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Group V secretory phospholipase A₂ is involved in macrophage activation and is sufficient for macrophage effector functions in allergic pulmonary inflammation

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Abstract

We reported that *Pla2g5*-null mice lacking group V secretory phospholipase A₂ (gV-sPLA₂) showed reduced eosinophilic pulmonary inflammation and Th2 cytokine generation when challenged with an extract (*Df*) from house dust mite *Dermatophagoides farinae*, compared to wild-type (WT) controls. Adoptive transfer studies suggested that gV-sPLA₂ in dendritic cells (DCs) was necessary for sensitization of *Pla2g5*-null mice, but was not sufficient to induce the effector phase of pulmonary inflammation. Here, we demonstrate that gV-sPLA₂ is inducibly expressed in mouse and human macrophages (Mφ activated by IL-4, and is required for the acquisition of Mφ effector functions that facilitate the effector phase of pulmonary inflammation. We demonstrate that gV-sPLA₂ expression in Mφ is sufficient for the development of pulmonary inflammation, even when inflammation is induced by intrapulmonary administration of IL-4. The concentrations of CCL22/CCL17 and effector T-cell recruitment are severely impaired in *Pla2g5*-null mice. Intratracheal transfers of enriched CD68⁺ cells isolated from the lungs of *Df*-challenged WT donor mice induce eosinophilia, chemokine production, and recruitment of T-cells into the lungs of *Pla2g5*-null recipients previously sensitized by WT *Df*-loaded DCs. Our studies identified a unique function of gV-sPLA₂ in activation of Mφ and in their capacity to recruit T-cells to amplify the effector phase of pulmonary inflammation.

Introduction

Phospholipases A₂ (PLA₂) are a family of enzymes that generate free fatty acids and lysophospholipids from membrane phospholipids (1). PLA₂-liberated arachidonic acid provides the substrate for the synthesis of eicosanoids (prostaglandins and leukotrienes), mediators of inflammation and allergic disorders (2, 3). Whereas the group IVA or cytosolic (c)PLA₂α is exclusively intracellular and absolutely required for the generation of

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arachidonic acid (4), the secretory (s)PLA₂s likely serve both intracellular and/or extracellular functions (1, 5) some in a cell-specific fashion (6–8). Studies in humans suggest that sPLA₂s can be inducibly expressed and activated during allergic pulmonary inflammation in asthma (9), mostly gV-sPLA₂ (10), gIIA- and gX-sPLA₂s (10, 11). Mouse models of pulmonary inflammation induced by systemic sensitization and inhalation challenge with OVA suggest that gV-sPLA₂ and gX-sPLA₂ may be functionally important in allergic airway disease (12, 13). However neither the cellular sources of the enzymes nor their distinguishing functions are known.

M ϕ are a heterogeneous cell lineage (14–17). Immune responses can direct macrophage development into classical activated (Ca) and alternative activated (AA) macrophages (M ϕ) (18) or M1 and M2 (19). CaM ϕ are associated with Th1-type immunity and are involved in killing of pathogens. AAM ϕ in vivo acquire functional characteristics to serve context-specific roles in homeostasis, inflammation, and fibrosis with reports of both proinflammatory and immunoregulatory effects (20–23). AAM ϕ are associated with Th2-biased immune responses such as those occurring in responses to helminths and allergens (20, 23–26), likely reflecting the polarizing effects of the microenvironment in which they develop (27, 28). The relative expression of molecules related to alternative activation of M ϕ may also be context dependent and not limited to M ϕ (25, 29, 30). Molecules linked to alternative activation of M ϕ (Resistin-like molecule- α (Relm- α), Arginase-1 (Arg-1), Ym1/Chitinase3-like3, are found in the lungs of OVA-sensitized and challenged mice (20, 25, 31, 32). Related molecules (Chitinase1, Chitinase3-like1, Stabilin1) are found in humans with asthma and COPD (33–35). However, the function of M ϕ especially in a complex disease like asthma is still controversial (26, 36).

We recently showed using a model of allergic pulmonary inflammation induced by an extract (*Df*) from the house dust mite *Dermatophagoides farinae* (37) that while DCs require the functions of gV-sPLA₂ for effective antigen uptake and Th2 priming ability, gV-sPLA₂ in other cell type(s) is required for the development of *Df*-induced pulmonary pathology. Because gV-sPLA₂ is critical for certain macrophage functions (38, 39) and M ϕ have been associated to antigen-induced Th2 responses, we hypothesized that endogenous expression of gV-sPLA₂ by M ϕ might contribute to the induction of *Df*-mediated pulmonary inflammation.

Indeed, we now demonstrate that gV-sPLA₂ expression by M ϕ can facilitate allergic pulmonary inflammation in *Df*-sensitized *Pla2g5*-null recipients, or in IL-4-treated *Pla2g5*-null mice. GV-sPLA₂ expression is induced during both human and mouse IL-4 mediated activation of M ϕ in vitro, and its absence impairs M ϕ activation in vivo, independently of effects on the adaptive immune response. Moreover, whereas gV-sPLA₂-expressing DCs can initiate Th2 responses in the lymph-nodes of *Pla2g5*-null recipients (37), gV-sPLA₂-expressing M ϕ are sufficient to induce the generation of CCL22 and CCL17, pulmonary T-cell recruitment and eosinophilic inflammation in response to *Df*. GV-sPLA₂ is thus a novel cross-species mediator of M ϕ activation and regulator of M ϕ effector functions, and may prove to be a fruitful therapeutic target in asthma.

Materials and Methods

Df- and IL-4-induced pulmonary inflammation

C57BL/6 WT and *Pla2g5*-null mice (40) (9– to 12-wk old males) received 3 μ g of *Df* extract (Greer Laboratories, Lenoir, NC) in 20 μ l NaCl 0.9% (containing <0.005 endotoxin/ml, Sigma-Aldrich, St. Louis, MO) or saline alone, intranasally on days 0, 4, 7, 11, 14 and 18 (37). For IL-4 induced pulmonary inflammation, WT and *Pla2g5*-null naïve mice were given mouse r-IL-4 (1 or 5 μ g/dose) (PeproTech, Rocky Hill, NJ) intratracheally (i.t.) on

days 1, 2, and 5 and mice were euthanized 36 h after the last dose (41). For blocking experiments mouse CCL22 and CCL17 polyclonal neutralizing antibodies (R&D Systems, Minneapolis, MN) were administered i.p. (20 µg/dose in 500 µl of NaCl 0.9%) 30 min before the last four doses of *Df*. BAL was performed with 0.7 ml of PBS (Sigma-Aldrich) containing 0.5 mM EDTA (three times). The BAL fluid was collected and the cell pellet was counted, cytopun and stained with Diff-Quick (Fisher Diagnostic, Middletown, VA) (37). ELISAs were used to measure CCL22 and CCL17 (R&D Systems), as well as IL-5 and IL-13 (eBiosciences) in cell-free BAL fluid.

All animal experiments were approved by the Animal Care and Use Committee of the Dana-Farber Cancer Institute (Boston, MA).

Immunohistochemical and immunofluorescence analysis

For immunohistochemical and immunofluorescence analysis, left lungs were collected from each mouse after BAL and fixed with 4% paraformaldehyde for at least 8 h. The lungs were washed twice with washing buffer (PBS containing 2% DMSO), and suspended in 50 mM of NH₄Cl overnight at 4°C. Then the lungs were embedded in paraffin or glycolmethacrylate. Lung sections were deparaffinized and rehydrated and antigen retrieval was performed with Target Retrieval Solution (Dako, Glostrup, Denmark) at 37°C for 30 min. After blocking with 10% chicken serum, samples were stained (1 h) with rabbit anti-mouse Relm-α (PeproTech), then washed and incubated (1 h) with Alexa-Fluor-594 chicken anti-rabbit IgG and Hoechst nuclear staining (Invitrogen, Carlsbad, CA). The sections were washed mounted and imaged with an Eclipse 80i microscope (Nikon Instruments, Melville, NY) (37).

