

Epstein-Barr Virus–induced Gene 2 Mediates Allergen-induced Leukocyte Migration into Airways

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Abstract

Rationale: Leukocyte recruitment to sites of allergic inflammation depends on the local production of priming cytokines, chemokines, and potentially other mediators. Previously, we showed that eosinophils (Eos) express numerous orphan G-protein–coupled receptors, including Epstein-Barr virus–induced gene 2 (*EBI2*). Despite its contribution to inflammatory diseases, the role of *EBI2* in pulmonary eosinophilia is unknown.

Objectives: To determine whether oxysterol ligands for *EBI2* are increased in asthma exacerbation, and if or how they promote Eos pulmonary migration.

Methods: *EBI2* ligands and pulmonary eosinophilia were measured in the bronchoalveolar lavage fluid from patients with mild asthma 48 hours after acute allergen challenge. *In vitro*, the ability of *EBI2* ligands alone or in combination with IL-5 priming to induce the migration of human blood Eos was assessed.

Measurements and Main Results: *EBI2* was constitutively and stably expressed in peripheral blood Eos. Eos treated with the *EBI2* ligands showed significantly increased transwell migration that was enhanced by priming with physiologic doses of IL-5. Migration was suppressed by inhibitors of the prolyl isomerase Pin1 or extracellular signal–regulated kinases (ERK) 1/2 or by pertussis toxin. *EBI2* signaling activated Pin1 isomerase activity through a cascade that was sensitive to ERK inhibitors and pertussis toxin. The concentration of *EBI2* ligands was significantly increased in the bronchoalveolar lavage fluid 48 hours after segmental allergen challenge and strongly

correlated with the increased numbers of Eos, lymphocytes, and neutrophils in the airways.

Conclusions: Oxysterols are increased in inflamed airways after allergen challenge and, through G-protein subunit α , ERK, and Pin1 signaling, likely participate in the regulation of Eos migration into the lung in people with asthma.

Keywords: asthma; eosinophils; migration; *EBI2*; Pin1

At a Glance Commentary

Scientific Knowledge on the Subject: Inhibition or ablation of chemokine or IL-5 function/expression has not eliminated eosinophilic inflammation in asthma. These data suggest the existence of additional, biologically relevant proinflammatory mediators that contribute to allergic inflammation in humans. These mediators could be targeted to further reduce inflammation and pulmonary pathology.

What This Study Adds to the Field: In this study, we found that segmental allergen challenge increases the level of airway oxysterol Epstein-Barr virus–induced gene 2 (*EBI2*) ligands, which was strongly correlated with leukocyte infiltration. *In vitro*, *EBI2* ligands induced eosinophil migration through a $G\alpha(i)$ -, extracellular signal–regulated kinase-, and Pin1-dependent process. Thus, the oxysterol–*EBI2* signaling is likely another mechanism for leukocyte and eosinophil recruitment to allergic airways, and *EBI2* antagonists combined with Pin1 inhibitors may have therapeutic potential.

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In patients with asthma, inhaled allergen triggers eosinophil (Eos) infiltration into the bronchial mucosa and airway lumen from the peripheral blood (1). This recruitment occurs in direct response to the pulmonary release of a variety of lipid mediators, chemoattractants, and cytokines. Eotaxins 1–3; regulated upon activation, normal T-cell expressed and secreted (RANTES); macrophage inflammatory protein (MIP)-1 α ; and monocyte-chemotactic-protein-3 are among the best-studied and most potent Eos attractants and are produced by pulmonary epithelial and endothelial cells soon after contact with allergen. Once secreted, they interact with CCR3 on the surface of Eos, triggering migration and the release of reactive oxygen species and cationic proteins (degranulation), escalating epithelial cell damage, acute airway inflammation, and pathology.

Although clearly important in rodent models, interference with chemokine function has yielded conflicting results in humans with asthma. For example, whereas genomic knockout of *CCR3* abrogated experimental asthma in rodents (2), treatment of patients with *CCR3* inhibitors only reduced pulmonary Eos accumulation by approximately 40% (3, 4). This may reflect the loss of surface *CCR3* expression by airway Eos, possibly after exposure to IL-3 and RANTES (5–8). Similarly, polymorphisms in *CCR3* were not associated with asthma susceptibility or eosinophilia (9). Moreover, although anti-IL-5 treatment significantly reduced the number of blood and sputum Eos, those effects were only observed in a subgroup of patients with steroid-refractory asthma (10, 11). These observations suggest that other mediators, in addition to the known chemokine–cytokine/receptor pairs, contribute to allergic inflammation in humans with asthma.

We have previously shown that peripheral blood Eos express numerous orphan G-protein-coupled receptors (GPRs), including Epstein-Barr virus-induced gene 2 (*EBI2*; also known as *GPR183*) (12). *EBI2*, a seven-transmembrane GPR, was initially implicated as a guidance receptor to position B cells within lymph nodes and spleen (13). Polymorphisms or knock-out of *EBI2* impairs the positioning of activated B cells and was associated with inflammatory autoimmune diseases (14, 15). Subsequent studies demonstrated that the receptor was also expressed on dendritic cells, neutrophils, T cells, and astrocytes after immune challenge (16–18).

EBI2 binds to a family of 7-, 25-, and/or 27-hydroxycholesterols (HCs) (oxysterols) that are produced and released by inflamed tissue, including the lung (19). The endogenous ligands for *EBI2* are oxidized derivatives of cholesterol and linked to multiple physiologic processes, such as fat metabolism, bile acid synthesis, and lipid transport. On ligand engagement, *EBI2* typically signals via inhibitory G-protein subunit α ($G\alpha[i]$) to induce intracellular calcium mobilization, mitogen-activated protein kinase (MAPK) activation, and cell proliferation (20, 21), but can also signal via G-protein-independent pathways, such as β -arrestin (17, 22). Although oxysterol-dependent activation of the liver X receptor has been linked to the acquired immune response (23), the aggregate effect of *EBI2* signaling also contributes to cell migration.

