell-derived PGD<sub>2</sub> is also an important mediator involved in eosinophilic airway inflammation $19,23$  and plays a substantial role in persistent eosinophil accumulation in airways repeatedly exposed to allergens. Current evidence showed that



Figure 6. Expression of CRTH2 in lung tissue. Western blot was performed on whole tissue extracts from the five groups: non-sensitized control group (NS); Aspergillus fumigatus spore-exposed group (NS + AF); ovalbumin (OVA) -sensitized and OVA-challenged group (OVA); OVA-sensitized, OVA-challenged and A. fumigatus spore-exposed group (OVA + AF); and CRTH2 antagonist OC00459-treated group (OVA + AF + Treat).; GAPDH was used as control.

A. fumigatus exposure significantly increased both chemoattractants (data not shown), but the levels of  $PGD<sub>2</sub>$  were also increased.

In the present study, we developed an experimental model that mimics the pathological and physiological changes observed in the airways of asthmatic patients during exacerbations induced by fungus exposure. The experiment focused on the effect of A. fumigatus on a Wistar rat model of chronic asthma in rats sensitized and challenged by OVA antigen. Herein, we show that the PGD<sub>2</sub>/CRTH2 pathway, which has been reported to be up-regulated in lung tissues of subjects with allergic asthma, has the ability to be up-regulated with A. fumigatus. We have shown, in this study, that (i) the administration of A. fumigatus into the airways increased the synthesis of  $PGD<sub>2</sub>$  in OVA-exposed rats only, (ii) inhalation delivery of A. fumigatus conidia recruited eosinophils into the airways of OVA-exposed rats and increased bronchial hyper-responsiveness, (iii) the pharmacological blockade of CRTH2, a  $PGD<sub>2</sub>$  receptor, prevented the increase in eosinophil accumulation and/or bronchial hyper-responsiveness induced by the administration of A. fumigatus in rats, and (iv) the administration of A. fumigatus into the airways increased the expression of CRTH2 in OVA-exposed lungs and its effect on the level of CRTH2 mRNA suggesting that the enhancing effect of A. fumigatus is mediated by an increase of transcription at the CRTH2 locus. All of these observations strongly indicate that  $PGD<sub>2</sub>$  and its receptor CRTH2 is the essential system for the exacerbation of eosinophilic airway inflammation induced by A. fumigatus.

Studies in pre-clinical models of allergic disease support the view that CRTH2 plays a central role in leucocyte recruitment, $30,54-57$  AHR<sup>57</sup> and the production of cytokines,<sup>58,59</sup> mucus<sup>54</sup> and IgE.<sup>30,58,60</sup> Independently, it has been shown that  $PGD<sub>2</sub>$  promotes chemotaxis of eosinophils through a receptor unrelated to DP1 and this was designated CRTH2.<sup>20</sup> The ability of PGD<sub>2</sub> to promote eosinophil accumulation in the airways of pre-clinical species is mimicked by selective CRTH2 agonists but not DP1 agonists<sup>28,29</sup> and is inhibited by the CRTH2 antagonist.<sup>28</sup> The effect of the mast cell supernatants on migration of T helper type 2 lymphocytes was blocked by a CRTH2 antagonist, $61$  as was the effect of mast cell supernatants on activation of eosinophils (S.L. Gyles, L. Xue, E.R. Townsend , F. Wettey, R. Pettipher, unpublished observations). Taken together, these data suggest that mast cell-dependent activation of eosinophils and T helper type 2 cells is mediated by  $PGD<sub>2</sub>$  acting on CRTH2.

The observations linking CRTH2 to the development of allergic inflammation has spurred interest in identifying more potent, selective and orally bioavailable antagonists of this receptor for treating asthma and related disorders. A number of chemical series have been described that antagonize CRTH2, including tetrahydroquinolone derivatives, carbazole derivatives, indole acetic acids, azaindole-3-acetic acids, phenoxyacetic acids, phenylacetic acids, thiazoleacetic acids, 3-indolyl sultams, and other series as reviewed in detail by Ulven and Kostenis.<sup>62</sup> OC000459, an indole-1-acetic acid derivative that is an orally active CRTH2 antagonist, which, in proof-of-concept phase IIb clinical trials has been shown to be highly potent and selective in inhibiting  $PGD_2$ -mediated activation of T helper type 2 cells and eosinophils and is an ideal tool to explore the clinical utility of CRTH2 antagonism.<sup>63</sup> The clinical efficacy findings with OC000459 are consistent with the results of pre-clinical studies in CRTH2-deficient mice and with small-molecule antagonists where reduction in eosinophil accumulation<sup>54–57</sup> has been observed. It is also effective in inhibiting mast cell-dependent activation of T helper type 2 cells and eosinophils and airway eosinophilia in rats and guinea-pigs. It is concluded that OC000459 shows promise in the treatment of asthma and warrants further investigation in this indication.

