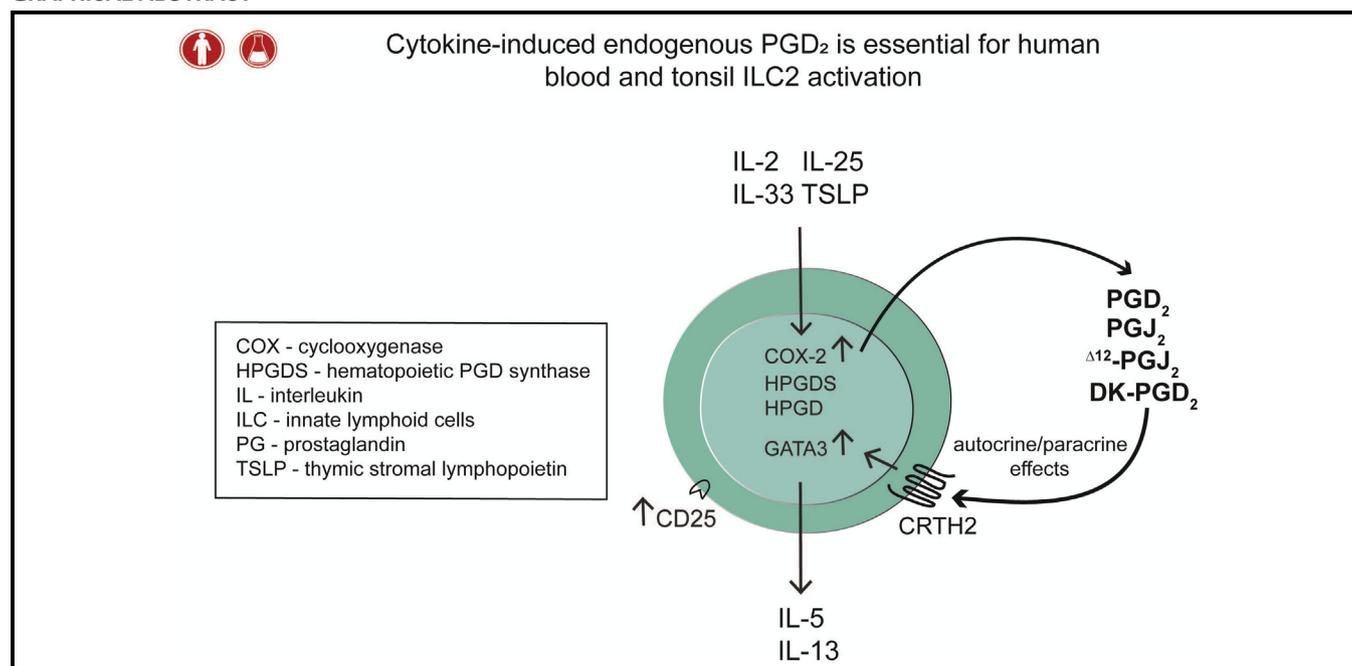


# Cytokine-induced endogenous production of prostaglandin D<sub>2</sub> is essential for human group 2 innate lymphoid cell activation

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## GRAPHICAL ABSTRACT



**Background:** Group 2 innate lymphoid cells (ILC2s) play a key role in the initiation and maintenance of type 2 immune responses. The prostaglandin (PG) D<sub>2</sub>-chemoattractant

receptor-homologous molecule expressed on T<sub>H</sub>2 cells (CRTH2) receptor axis potently induces cytokine production and ILC2 migration.

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**Objective:** We set out to examine PG production in human ILC2s and the implications of such endogenous production on ILC2 function.

**Methods:** The effects of the COX-1/2 inhibitor flurbiprofen, the hematopoietic prostaglandin D<sub>2</sub> synthase (HPGDS) inhibitor KMN698, and the CRTH2 antagonist CAY10471 on human ILC2s were determined by assessing receptor and transcription factor expression, cytokine production, and gene expression with flow cytometry, ELISA, and quantitative RT-PCR, respectively. Concentrations of lipid mediators were measured by using liquid chromatography–tandem mass spectrometry and ELISA.

**Results:** We show that ILC2s constitutively express HPGDS and upregulate COX-2 upon IL-2, IL-25, and IL-33 plus thymic stromal lymphopoietin stimulation. Consequently, PGD<sub>2</sub> and its metabolites can be detected in ILC2 supernatants. We reveal that endogenously produced PGD<sub>2</sub> is essential in cytokine-induced ILC2 activation because blocking of the COX-1/2 or HPGDS enzymes or the CRTH2 receptor abolishes ILC2 responses.

**Conclusion:** PGD<sub>2</sub> produced by ILC2s is, in a paracrine/autocrine manner, essential in cytokine-induced ILC2 activation. Hence we provide the detailed mechanism behind how CRTH2 antagonists represent promising therapeutic tools for allergic diseases by controlling ILC2 function. (J Allergy Clin Immunol 2019;■■■:■■■-■■■.)

**Key words:** Group 2 innate lymphoid cells, prostaglandin D<sub>2</sub>, chemoattractant receptor–homologous molecule expressed on T<sub>H</sub>2 cells, allergy

Group 2 innate lymphoid cells (ILC2s) play important roles in the initiation and amplification of type 2 immune responses.<sup>1</sup> Under inflammatory conditions, ILC2s are stimulated by the innate cytokines IL-33, IL-25, and thymic stromal lymphopoietin (TSLP), which are products of activated epithelial cells and macrophages.<sup>2</sup> Optimal activation of human ILC2s requires a combination of these 3 cytokines, as well as IL-2.<sup>3,4</sup>

Furthermore, ILC2s can be activated by proinflammatory lipid mediators.<sup>5</sup> Leukotriene (LT) D<sub>4</sub>, LTE<sub>4</sub>,<sup>6,7</sup> and prostaglandin (PG) D<sub>2</sub> are potent stimulators of mouse and human ILC2s.<sup>8-10</sup> PGD<sub>2</sub> has been shown to induce IL-5 and IL-13 production, as well as migration of ILC2s.<sup>8,10,11</sup> Human ILC2s were initially identified based on the expression of one of the receptors for PGD<sub>2</sub>.<sup>12</sup> The PGD<sub>2</sub> receptor chemoattractant receptor–homologous molecule expressed on T<sub>H</sub>2 cells (CRTH2) now serves as an important cell-surface marker of human ILC2s. CRTH2 is a Gi protein–coupled receptor, and in type 2 immune cells, such as T<sub>H</sub>2 cells, eosinophils, basophils, and ILC2s, CRTH2 activation increases free intracellular calcium levels.<sup>13</sup> Activation of these cells is crucial in the development of allergic inflammatory processes.

PGD<sub>2</sub>, belonging to the eicosanoid family, is derived from arachidonic acid, which is liberated from membrane phospholipids by phospholipase A<sub>2</sub>. Activation of COX-1, COX-2, or both leads to a 2-step metabolism of arachidonic acid into PGG<sub>2</sub> and subsequently to PGH<sub>2</sub>, which is the substrate for enzymes specialized in generating different PGs. There are 2 types of PGD<sub>2</sub> synthases: hematopoietic prostaglandin D<sub>2</sub>

#### Abbreviations used

CBMC:	Cord blood–derived mast cell
CRTH2:	Chemoattractant receptor–homologous molecule expressed on T <sub>H</sub> 2 cells
DK-PGD <sub>2</sub> :	13, 14-Dihydro-15-keto prostaglandin D <sub>2</sub>
DPI:	Prostaglandin D <sub>2</sub> receptor 1
15d-PGJ <sub>2</sub> :	15-Deoxy-delta-12, 14-PGJ <sub>2</sub>
HPGD:	15-Hydroxyprostaglandin dehydrogenase
HPGDS:	Hematopoietic prostaglandin D <sub>2</sub> synthase
ILC2:	Group 2 innate lymphoid cell
IMDM:	Iscove modified Dulbecco medium
LC-MS/MS:	Liquid chromatography–tandem mass spectrometry
LPGDS:	Lipocalin prostaglandin D <sub>2</sub> synthase
LT:	Leukotriene
NHS:	Normal human serum
PE:	Phycoerythrin
PGD <sub>2</sub> :	Prostaglandin
PPAR-γ:	Peroxisome proliferator–activated receptor γ
RT-qPCR:	Quantitative RT-PCR
TSLP:	Thymic stromal lymphopoietin
TXB <sub>2</sub> :	Thromboxane B <sub>2</sub>

synthase (HPGDS), which is expressed on immune cells,<sup>14-16</sup> and lipocalin prostaglandin D<sub>2</sub> synthase (LPGDS), which is found mainly in nervous system, heart, and adipose tissue.<sup>17,18</sup> Mast cells are the major source of PGD<sub>2</sub>,<sup>19</sup> whereas dendritic cells, macrophages,<sup>20</sup> eosinophils,<sup>21</sup> and T<sub>H</sub>2 cells release smaller amounts of PGD<sub>2</sub>.<sup>22</sup> In the metabolism of PGD<sub>2</sub>, 15-hydroxyprostaglandin dehydrogenase (HPGD) converts PGD<sub>2</sub> into 13, 14-dihydro-15-keto prostaglandin D<sub>2</sub> (DK-PGD<sub>2</sub>),<sup>23,24</sup> a natural selective agonist of CRTH2 receptor.<sup>25</sup> Furthermore, PGD<sub>2</sub> can be nonenzymatically converted into PGJ<sub>2</sub> and its derivatives, metabolites with biological activity.<sup>26</sup>

Because CRTH2 is a key receptor in allergic inflammatory processes and mediates the PGD<sub>2</sub>-induced activation of ILC2s and other type 2 immune cells, the PGD<sub>2</sub>-CRTH2 axis is an emerging therapeutic target to control allergic diseases.<sup>22,27,28</sup> The role of exogenous PGD<sub>2</sub> in ILC2 function has been clearly established; however, it has not been addressed whether endogenously produced PGD<sub>2</sub> plays a role in ILC2 activation.

Our study shows that human ILC2s express HPGDS and that IL-2, IL-25, and IL-33 plus TSLP activation of human ILC2s leads to upregulation of the rate-limiting COX-2 enzyme. This renders ILC2s capable of producing PGD<sub>2</sub> and its metabolites. Endogenously produced PGD<sub>2</sub> is essential in cytokine-induced activation of human ILC2s because a COX-1/2 blocker potently prevents IL-5 and IL-13 production from the cells. Similarly, selective antagonism of CRTH2 on cytokine stimulation inhibits ILC2 activation. Thus we demonstrate that human ILC2s critically depend on endogenously produced PGD<sub>2</sub> for activation by cytokines. Our results have implications for the treatment of allergic inflammation.

## METHODS

A detailed description of the methods and materials used in this study is provided in the [Methods](#) section in this article's Online Repository at [www.jacionline.org](http://www.jacionline.org).

## Isolation and flow cytometric sorting of tonsillar and blood ILC2s

Human ILC2s were obtained from tonsils and buffy coats. Cells were isolated by means of flow cytometric sorting with antibodies against specific cell-surface markers, as previously described.<sup>29</sup> Tonsils were obtained from both children and adults. Because we did not observe any differences in the results depending on age or sex, we present data pooled from children and adults of both sexes. A detailed description can be found in the [Methods](#) section in this article's Online Repository.