Pin1, a *cis-trans* peptidyl prolyl-isomerase, plays a key role in the regulation of innate and adaptive immunity and the expression and signaling of cytokines (24). *Pin1* was originally implicated in cell proliferation in part through control of cyclin D1 phosphorylation and stability (25). *Pin1* specifically binds to and isomerizes pSer/pThr-Pro peptide bonds, altering target protein conformation, location, levels, and/or function. Previously, we have shown that pharmacologic blockade or genomic deletion of *Pin1* prevents the development of experimental asthma in rodents (26, 27).

In the present study, we tested the hypothesis that *EBI2* directs Eos migration through a signaling cascade that requires *Pin1*. We show that segmental allergen challenge increases the level of *EBI2* ligands (*EBI2*-Ls) in the airway that was strongly correlated with leukocyte infiltration into lung. *In vitro*, *EBI2*-Ls induced Eos migration through a $G\alpha(i)$ -, extracellular signal-regulated kinase (ERK)-, and *Pin1*-dependent process. Therefore, the oxysterol-*EBI2* signaling is likely another mechanism for leukocyte and Eos recruitment to allergic airways, and *EBI2* antagonists combined with *Pin1* inhibitors may have therapeutic potential.

Methods

Research Volunteers for the Bronchoalveolar Lavage Study

Bronchoscopy with bronchoalveolar lavage (BAL) was approved by the University

of Wisconsin-Madison Center for Health Sciences Human Subjects Committee and each participant provided written informed consent. Subjects had a history of mild asthma, positive skin prick test, a reversibility to albuterol (180 μ g) of greater than 12%, and/or airway hyperresponsiveness to methacholine (the provocative concentration of methacholine producing a 20% fall in FEV₁ was <8 mg/ml). Subjects were nonsmokers, were not receiving asthma controller medications, and did not have evidence of a respiratory infection or asthma exacerbation within the previous 4 weeks. Segmental bronchial provocation with allergen (SBP-Ag) and preparation of BAL fluid was performed as described previously (28) and with a brief summary in the online supplement.

Detection of Oxysterols in BAL Fluid by Mass Spectrometry

Detailed methods have previously been described (29). Crude sterols in BAL fluids were extracted with dichloromethane/methanol (1:1) and 10N KOH before isolation using solid-phase extraction columns. Quantitative analysis of oxysterols was performed using a tertiary Shimadzu LC-20XR HPLC system (Shimadzu Corporation, Kyoto, Japan) equipped with an ultrasonic degasser, column oven, and autosampler. Oxysterols were resolved with a binary solvent gradient after injection of 30 μ l extract. Integration of areas under elution curves was done using ChemStation software (CSS Analytical Co Inc., Shawnee, KS). Quantitative values were calculated using isotope dilution and single-point calibration through a relative response factor calculation. The relative response factor standard was run at the beginning, middle, and end of each sample set; typically, 54 samples per set.

Cell Migration Assay

CytoSelect 24-Well Cell Migration Assay kit (Cell Biolabs, Inc., San Diego, CA) was used. Cells were seeded in the top of the insert in a medium with or without 10% fetal bovine serum, while chemoattractants were placed in the well below. Inhibitors and IL-5 (for priming) were added directly to the cell suspension before stimulation with chemoattractants. Cells migrated through the pores (5 μ m) toward the chemoattractants and were lysed and

stained with CyQuant GR dye (from the assay kit) before fluorescence measurement in a plate reader at 480 nm/520 nm.

Pin1 Activity Assay

Activity was measured as described previously (30) with slight modifications.

Statistics

All *P* values were calculated using GraphPad Prism 7.01 (GraphPad Software, La Jolla, CA). For comparing multiple groups, we performed nonparametric

one-way analysis of variance followed by *post hoc* Tukey test. Data are represented as mean \pm SD; *P* less than 0.05 was considered statistically significant.

Results

EBI2 Ligands Are Increased in Human Asthma

EBI2 oxysterol ligands are up-regulated in a variety of human inflammatory diseases, including atherosclerosis, autoimmunity,

diabetes, and cancer (20, 31). Thus, we hypothesized that oxysterol production would be elevated in the BAL fluid of patients with allergic pulmonary inflammation. Therefore, seven patients with mild asthma (see Table E1 in the online supplement) were subjected to SBP-Ag (see online supplement). After 48 hours, BAL fluid was collected, and cellular infiltrate before and after provocation was analyzed. Consistent with previous observations (32), total BAL cell counts increased by approximately fourfold from a