Enrichment of *Df*-induced lung Mφ

After removing airway cells and alveolar Mφ by extensive BAL with PBS/0.5mM EDTA, lungs were homogenized through a 70-µm mesh in ice-cold complete medium (37) (RPMI 1640, 10% FBS, 100 U/ml penicillin, 100 µg/ml streptomycin, 0.1 mM non-essential amino acids, 2 mM L-glutamine, 0.05 µM 2-ME). The homogenate was then washed and incubated (30 min, 37°C) with 428 U/ml Collagenase IV (Worthington Lakewood, NJ) and 20 µg/ml DNase I (Roche, Mannheim, Germany). The obtained cell suspension from pooled mice was washed and layered onto a Percoll® gradient (Sigma-Aldrich) (60% and 40%). After centrifugation at 600 g for 20 min at 37°C, the cells at each interface were collected, washed, and centrifuged again onto a second density gradient. At the end of the procedure, the obtained cells represented an enriched (70–80%) population of lung Mφ (CD68^{enr}) as determined by flow cytometry. Total RNA was isolated from the cell lysate with TRI Reagent (Sigma-Aldrich), reverse transcribed into cDNA (High-Capacity cDNA Reverse Transcription Kit, Applied Biosystems) and measured by real-time quantitative PCR (qPCR) with the use of SYBR Green/ROX master mix (SABiosciences, Frederick, MD) on an Mx3005P thermal cycler (Stratagene, Santa Clara, CA) (37). The ratio of each mRNA relative to the GAPDH mRNA was calculated with the $\Delta\Delta C_T$ method. The primers are described in Table I.

Bone marrow-derived Mφ

WT and *Pla2g5*-null or C57BL/6 and IL-2Rγ-null bone marrow (BM) cells were collected from femurs and tibiae of mice. The disaggregated cells were counted and suspended in complete medium containing 50 ng/ml murine r-M-CSF (PeproTech) at a concentration of 4.0×10^5 cells/ml, 10 ml/Petri dish. On day 3, 10 ml of medium containing r-M-CSF were added. On day 7, the cells were pulsed with 20 ng/ml of IL-4 (PeproTech) for BM-IL-4-Mφ (a surrogate of AAMφ), LPS (100 ng/ml) (Sigma-Aldrich) and IFN-γ (200 U/ml) (PeproTech) for BM-LPS/IFNγ-Mφ (a surrogate of CaMφ), cultured in medium (BM-

Immature-M ϕ , BM-IM-M ϕ), or IL-4 and/or IL-13, IL-33 and GM-CSF (PeproTech). After 24 h, cells were harvested with PBS containing Lidocaine (4 mg/ml) and EDTA (5 mM) (15 min, 37°C) (39) and analyzed by flow cytometry. For adoptive transfer, 1×10^5 WT and *Pla2g5*-null BM-IM-M ϕ were transferred i.t. into naïve WT and *Pla2g5*-null mice at day 0, followed by 5 μ g/dose of IL-4 i.t. on days 1, 2, and 5.

Human monocyte-derived M ϕ

Leukocyte-enriched buffy coat from healthy donors was overlaid on Ficoll-Paque Plus (GE Healthcare, Buckinghamshire, UK) and centrifuged at 600 *g* for 20 min. The mononuclear layer at the interface was collected, washed, and counted. Monocytes were isolated by negative selection (Miltenyi Biotec, Auburn, CA) and plated at $1-1.5 \times 10^6$ cells/ml in 30-mm Petri dishes. The cells were cultured for 7 days in complete medium supplemented with human r-M-CSF (50ng/ml) (R&D, Minneapolis, MN). To activate monocyte-derived M ϕ (42), cells were polarized for 24h in complete medium, without M-CSF, supplemented with IL-4 (R&D) (20ng/ml) (h-IL-4-M ϕ) as a surrogate of human AAM ϕ or LPS (Sigma Aldrich) (100ng/ml) and INF- γ (R&D) (20ng/ml) (h-LPS/INF γ -M ϕ) as a surrogate of human classical activated M ϕ .

To knock down group V sPLA₂, after culturing the monocytes for 7 days in M-CSF, the cells were harvested by incubation with Lidocaine/EDTA, washed, and counted. M ϕ were transfected with human *Pla2g5* ON-TARGET Plus siRNA or non-targeting vector control (Dharmacon, 100nM and 1000nM) using the Amaxa® Human Macrophage Nucleofector® kit (Lonza, Germany), according to the manufacturer's instructions. After 24 h, the transfection medium was replaced by complete medium with or without IL-4 (20ng/ml) for polarization. Twenty-four hours later, the adherent cells were harvested by incubation with Lidocaine/EDTA and analyzed by qPCR and flow cytometry. QPCR was performed as described above. The primers are described in Table I.

Flow cytometry

Lung cells were fixed with 4% paraformaldehyde (7 min, 21°C), washed, and permeabilized with 0.1% saponin (7 min, 21°C). After washing, cells were blocked (1 h, 4°C) with 1% rat anti mouse CD16/CD32 (BD Biosciences, San Jose, CA) and 10% donkey serum, and then stained (1 h, 4°C) with appropriate antibodies. Mouse cells were stained with Dectin1-FITC (clone 2A11), CD68-APC (clone FA11), CD206-FITC (clone MR5D3) (AbD Serotec, Raleigh, NC), CD11c-PECy7 (clone N418), CD3-PECy7 (clone 145-2C11), B220-PerCP-Cy5.6 (clone RA3-6B2) (eBioscience, San Diego, CA), CD11b-FITC or -PE-Cy7 (clone M1/70), Ly6G-PE (clone 1A8), Ly6C-APC-Cy7 (clone HK1.4), SiglecF-PE (clone E50-2440), CD36-FITC (clone HM36), F4/80-APC-Cy7 (clone BM8) (BioLegend, San Diego, CA), CD4-PECy7 (clone RM4-5) (BD Biosciences) and corresponding isotypes as controls. Cells stained with rabbit antibody Relm- α (PeproTech) or rabbit IgG as control were washed and stained with Alexa488 donkey anti-rabbit IgG (1:200) (Jackson ImmunoResearch, West Grove, PA). Human cells were stained with: CD11b-PE-Cy7 (clone ICRF44), CD209 (clone 9E9A8) (BioLegend), CD206-FITC (clone 19.2) (BD Biosciences), CD36-FITC (clone SM ϕ) (AbD Serotec), and corresponding isotypes as controls. Acquisition was performed on a FACSCanto flow cytometer with FACSDiva software (BD Biosciences) and data were analyzed with FlowJo (Tree Star, Ashland, OR).

Generation of chimeric mice

Three-wk-old recipient mice were lethally irradiated with two doses of 600 rad each (separated by 4 h). BM from donor mice was harvested from femora and tibiae, and 10×10^6 cells in 0.2 ml PBS were injected i.v. into recipient mice. BM transplantation was performed in four groups of mice: BM from *Pla2g5*-null into WT (expressing group V

sPLA₂ on non-hematopoietic cells only); BM from WT into *Pla2g5*-null mice (expressing group V sPLA₂ on hematopoietic cells only); BM from *Pla2g5*-null to *Pla2g5*-null (expressing no group V sPLA₂); and BM from WT to WT. Mice in the latter two groups served as negative and positive controls for possible radiation effects. To permit complete chimerism, we allowed 10 weeks of reconstitution time before we started experiments (43).