## Culture, expansion, and treatments of ILC2s

Freshly sorted tonsillar ILC2s were expanded for 2 weeks in culture with a mixture of feeder cells, IL-2 (100 U/mL), PHA, and IL-4, as previously described.<sup>12</sup> After 2 weeks of expansion, the majority of the feeder cells were engulfed by the ILC2s, rendering ILC2s greater than 85% pure, and the remaining feeder cells were not viable, as determined by means of dead cell marker discrimination in flow cytometry (data not shown). ILC2s were preincubated for 40 minutes with flurbiprofen (1  $\mu$ mol/L) or a specific HPGDS inhibitor developed by Sanofi-Aventis (Paris, France; 2-phenyl-pyrimidine-5-carboxylic acid [2,3-dihydro-indol-1-yl]-amide [United States patent 8,202,863], which for this research study was synthesized by Cayman Chemical [Ann Arbor, Mich], given the internal code KMN698, and used at a concentration of 100 nmol/L) or for 20 minutes with the CRTH2 antagonist CAY10471 (1  $\mu$ mol/L) before IL-2 (10 U/mL) with or without IL-33, TSLP, and IL-25 (50 ng/mL each). Expanded ILC2s were incubated for 6 and 24 hours. Freshly sorted blood and tonsillar ILC2s were cultured for 5 days.

## Flow cytometric staining and analysis of intracellular cytokines

Cell-surface expression of CD25 and CRTH2 was detected by using specific mAbs. Intracellular IL-5 and IL-13 protein, as well as the transcription factor GATA-3, were detected after fixation and permeabilization of the cells with appropriate antibodies (see the [Methods](#) section in this article's Online Repository). After staining, cells were immediately analyzed on an LSR Fortessa flow cytometer (BD Biosciences, San Jose, Calif).

## Mast cell culture and stimulation

Human cord blood-derived mast cells (CBMCs) were isolated and then maintained, as previously described.<sup>30,31</sup> CBMCs were preincubated with 10 ng/mL human IL-4 for 4 days and received 1  $\mu$ g/mL IgE overnight before activation. CBMCs were activated in Iscove modified Dulbecco medium (IMDM)/1% normal human serum (NHS) by using either 2  $\mu$ g/mL anti-human IgE for 6 hours or 2  $\mu$ mol/L of the calcium ionophore A23187 for 2 hours. Supernatants were collected and stored at  $-80^{\circ}\text{C}$ .

## ELISA

Concentrations of IL-5 and IL-13 in cell supernatants were analyzed by using an IL-5 ELISA Duo-Set kit (R&D Systems, Minneapolis, Minn) or an IL-13 ELISA kit. Concentrations of PGD<sub>2</sub> in ILC2 supernatants in some experiments were determined by using the PGD<sub>2</sub>-MOX (methoxime) ELISA Kit (Cayman Chemical).

## Liquid chromatography–tandem mass spectrometry

Lipid mediators in ILC2 supernatants were detected after indicated treatments for 24 hours by using liquid chromatography–tandem mass spectrometry (LC-MS/MS). LC-MS/MS was carried out after the supernatants were spiked with the respective stable isotopes as internal standards and extracted by using ethyl acetate. A detailed description can be found in the [Methods](#) section in this article's Online Repository.

## Quantitative RT-PCR

Total RNA from expanded tonsillar ILC2 cells was isolated by using the RNeasy Micro Kit (Qiagen, Hilden, Germany). cDNA was synthesized by using the iScript cDNA Synthesis Kit, and quantitative PCR reactions were performed with SsoAdvanced Universal SYBR Green Supermix (Bio-Rad Laboratories, Hercules, Calif). Primers for human transcripts for molecules important for PGD<sub>2</sub> metabolism and action, *COX1*, *COX2*, *HPGDS*, *LPGDS*, *HPGD*, prostaglandin D<sub>2</sub> receptor 1 (*DPI1*), and *CRTH2* and the housekeeping gene *RPS18*, were from Bio-Rad Laboratories. Quantitative RT-PCR (RT-qPCR) was performed in the CFX Connect Real-Time PCR Detection System (Bio-Rad Laboratories).

## Statistical analyses

Differences between 2 groups were analyzed by using the matched-pairs *t* test, and 3 or more groups were compared by using 1-way ANOVA for repeated measurements and the Dunnett multiple comparisons test. Analyses were performed by using GraphPad Prism 6 software (GraphPad Software, La Jolla, Calif).

## RESULTS

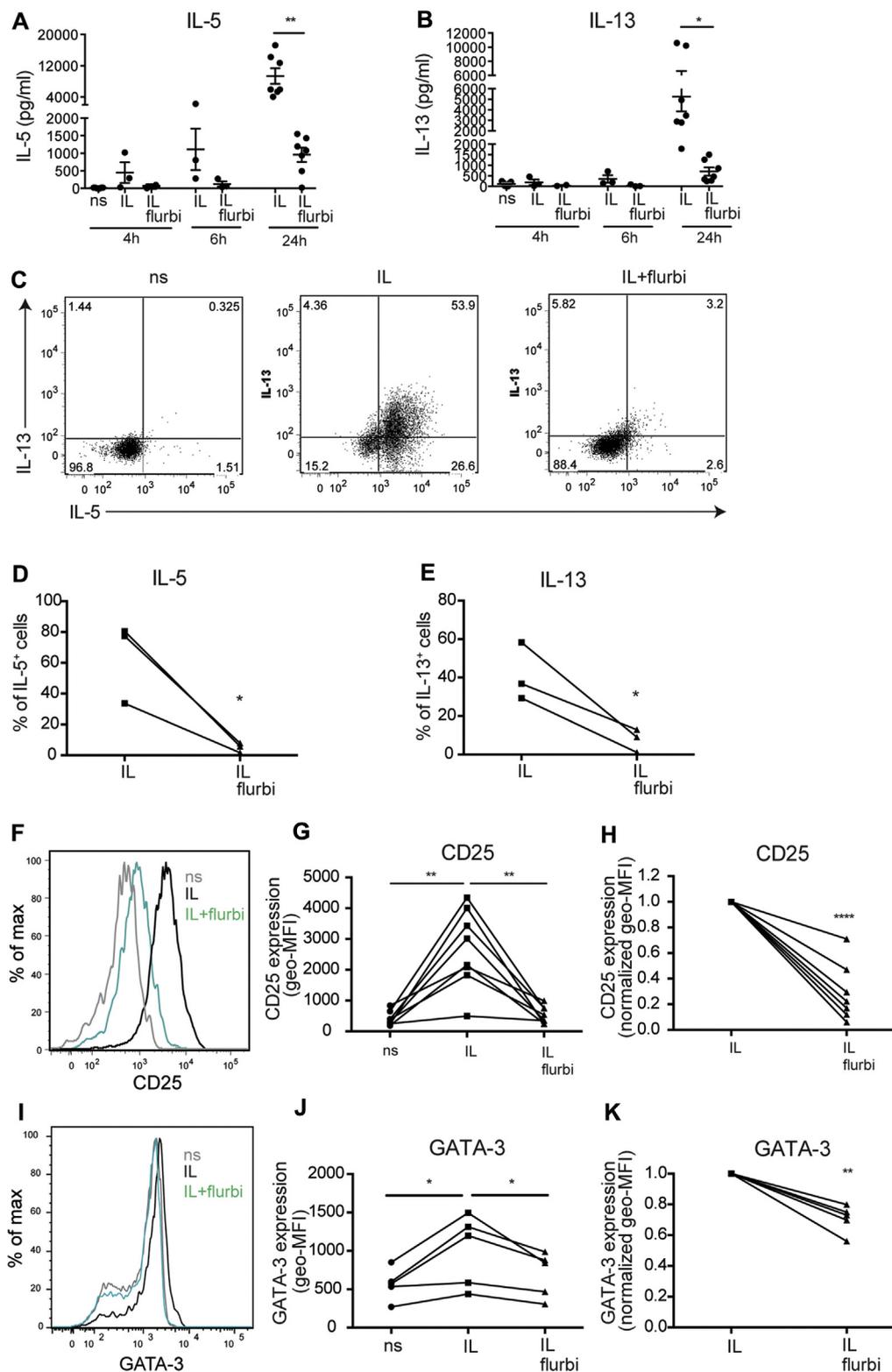
### Production of IL-5 and IL-13 by ILC2s is dependent on endogenous PGs

Recently, we showed that human ILC2s express HPGDS<sup>32</sup> and therefore hypothesized that ILC2s are able to produce PGs, which might be important for ILC2 activation. Because ILC2s are typically activated by cytokines, we sought to investigate the role of endogenous PGs in the context of cytokine-induced ILC2 activation. We incubated human tonsillar ILC2s with cytokines known to activate ILC2s: IL-33, TSLP, and IL-25 (each 50 ng/mL) plus IL-2 (10 U/mL). To investigate the importance of endogenous PGs in ILC2 activation, we pretreated ILC2s with flurbiprofen. Inhibition of the COX-1/2 enzymes almost completely prevented IL-5 and IL-13 secretion from cytokine-stimulated ILC2s ([Fig 1, A and B](#)). Suppression of type 2 cytokine production on a per-cell basis was confirmed by analyzing intracellular cytokine production, demonstrating that flurbiprofen inhibited the synthesis of IL-5 and IL-13 in ILC2s ([Fig 1, C-E](#)) without hampering cell viability (data not shown). Because IL-2 is an important costimulator of ILC2s,<sup>4</sup> we determined how CD25 (IL-2 receptor  $\alpha$ ) expression is affected by ILC2-derived PGs. Although stimulatory cytokines upregulated CD25 expression on ILC2s ([Fig 1, F-H](#)), coincubation of ILC2s with flurbiprofen largely abolished the cytokine-triggered CD25 upregulation ([Fig 1, F-H](#)). Because the transcription factor GATA-3 is important for maintenance and activation of ILC2s, we examined whether its expression is affected by inhibition of PG production. Indeed, inhibition of COX-1/2 by flurbiprofen reduced cytokine-induced GATA-3 upregulation ([Fig 1, I-K](#)).

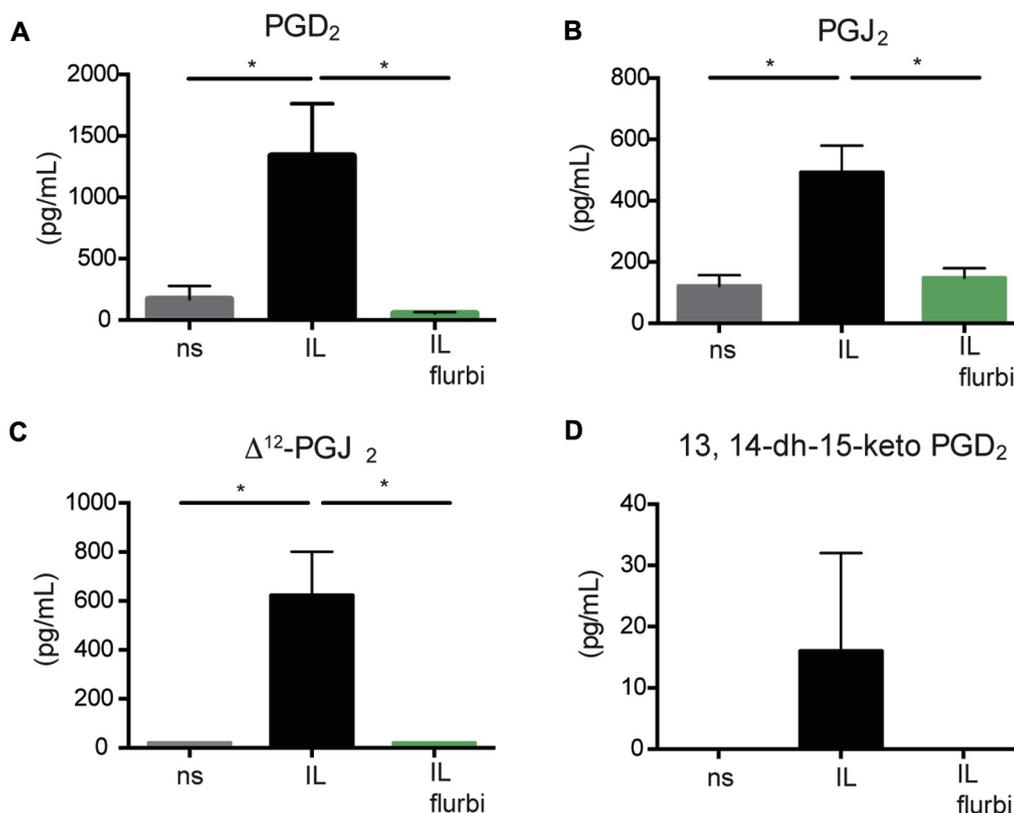
These data suggest that PGs produced by ILC2s on cytokine stimulation are necessary for optimal ILC2 activation.