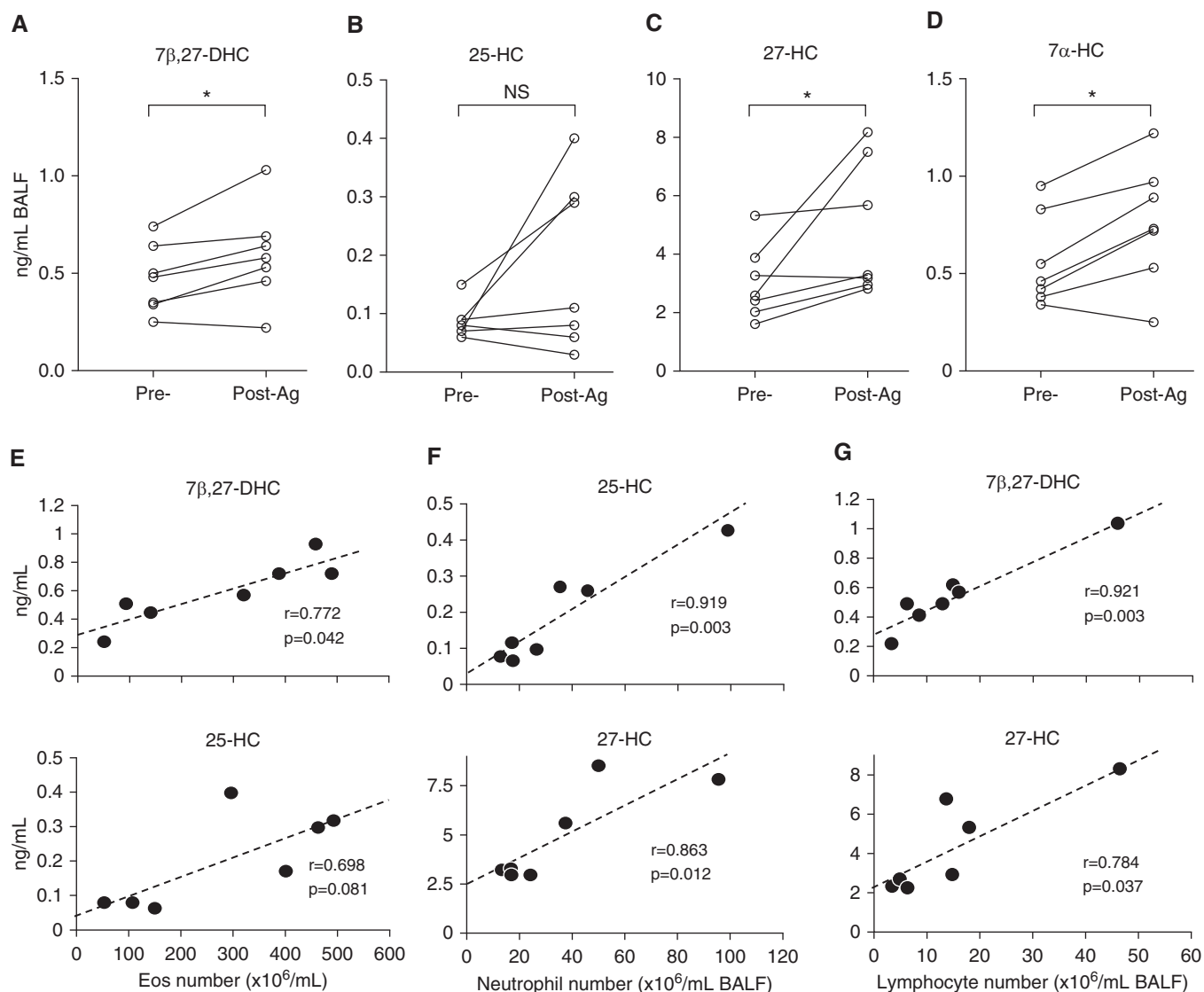


Figure 1. Epstein-Barr virus–induced gene 2 ligands are increased in bronchoalveolar lavage fluid (BALF) after segmental allergen challenge. (A–D) BALF was collected before and 48 hours after segmental allergen challenge of patients (*n* = 7) with mild asthma. After removal of cells, the supernatants were subjected to mass spectrometry. (E–G) Pearson correlation coefficients (*r*) were measured to evaluate the linear relationship between ligand concentration and cell numbers in BAL from after segmental allergen challenge. **P* < 0.05 by paired Student's *t* test. Ag = allergen challenge; DHC = dihydroxycholesterol; Eos = eosinophil; HC = hydroxycholesterol; NS = nonsignificant.

baseline of $126 \pm 59 \times 10^4$ to $516 \pm 200 \times 10^4$ per milliliter of BAL fluid (see Table E2). Absolute Eos numbers in BAL showed the greatest change and were increased by approximately 40-fold. We then analyzed BAL fluid from these patients for EB12-Ls (17, 21) by mass spectrometry (29, 33). Of the eight known EB12-Ls, 25-HC showed a trend to increase after SBP, whereas 7 β ,27-dihydroxycholesterol (7 β ,27-DHC), 27-HC, and 7 α -HC were significantly increased (Figures 1A–1D). Interestingly, the BAL concentration of 7 β ,27-DHC, 25-HC, and 27-HC was strongly positively correlated with inflammatory cell counts (Eos, neutrophils, and lymphocytes) with 7 β ,27-DHC correlating best with Eos and lymphocyte counts and 25-HC correlating best with neutrophilia (Table 1, Figures 1E–1G). These data show that multiple EB12-Ls are increased in the BAL after acute allergic stimulation and likely participate in Eos, neutrophil, and lymphocyte recruitment/retention in allergic airway disease.

EB12 Is Expressed by Eos and Directs Migration

These data suggested that EB12 and its ligands play a role in allergic airway inflammation. EB12 (GPR183) is highly expressed in lung (34) and most immune cells after activation (16, 17), although we are unaware of any reports regarding its expression by Eos. RNA Seq revealed 24 GPRs, including EB12, were expressed above baseline by human peripheral blood Eos (see Table E3) (12). Western blotting confirmed that EB12 was present in resting Eos or those treated with the prosurvival cytokine IL-5 or EB12-L (Figure 2A and data not shown). Therefore, EB12–oxysterol signaling apparatus is present in Eos and based on our *in vivo* data above, likely

functional and physiologically relevant for the migration of human Eos into allergen-challenged lung.