Transfer of CD68^{enr}/*Df*-lung-cells into DC-sensitized recipients

At day 0 recipient mice were sensitized to *Df* by transferring intranasally WT *Df*-pulsed BM-derived DCs (*Df*-BMDCs) (5×10^4 i.t.) (37). At day 8, lung cells were harvested from *Df*-challenged (3 μ g/dose, 6 doses) WT donor mice. Lung cells were enriched over three Percoll® gradients to have a population of >85% CD68⁺ M ϕ (Supplemental Fig. 3) and transferred (1×10^5) i.t. into sensitized WT and *Pla2g5*-null recipient mice. The recipient mice were challenged with *Df* intranasally at day 9 and 12, and euthanized 36 h later (Fig. 6 inset).

Intracellular cytokine generation by T-cells

Lungs and lymph nodes were mechanically dispersed and filtered through a 70- μ m cell strainer. Mononuclear cells were isolated on a NycoPrep density gradient (Axis-Shield) at 600 g for 20 min. The mononuclear cells were washed and plated at 1×10^6 cells/ml in 24-well plates in RPMI containing 10% FBS, 100U/ml penicillin, 100 μ g/ml streptomycin, 0.1mM non-essential amino acids, 2mM L-glutamine, 0.043 mM 2-ME, 0.025M Hepes buffer, and 1mM sodium Pyruvate. The cells were stimulated for 6 h at 37°C with PMA (50 ng/ml) and ionomycin (1 μ M) to induce cytokine production. To stop exocytosis we added monensin (2.5 μ M) to the culture 4 h before collecting the cells. The cells were then treated with DNase I (60 μ g/ml) for 15 min at 37°C. After washing, the cells were resuspended in fixation buffer (eBioscience) for 15 min, then washed and blocked in permeabilization buffer (eBioscience) containing 1% mouse IgG (Sigma-Aldrich) and 1% rat anti mouse CD16/CD32 (BD) for 20 min. Cells were stained for FACS analysis with anti-mouse CD4-PE-Cy7 (clone RM4-5), CD8 α -allophycocyanin (clone 53-6.7), IL-4-PE (clone 11B11), IL-5-PE (clone TRFK5), IL-17A-PE (clone TC11-18H10) (BD), IFN- γ -PE (clone XMG1.2) (eBioscience), or rat IgG1-PE isotype Ab (BD) (44). The acquisition was performed on a FACSCanto flow cytometer, and data were analyzed with FlowJo.

Statistical analysis

To compare dose-dependent induction of pulmonary inflammation by IL-4 in WT and *Pla2g5*-null mice, we performed two-way ANOVA with Tukey correction for multiple comparisons. To compare expression of gV-sPLA₂ mRNA in in vitro-derived M ϕ (three groups), we used one-way ANOVA with Dunnett's correction for multiple comparisons. Both comparisons were performed with Prism software (GraphPad Software, La Jolla, CA). Comparisons between two groups were made using unpaired Student's *t* test. Data are expressed as mean \pm SEM, and $P < 0.05$ was considered significant.

Results

GV-sPLA₂ contributes to activation of M ϕ in vivo during *Df*-induced pulmonary inflammation

Our previous study showed that gV-sPLA₂ was expressed in the epithelium and in cells infiltrating the lung of *Df*-treated mice, including cells with the morphology of M ϕ (37). To start to investigate whether gV-sPLA₂ is involved in the activation and functions of M ϕ generated during *Df*-induced pulmonary inflammation, we analyzed the expression of Relm- α by immunofluorescence. The lungs of *Df*-treated WT mice showed strong staining for

Relm- α on epithelial cells (Fig. 1Aii, thick arrow) and on infiltrating cells with the morphologies typical of M ϕ and eosinophils (Fig. 1Bi, large and small arrowhead, respectively). In contrast, Relm- α staining was not observed on cells infiltrating the lungs of *Df*-treated *Pla2g5*-null mice (Fig. 1Bii), but did localize to the epithelium with a distribution and intensity similar to the WT controls (Fig. 1Aiv, thin arrow). Neither genotype showed Relm- α staining in the lungs of saline-treated mice (Fig. 1Ai,iii).

To confirm that M ϕ from the lungs of *Df*-treated *Pla2g5*-null mice lacked Relm- α , and to determine whether they also differed in their expression of other molecules related to macrophage activation, M ϕ were enriched from dispersed lung cells of *Df*-treated WT and *Pla2g5*-null mice (CD68^{enr}/*Df*-lung-cells) and analyzed by flow cytometry. Forward and side scatter analysis showed three or two distinct populations respectively in WT and *Pla2g5*-null mice (Supplemental Fig. 1, respectively upper inset and lower inset): both genotypes showed population (a) of which more than 90% were CD68⁺ and expressed CD11c (Supplemental Fig. 1A,D). In both genotypes, less than 10% of the CD68⁺ cells were in population (b) showing small forward and small side scatter (Supplemental Fig. 1B, E); of the CD68⁻ cells less than 5% expressed CD3 or B220 (data not shown). Only WT mice showed population (c) with homogeneous forward scatter and high side scatter, characteristic of granulocytes (Supplemental Fig. 1C). Similar numbers of CD68⁺ cells were recovered from the lungs of *Df*-challenged WT and *Pla2g5*-null mice (respectively 1.52 ± 0.04 and $1.02 \pm 0.06 \times 10^5$ from four experiments, data not shown) expressing similar levels of CD11c ($67.78 \pm 6.8\%$ and $66.29 \pm 0.41\%$, respectively) (data not shown). The CD68⁺ cells of *Df*-treated WT mice expressed significantly more Relm- α protein than CD68⁺ cells of *Df*-treated *Pla2g5*-null mice (Fig. 1D); Dectin1 also showed a similar but more modest trend. The expression of CD206 was similar in both genotypes (data not shown). The qPCR of the CD68^{enr}/*Df*-lung-cells from *Pla2g5*-null mice showed significantly lower expression of mRNAs encoding CCL22 and Arg-1 compared to CD68^{enr}/*Df*-lung-cells from WT mice (Fig. 1E), with similar trends for matrix metalloproteinase-12 (MMP12), CCL17, Ym1, and CCL11. There were no differences in mRNA encoding inducible nitric oxide synthase (iNOS) and CXCL10 (Fig. 1E).

The expression of gV-sPLA₂ in hematopoietic cells is sufficient to develop major features of *Df*-induced pulmonary inflammation

Because gV-sPLA₂ is expressed on epithelial cells and cells infiltrating the lungs of *Df*-treated mice (37), we generated chimeric mice to determine whether gV-sPLA₂ expression by hematopoietic cells or by resident cells was more essential for development of *Df*-induced pulmonary inflammation and/or the activation of M ϕ . Irradiated WT and *Pla2g5*-null mice were reciprocally transferred BM from both genotypes and after 10 weeks exposed to *Df*. WBC counts showed equivalent engraftment efficiencies among the groups (data not shown). *Df*-treated WT mice receiving *Pla2g5*-null-BM did not develop BAL fluid eosinophilia (Fig. 2A). In contrast, *Df*-treated *Pla2g5*-null mice engrafted with WT-BM developed pulmonary inflammation and BAL fluid eosinophilia to a level that exceeded that of the WT mice reconstituted with WT-BM (Fig. 2B). The *Pla2g5*-null mice engrafted with *Pla2g5*-null-BM did not develop inflammation in response to *Df* (Fig. 2B). CD68^{enr}/*Df*-lung-cells from engrafted mice that received WT-BM showed increased expression of Relm- α and Dectin1 following *Df*-treatment, independently of the genotype of the host (Fig. 2C–D). In marked contrast, CD68⁺ cells in mice receiving *Pla2g5*-null-BM lacked Relm- α and expressed less Dectin-1 than in mice receiving WT BM, regardless of host genotype.