### The PGD<sub>2</sub> pathway is turned on in stimulated human ILC2s

Based on the dramatically reduced ILC2 function on inhibition of PG production, we aimed to investigate which PGs can be released by activated ILC2s. It has already been shown that exogenous PGD<sub>2</sub> acts as a strong stimulator of ILC2s through engaging the CRTH2 receptor.<sup>8</sup> However, it has not yet been determined whether ILC2s themselves can produce PGD<sub>2</sub>.



**FIG 1.** Production of IL-5 and IL-13 by ILC2s is dependent on endogenously produced PGs. **A** and **B**, Sorted and expanded ILC2s in the presence of IL-2 (10 U/mL) were stimulated with IL-33, IL-25, and TSLP (50 ng/mL each) or left nonstimulated for 4, 6, and 24 hours. Flurbiprofen (1  $\mu$ mol/L) was used at 40 minutes before treatment. Concentrations of IL-5 (Fig 1, A) and IL-13 (Fig 1, B) in ILC2 supernatants were detected by using ELISA; graphs show individual concentrations and means  $\pm$  SEMs. \* $P$  < .05 and \*\* $P$  < .01, 3 donors for the 4- and 6-hour time points and 7 donors for 24 hours. **C**, Intracellular expression of IL-5 and IL-13 in ILC2s was assessed after 24 hours by using flow cytometry. **D** and **E**, Graphs show individual percentages of IL-5-producing (Fig 1, D) and IL-13-producing (Fig 1, E) ILC2s. \* $P$  < .05, 3 donors. **F** and **I**, Expression of CD25 (Fig 1, F)



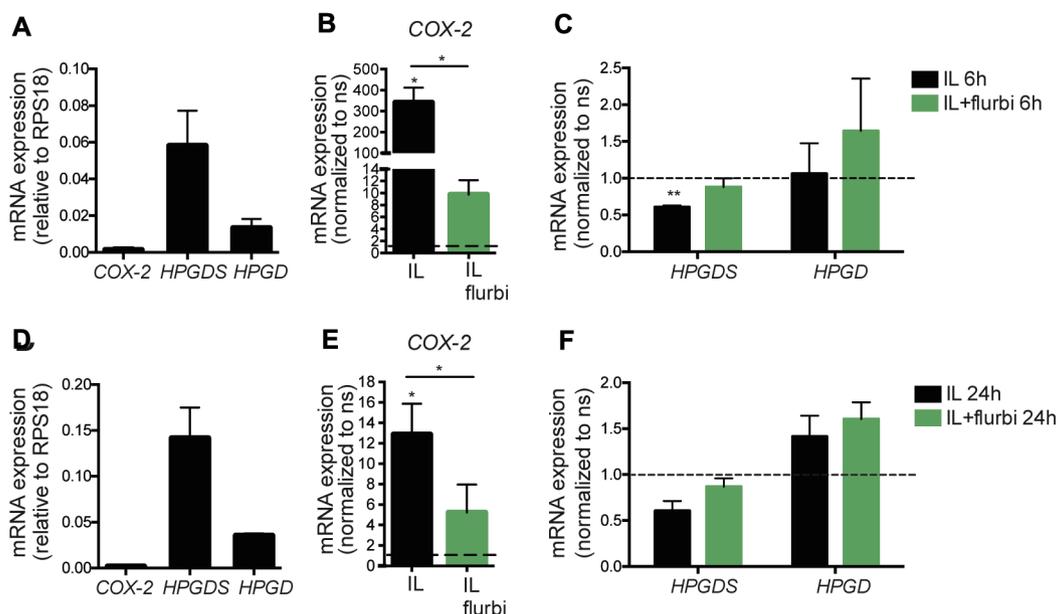
**FIG 2.** Human tonsillar ILC2s produce PGD<sub>2</sub> and its metabolites on cytokine stimulation. Sorted and expanded tonsillar ILC2s were stimulated as indicated for 24 hours with or without flurbiprofen (1 μmol/L, 40 minutes before treatment). Mediators of the PGD<sub>2</sub> pathway were detected in ILC2 supernatants by using LC-MS/MS. Graphs show mean + SEM concentrations of PGD<sub>2</sub> (\**P* < .05, 5 donors for the nonstimulated [*ns*] and IL-33, IL-25, and TSLP [*IL*] groups and 4 donors for the IL plus flurbiprofen group; **A**), PGJ<sub>2</sub> (\**P* < .05, 7 donors; **B**), Δ<sup>12</sup>-PGJ<sub>2</sub> (\**P* < .05, 7 donors; **C**), and DK-PGD<sub>2</sub> (**D**) from 4 donors. Samples were compared by using ANOVA matched-pair measurements with the Dunnett posttest, in which other groups were compared with the IL group, except for PGD<sub>2</sub> data, for which ANOVA unpaired measurement was performed. *flurbi*, Flurbiprofen; *IL*, IL-33, IL-25, and TSLP; *ns*, nonstimulated.

Therefore we collected supernatants of *in vitro*-expanded human tonsillar ILC2s and performed a highly sensitive mass spectrometric analysis for detection of lipid mediator release. Nonstimulated ILC2s did not constitutively release any lipids, whereas upon cytokine stimulation, we were able to detect PGD<sub>2</sub> and its metabolites PGJ<sub>2</sub>, Δ<sup>12</sup>-PGJ<sub>2</sub>, and, in lower concentrations, DK-PGD<sub>2</sub> (Fig 2). Coincubation with flurbiprofen completely inhibited the release of these lipid mediators. Additionally, we detected thromboxane B<sub>2</sub> (TXB<sub>2</sub>) in ILC2 supernatants, which was also completely inhibited by flurbiprofen (see Fig E1 in this article's Online Repository at [www.jacionline.org](http://www.jacionline.org)). Upon cytokine stimulation, we additionally detected the PGD<sub>2</sub> metabolite 15-deoxy-Δ<sup>12,14</sup>-PGJ<sub>2</sub> (15d-PGJ<sub>2</sub>; data not shown). Other prostanoids, such as PGE<sub>2</sub>, PGI<sub>2</sub>, and metabolites, were not detected in the supernatants of stimulated ILC2s (data not shown).

To address the relative contribution of ILC2s to the PGD<sub>2</sub> pool, we analyzed lipid mediators produced by mast cells, which are known to be the richest source of PGD<sub>2</sub>.<sup>33-36</sup> We detected high amounts of PGD<sub>2</sub> on IgE receptor activation of human CBMCs (see Fig E2, A, in this article's Online Repository at [www.jacionline.org](http://www.jacionline.org)). Compared with CBMCs, ILC2s clearly produced less PGD<sub>2</sub>, suggesting most likely an autocrine but potentially also paracrine role of PGD<sub>2</sub> in ILC2 activation rather than contributing to the systemic levels of PGD<sub>2</sub>.

Furthermore, in addition to detecting PGD<sub>2</sub> and its metabolites in ILC2 supernatants, we analyzed gene expression of the enzymes involved in PGD<sub>2</sub> metabolism: COX-1, COX-2, HPGDS, and HPGD (Fig 3). Although nonstimulated ILC2s constitutively expressed *HPGDS* mRNA, they did not express *COX1* or *COX2* mRNA, providing an explanation for the lack of PG production

and GATA-3 (Fig 1, I) was analyzed by using flow cytometry; representative histograms are shown. **G, H, J,** and **K,** Graphs show individual geometric mean fluorescence intensity of CD25 expression (\*\**P* < .01, 8 donors; Fig 1, G) and GATA-3 expression (\**P* < .05, 5 independent experiments, 4 donors; Fig 1, J), and geometric mean fluorescence intensity normalized to stimulated cells of CD25 (\*\*\*\**P* < .0001, 8 donors; Fig 1, H), and GATA-3 expression (\*\**P* < .01, 5 independent experiments, 4 donors; Fig 1, K) after 24 hours of culture. *flurbi*, Flurbiprofen; *IL*, IL-33, IL-25, and TSLP; *ns*, nonstimulated.



**FIG 3.** ILC2s express enzymes necessary for PGD<sub>2</sub> production and metabolism. Sorted and expanded tonsillar ILC2s were stimulated as indicated for 6 (A-C) and 24 (D-F) hours, with or without flurbiprofen (1 μmol/L, 40 minutes before treatment). Basal mRNA expression of *COX2*, *HPGDS*, and *HPGD* in nonstimulated ILC2s after 6 (Fig 3, A) and 24 (Fig 3, D) hours of culture were analyzed by using quantitative PCR. Data are shown as means + SEMs (data are from 4 donors). Changes of gene expression normalized to nonstimulated cells after 6 hours of stimulation (\**P* < .05 between IL-33, IL-25, and TSLP [IL] treatment and the nonstimulated group and between IL plus flurbiprofen and IL for *COX2* and \*\**P* < .01 between the IL treatment and nonstimulated groups for *HPGDS*, 4 donors for *COX-2* data and 3 donors for *HPGDS* and *HPGD*; Fig 3, B and C) and after 24 hours (\**P* < .05 between the IL treatment and nonstimulated groups and between the IL plus flurbiprofen and IL groups, 4 donors; Fig 3, E and F) are shown as means + SEMs. *flurbi*, Flurbiprofen; *IL*, IL-33, IL-25, and TSLP; *ns*, nonstimulated.

from resting ILC2s. However, after cytokine stimulation, ILC2s upregulated *COX2* expression approximately 300-fold (Fig 3, A, B, D, and E). *COX1* expression was not induced on cytokine stimulation (data not shown). Interestingly, the COX-1/2 inhibitor flurbiprofen partially prevented upregulation of *COX2* mRNA, indicating a possible regulatory role of COX-2 activation in transcription of *COX-2* in ILC2s (Fig 3, B and E). Although ILC2s expressed *HPGDS* transcripts under basal conditions (Fig 3, A and D), they lacked *LPGDS* expression (data not shown). These findings are in line with our previous observation that freshly isolated human ILC2s express *HPGDS* transcripts, as detected by using single-cell RNA sequencing.<sup>32</sup> Additionally, we revealed here that nonstimulated ILC2s express *HPGD* (Fig 3, A and D), the enzyme responsible for conversion of PGD<sub>2</sub> to DK-PGD<sub>2</sub>.<sup>23</sup> Interestingly, DK-PGD<sub>2</sub> is a natural selective agonist of the CRTH2 receptor,<sup>25</sup> and we were able to detect small amounts of this PGD<sub>2</sub> metabolite in ILC2 supernatants (Fig 2, D). Cytokine stimulation caused downregulation of *HPGDS* expression after 6 hours (Fig 3, C) and slightly but not significantly reduced *HPGDS* expression after 24 hours (Fig 3, F). Coincubation with flurbiprofen did not influence *HPGDS* expression, and ILC2 stimulation did not affect the expression of *HPGD* (Fig 3, C and F).