To verify this, we performed cell migration assays using transwell chambers. Peripheral blood Eos was added to the upper chamber and EB12-L was added to the lower chamber. After incubation for 18 hours, migrated cells were lysed and quantitated (see METHODS). In the presence of 10% fetal bovine serum, the EB12-L 7 α ,25-DHC (100 nM = 40.3 ng/ml) induced significant migration into the lower chamber (Figure 2B). We observed similar results after migration for 3 hours, indicating that most of cell migration occurs in first 2–3 hours. The difference in lower chamber cell counts was not caused by variation in cell viability because at 96 hours of culture, daily supplementation with 7 α ,25-DHC (10–100 nM) had no effect on Eos survival either alone or in combination with IL-5 (5–150 pM) (Figure 2C and data not shown). Because BAL IL-5 increases 1.7-fold (5 pM = 13 pg/ml) in patients with asthma (35) and also acts as an independent Eos chemoattractant at high concentration (>150 pM) (36, 37), we primed the cells with different doses of IL-5 for 10 minutes before the induction of migration for 3 hours. High-dose IL-5 (20–100 pM) had similar chemoattractant activity as the EB12-L alone (100 nM) ($EC_{50} = 2$ nM for the receptor activation) or eotaxin (50 ng/ml) (Figure 2D). Interestingly, the EB12-induced migration was augmented when cells were primed with pathophysiologically relevant doses of IL-5 (5–10 pM).

We have obtained similar data with other EB12-Ls, including 7 β ,27-DHC (6 μ M) and 25-HC (152 μ M) (Figures 1 and 2E) but at higher doses than their EC_{50} (17, 21). These results are consistent with

postchallenge *in vivo* events in which Eos would be expected to encounter both elevated IL-5 and EB12-L in the blood and airway after pulmonary allergen exposure. To understand the specificity of the ligands, we isolated human peripheral blood mononuclear cells and neutrophils, and tested their response to 7 α ,25-DHC, 7 β ,27-DHC, and 27-HC (Figure 1). All freshly isolated, resting cells had poor response to the ligands. However, prior activation of T cells with anti-CD3/CD28 (see Figure E1) or phorbol myristate acetate/ionomycin (data not shown) increased cell migration to 7 α ,25-DHC and 7 β ,27-DHC, as did neutrophils primed with granulocyte-macrophage colony-stimulating factor (GM-CSF) (see Figure E1), suggesting EB12 also regulates the trafficking of other inflammatory cell types (Figures 1F and 1G; see Table E2) in addition to Eos.

G α (i), cAMP, and ERK Signaling Are Required for EB12-directed Eos Migration

EB12 signaling can drive MAPKs activation through G α (i)-dependent or –independent mechanisms (18, 20). G α (i) blocks G-protein α stimulatory (G α [s]) function, decreasing the activity of adenylyl cyclases (ACs) 1, 5, and 6 (38). G β /G γ subunits have the opposite effect and can directly stimulate ACs 2, 4, and 7. Thus, the overall intracellular cAMP levels are an aggregate reflection of the levels and activity of G α (i), G α (s), and G β /G γ . To identify which GPR pathway EB12 signaling used (39), we used purified Eos, which were preincubated with either pertussis toxin (PT) (inhibitor of G α [i]) (40) or the pan-AC inhibitor MDL-12330A (MDL) (41, 42). Interestingly, both inhibitors had no effect on Eos only treated with IL-5 but suppressed cell migration induced by IL-5-priming and

Table 1. Pearson Correlation Coefficients and *P* Values for Comparisons of BAL Cells and EB12 Oxysterol Ligands (n = 7)

Ligands	Total BAL Cells		Eosinophils		Neutrophils		Lymphocytes		Macrophages	
	<i>r</i>	<i>P</i> Value	<i>r</i>	<i>P</i> Value	<i>r</i>	<i>P</i> Value	<i>r</i>	<i>P</i> Value	<i>r</i>	<i>P</i> Value
7 β ,27-DHC	0.689	0.087	0.772	0.042	0.225	0.628	0.921	0.003	–0.133	0.777
25-HC	0.649	0.115	0.698	0.081	0.919	0.003	0.648	0.116	–0.333	0.466
27-HC	0.667	0.102	0.681	0.092	0.863	0.012	0.784	0.037	–0.257	0.578
7 α -HC	0.342	0.453	0.430	0.335	–0.250	0.589	0.523	0.229	–0.081	0.863

Definition of abbreviations: BAL = bronchoalveolar lavage; DHC = dihydroxycholesterol; EB12 = Epstein-Barr virus–induced gene 2; HC = hydroxycholesterol. *r*, Pearson correlation coefficient; cells are per milliliter of BAL fluid recovered; neutrophils and lymphocytes (but not eosinophils) are logarithmically transformed.

Bold indicates *P* < 0.05.