GV-sPLA₂ contributes to IL-4-induced pulmonary eosinophilia and activation of M ϕ in vivo

In the absence of gV-sPLA₂, *Df*-treated mice showed deficient expression of Th2 cytokines in the lung (37). Because M ϕ are activated in Th2 environments, we sought to determine

whether endogenously expressed gV-sPLA₂ contributes to activation of M ϕ in vivo independently of the adaptive immune system (41). We instilled r-IL-4 intratracheally (1 and 5 μ g/dose) to directly induce the effector pathways associated with a Th2 response without requirement of sensitization or T-cell activation. In WT mice, both doses of IL-4 significantly increased the number of cells recruited into the airways, as well as eosinophils (5 μ g/dose) and mononuclear cells (1 μ g/dose). These responses were significantly diminished in the *Pla2g5*-null mice (Fig. 3A). The numbers of neutrophils and M ϕ were similar in the two genotypes. To verify the effect of IL-4-activation on M ϕ in vivo, we analyzed the expression of Relm- α and Dectin1 in WT and *Pla2g5*-null lung CD68⁺ cells. IL-4 at 5 μ g induced the expression of Relm- α by WT CD68⁺ cells (Fig. 3B), but not by *Pla2g5*-null CD68⁺ cells. WT and *Pla2g5*-null CD68⁺ cells from saline- and IL-4- (1 μ g) treated mice did not express Relm- α while the expression of Dectin1 was similar in IL-4-treated WT and *Pla2g5*-null CD68⁺ cells (data not shown). Both genotypes showed minimal induction and similar amounts of IL-5 and IL-13 in the BAL after IL-4 treatment (data not shown).

GV-sPLA₂ is directly induced by IL-4 in human and mouse in vitro-derived M ϕ , and is involved in M ϕ activation in vitro

Next we asked whether the in vivo phenotype of *Df*-induced macrophages could be replicated in vitro. WT and *Pla2g5*-null BM-M ϕ cultured with r-M-CSF (BM-IM-M ϕ), were polarized with IL-4 (BM-IL-4-M ϕ , a surrogate of AAM ϕ) or LPS and IFN- γ (BM-LPS/IFN γ -M ϕ , a surrogate of CaM ϕ) (Fig. 4 inset). As determined by qPCR, gV-sPLA₂ mRNA was virtually undetectable in WT BM-IM-M ϕ and WT BM-LPS/IFN γ -M ϕ , but was sharply upregulated in WT BM-IL-4-M ϕ (Fig. 4A). To determine whether the lack of gV-sPLA₂ in in vitro-derived M ϕ altered the ability of IL-4 to induce the expression of markers of alternative activation (18, 19), we analyzed WT and *Pla2g5*-null BM-IL-4-M ϕ by flow cytometry. The expression of CD206 was significantly higher on WT than on *Pla2g5*-null BM-IL-4-M ϕ , with similar trends for CD36 and Dectin1 (Fig. 4B). Relm- α expression was weak and inconsistently detected in BM-IL-4-M ϕ while other markers showed similar expression (data not shown). Stimulation of WT BM-IM-M ϕ with IL-13 induced gV-sPLA₂ expression, but less potently than IL-4 (Supplemental Fig. 2A). The induction of gV-sPLA₂ expression by IL-4 was absent in BM-IM-M ϕ lacking the common γ -chain of the IL-2 receptor (IL-2R γ -null mice) (Supplemental Fig. 2B, left panel). The expression of CD206 was also significantly less inducible in the IL-2R γ -null BM-IL-4-M ϕ than in the WT control cells (Supplemental Fig. 2B, right panel). Expression of the M ϕ associated chemokines CCL17 and CCL22 mRNA was not induced by IL-4 alone in either genotype. However, the addition of IL-33 (20), in combination with GM-CSF (45) during the IL-4-induced polarization step resulted in robust induction of CCL22 mRNA expression. This induction was substantially blunted in the *Pla2g5*-null BM-IL-4-M ϕ treated with GM-CSF and IL-33 (Fig. 4C). IL-4/IL-33/GM-CSF did not increase the expression of other markers of alternative activation (including CCL17), nor did it further increase the expression of group V sPLA₂ mRNA compared with IL-4 alone (data not shown).

To determine whether the expression of gV-sPLA₂ was induced by IL-4-polarization of human- (h)-monocyte-derived M ϕ , M ϕ were activated by IL-4 (h-IL-4-M ϕ) or LPS/IFN- γ (h-LPS/IFN γ -M ϕ) and analyzed for the expression of gV-sPLA₂ by qPCR. Freshly isolated monocytes and h-monocyte-derived M ϕ lacked gV-sPLA₂ mRNA (Fig. 4D and data not shown). However, gV-sPLA₂ mRNA was significantly induced in h-IL-4-M ϕ but not in h-LPS/IFN γ -M ϕ (Fig. 4C). IL-13 polarization of h-monocyte-derived M ϕ modestly induced the expression of gV-sPLA₂ (Supplemental Fig. 2C). KD of gV-sPLA₂ in h-IL-4-M ϕ (Supplemental Fig. 2D) significantly reduced the expression of CD209 (or Dendritic Cell-Specific Intercellular adhesion molecule-3-Grabbing Non-integrin (DC-SIGN), a molecule

associated with IL-4 activation of human macrophages) (42) compared to vector-treated cells, while the effects on CD206 and CD36 were not significant (Fig. 4E).

The lack of group V sPLA₂ results in reduced production of Mφ-associated CCL17 and CCL22 that are necessary for effector T-cell recruitment

Next we wanted to determine whether in vivo group V sPLA₂ would permit the generation of mediators that could amplify pulmonary inflammation. Because *Pla2g5*-null CD68^{enr}/*Df*-lung-cells, had reduced mRNA expression for CCL22 and CCL17 (Fig. 1 E), two Th2-active chemokines that can be produced by Mφ in vivo (19), and because this phenotype could be partially replicated in vitro (Fig. 4C), we measured CCL22 and CCL17 levels in the BAL fluid of WT and *Pla2g5*-null mice after 6 challenges with *Df*. The levels of CCL22 and CCL17 were significantly higher in the BAL fluid of *Df*-treated WT mice compared to *Df*-treated *Pla2g5*-null mice (Fig. 5A). Because these chemokines recruit Th2-cells (46, 47) we quantified the number and type of T-cells present in the lungs of WT and *Pla2g5*-null mice treated with *Df*. The WT but not the *Pla2g5*-null mice showed a significant increase in the number of T-cells recruited into the lung in response to *Df* (Fig. 5B). Compared to WT mice, *Pla2g5*-null mice had significant reductions in lung T-cells producing IL-4, IL-5, IL-17, and IFN-γ. T-cells from parabronchial lymph nodes showed a similar trend, possibly reflecting the absence of group V sPLA₂ in DC (data not shown). To ascertain the contributions of CCL22 and CCL17 to the pathologic features of the *Df* model of pulmonary inflammation, we administered blocking antibody against CCL22 or CCL17 to WT and *Pla2g5*-null mice 30 min before each of the last 4 doses of *Df*. Administration of CCL22 Ab to *Df*-treated WT mice reduced the total numbers of cells and lymphocytes in the BAL fluid to levels comparable to those observed in *Df*-treated *Pla2g5*-null mice (Fig. 5C). The blocking of CCL17 did not reduce BAL cellularity (Fig. 5D).