These data demonstrate an important role for CRTH2 in the pathogenesis of allergic inflammation. Combined with our observation that exacerbation of allergic airway inflammation and AHR induced by A. fumigatus is reduced in rats treated with OC000459, these findings suggest that the CRTH2 antagonist may find utility in the prevention or treatment of fungal exacerbations in asthmatic subjects. It is tempting to speculate that CRTH2 receptor antagonism might be useful as a preventive or therapeutic agent in the exacerbations of asthma. Although CRTH2 receptor antagonists have been found to be effective to suppress inflammation, CRTH2-deficient mice developed more severe inflammation of the airways compared with wild-type mice, $64$  which makes the role of the CRTH2 receptor on the pathogenesis of allergic inflammation controversial. The mechanisms are still not well understood and await further investigation.

One limitation of the model is that we did not directly measure bronchoconstriction, but we assessed changes in ventilation by unrestrained whole body plethysmography. Penh (Enhanced pause), which derived from unrestrained whole body plethysmography, is being widely used as a method to study bronchial responsiveness in various animal models of lung disease.<sup>65–72</sup> This technique, and particularly the use of Penh as an index of airway resistance, has been heavily criticized recently.<sup>73–75</sup> Whether an increased Penh is a consequence of changes in the mechanical properties of the respiratory system has not been investigated thoroughly; nevertheless, there is little doubt that increases in Penh do reflect alterations in ventilatory pattern that are compatible with airway obstruction. Other observations in our laboratory support correlation between the Penh and pulmonary resistance  $(R_L - lung resistance, a well recognized pulmonary func$ tion parameter; Figs S1 and S2). So it is very difficult to interpret the correlation between airway obstructive character and Penh. Independently of the physiological meaning of Penh, we infer that some degree of correlation between Penh and bronchoconstriction is expected. This index was indeed measuring changes in the pulmonary function, at least in our experimental conditions. However, whether Penh can be used as a surrogate for  $R_L$  is unclear and requires further research.

Our study showed that prolonged exposure of OVAsensitized and OVA-challenged rats to A. fumigatus spores resulted in enhanced levels of pulmonary eosinophils and bronchial hyper-responsiveness, pharmacological blockade of CRTH2 successfully suppressed pulmonary eosinophilia and AHR. Although eosinophils tend to parallel AHR, they did not appear to be the definitive causative process in this response. It is generally accepted that airway inflammation contributes to the presence and severity of AHR, but the association between allergen-induced pulmonary eosinophilia and the development of AHR has been, at best, a collection of confusing and often contradictory observations. A dissociation between eosinophil numbers and AHR has been shown in guinea-pigs, $^{76}$ 

Brown Norway rats<sup>77</sup> and mice.<sup>78</sup> Clinical studies in human allergic asthma have also shown dissociation between airway inflammation, AHR and the late asthmatic reaction.<sup>79</sup> These studies are not definitive; however, they do cast a shadow on the (cellular) inflammation AHR link. Some researchers point out that the lack of symptom improvement in asthma patients after administration of antibodies to interleukin-5 exemplifies the ambiguous character of clinical studies that attempt to ablate eosinophils. Mouse models purporting to ablate eosinophils are also ambiguous, as they either do not completely eliminate pulmonary eosinophils or they elicit the loss of eosinophils by mechanisms that do not differentiate between effects on eosinophils and other potentially important cellular targets. In each of these studies,  $76-78$  eosinophil numbers in the experimental mice are low but elevated relative to control animals, the small numbers of eosinophils present may actually be sufficient to elicit AHR. And measurements of lung function after OVA sensitization/aerosol challenge of eosinophil-deficient mice (a transgenic line of mice that are specifically devoid of eosinophils but otherwise have a full complement of hematopoietically derived cells, PHIL mice) showed that methacholine-induced AHR was dependent on the presence of eosinophils.<sup>36</sup> Therefore, to identify the contributions of inflammation to AHR, it is important to consider the many aspects of this process, including the location of the cells (i.e. lumen versus bronchial wall) and cell type as well as the many mediators associated with the injury to the airway. The present data are not sufficient to make a definitive conclusion, cellular mechanisms involved in the development of AHR need further study.

In conclusion, OVA-induced inflammation led to an increased susceptibility of the airways to A. fumigatus and increased their likelihood to develop eosinophilic inflammation and bronchial hyper-responsiveness via the activation of the  $PGD<sub>2</sub>/CRTH2$  pathway. The responsiveness to A. fumigatus that we observed appears enhanced in the inflamed airways, because (i) they have a higher capacity to produce  $PGD<sub>2</sub>$  in response to A. fumigatus and (ii) eosinophils are more likely to be recruited into the inflamed airways in response to  $PGD<sub>2</sub>$  through a CRTH2 receptor. The immunomodulatory effects of A. fumigatus and the relationship between allergen dosage and this effect pose interesting problems for future study.

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#### **Disclosures**

The authors have no conflicts of interest to disclose.

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## Supporting Information

Additional Supporting Information may be found in the online version of this article:

Figure S1. Airway responsiveness measured by barometric plethysmography in spontaneously breathing animals (Penh) (A) and plethysmography in mechanically ventilated animals (RL) (B).

Figure S2. Comparison of RL and Penh in Wistar rats challenged with methacholine (MCh) (0.025, 0.05, 0.1, 0.2, 0.4 mg/kg, i.v.). Data are presented as % increase from base level. RL and Penh are correlative measurements of airway reactivity ( $r^2 = 0.842$ ). Solid line: regression line.