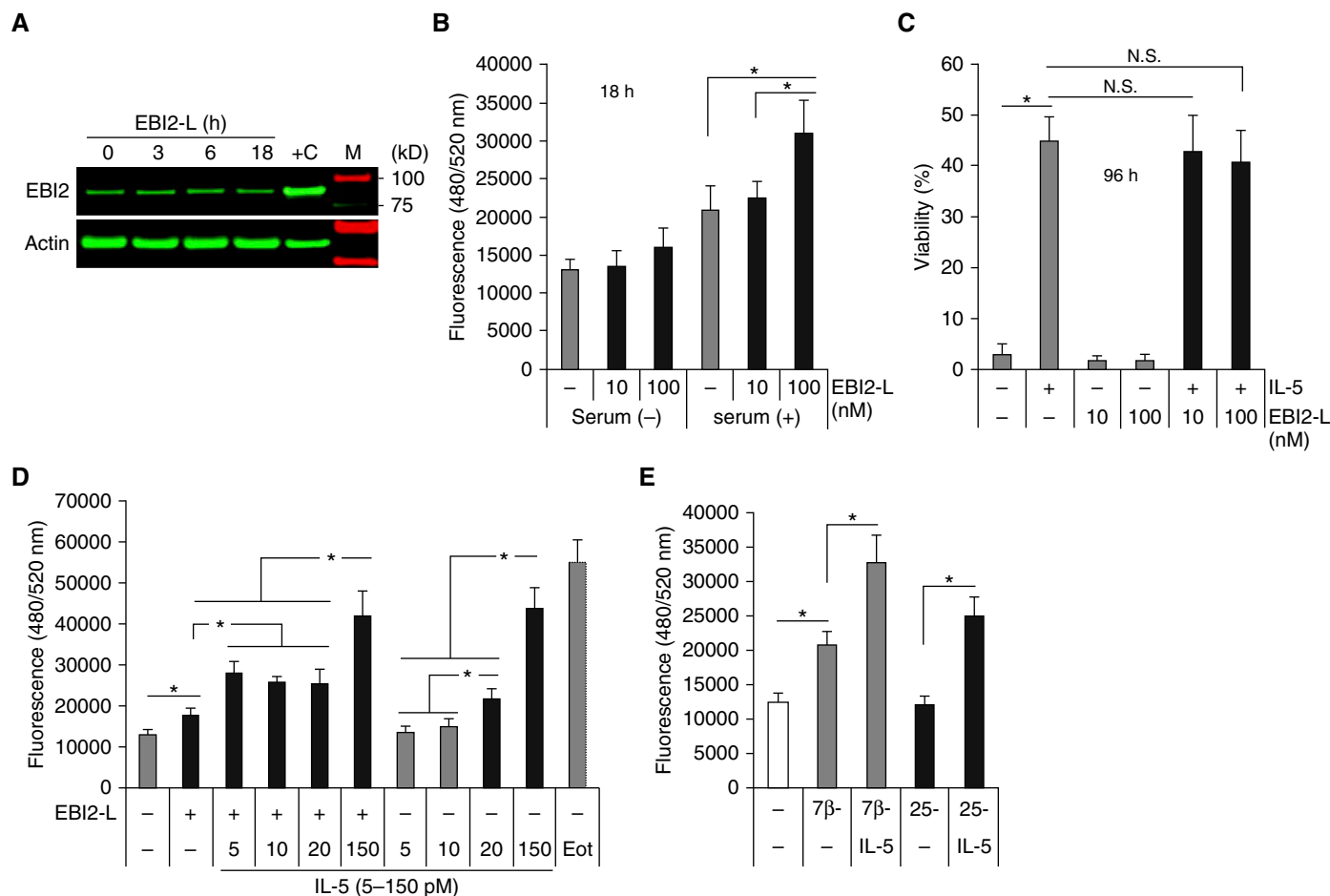


Figure 2. Epstein-Barr virus–induced gene 2 (*EBI2*) is expressed by eosinophil (Eos) and directs migration. (A) Freshly purified blood Eos were treated with EBI2 ligand (EBI2-L) (7α , 25-dihydroxycholesterol [DHC]) (100 nM = 40.3 ng/ml) for the indicated times. Whole-cell lysates were subjected to immunoblots with anti-EBI2 and β -actin as shown. +C: Ramos cell lysate served as positive control. (B) Cell migration assay was performed (see METHODS) in the presence (+) or absence (–) of 10% fetal bovine serum and EBI2-L. Migrated cells were lysed and stained with DNA dye. (C) Cells were cultured for 96 hours in the presence or absence of EBI2-L (10–100 nM) and IL-5 (150 pM) in the medium containing 10% fetal bovine serum before viability assay. (D) Cells were primed in the presence of different doses of IL-5 (5–150 pM) for 10 minutes before induction of migration with EBI2-L (100 nM) for 3 hours. Eotaxin (50 ng/ml) used as a positive control. (E) Cells were primed with IL-5 (10 pM) before induction of migration with other EBI2-L (6 μ M of 7β , 27-DHC, or 152 μ M of 25-DHC) for 3 hours. Data are expressed as mean \pm SD. * P < 0.05. The data are from at least four independent experiments. Eot = eotaxin; N.S. = nonsignificant.

EBI2-L (Figures 3A and 3B). These data suggest that EBI2 signaling is dually regulated by $G\alpha(i)$ and AC, presumably with AC downstream of $G\alpha(i)$ independent of IL-5 signaling.

EBI2 engagement is known to activate MAPK pathways in diverse cell types, including transformed cells and glia (20). Therefore, before transwell migration assays, Eos were preincubated with a variety of kinase inhibitors. Although inhibitors for c-Jun N-terminal kinases, p38, phosphatidylinositol-3 kinases (PI3K), protein kinase C- α/β 2, and ribosomal S6 kinase had no effect, the MAPK/ERK kinase (MEK) 2 inhibitor (PD98059)

reproducibly blocked cell migration induced by IL-5 and EBI2-L (Figure 4A). Western blots and flow cytometry analysis (Figure 4B) revealed the activation of ERK1/2 after EBI2 signaling. Therefore, EBI2 triggers a cascade involving $G\alpha(i)$, AC, and ERK, all of which are required for Eos migration.

Pin1 Is Required for EBI2-directed Eos Migration

We have shown that pharmacologic blockade of or genetic deletion of Pin1 significantly reduced Eos inflammation and airway fibrosis in experimental rodent models of asthma (26, 27). Although

some of the reductions reflected increased apoptosis of pulmonary Eos, these changes did not account for all of the observed differences. Thus, we hypothesized that Pin1 may also play a role in human Eos migration induced by EBI2-L. To test this possibility, we incubated Eos with a specific Pin1 inhibitor before measuring transwell cell migration. As shown (Figure 5A), dominant negative WW peptide of Pin1 alone had no effect on random cell migration; however, it significantly suppressed EBI2-L–induced and IL-5–enhanced cell migration at multiple time points (3 h and 24 h). Control peptides (W34A nonfunctional