Transfer of gV-sPLA₂-expressing CD68^{enr}/*Df*-lung-cells can amplify *Df*-induced pulmonary inflammation in sensitized *Pla2g5*-null recipients

Because Mφ activated in vitro by IL-4 (Fig. 4) only partially replicated the phenotype of Mφ developed in vivo after *Df* exposure (Fig. 1), we adoptively transferred lung Mφ derived in vivo to verify that the presence of gV-sPLA₂ in Mφ was important to amplify inflammation in sensitized mice. We sensitized WT and *Pla2g5*-null mice to *Df* using WT *Df*-BMDCs (containing less than 1% eosinophils, data not shown), which are sufficient in both recipient strains to elicit an antigen-specific recall response in regional lymph nodes (37). Then, we transferred WT CD68^{enr}/*Df*-lung-cells into the sensitized mice (Fig. 6 inset). The CD68^{enr}/*Df*-lung-cells were enriched to >85% CD68⁺ cells using three consecutive gradients before the transfer (Supplemental Fig. 3). Less than 5% of the CD68⁻ cells expressed CD3 or B220 (data not shown). Control *Pla2g5*-null mice not receiving WT CD68^{enr}/*Df*-lung-cells failed to increase cell numbers or eosinophils in the BAL fluid after *Df* challenge, whereas identically treated WT control mice did (Fig. 6A, bottom row, middle columns). The transfer of WT CD68^{enr}/*Df*-lung-cells into DC-sensitized *Pla2g5*-null mice significantly increased the total cell number, eosinophils and lymphocytes in the BAL fluid (Fig. 6A, bottom row, left columns), albeit with lower values compared to control WT mice. The transfer WT CD68^{enr}/*Df*-lung-cells into WT mice did not significantly increase the recruitment of eosinophils, lymphocytes or total BAL fluid cells (Fig. 6A, top row).

To confirm that the amplification of the effector phase of pulmonary inflammation by WT CD68^{enr}/*Df*-lung-cells was associated with increased generation of CCL22 and CCL17, we measured these chemokines in the BAL fluid of transferred mice. There was a significant increase in the amount of CCL22 and CCL17 produced by *Pla2g5*-null mice transferred with WT *Df*-BMDCs and WT CD68^{enr}/*Df*-lung-cells compared to the *Pla2g5*-null mice transferred only with WT *Df*-BMDCs (Fig. 6B, bottom row), although with values lower

than control WT mice. To assess whether the transfer of gV-sPLA₂-expressing WT CD68^{enr}/*Df*-lung-cells into *Pla2g5*-null-sensitized mice increased the recruitment of CD4⁺ T-cells into the lung, we measured the number of lung CD4⁺ T-cells by flow cytometry. *Pla2g5*-null mice receiving *Df*-BMDCs and WT CD68^{enr}/*Df*-lung-cells had significant more CD4⁺ T-cells in the lung than *Pla2g5*-null mice receiving only *Df*-BMDCs (Fig. 6C, bottom row).

Transfer of WT BM-IM-Mφ amplify IL-4-induced eosinophilic inflammation in *Pla2g5*-null mice

Because the CD68^{enr}/*Df*-lung-cells contain non-Mφ which may also influence the effector phase of *Df*-induced pulmonary inflammation, we used a second approach to verify that endogenous gV-sPLA₂ in Mφ is required for amplification of the effector phase of pulmonary inflammation. We cultured BM-IM-Mφ from WT and *Pla2g5*-null BM and transferred them into WT and *Pla2g5*-null mice before exposing the mice to IL-4 to bypass the requirement for DC sensitization and to directly induce activation of the Mφ in vivo. The BM-IM-Mφ from WT and *Pla2g5*-null were 94.68 ± 2.82% and 92.48 ± 3.2% CD68⁺ respectively (Supplemental Fig. 4 and not depicted). The WT and *Pla2g5*-null CD68⁺ BM-IM-Mφ expressed similar levels of CD11b (respectively 95.48 ± 1.8% and 95.97 ± 0.67%) and F4/80 (respectively 48.21 ± 6.01% and 48.65 ± 3.84%, Supplemental Fig. 4 and not depicted). Transfers of WT BM-IM-Mφ into *Pla2g5*-null mice significantly increased total number of cells in the BAL, and numbers of eosinophils, lymphocytes/monocytes, and neutrophils in response to IL-4 compared to transfers of *Pla2g5*-null BM-IM-Mφ, with values similar to equally treated WT recipients (Fig. 7). WT BM-IM-Mφ also tended to increase pulmonary inflammation in IL-4-treated WT mice, compared to WT mice transferred with *Pla2g5*-null BM-IM-Mφ.

Discussion

Several pathologic features of asthma, including eosinophilia, airway remodeling, and goblet cell metaplasia, are linked to the development and persistence of Th2-associated effector pathways. Using a model of allergic pulmonary inflammation induced by *Df*, a complex allergen with endogenous adjuvant properties that commonly sensitizes individuals with asthma, we previously reported that gV-sPLA₂ expression by DCs was necessary, but not sufficient, to cause the development of pulmonary Th2 responses (37). The current study establishes that although gV-sPLA₂ is expressed by eosinophils and epithelial cells, its expression is induced in Mφ during IL-4 activation and that the presence of gV-sPLA₂ in Mφ is sufficient to generate critical chemokines for the recruitment of effector T-cells in this model. Indeed, Mφ appear to be intermediaries between DC-dependent sensitization and the amplification of a *Df*-directed effector response, with gV-sPLA₂ being essential for both steps. Moreover, interference with the functions of gV-sPLA₂ has the potential to interrupt this pathway for therapeutic purposes.

Several sPLA₂s (group V, IIA and X) have been identified in the airways of patients with asthma (10, 11) and OVA models of pulmonary inflammation have established central roles for group V and group X sPLA₂ (12, 13). However the cellular sources and the particular functions of each sPLA₂ are largely unknown. GV-sPLA₂ expression in the lungs of naïve mice is weak or undetectable (37), but our previous studies showed that the enzyme localized prominently to Mφ as well as eosinophils and epithelial cells following exposure to *Df* (37). Since gV-sPLA₂ is essential for normal function of Mφ and for immune responses in a model of candidiasis (38, 39) in which Mφ have the characteristics of AAMφ (48, 49), and because allergen-induced pulmonary inflammation typically results in the

expression of AAM ϕ markers such as Relm- α (20, 25, 31, 32), we suspected that the absence of gV-sPLA₂ might affect the phenotype of M ϕ and their function in the *Df* model.

We found that *Df*-challenged mice lacking gV-sPLA₂ showed induced expression of Relm- α in epithelial cells, but complete absence of this marker on cells infiltrating the lung, including M ϕ (Fig. 1A). We verified that WT and *Pla2g5*-null mice had similar total numbers of lung CD68⁺ M ϕ after *Df* challenge, but that the CD68⁺ M ϕ from the lungs of *Df*-treated *Pla2g5*-null mice showed very weak expression of molecules related to macrophage alternative activation at both the protein (Fig. 1D) and mRNA levels (Fig. 1E). These experiments suggested that the absence of gV-sPLA₂ markedly impairs the activation of M ϕ in vivo.