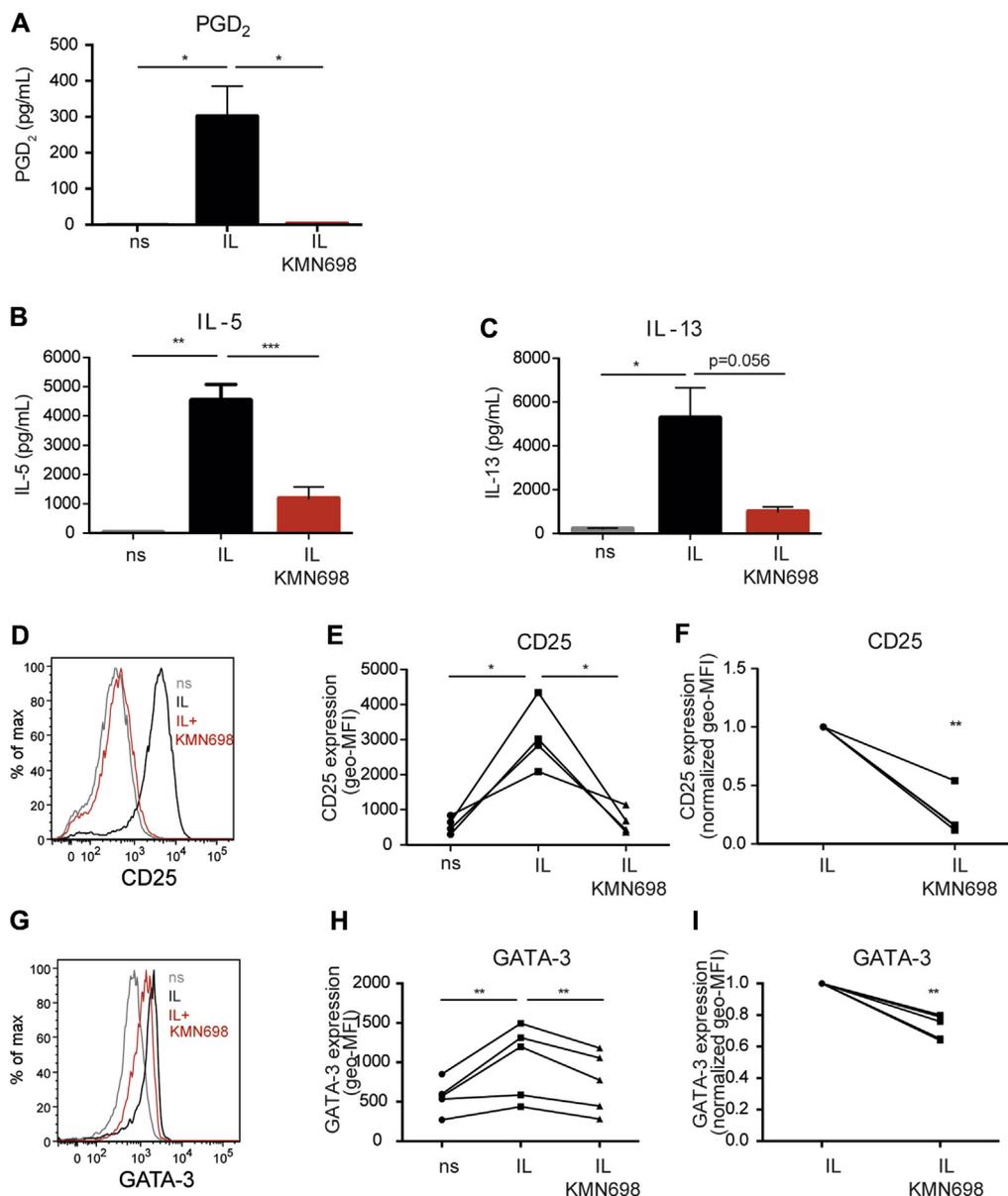
Overall, we demonstrated that ILC2s express the enzymes required for PGD<sub>2</sub> synthesis, *HPGDS* and *COX2*, with the latter being highly upregulated on cytokine stimulation. As a consequence, cytokine activation of ILC2s leads to endogenous PG synthesis, as demonstrated by our detection of PGD<sub>2</sub> and its metabolites in the supernatant of stimulated ILC2s. Additionally, we

showed that ILC2s express *HPGD* and that its product, the selective CRTH2 agonist DK-PGD<sub>2</sub>, can be detected in ILC2 supernatants.

### Activation of ILC2s on cytokine stimulation depends on endogenous PGD<sub>2</sub>

We found that PGD<sub>2</sub> is the major PG released by ILC2s. To confirm that the observed inhibitory effects of flurbiprofen are due to abolished PGD<sub>2</sub> production and thereby reduced paracrine/autocrine action of PGD<sub>2</sub>, we incubated ILC2s with the selective *HPGDS* inhibitor KMN698 and the CRTH2 antagonist CAY10471 before cytokine treatment. The *HPGDS* inhibitor blocked release of PGD<sub>2</sub> from stimulated ILC2s, showing its specific effects (Fig 4, A). Inhibition of PGD<sub>2</sub> production reduced IL-5 and IL-13 release from cytokine-stimulated ILC2s, confirming a crucial role of PGD<sub>2</sub> production in ILC2 stimulation (Fig 4, B and C). Furthermore, we observed that the *HPGDS* inhibitor reduced cytokine-induced upregulation of CD25 (Fig 4, D-F) and GATA-3 expression (Fig 4, G-I).

Our data also revealed that the selective CRTH2 antagonist almost completely abolished IL-5 and IL-13 secretion from cytokine-stimulated ILC2s (Fig 5, A and B). Furthermore, blocking of the CRTH2 receptor prevented the increase in intracellular IL-5 and IL-13 production in activated ILC2s (Fig 5, C-E). CRTH2 antagonism also prevented CD25 (Fig 5, F-H) and GATA-3 upregulation (Fig 5, I-K) of stimulated ILC2s. These data clearly show that endogenous PGD<sub>2</sub> produced by ILC2s on



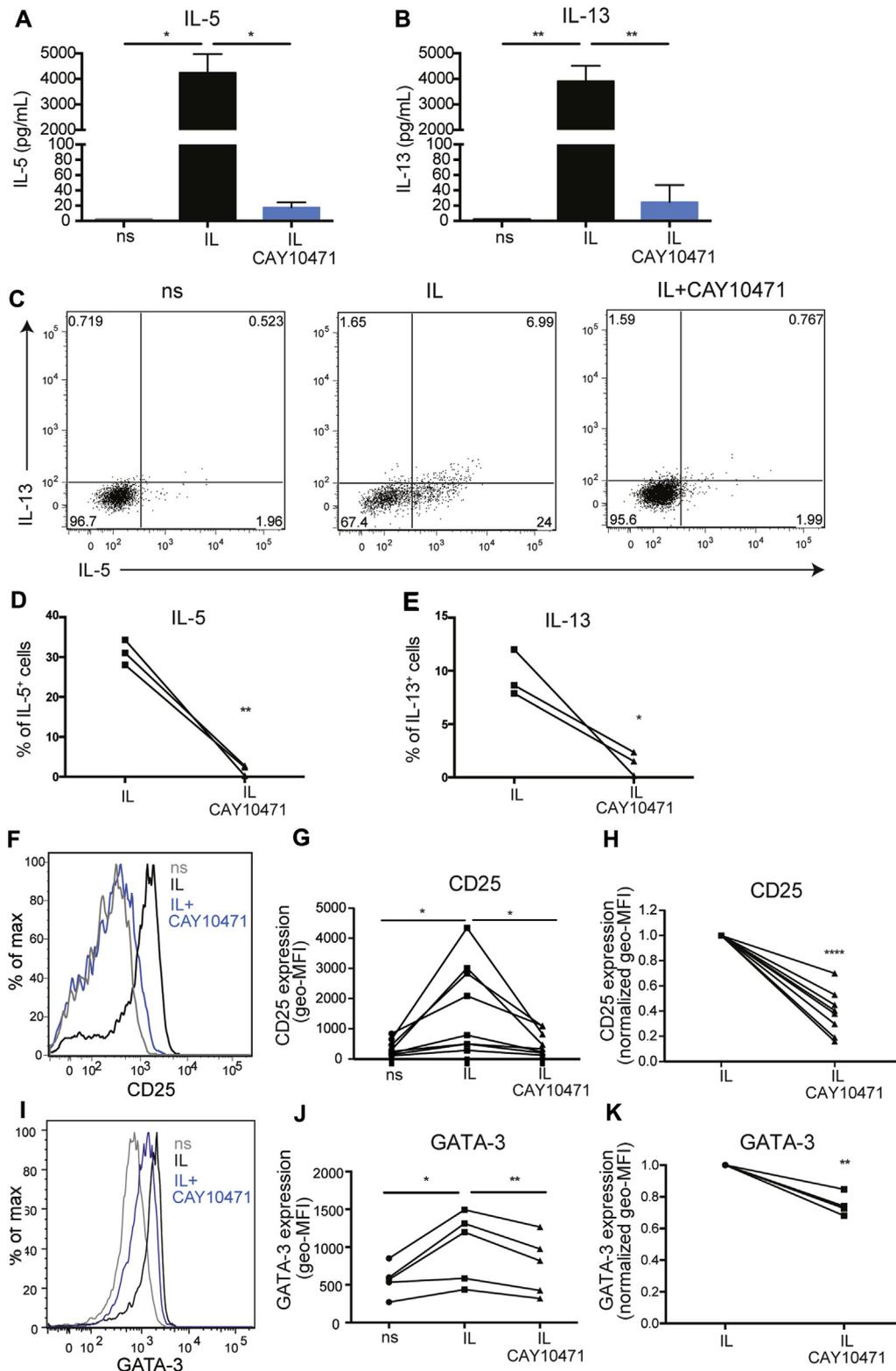
**FIG 4.** Endogenously produced PGD<sub>2</sub> is essential for ILC2 stimulation. Sorted and expanded tonsillar ILC2s were stimulated as indicated with or without the HPGDS inhibitor KMN698 (100 nmol/L, 40 minutes before treatment) for 24 hours. **A**, Concentration of released PGD<sub>2</sub> was measured by using ELISA; graph shows means + SEMs (\**P* < .05, data are from 2 donors, 4 independent experiments). **B** and **C**, Concentrations of released IL-5 (Fig 4, **B**) and IL-13 (Fig 4, **C**) were measured by using ELISA; graphs show means + SEMs (\**P* < .05, \*\**P* < .01, and \*\*\**P* < .001, 4 donors, 5 independent experiments). **D** and **G**, Expression of CD25 (Fig 4, **D**) and GATA-3 (Fig 4, **G**) was analyzed by using flow cytometry; representative histograms are shown. **E**, **F**, **H**, and **I**, Graphs show individual geometric mean fluorescence intensity of CD25 (\**P* < .05, 3 donors, 4 independent experiments; Fig 4, **E**) and GATA-3 expression (\*\**P* < .01, 4 donors, 5 independent experiments; Fig 4, **H**) and geometric mean fluorescence intensity normalized to stimulated cells of CD25 (\*\**P* < .01, 3 donors, 4 independent experiments; Fig 4, **F**) and GATA-3 expression (\*\**P* < .01, 4 donors, 5 independent experiments; Fig 4, **I**). *IL*, IL-33, IL-25, and TSLP; *ns*, nonstimulated.

cytokine stimulation is essential for activation of ILC2s acting through the CRTH2 receptor.