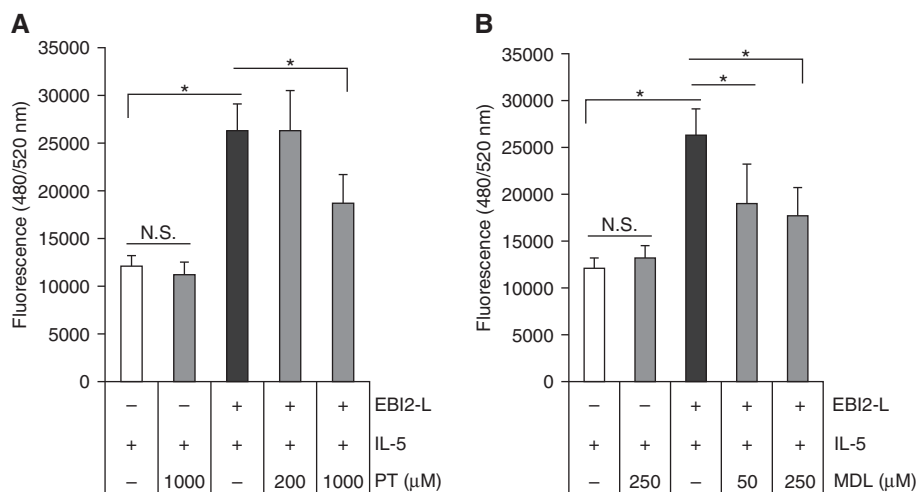


Figure 3. G-protein α and cAMP mediate Epstein-Barr virus-induced gene 2 (*EBI2*) signaling. (A and B) Eosinophils were pretreated for 10 minutes with pertussis toxin or adenylyl cyclase inhibitor (MDL-12330A) before priming with IL-5 (10 pM, 10 min) and subsequent induction of migration with EBI2 ligand (100 nM) for 3 hours. * $P < 0.05$. The data are from at least four independent experiments. EBI2-L = Epstein-Barr-virus-induced gene 2 ligand; MDL = MDL-12330A; N.S. = nonsignificant; PT = pertussis toxin.

mutant peptide) unable to compete with endogenous Pin1 also had no effect on migration (Figure 5B).

These data suggested that Pin1 isomerase activity might be regulated by EBI2 signaling. Thus, Eos were primed and/or treated with EBI2-L before cell lysis and Pin1 activity assay. As shown (Figure 5C), 10-minute exposure to the ligand markedly

increased Pin1 *cis-trans* isomerase activity that was further enhanced when cells were primed with low dose IL-5. The specificity of the assay for Pin1 was confirmed by the inhibitory effect of dominant negative WW peptide (Figure 5C). Collectively, these data suggest that Pin1 isomerase activity is required for and regulates cell migration directed by the ligands.

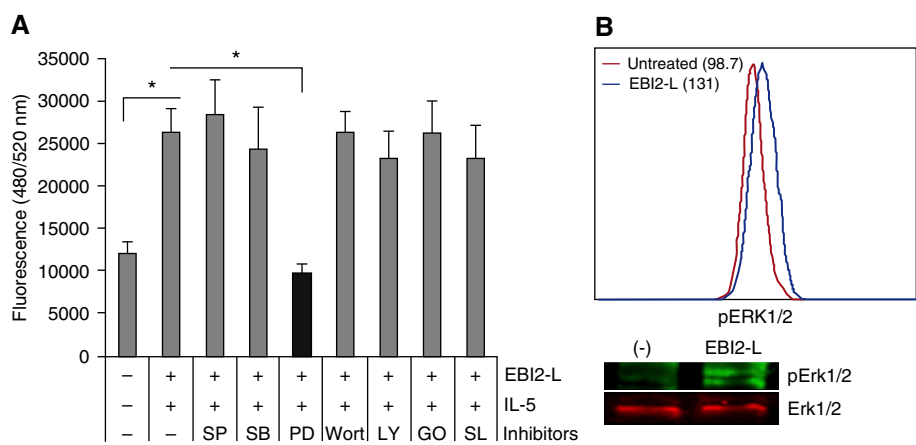


Figure 4. Extracellular signal-regulated kinase (ERK) mitogen-activated protein kinase (MAPK) mediates Epstein-Barr virus-induced gene 2 (*EBI2*) signaling. (A) Eosinophils were pretreated with kinase inhibitors (5 μ M SP600125 [SP] for c-Jun N-terminal kinases, 5 μ M SB203580 [SB] for p38 MAPK, 50 μ M PD98059 [PD] for ERK MAPK, 200 nM wortmannin [Wort] for phosphatidylinositol-3 kinases, 10 μ M LY294002 [LY] for phosphatidylinositol-3 kinases, 10 nM Gö6976 [GO] for protein kinase C- α/β , 100 nM SL0101-1 [SL] for p90 ribosomal S6 kinase) before priming with IL-5 (10 pM) and subsequent induction of migration with EBI2 ligand (EBI2-L) (100 nM) for 3 hours. (B) Cells were treated with EBI2-L alone for 10 minutes and subjected to flow cytometry and immunoblots for the activation of ERK1/2. * $P < 0.05$. The data are from at least four independent experiments.

If Pin1 was required for directed Eos migration, we reasoned that other inhibitors of EBI2 signaling, namely PT and MDL, would also negatively impact Pin1 activity. Thus, cells were pretreated with PT or MDL before IL-5 and EBI2-L. As shown (Figure 5D), although PT and PD98059 (MEK/ERK inhibitor) (data not shown) significantly attenuated Pin1 activity, MDL had no effect. This suggests that AC signaling does not impact Pin1 and that ERK activation needed for both migration and Pin1 activation must be triggered by $G\alpha(i)$ independently of AC.

Discussion

In this study, we demonstrated that human blood Eos constitutively express EBI2 and its ligands strongly induce cell migration in concert with IL-5 priming. The chemotactic effects of EBI2-Ls were also observed in activated human T cells and neutrophils, although their responses were relatively poor. Relevant to allergic diseases, the endogenous EBI2 oxysterol ligands were significantly increased in the lung after segmental allergen challenge, an accepted experimental surrogate for acute asthma. At the molecular level, EBI2 receptor signaling was mediated via $G\alpha(i)$, cAMP, and ERK1/2 MAPK. Pin1 was also required for Eos migration, functioning downstream of $G\alpha(i)$ and ERK, although the molecular targets regulated by the isomerase that are involved in migration remain to be elucidated. Thus, along with established chemokines, EBI2-Ls may facilitate Eos and other immune cell migration and airway inflammation (see Table E2).