The expression of gV-sPLA₂ in the lungs of *Df*-challenged mice includes both resident (epithelium) and hematopoietic cells (M ϕ and eosinophils), all of which play a role in the pathophysiology of inflammation and Th2 responses. Several lines of evidence now indicate that the expression of gV-sPLA₂ in M ϕ *per se* is required for the activation of M ϕ in our model. First, studies using chimeric mice we demonstrate that the expression of gV-sPLA₂ by hematopoietic cells, but not by resident cells, contributes to the Th2-driven inflammatory response to *Df*, and to the activation of M ϕ in vivo (Fig. 2). Second, the activation of lung M ϕ (along with other features of allergic pulmonary inflammation) is markedly impaired in *Pla2g5*-null mice intratracheally instilled with IL-4 (a model that induces Th2-like pathology independently of the adaptive immune system) (41) compared to WT controls (Fig. 3). Thus, although we cannot completely exclude contributions from gV-sPLA₂ in cells other than macrophages, the presence of the enzyme in hematopoietic cells contributes to activation of M ϕ by a pathway downstream of the adaptive immune response and the Th2-type cytokines IL-4 and IL-13. This is further validated by the impaired IL-4-IL-4R γ -chain-mediated expression of gV-sPLA₂ and other markers of macrophage activation (Supplemental Fig. 3B). Moreover gV-sPLA₂ expression is upregulated by IL-4 (and to a lesser extent by IL-13) in mouse and human M ϕ (Fig. 4A,D; Supplemental Fig. 3A,C), and the elimination or depletion of gV-sPLA₂ from the M ϕ of both species selectively reduces the expression of some molecules related to macrophage activation (Fig. 4B, C, E). Collectively, these findings suggest that gV-sPLA₂ contributes to the Th2-dependent pathway for activation of M ϕ , and that this function is conserved across species.

M ϕ are prominent in various models of antigen- or parasite-induced Th2-type pulmonary inflammation, although their specific function(s) are incompletely understood (22, 23, 26, 36, 50, 51). Because our previous studies suggested that *Df*-driven pulmonary pathology required the function(s) of gV-sPLA₂ in at least one cell type in addition to DCs (37), we focused on potential pathways and mechanisms by which gV-sPLA₂-expressing M ϕ might amplify the response to *Df*. A previous study revealed that lung myeloid cells were the most abundant source of the Th2-active chemokines CCL22 and CCL17 in a model of OVA-induced pulmonary disease using mice with an OVA-specific T-cell receptor (52). We found that the expression of both chemokines was profoundly decreased at the mRNA level in *Pla2g5*-null CD68^{enr}/*Df*-lung-cells compared to WT controls (Fig. 1E), a phenotype that could be partially replicated in vitro (Fig. 4C), and that the levels of the corresponding proteins were significantly lower in the BAL fluid of *Df*-treated *Pla2g5*-null mice as well (Fig. 5A). The importance of one of these chemokines (19), CCL22, was verified by Ab-mediated blockade, which markedly suppressed the accumulation of eosinophils and lymphocytes in the BAL fluid (Fig. 5C–D). The transfer of WT CD68^{enr}/*Df*-lung-cells increased total cells, eosinophils and lymphocytes in the BAL fluid of DC-sensitized *Pla2g5*-null mice (Fig. 6A) while also amplifying CCL22/CCL17 generation and the recruitment of CD4⁺ lung T-cells (Fig. 6B–C). Finally, transfers of WT BM-IM-M ϕ , but not *Pla2g5*-null BM-IM-M ϕ , followed by intratracheal administration of IL-4 to directly induce

the effector phase of pulmonary inflammation and in vivo activation of M ϕ , increased pulmonary inflammation in *Pla2g5*-null mice (Fig. 7).

Although these findings do not preclude important functions of gV-sPLA₂ in other cell types, they do suggest that gV-sPLA₂-expressing M ϕ can amplify lung T-cell recruitment and downstream eosinophilia (21) by generating T-cell-attractive chemokines. Finer resolution of the roles of gV-sPLA₂ in M ϕ vs. other BM-derived cell types awaits the creation of mice with conditional deletion of the *Pla2g5* allele.

Our studies suggest a novel and unique role for gV-sPLA₂ in regulating the activation and effector function of M ϕ in a model of pulmonary inflammation caused by *Df*, a clinically relevant allergen. Our data support a role of M ϕ in amplification of pulmonary inflammation through T-cell recruitment and production of Th2-attractive chemokine (CCL22 and CCL17), molecules increased in the lung and/or serum of patients with severe asthma and COPD (20, 32–34, 53). Because gV-sPLA₂ is induced by IL-4 in both mouse and human M ϕ and is sufficient for the effector functions of M ϕ in pulmonary inflammation, our study suggests potential therapeutic implications of targeting this enzyme in severe asthma, COPD, or other diseases in which Th-2 activated M ϕ may play an effector role.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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Abbreviations used

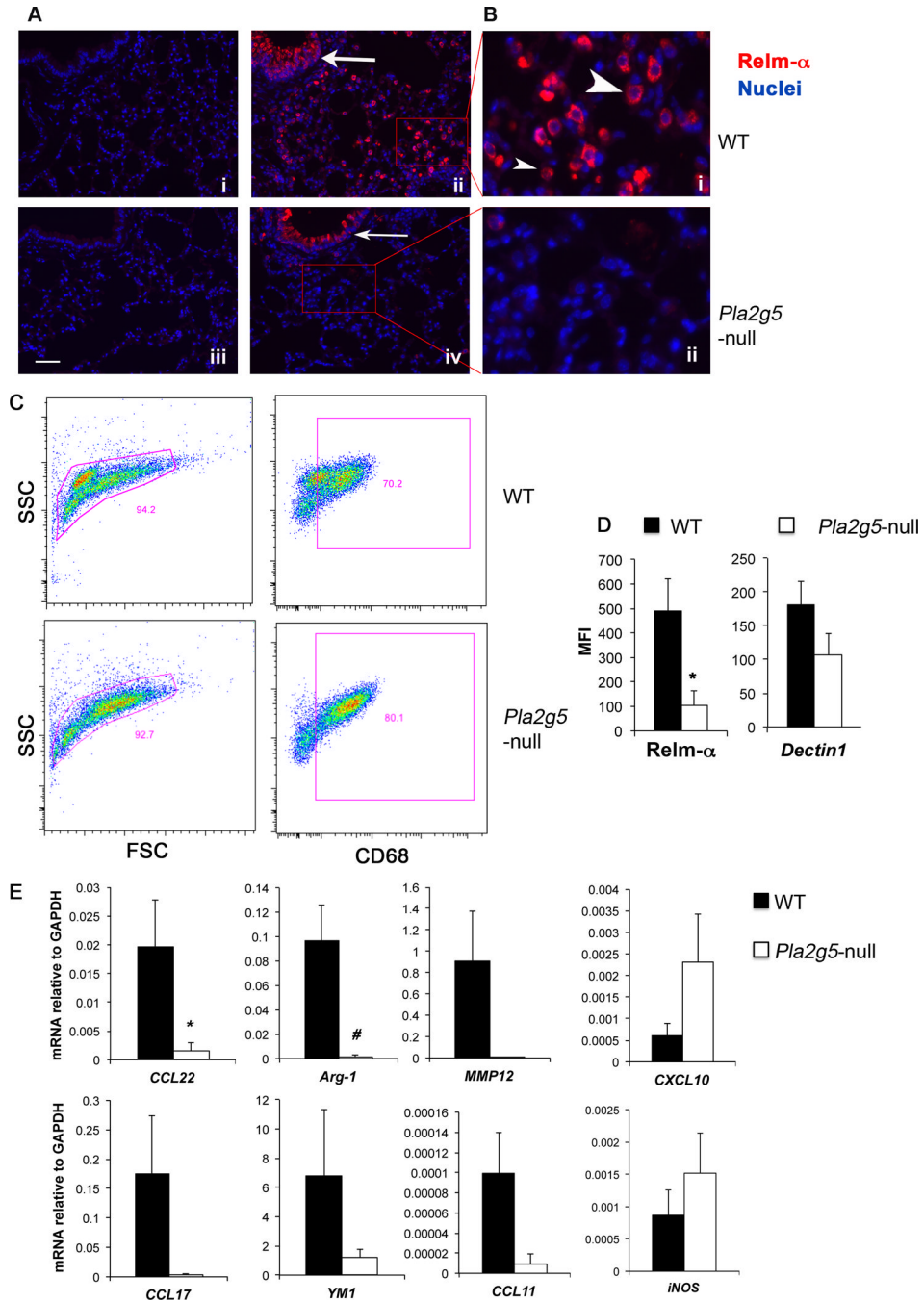
AAMϕ	alternatively activated macrophages
Arg-1	Arginase-1
BM	bone marrow
BAL	bronchoalveolar lavage
CD68^{enr}/<i>Df</i>-lung-cells	<i>Df</i> -induced lung cells enriched for CD68 ⁺ cells
<i>Df</i>	<i>Dermatophagoides farina</i>
<i>Df</i>-BMDC	<i>Df</i> -pulsed BMDC
iNOS	inducible nitric oxide synthase
i.t.	intratracheally
Mϕ	macrophages
MFI	mean fluorescence intensity
Relm-α	Resistin-like molecule- α
sPLA₂	secretory phospholipase A ₂
WT	wild-type.