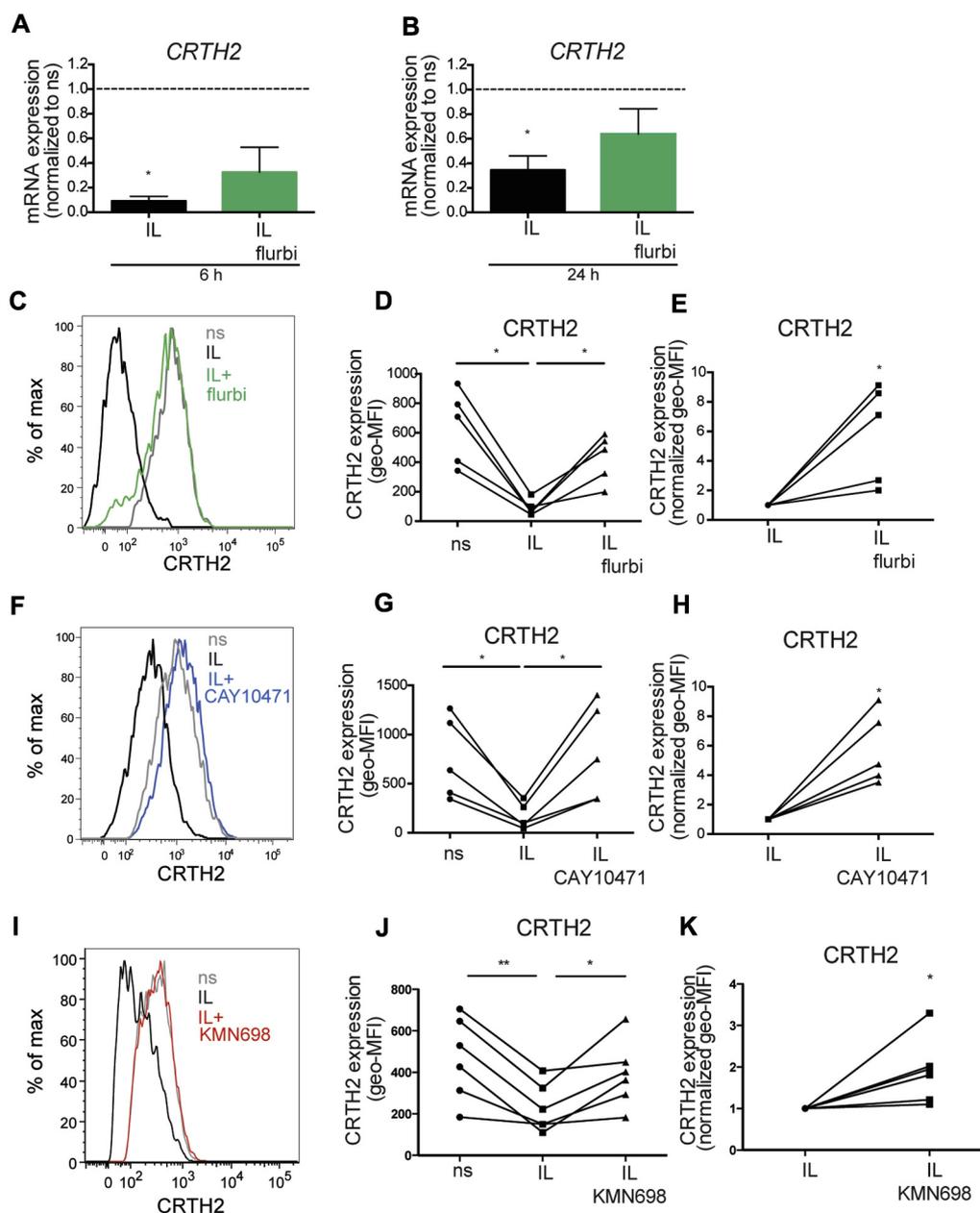
It has been demonstrated that exogenous PGD<sub>2</sub> directly stimulates ILC2s through binding to the CRTH2 receptor and that such engagement leads to downregulation of CRTH2 expression at both the mRNA and protein levels.<sup>8</sup> Thus we set out to investigate

the role of endogenous PGs in regulating expression of CRTH2 in ILC2s.

Although we detected high levels of *CRTH2* transcripts in non-stimulated ILC2s, only low expression of the *DPI1* receptor could be found (see Fig E3, **A** and **B**, in this article's Online Repository at [www.jacionline.org](http://www.jacionline.org)). Cytokine stimulation significantly

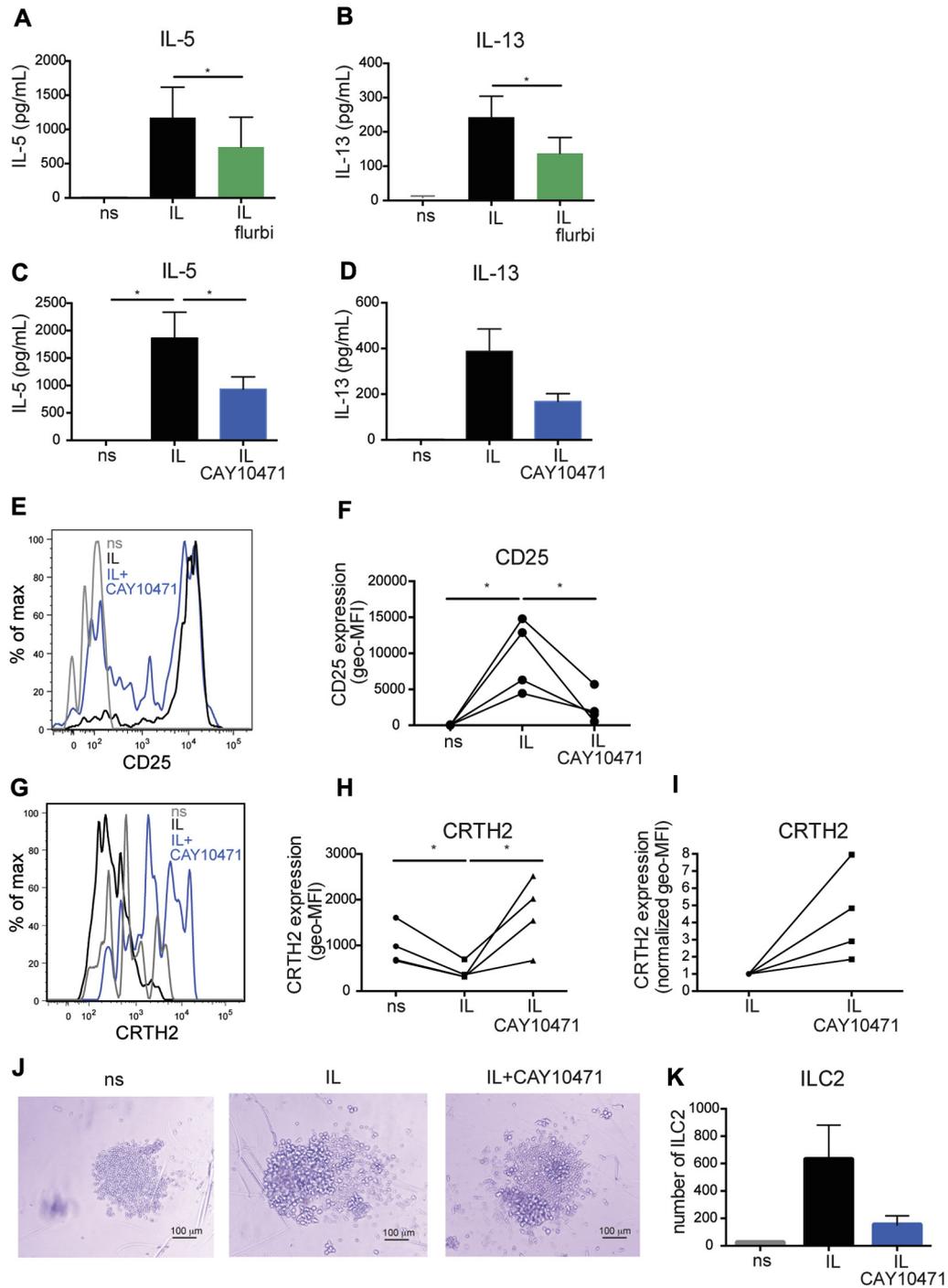


**FIG 5.** Endogenously produced PGD<sub>2</sub> acts through the CRTH2 receptor on ILC2s. **A** and **B**, Sorted and expanded ILC2s were treated as indicated in the presence or absence of the CRTH2 selective antagonist CAY10471 (1  $\mu$ mol/L, 20 minutes before treatment) for 24 hours. **A** and **B**, Concentrations of IL-5 (Fig 5, **A**) and IL-13 (Fig 5, **B**) in ILC2 supernatants were detected by using ELISA; graphs show means + SEMs (\* $P$  < .05 and \*\* $P$  < .01, 4 donors). **C-E**, Intracellular expression of IL-5 and IL-13 in ILC2s was assessed after 24 hours by using flow cytometry; graphs show individual percentages of IL-5-producing (Fig 5, **D**) and IL-13-producing (Fig 5, **E**) ILC2s (\* $P$  < .05 and \*\* $P$  < .01, 3 donors). **F** and **I**, Expression of CD25 (Fig 5, **F**) and GATA-3 (Fig 5, **I**) was analyzed by using flow cytometry; representative histograms are shown. **G**, **H**, **J**,



**FIG 6.** Cytokine-induced downregulation of CRTH2 is prevented by inhibition of endogenous PG production or antagonism of CRTH2 receptor. Tonsillar ILC2s were stimulated as indicated for 6 or 24 hours, with pretreatment of flurbiprofen (1  $\mu\text{mol/L}$ , 40 min; **A-E**), CAY10471 (1  $\mu\text{mol/L}$ , 20 minutes; **F-H**), or the HPGDS inhibitor KMN698 (100 nmol/L, 40 minutes; **I-K**). Changes in *CRTH2* mRNA expression normalized to nonstimulated cells after 6 hours of stimulation ( $*P < .05$ , 3 donors; Fig 6, **A**) and after 24 hours ( $*P < .05$ , 4 donors; Fig 6, **B**) are shown as means  $\pm$  SEMs. Fig 6, **C**, **F**, and **I**, CRTH2 protein expression was analyzed by using flow cytometry. Graphs show individual geometric mean fluorescence intensity (Fig 6, **D**, **G**, and **J**) and individual geometric mean fluorescence intensity normalized to stimulated cells of CRTH2 expression after 24 hours of culture (Fig 6, **E**, **H**, and **K**;  $*P < .05$  and  $**P < .01$ , for flurbiprofen and CAY10471 data are from 5 donors and for KMN698 from 4 donors, 6 independent experiments). *flurbi*, Flurbiprofen; *IL*, IL-33, IL-25, and TSLP; *MFI*, mean fluorescence intensity; *ns*, nonstimulated.

and **K**, Graphs show individual geometric mean fluorescence intensity of CD25 ( $*P < .05$ , 7 donors, 8 independent experiments; Fig 5, **G**) and GATA-3 expression ( $*P < .05$  and  $**P < .01$ , 5 independent experiments, 4 donors; Fig 5, **J**) and geometric mean fluorescence intensity normalized to stimulated cells of CD25 ( $***P < .0001$ , 7 donors, 8 independent experiments; Fig 5, **H**) and GATA-3 expression ( $**P < .01$ , 4 donors, 5 independent experiments; Fig 5, **K**) after 24 hours of culture. *IL*, IL-33, IL-25, and TSLP; *MFI*, mean fluorescence intensity; *ns*, nonstimulated.