Much evidence implicates C-C chemokines in allergic inflammation. RANTES, eotaxins, MIP-1, and monocyte-chemotactic protein 3/4 are dramatically increased in rodent models and humans with asthma (1, 2). Chemoattractants are released by activated Th2 lymphocytes, macrophages, epithelial cells, and mast cells (1, 2), and show variable specificity and potency on Eos. These molecules bind to and signal through a common receptor, CCR3, and function as powerful chemoattractants, *in vitro*. Despite this, the role of these molecules in the induction of eosinophilic inflammation in patients with asthma remains unclear. For example, BAL Eos showed significantly reduced CCR3 levels after exposure to airway

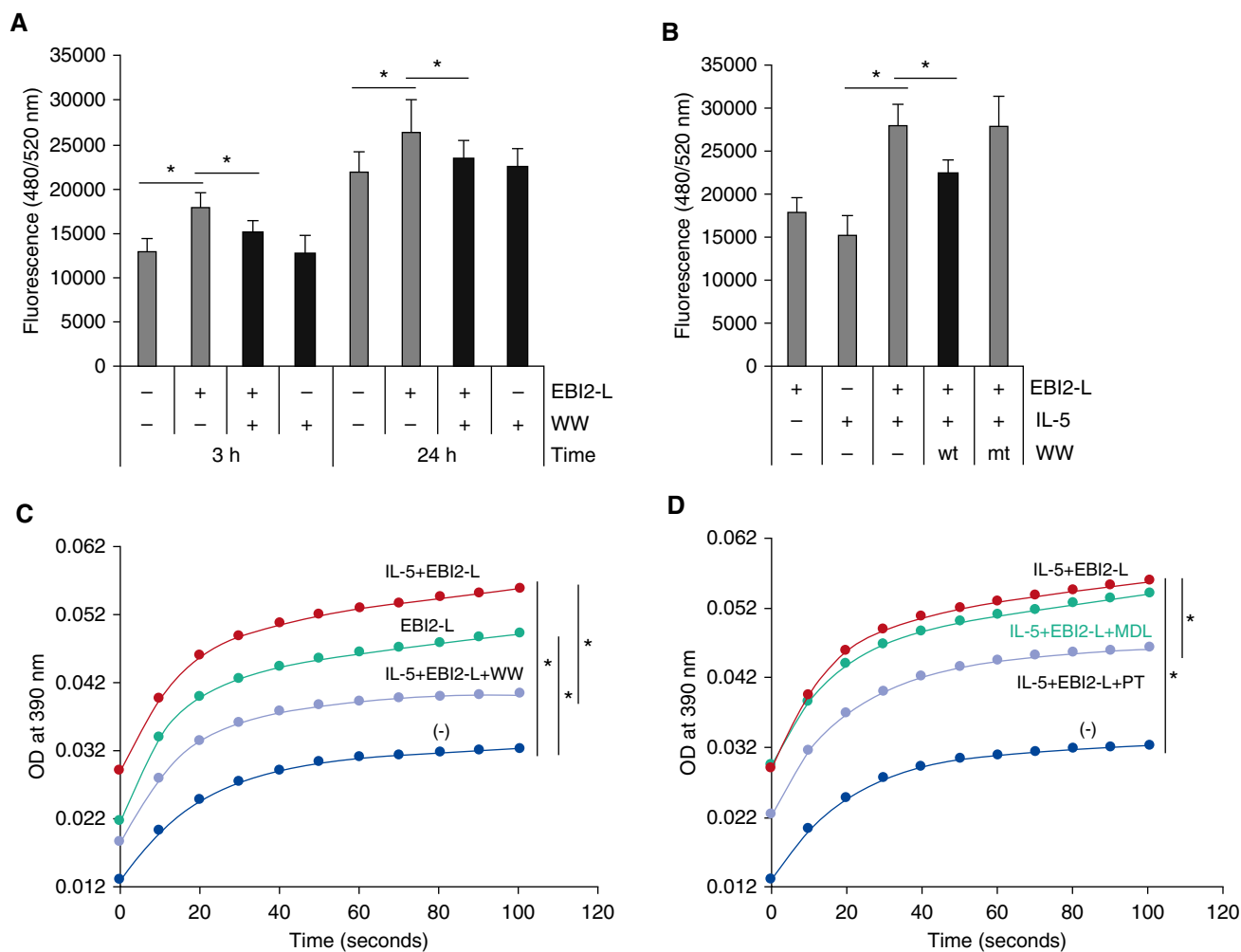


Figure 5. Pin1 is required for Epstein-Barr virus–induced gene 2 (*EBI2*)-directed eosinophil migration. (A) Eosinophils were pretreated with dominant negative TAT-WW of Pin1 peptide (WW, 10 nM) for 10 minutes before induction of migration with *EBI2* ligand (*EBI2*-L) (100 nM) alone for the indicated times. (B) Cells were pretreated with TAT-WW (wt) or with its nonfunctional mutant (mt) before priming with IL-5 (10 pM) and subsequent induction of migration with *EBI2* ligand for 3 hours. (C and D) Cells were treated as in B for 10 minutes. Cell lysates were subjected to Pin1 isomerase activity assay as described in METHODS. In D, 1,000 μ M of pertussis toxin and 250 μ M of MDL were used. * $P < 0.05$. The data are from at least four independent experiments. MDL = MDL-12330A; OD = optical density; PT = pertussis toxin.