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Pla2g5-null (open bars) cells. (E) Expression of mRNA encoding alternatively (CCL22, Arg-1, MMP-12, CCL17, YM1, and CCL11) and classically activated (CXCL10 and iNOS) M ϕ markers measured by qPCR. Data are expressed as ratio of the indicated mRNA expression relative to GAPDH. (A, B) Images are from one representative mouse per group from one of two experiments. (C) The lung cells are pooled from 7–9 mice per group, and representative plots are from one of six experiments. (D and E) Values are mean \pm SEM of four to six independent experiments. *, $p < 0.05$; #, $p < 0.02$

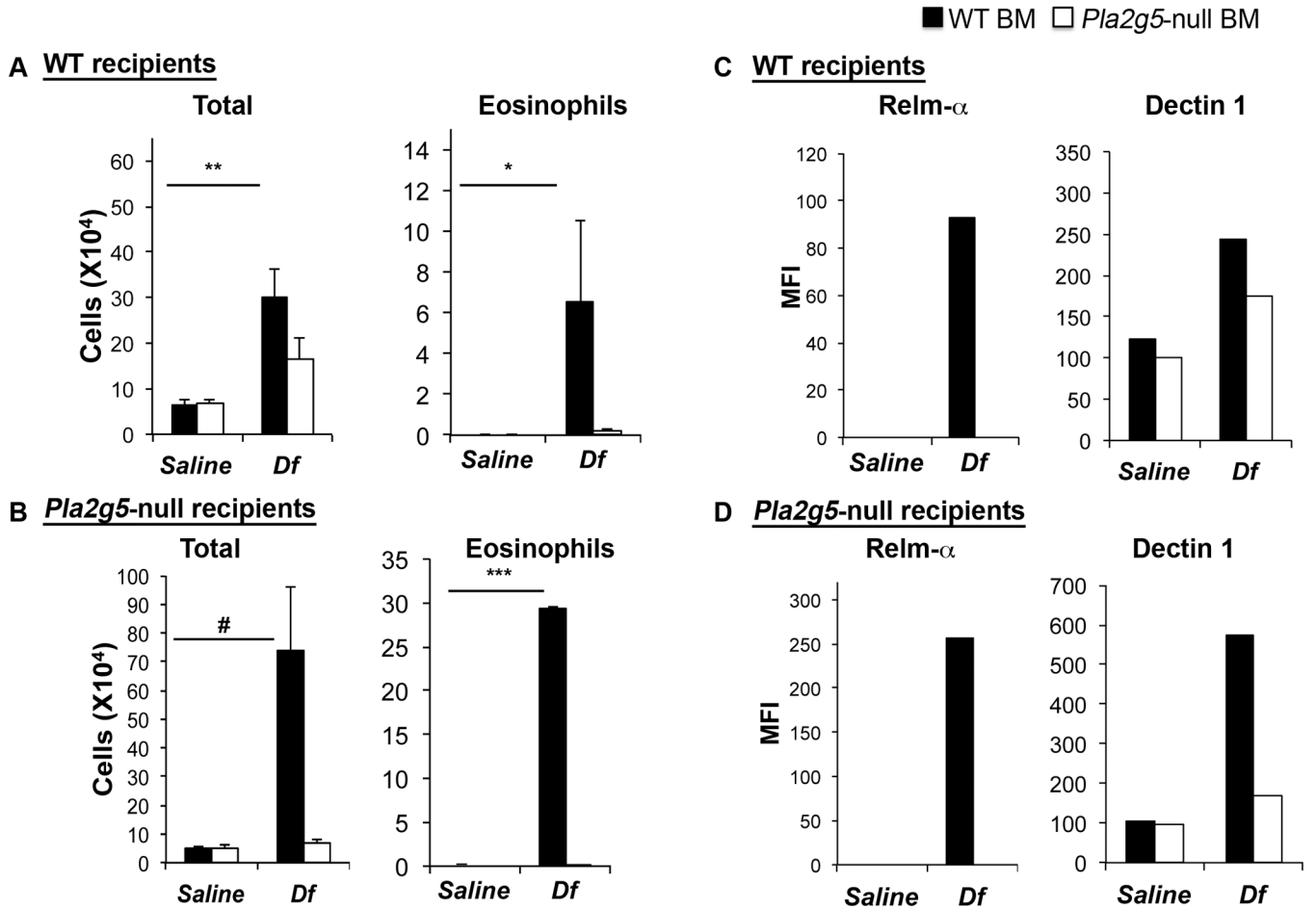


Figure 2. *Df*-induced pulmonary inflammation in WT and *Pla2g5*-null chimeric mice depends on the presence of gV-sPLA₂ in hematopoietic cells

Total and eosinophil cell counts from BAL fluid of WT (A) and *Pla2g5*-null mice (B) receiving WT (filled bars) or *Pla2g5*-null (open bars) BM cells. Net (isotype control subtracted) MFI of Dectin1 and Relm- α on gated CD68⁺ lung cells from pooled lungs of WT (C) and *Pla2g5*-null mice (D) receiving WT- (filled bars) or *Pla2g5*-null- (open bars) BM cells evaluated by flow cytometry. (A, B) Values are mean \pm SEM of two independent experiments with 7–13 mice per group. (C, D) Results represent pooled cells from multiple mice in one of the two experiments, which showed similar results but different magnitude of responses. *, $p < 0.05$; #, $p < 0.03$; **, $p < 0.02$; ***, $p < 0.01$