**FIG 7.** Endogenously produced PGD<sub>2</sub> contributes to activation of freshly sorted tonsillar and blood ILC2s on cytokine stimulation. Freshly sorted human tonsillar (**A** and **B**) and blood (**C-K**) ILC2s were incubated as indicated for 5 days in the presence or absence of flurbiprofen (1 μmol/L; Fig 7, **A** and **B**) or CAY10471 (1 μmol/L; Fig 7, **C-K**). Concentrations of released IL-5 from tonsillar ILC2s (Fig 7, **A**) and blood ILC2s (Fig 7, **C**) and IL-13 from tonsillar ILC2s (Fig 7, **B**) and blood ILC2s (Fig 7, **D**) were measured by means of ELISA; graphs show means + SEMs (\**P* < .05, 4 donors for blood ILC2s and 3 donors for tonsillar ILC2s). Fig 7, **E** and **G**, Expression of CD25 and CRTH2 in freshly sorted blood ILC2s was analyzed by using flow cytometry; representative histograms are shown. Graphs show individual geometric mean fluorescence intensity of CD25 expression in blood ILC2s after 5 days of culture (\**P* < .05, 4 donors; Fig 7, **F**) and of CRTH2 expression (\**P* < .05, 4 donors; Fig 7, **H**) and individual geometric mean fluorescence intensity of CRTH2 expression normalized to stimulated cells (4 donors; Fig 7, **I**). Fig 7, **J**, Images of freshly sorted blood ILC2s after 5 days of treatments (×10 magnification, scale bar = 100 μm). Fig 7, **K**, Numbers of freshly sorted blood ILC2s acquired by using flow cytometry from 4 donors. *flurbi*, Flurbiprofen; *IL*, IL-33, IL-25, and TSLP; *MFI*, mean fluorescence intensity; *ns*, nonstimulated.

reduced *CRTH2* mRNA expression in ILC2s, which was partially but not significantly reversed on inhibition of COX-1/2 by using flurbiprofen (Fig 6, A and B). *DPI* expression was not modulated by cytokine stimulation after 6 hours but was downregulated after 24 hours of stimulation (see Fig E3, C and D). Moreover, we showed that stimulatory cytokines also reduce CRTH2 protein expression on the cell surface (Fig 6). Of note, pretreatment of ILC2s with either flurbiprofen (Fig 6, C-E), CRTH2 antagonist (Fig 6, F-H), or a HPGDS inhibitor (Fig 6, I-K) reversed cell-surface downregulation of CRTH2 to the expression level of nonstimulated ILC2s. These findings strongly suggest that endogenously released PGD<sub>2</sub> is responsible for cytokine-induced reduction of CRTH2 expression.

### Endogenously produced PGD<sub>2</sub> is crucial for cytokine-induced activation of freshly isolated blood and tonsillar ILC2s

Next, we aimed to confirm our findings based on *in vitro*-expanded tonsillar ILC2s by using freshly isolated and sort-purified ILC2s. Additionally, we extended our studies to include not only tissue (tonsillar) ILC2s but also blood ILC2s, which display a more naive phenotype.<sup>37</sup> ILC2s freshly isolated from tonsils (Fig 7, A and B) and blood (Fig 7, C-K) elaborated IL-5 and IL-13 after 5 days of *in vitro* culture with IL-2, IL-25, and IL-33 plus TSLP. As observed in *in vitro*-expanded ILC2s, cotreatment with flurbiprofen (Fig 7, A and B) or the CRTH2 antagonist (Fig 7, C and D) significantly reduced IL-5 and IL-13 secretion.

We also determined changes in CD25 expression in freshly isolated blood ILC2s. Cytokine stimulation markedly upregulated CD25 expression in ILC2s, which was prevented by means of preincubation with the CRTH2 antagonist (Fig 7, E and F). Similar to our observation with *in vitro*-expanded ILC2s, cytokine-induced downregulation of CRTH2 in freshly isolated ILC2s was reversed by coinubation with the CRTH2 antagonist (Fig 7, G-I). Furthermore, we captured microscopic images of freshly isolated ILC2s that were cultured for 5 days. These images revealed that specific inhibition of the CRTH2 receptor allowed less prominent cell growth compared with the treatment with only stimulatory cytokines (Fig 7, J). The microscopic analysis was confirmed by quantification of the number of viable ILC2s by using flow cytometry (Fig 7, K), which showed that blocking of CRTH2 prevented cytokine-induced increase of ILC2 numbers.

Altogether, we show here that endogenously produced PGD<sub>2</sub> is essential in complete activation of freshly isolated human blood and tissue ILC2s.

## DISCUSSION

ILC2s require several signals for optimal activation. Synergy between the signal transducer and activator of transcription activators TSLP and IL-2, as well as the nuclear factor  $\kappa$ B stimulators IL-33 and IL-25, potently activates human ILC2s.<sup>3,38</sup> Interestingly, ILC2s can also be activated by the lipid mediators PGD<sub>2</sub>, LTD<sub>4</sub>, and LTE<sub>4</sub>.<sup>6-8</sup> The PGD<sub>2</sub>-CRTH2 axis is described to be important in identification and function of ILC2s.<sup>7,8,39</sup> Because ILC2s are crucial in the initial steps and maintenance of type 2 immune responses during allergic inflammation,<sup>1,40</sup> better understanding of regulation of these cells is required for

development of novel therapeutic approaches. Although recent studies highlight the role of lipid mediators in ILC2 function,<sup>5</sup> it is still unknown whether ILC2s are able to produce PG levels on their own. We demonstrate here that upon cytokine stimulation, ILC2s, which constitutively express *HPGDS*, upregulate *COX2* and start producing PGD<sub>2</sub>, which in turn leads to engagement of the CRTH2 receptor. We prove that this endogenous PGD<sub>2</sub>-CRTH2 pathway is essential for cytokine-induced ILC2 activation.

In our study we detected PGD<sub>2</sub> and its metabolites PGJ<sub>2</sub>,  $\Delta^{12}$ -PGJ<sub>2</sub>, and 15d-PGJ<sub>2</sub>, as well as TXB<sub>2</sub>, in supernatants of stimulated ILC2s. Importantly, we observed that ILC2s exclusively produce these lipid mediators. The COX-1/2 inhibitor flurbiprofen inhibited prostanoid production of ILC2s. To our surprise, flurbiprofen almost completely abolished IL-5 and IL-13 production of ILC2s on cytokine stimulation, suggesting an interesting mechanism of ILC2 activation by endogenously produced PGs.

We were able to confirm *HPGDS* mRNA expression in *in vitro*-expanded tonsillar ILC2s, as previously shown in freshly isolated ILC2s.<sup>32</sup> *HPGDS* is also expressed by other immune cells, such as mast cells, T<sub>H</sub>2 and dendritic cells, macrophages, and eosinophils, which are all able to produce PGD<sub>2</sub>.<sup>14,16,22</sup> Even though *HPGDS* is constitutively expressed by ILC2s, nonstimulated ILC2s do not produce PGD<sub>2</sub>, which is most probably due to the lack of *COX1* and *COX2* expression. In contrast, mast cells express COX-1 and release PGD<sub>2</sub> already under basal conditions, whereas COX-2 can be induced for instance on immunologic activation with IgE crosslinking.<sup>15</sup> Here we show that stimulation of ILC2s with the cytokines IL-33, TSLP, and IL-25 plus IL-2 induces approximately 300-fold upregulation of *COX2* mRNA expression. COX-2 is the rate-limiting enzyme in PGD<sub>2</sub> production because it provides the substrate PGH<sub>2</sub> for the already highly expressed *HPGDS* in ILC2s. We used a cytokine mixture for activating ILC2s because this approach ensures full stimulation of human ILC2s.<sup>3,12</sup>

We have not investigated the effect of single cytokines on COX-2 induction. However, IL-33, a member of the IL-1 family, has previously been shown to induce PGD<sub>2</sub> synthesis in mast cells,<sup>41</sup> and a recent study demonstrates that TSLP synergistically with IL-33 contributes to PGD<sub>2</sub> production by human mast cells.<sup>36</sup> Similar to ILC2s, T<sub>H</sub>2 cells can also upregulate COX-2 on different stimuli and hence are capable of PGD<sub>2</sub> production.<sup>16,42</sup> Interestingly, the COX inhibitor flurbiprofen also attenuated upregulation of *COX2* mRNA in stimulated ILC2s. It is likely that endogenous PGD<sub>2</sub>, acting through a positive feedback loop, further contributes to *COX2* upregulation and, consequently, inhibition of its production reduces *COX2* expression in ILC2s. Positive feedback regulation of PGs on *COX2* expression has been shown.<sup>43</sup>

Importantly, we demonstrate that endogenous PGD<sub>2</sub> is responsible for ILC2 activation on cytokine stimulation by using an *HPGDS* inhibitor and a selective antagonist of CRTH2. It was previously shown that exogenous PGD<sub>2</sub> treatment of ILC2s led to downregulation of CRTH2 expression both at the mRNA and protein levels.<sup>8</sup> Furthermore, PGD<sub>2</sub> was reported to induce internalization and desensitization of CRTH2 in HEK293 cells and in human eosinophils.<sup>25,44</sup> We observed a reduction in CRTH2 mRNA and protein expression on cytokine stimulation of ILC2s. Most likely, restored cell-surface expression of CRTH2 after flurbiprofen, CRTH2 antagonist, and *HPGDS* inhibitor treatment is due to hampering of endogenous PGD<sub>2</sub> production and

CRTH2 binding. Thus in addition to the positive regulatory role of ILC2-derived PGs in COX-2 upregulation, there is an additional self-limiting mechanism in the PGD<sub>2</sub>-caused reduction of CRTH2 expression in activated ILC2s. Furthermore, because reduced *CRTH2* mRNA expression on cytokine stimulation was not completely reversed by flurbiprofen, it is likely that ILC2-stimulatory cytokines can also directly affect *CRTH2* gene expression in a PGD<sub>2</sub>-independent manner.

We show that ILC2s also express *HPGD*, and we detected the product of the enzyme DK-PGD<sub>2</sub> in ILC2 supernatants. DK-PGD<sub>2</sub> is a naturally occurring selective CRTH2 agonist that can activate human T<sub>H</sub>2 cells and eosinophils.<sup>13,25</sup> Additionally, other PGD<sub>2</sub> metabolites present in ILC2 supernatants, such as PGJ<sub>2</sub>, Δ<sup>12</sup>-PGJ<sub>2</sub>, and 15d-PGJ<sub>2</sub>, can also mediate their effects through the CRTH2 receptor.<sup>26,27</sup> These metabolites, along with PGD<sub>2</sub>, could be additionally responsible for CRTH2 receptor-dependent ILC2 activation. 15d-PGJ<sub>2</sub>, in addition to exerting its effects through the CRTH2 receptor, also binds to the peroxisome proliferator-activated receptor γ (PPAR-γ). Although PPAR-γ activation can lead to anti-inflammatory effects,<sup>45</sup> other studies showed that it triggered human eosinophil chemotaxis<sup>46</sup> and the type 2 response in mouse T<sub>H</sub>2 cells and ILC2s.<sup>47</sup> Hence the 15d-PGJ<sub>2</sub>-PPAR-γ axis might also be partly involved in ILC2 activation. We show that ILC2s also express low levels of *DP1* receptor. However, because of the powerful inhibitory effect of the CRTH2 antagonist, it is unlikely that the DP1 receptor is involved in ILC2 activation, which is in agreement with earlier studies.<sup>11</sup>

In addition to metabolites of the PGD<sub>2</sub> pathway, we also detected low levels of TXB<sub>2</sub>; although there is no evidence yet for thromboxane receptor expression on ILC2s, thromboxane metabolites still could play some role in ILC2 activation because they have been shown to act as CRTH2 agonists.<sup>48</sup>

Importantly, we confirmed the role of endogenous PGD<sub>2</sub> in freshly isolated ILC2s by culturing these cells with cytokines and COX-1/2 inhibitor or CRTH2 antagonists for 5 days. We chose this rather long incubation time because we have observed that freshly isolated, compared with *in vitro*-expanded, ILC2s require that for activation. In comparison with *in vitro*-expanded ILC2s, the effect of COX-1/2 inhibition and CRTH2 antagonism on freshly isolated cells was more modest, indicating that *in vitro* expansion of ILC2s leads to a greater dependence on PGD<sub>2</sub> rather than the cytokines themselves. An alternative explanation could be that during this long incubation time, the reversible inhibitory effects of the CRTH2 antagonist and flurbiprofen were less pronounced. Nevertheless, we confirmed that also freshly isolated ILC2s use endogenous PGD<sub>2</sub> as a mechanism for cytokine-induced activation.

As an intermediate step in ILC2 activation, we investigated IL-2 receptor α (CD25) receptor expression because IL-2 is an important costimulator of ILC2s.<sup>4,7</sup> Suppression of cytokine-induced CD25 upregulation by the CRTH2 antagonist and flurbiprofen indicates that regulation of CD25 expression is one of the downstream mechanisms of the PGD<sub>2</sub>-CRTH2 pathway. Furthermore, we showed that endogenous PGD<sub>2</sub> is of importance for GATA-3 upregulation.