IL-3/RANTES (5–7), attenuating CCR3-mediated signaling. Although the CCR3 agonist MIP-1 is a powerful Eos chemoattractant in mice (43), another agonist (eotaxin-3) shows limited activity in humans (8). Prosurvival cytokines IL-5 and GM-CSF show modest chemotactic activity but very potently prime Eos for enhanced chemotactic responsiveness to suboptimal concentrations of chemokines (44) and oxysterols (Figure 2D). In aggregate these data suggest Eos respond to a complex combination of signals that are only partially predictable based on the cell-surface receptor they engage.

The bioactive oxysterols that were increased after allergen challenge (Figure 1) are potentially a new class of Eos chemoattractants

that also show biologic activity toward other inflammatory cells that express *EBI2*, including neutrophils and lymphocytes (see Table E2 and Figure E1). The most potent physiologic *EBI2*-L is 7 α ,25-DHC (17, 21). The short half-life of these compounds quenches their signaling in normal tissues (36). However, during pathology involving tissue damage, such as atherosclerosis, cataracts, and Alzheimer disease (45), *EBI2*-Ls dramatically increase as we have seen after segmental allergen challenge (Figure 1). Blockade or knock-out (13) of *EBI2*-L–mediated signaling greatly reduced innate and adaptive immune responses (45), processes essential for responses to infectious, allergic, or inflammatory diseases and cancer. These

observations suggest a diverse array of chemotactic signals contribute to immune cell trafficking and activation.

We propose that oxysterols and chemokines are nonredundant signaling molecules because of the differences in their production and metabolism. Oxysterols likely are more short-acting and thus locally relevant because of their rapid production by many cell types and their relatively short half-life. In contrast, chemokines secretion is slower and restricted to fewer cell types but capable of communicating over greater distances (e.g., between lung and bone marrow). Thus, both types of chemoattractants likely function in a spatial and temporal cooperative manner.

It remains to be determined if oxysterol production and/or levels can be used as a biomarker for asthma severity or Eos inflammation during disease exacerbations.

In addition to EBI2, Eos expressed a large number of orphan GPRs (see Table E3) (12). Although some ligand-receptor pairs have been identified, their functional roles remain poorly characterized. We have recently shown that many orphan GPRs, including *EBI2*, are markedly increased in total BAL cells 48 hours after allergen challenge (46), although the cellular source was not definitively determined. These included *GPR35*, *GPR65*, and *GPR77* (see Table E3), all of which are highly expressed by untreated blood Eos and have been associated with asthma and allergic disease (47–49). Eos lacking *GPR65* failed to increase cAMP production or prolong survival after acid treatment (47). *GPR77/C5L2* (coded by *C5AR2* gene in human) knockout mice were protected from eosinophilic inflammation and airway hyperresponsiveness (49). The latter may reflect the function of GPRs on airway smooth muscle cells that interact with integrins (50). Whether *GPR183*-deficient mice are similarly protected from challenge-induced eosinophilic pulmonary inflammation is unknown. Taken together, these data suggest that orphan GPR signaling plays a variety of roles in Eos function and asthma pathogenesis.

The signaling that regulates Eos migration in response to various chemoattractants is complex and includes kinase cascades involving MAPKs (51–53) and PI3Ks (54–56). Other activated kinases during Eos migration include protein kinase C (57), Src (58), and Janus kinase 2 (59). Eotaxin, an extremely potent and Eos-selective chemokine rapidly induced

tyrosine phosphorylation of multiple intracellular proteins (including ERK1/2), calcium mobilization, actin reorganization, and changes in cell shape (51). In addition to recruitment, eotaxin induces Eos degranulation (60). Coimmunoprecipitation studies revealed that eotaxin binding triggered internalization of the GPR CCR3 and intracellular association with Src family kinases (Hck and c-Fgr) (61). Kinetically similar, oxysterol–GPR EBI2 interactions induced the rapid (1–5 min) recruitment of Gα(i), with subsequent cAMP activation, calcium mobilization (19, 20), and ERK1/2 MAPK phosphorylation (Figures 3B and 4). These data suggest that oxysterol-induced signaling uses a distinct, initial mechanism from CCR3-bound chemokines, but one that eventually integrates into the common pathways required for cell migration.

A role for Pin1 in EBI2 signaling was not unexpected given prior data (24, 26, 27). Pin1 is required for enhanced cytokine gene expression by immune cells and fibroblasts in response to tissue injury or damage in asthma, fibrosis, and infection. Pin1 blockade suppressed GM-CSF or IL-5 production and their prosurvival signaling in activated human Eos as well as in rodent model of asthma (26, 27). At a molecular level, ERK MAPK activation was critical for *in vitro* survival cytokine expression and *in vivo* prosurvival signaling in Eos (62–64). The increased level of pulmonary oxysterols (Figure 1) and their activation of ERK1/2 (Figure 4B) via EBI2 provide an additional dimension for defining and understanding of eosinophilic accumulation in the airway of subjects with asthma. Pin1 is required for the EBI2 signaling but not for eotaxin signaling (Chu and colleagues, unpublished data). The

molecular mechanism by which Pin1 selectively modulates oxysterol–EBI2 signaling remains elusive. Potential targets include β-arrestin-1/2, protein kinase A, ERK1/2, regulator of G-protein signaling 14, p21-activated kinases, MAPK/ERK kinase kinase 2, and PI3K, all of which have been implicated in Gα(i) signaling and cytoskeleton reorganization (17, 21, 65–67). Nevertheless, we were unable to coimmunoprecipitate these molecules with Pin1 (not shown) despite Eos activation with high-dose EBI2-L. Future studies using mass spectrometry-based proteomics approach should help to delineate the molecule mechanisms.

In summary, we have shown that oxysterol–EBI2 signaling is relevant to human asthma through a complex signaling cascade involving Gα(i), cAMP, ERK, and Pin1. Oxysterols can be pharmacologically manipulated through drugs directed at rate-limiting enzymes in their synthetic pathway that could provide additional approaches to reduce inflammation in allergic diseases. ■

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