In our *in vitro* system, we analyzed purified tonsillar and blood ILC2s. It is possible that *in vivo* PGD<sub>2</sub> released from other cellular sources, such as mast cells, can have additional effects on ILC2s. However, in early innate immune responses, the paracrine/auto-crine effects of PGD<sub>2</sub> produced by stimulated ILC2s might be crucial.

In conclusion, we revealed that cytokine-stimulated ILC2 activation largely depends on endogenous PGD<sub>2</sub> production. We showed that human ILC2s express *HPGDS*; upregulate *COX2* on stimulation with IL-33, TSLP, and IL-25 plus IL-2; and consequently produce PGD<sub>2</sub>. We could also demonstrate that endogenously produced PGD<sub>2</sub> and its metabolites, acting through the CRTH2 receptor, are essential for CD25 upregulation and subsequent production of IL-5 and IL-13 in cytokine-stimulated ILC2s. Our data reveal a novel mechanism by which ILC2s critically depend on endogenously produced PGD<sub>2</sub> induced by stimulating cytokines for optimal activation. Hence our data highlight the importance of therapeutic CRTH2 antagonists in control of allergic inflammation.

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### Key messages

- ILC2s express *HPGDS*, upregulate *COX-2*, and consequently produce PGD<sub>2</sub> upon cytokine stimulation
- Endogenously produced PGD<sub>2</sub>, acting through the CRTH2 receptor, is responsible for cytokine-induced activation of ILC2s.
- CRTH2 antagonists prevent cytokine-induced ILC2 activation and might serve as promising therapeutic tools.

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## METHODS

### Isolation and flow cytometric sorting of tonsillar and blood ILC2s

Human tonsils were freshly received from tonsillectomies of patients with obstructive sleep apnea syndrome at the Ear Nose Throat Clinic at the Karolinska University Hospital, Huddinge, Sweden. Collection and use of tonsils were approved by the regional ethical board at the Karolinska Institutet. Buffy coats were provided by the blood bank at the Karolinska University Hospital Huddinge and were collected according to the approval of the regional ethical board at Karolinska Institutet. Isolation of ILCs from tonsils was performed, as previously described.<sup>E1</sup> Briefly, tonsils were dissected into small pieces, ground through a cell strainer, and centrifuged. After Lymphoprep centrifugation (STEMCELL Technologies, Vancouver, British Columbia, Canada), layers of mononuclear cells were collected, and cells were depleted from lineage-positive cells through magnetic separation by using directly conjugated anti-CD3, anti-CD19, and anti-CD14 microbeads (markers for T cells, B cells, and monocytes, respectively) and LD MACS columns (Miltenyi Biotec, Bergisch Gladbach, Germany). Lineage-depleted PBMCs from buffy coats were additionally enriched by using the CD127 MicroBead Kit (Miltenyi Biotec). Lineage-depleted cells were incubated with the following specific mAbs: fluorescein isothiocyanate-conjugated anti-CD14 (TÜK4; Dako, Glostrup, Denmark); anti-CD3 (SK7), anti-FcεR1α (AER-37 CRA-1), anti-CD34 (581), anti-CD123 (6H6), anti-CD1a (HI149), anti-T-cell receptor αβ (IP26), anti-T-cell receptor γδ (B1), and anti-CD94 (DX22; all from BioLegend, San Diego, Calif); anti-BDCA2 (AC144; Miltenyi Biotec) and anti-CD19 (4G7; BD Biosciences). Additionally, we used Brilliant Violet 605-conjugated anti-CD161 (HP-3G10), Brilliant Violet 711-conjugated anti-CD56 (HCD56; both from BioLegend); phycoerythrin (PE)-CF594-conjugated anti-CRTH2 (BM16) and V500-conjugated anti-CD45 (HI30; both from BD Biosciences); and PE-Cy7-conjugated anti-CD127 (R34.34), PE-Cy5.5-conjugated anti-CD117 (104D2D1), and PE-Cy5-conjugated anti-NKp44 (Z231; all from Beckman Coulter, Fullerton, Calif).

In addition, cells were stained with the LIVE/DEAD Fixable Green Dead Cell Stain Kit (Life Technologies, Grand Island, NY). ILC2s were sort-purified based on their specific phenotype as Lin<sup>-</sup>CD127<sup>+</sup>CD161<sup>+</sup>CRTH2<sup>+</sup> cells by using the FACS Aria Fusion cell sorter (BD Bioscience) equipped with the FACSDiva software, version 8.

### Culture and expansion of ILC2s

Freshly sorted ILC2s were cultured in IMDM (Gibco, Carlsbad, Calif) supplemented with Yssel medium and 1% NHS (Human AB serum; Invitrogen, Carlsbad, Calif) with penicillin (100 U/mL; HyClone, San Angelo, Tex) and streptomycin (0.1 mg/mL; HyClone). ILC2s were expanded for 2 weeks in culture with a mixture of feeder cells containing PBMCs irradiated at 25 Gy and JY cells irradiated at 50 Gy, with IL-2 (100 U/mL), PHA (1 μg/mL), and addition of IL-4 (25 ng/mL; R&D Systems).

Freshly sorted ILC2s were kept in Yssel medium/1% NHS with IL-2 (10 U/mL) in a 96-well plate (2000 cells per sample) and treated the following day.

### Treatments of ILC2s

After expansion for 2 weeks, ILC2s were rested in Yssel medium/1% NHS with IL-2 (2 U/mL) for 3 days before the experiments. Cells were seeded in 96-well U-bottom plates at  $5 \times 10^4$  cells/well. ILC2s were incubated for 40 minutes with flurbiprofen (1 μmol/L) or 20 minutes with the CRTH2 antagonist CAY10471 (1 μmol/L) before adding IL-2 (10 U/mL) with or without IL-33 (PeproTech, Rocky Hill, NJ), TSLP (PeproTech), and IL-25 (R&D Systems) at 50 ng/mL each.

For LC-MS/MS and quantitative PCR experiments, expanded ILC2s were seeded at a density of  $2.5 \times 10^5$  cells/well in IMDM/1% NHS without Yssel supplement and incubated for 6 and 24 hours with IL-2 (10 U/mL) and with or without combination of IL-33, TSLP, and IL-25 (50 ng/mL each). Where indicated, flurbiprofen (1 μmol/L) was used 40 minutes before treatment.

Freshly sorted blood and tonsillar ILC2s were kept in the presence of IL-2 (10 U/mL) and nonstimulated or treated with the combination of IL-33, TSLP,

and IL-25 (50 ng/mL each) for 5 days. As pretreatments, flurbiprofen (1 μmol/L) and a specific HPGDS inhibitor developed by Sanofi-Aventis (2-phenyl-pyrimidine-5-carboxylic acid [2,3-dihydro-indol-1-yl]-amide [US patent 8,202,863], which for this research study was synthesized by Cayman, given the internal code KMN698, and used at a concentration of 100 nmol/L) were added at 40 minutes, whereas the CRTH2 antagonist CAY10471 (1 μmol/L) was added at 20 minutes before the cytokines. Flurbiprofen and CAY10471 treatments were repeated on days 2 and 4 of incubation. Microscopic images of freshly sorted blood ILC2s were captured after 5 days of culture at  $\times 10$  magnification by using a camera and the TouView (TouView Photonics, Zhejiang, China) program.

### Flow cytometry and analysis of intracellular cytokines

Cell-surface staining of ILC2s was determined by using PE-Cy7-conjugated anti-CD127, PE-Cy5.5-conjugated anti-CD117, Brilliant Violet 605-conjugated anti-CD161, PE-CF594-conjugated anti-CRTH2, and Brilliant Violet 711-conjugated anti-CD56. Dead cells were excluded by using the LIVE/DEAD Fixable Green Dead Cell Stain Kit (Life Technologies). CD25 receptor expression was assessed by using Brilliant Violet 650-conjugated anti-CD25 (BC96; BioLegend).

For intracellular detection of IL-5 and IL-13, ILC2s were stimulated for 24 hours, as already described, with Golgi Plug (containing Brefeldin A) and Golgi Stop (containing monensin) added after 20 hours. Thereafter, ILC2s were fixed (2% paraformaldehyde), permeabilized (FACS Permeabilizing Solution 2; BD Bioscience), and stained with BD Biosciences antibodies for intracellular IL-13 (allophycocyanin-conjugated anti-IL-13; JES10-5A2) and IL-5 (PE-conjugated anti-IL-5; TRFK5). Intracellular GATA-3 expression was assessed by using allophycocyanin-conjugated antibody (clone TWAJ; eBioscience) after fixation and permeabilization (Foxp3/Transcription factor staining buffer set; eBioscience).

All samples were measured on an LSR Fortessa flow cytometer (BD Biosciences) equipped with FACSDiva software version 8, and data were analyzed by using FlowJo software (TreeStar, Ashland, Ore).

### Mast cell culture and stimulation

CBMCs were derived and then maintained, as previously described.<sup>E2,E3</sup> CBMCs were preincubated with 10 ng/mL human IL-4 (PeproTech) for 4 days and received 1 μg/mL IgE (Calbiochem, Merck Millipore, Darmstadt, Germany) overnight before activation. For activation, cells (250,000 cells/well; equivalent to  $1.25 \times 10^6$  cells/mL) were activated in IMDM supplemented with 1% normal human serum (NHS) and penicillin (100 U/mL) and streptomycin (0.1 mg/mL) by using either 2 μg/mL anti-human IgE (Sigma-Aldrich) for 6 hours or 2 μmol/L calcium ionophore A23187 (Sigma-Aldrich) for 2 hours. Supernatants were collected and stored at  $-80^\circ\text{C}$ .

### ELISA

Concentrations of released IL-5 and IL-13 in ILC2 supernatants were assessed by means of ELISA. We used the human IL-5 ELISA Duo-Set Kit (R&D Systems) and the human IL-13 ELISA kit (Sanquin, Amsterdam The Netherlands).

In some experiments PGD<sub>2</sub> concentrations in ILC2 supernatants were measured by using the ELISA Prostaglandin D<sub>2</sub>-MOX ELISA Kit (Cayman Chemicals). PGD<sub>2</sub> was converted to a stable methoxime derivative by mixing cell supernatants or standards with methyl oximating reagents (1:1) and heated to 60°C for 30 minutes. Afterward, PGD<sub>2</sub>-methoxime and standards were used for ELISA, and their concentrations were determined spectrophotometrically.

### LC-MS/MS analysis of prostanoids and metabolites

Analysis of cell-culture supernatants was performed by using LC-MS/MS. All analytic and internal standards were purchased from Cayman Chemical. Solvents (LC-MS grade) were obtained from Carl Roth (Karlsruhe, Germany).

For extraction of prostanoids and metabolites, 200 μL of cell-culture supernatant were spiked with isotopically labeled internal standards (PGE<sub>2</sub>-d<sub>4</sub>, PGD<sub>2</sub>-d<sub>4</sub>, TXB<sub>2</sub>-d<sub>4</sub>, PGF<sub>2α</sub>-d<sub>4</sub>, 6-keto PGF<sub>1α</sub>-d<sub>4</sub>, 20-HETE-d<sub>6</sub>, and

13,14-dh-15-keto PGE<sub>2</sub>), 100  $\mu$ L of EDTA solution (0.15 mol/L), and 600  $\mu$ L of ethyl acetate. Samples were vortexed and centrifuged at 20,000g for 5 minutes. The organic phase was removed, and the extraction was repeated with 600  $\mu$ L of ethyl acetate. Organic fractions were combined and evaporated at a temperature of 45°C under a gentle stream of nitrogen. Residues were reconstituted with 50  $\mu$ L of acetonitrile/water/formic acid (20:80:0.0025 vol/vol/vol) and transferred to glass vials.

The LC-MS/MS analysis was carried out with an Agilent 1290 Infinity LC system (Agilent, Waldbronn, Germany) coupled to a hybrid triple quadrupole linear ion trap mass spectrometer QTRAP 6500+ (Sciex, Darmstadt, Germany) equipped with a Turbo-V-source operating in negative electrospray ionization mode. Chromatographic separation was carried out with a Synergi Hydro-RP Column (150  $\times$  2 mm, 4- $\mu$ m particle size, and 80-Å pore size; Phenomenex, Aschaffenburg, Germany). A linear gradient was used at a flow rate of 300  $\mu$ L/min. Mobile phase A was water/formic acid (100:0.0025 vol/vol), and mobile phase B was acetonitrile/formic acid (100:0.0025 vol/vol). Analytes were separated under gradient conditions within 16 minutes. The injection volume was 20  $\mu$ L. The gradient program started with 90% mobile phase A for 1 minute, and then mobile phase A was decreased to 60% within 1 minute, held for 1 minute, further decreased to 50% within 1 minute, and held for 2 minutes. Within 2 minutes, mobile phase A was further decreased to 10% and held for 1 minute. Within 1 minute, the initial conditions were restored and the column was re-equilibrated for 6 minutes. Mass spectrometric parameters were set as follows: ionspray voltage, -4500 V; source temperature, 500°C, curtain gas, 40 psi; nebulizer gas, 40 psi; and Turbo heater gas, 60 psi. Both quadrupoles were running at unit resolution.

For analysis and quantification, Analyst Software 1.6 and MultiQuant Software 3.0 (both from Sciex) were used. The following precursor-to-product ion transitions were used for quantification: m/z 351.2  $\rightarrow$  m/z 315.0 for PGE<sub>2</sub>, m/z 351.2  $\rightarrow$  m/z 233.3 for PGD<sub>2</sub>, m/z 353.2  $\rightarrow$  m/z 193.0 for PGF<sub>2 $\alpha$</sub> , m/z 369.2  $\rightarrow$  m/z 163.0 for 6-keto PGF<sub>1 $\alpha$</sub> , m/z 369.2  $\rightarrow$  m/z 195.0 for TXB<sub>2</sub>, m/z 279.1  $\rightarrow$  m/z 178.9 for 12-hydroxyheptadecatrienoic acid, m/z 353.2  $\rightarrow$  m/z 309.2 for 11 $\beta$ -PGF<sub>2 $\alpha$</sub> , m/z 351.2  $\rightarrow$  m/z 333.2 for 13,14-dh-15-keto-PGD<sub>2</sub>, m/z 333.2  $\rightarrow$  m/z 233.3 for PGJ<sub>2</sub>, and m/z 333.2  $\rightarrow$  m/z 233.3 for  $\Delta$ <sup>12</sup>-PGJ<sub>2</sub>. The peak area of each analyte was corrected by the peak area of the corresponding internal standard. Calibration curves were constructed

by using linear regression with  $1/x^2$  weighting. The coefficient of correlation was at least 0.99. Variations in accuracy were less than 15% over the whole range of calibration, except for the lowest limit of quantification, where a variation in accuracy of 20% was accepted. The calibration ranges were 0.02 to 10 ng/mL plasma for PGE<sub>2</sub>, PGD<sub>2</sub>, TXB<sub>2</sub>, 13,14-dh-15keto-PGD<sub>2</sub>, PGJ<sub>2</sub>, and  $\Delta$ <sup>12</sup>-PGJ<sub>2</sub>; 0.04 to 20 ng/mL plasma for PGF<sub>2 $\alpha$</sub> ; 0.08 to 40 ng/mL plasma for 6-keto PGF<sub>1 $\alpha$</sub> ; 0.1 to 10 ng/mL plasma for 11 $\beta$ -PGF<sub>2 $\alpha$</sub> ; and 0.2 to 10 ng/mL plasma for 12-hydroxyheptadecatrienoic acid.

## RT-qPCR

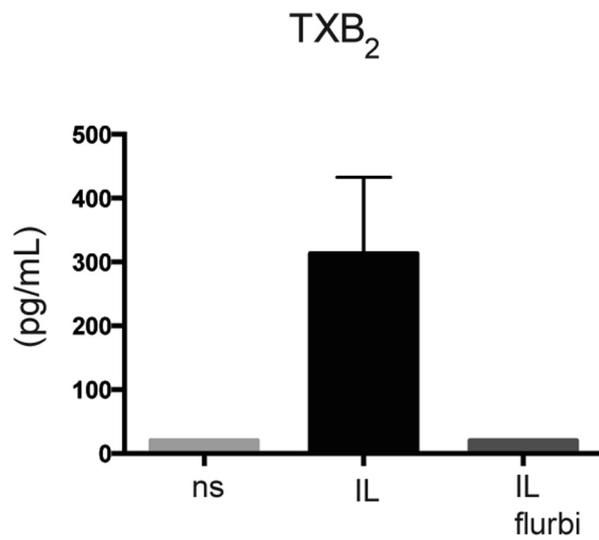
Total RNA from expanded tonsillar ILC2s was isolated with the RNeasy Micro Kit (Qiagen). The iScript cDNA Synthesis Kit (Bio-Rad Laboratories) was used for cDNA synthesis, and SsoAdvanced Universal SYBR Green Supermix (Bio-Rad Laboratories) was used for PCR reactions. Primers were the PrimePCR SYBR Green assay for human *PTGS2* (for COX-2), *HPGDS*, *HPGD*, *PTGDR* (for DP1), *GPR44* (for CRTH2), and *RPS18*; all primers were from Bio-Rad Laboratories. RT-qPCR was performed in the CFX Connect Real-Time PCR Detection System (Bio-Rad Laboratories).

## Statistical analyses

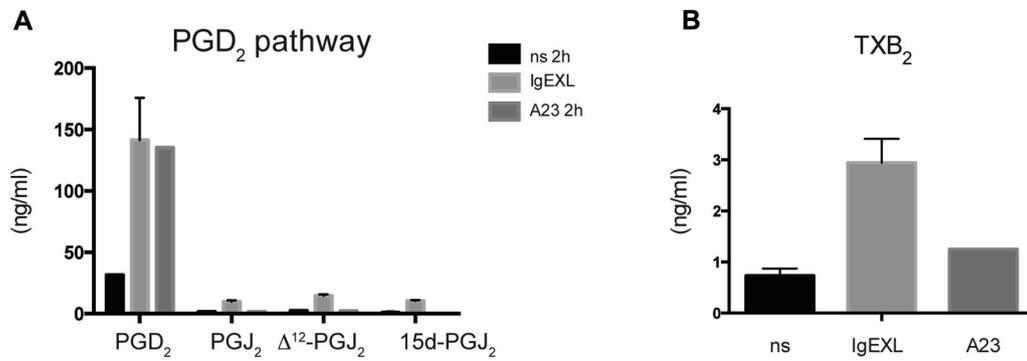
Differences between 2 groups were analyzed by using the matched-pairs *t* test, and 3 or more groups were compared by using 1-way ANOVA for repeated measurements and the Dunnett multiple comparisons test. Analyses were performed with GraphPad Prism 6 Software.

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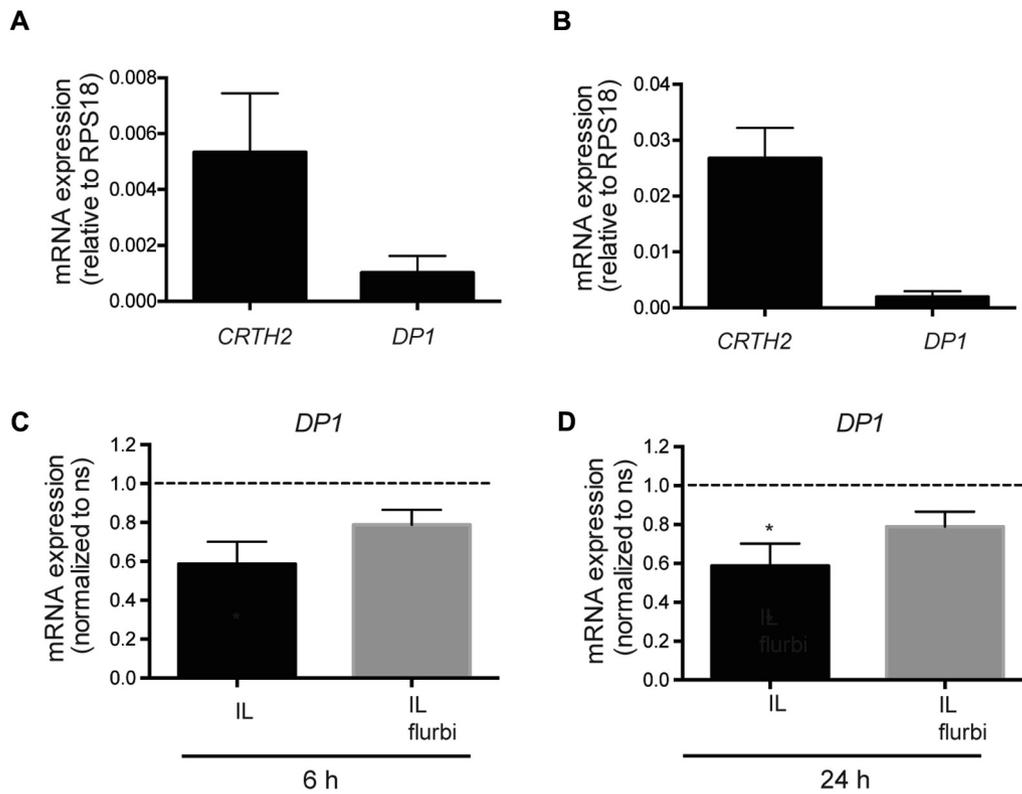
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**FIG E1.** Stimulated ILC2s produce TXB<sub>2</sub>. Sorted and expanded tonsillar ILC2s were stimulated as indicated for 24 hours. TXB<sub>2</sub> was detected in ILC2 supernatants by using LC-MS/MS. The graph shows mean + SEM concentrations (4 donors). *flurbi*, Flurbiprofen; *IL*, IL-33, IL-25, and TSLP; *ns*, nonstimulated.



**FIG E2.** Metabolites of the PGD<sub>2</sub> pathway and TXB<sub>2</sub> are released by activated human mast cells. CBMCs were stimulated with IgEXL for 6 hours or A23 for 2 hours. Mediators of the PGD<sub>2</sub> pathways (**A**) and TXB<sub>2</sub> (**B**) were detected in mast cell supernatants by using LC-MS/MS. Graphs show mean + SEM concentrations (n = 1-3). *ns*, Nonstimulated.



**FIG E3.** ILC2s express lower mRNA levels of *DP1* than *CRTH2* receptor. Sorted and expanded tonsillar ILC2s were stimulated as indicated for 6 (**A** and **C**) and 24 (**B** and **D**) with or without flurbiprofen (1  $\mu\text{mol/L}$ , 40 minutes before treatment). Basal mRNA expression of *CRTH2* and *DP1* after 6 (Fig E3, **A**) and 24 (Fig E3, **B**) hours of culture were analyzed by using PCR. Data are shown as means + SEMs (2 donors for *DP1* data for the 6-hour time point, 3 donors for *CRTH2* data for 6 hours, and 4 donors for 24 hours of culture for both receptors). Changes of *DP1* mRNA expression normalized to nonstimulated cells after 6 (2 donors; Fig E3, **C**) and after 24 (\* $P < .05$ , 4 donors; Fig E3, **D**) hours of stimulation are shown as means + SEMs. *flurbi*, Flurbiprofen; *IL*, IL-33, IL-25, and TSLP; *ns*, nonstimulated.