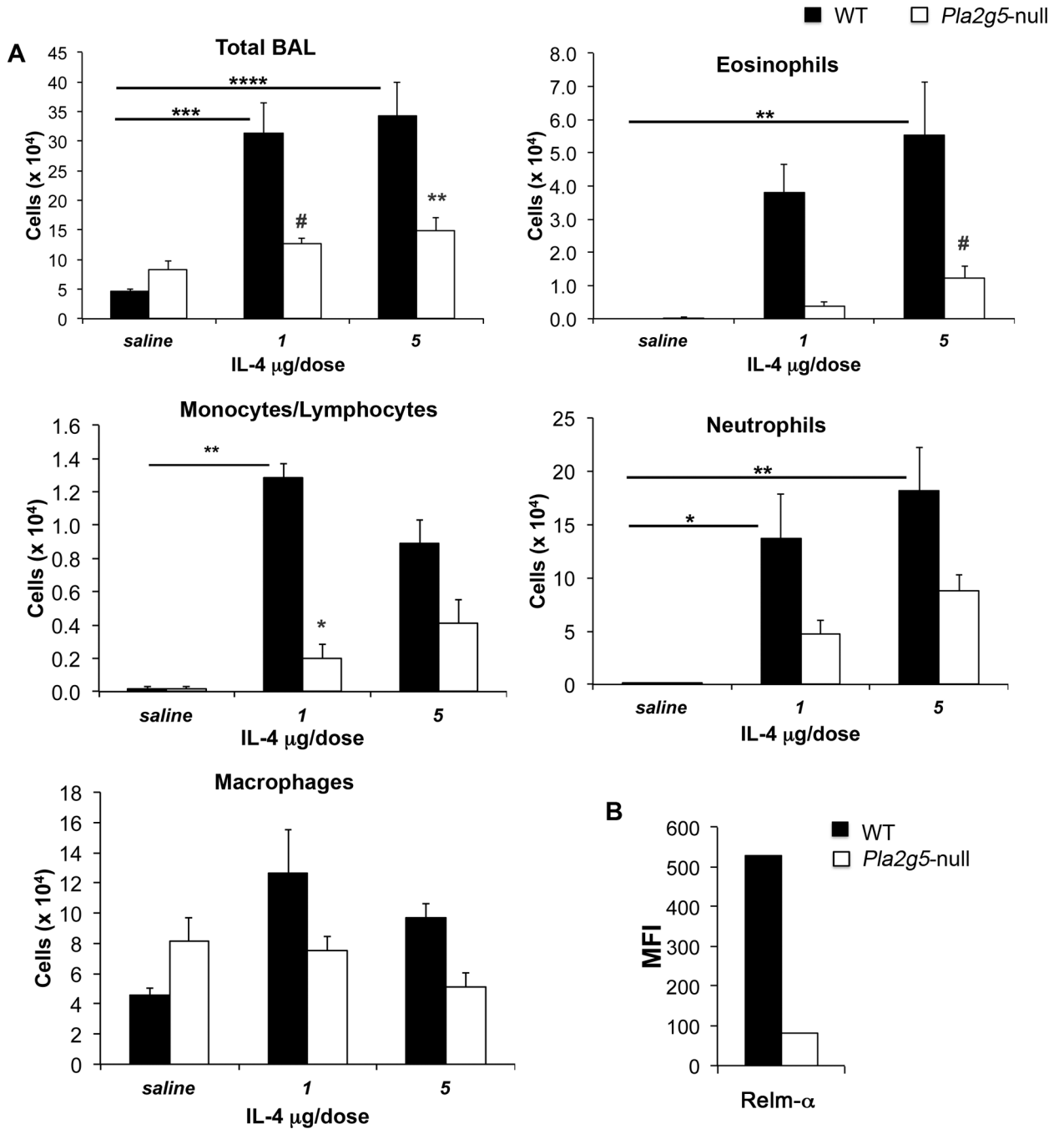


Figure 3. IL-4-induced pulmonary inflammation requires gV-sPLA₂

(A) Total and differential cell counts from BAL fluid of WT (filled bars) and *Pla2g5*-null (open bars) mice exposed to IL-4 1 or 5 $\mu\text{g}/\text{dose}$ intratracheally. (B) Net (isotype control subtracted) MFI of Relm- α on gated CD68⁺ lung cells of WT (filled bars) and *Pla2g5*-null (open bars) mice exposed to IL-4 5 $\mu\text{g}/\text{dose}$ intratracheally and evaluated by flow cytometry. (A) Values are mean \pm SEM from two independent experiments with 10–17 mice per group. (B) Data are representative of pooled cells from multiple mice in one of the two experiments with 7–9 mice per group. *, $p < 0.05$; #, $p < 0.02$; **, $p < 0.01$; ***, $p < 0.001$; ****, $p < 0.0001$

monocyte-derived M ϕ (h-IL-4-M ϕ and h-LPS/IFN γ -M ϕ , respectively) measured by qPCR. (E) Transfection of h-IL-4-M ϕ with human *Pla2g5*-siRNA (light gray columns) or non-targeting vector (dark gray columns) (1000 nM) and net (isotype control subtracted) MFI of CD209, CD206, and CD36 on gated CD11b⁺ cells evaluated by flow cytometry. Values are mean \pm SEM of three (A, C, D and E), or four to six (B) independent experiments. *, p<0.05; #, p<0.002; **, p<0.001

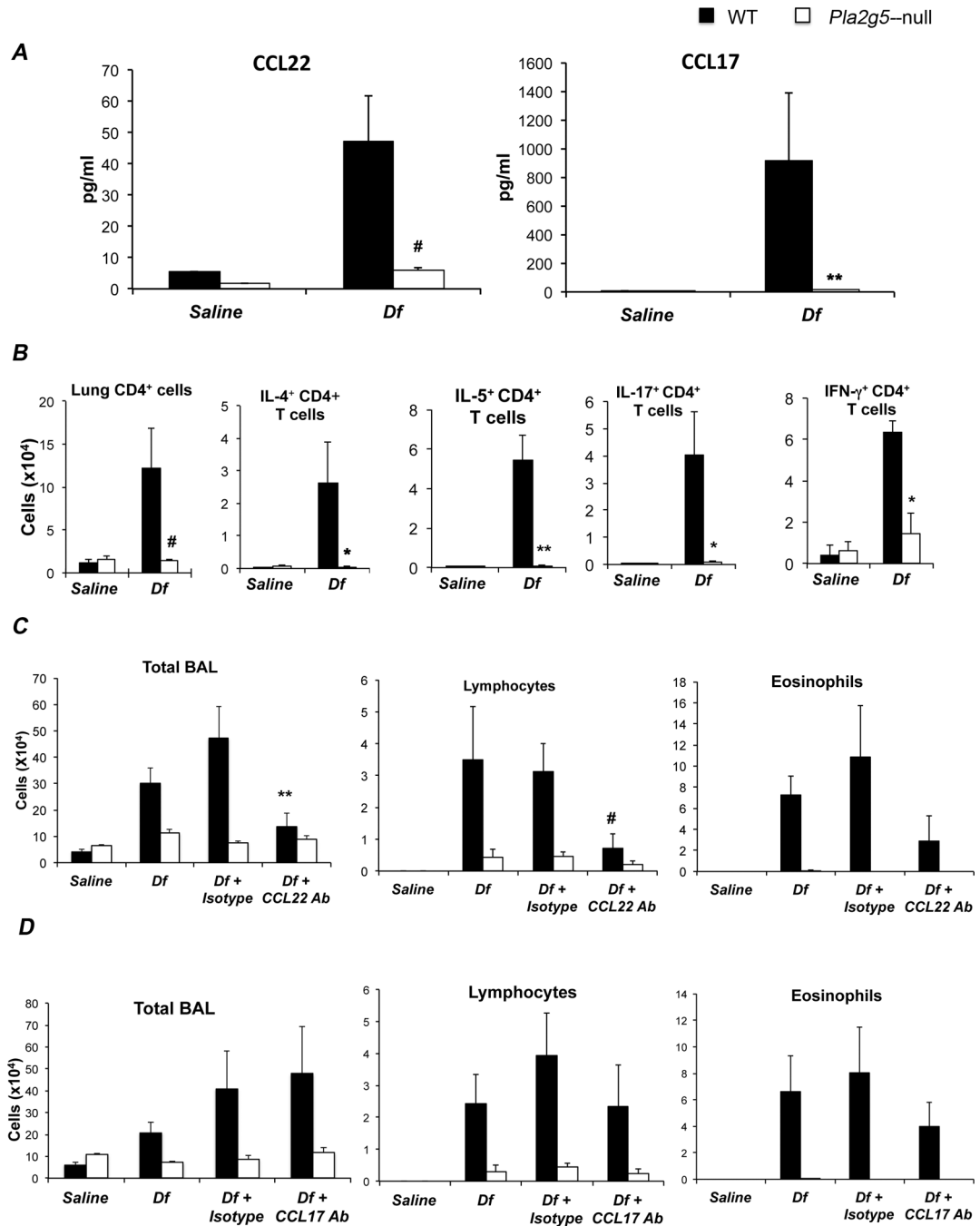
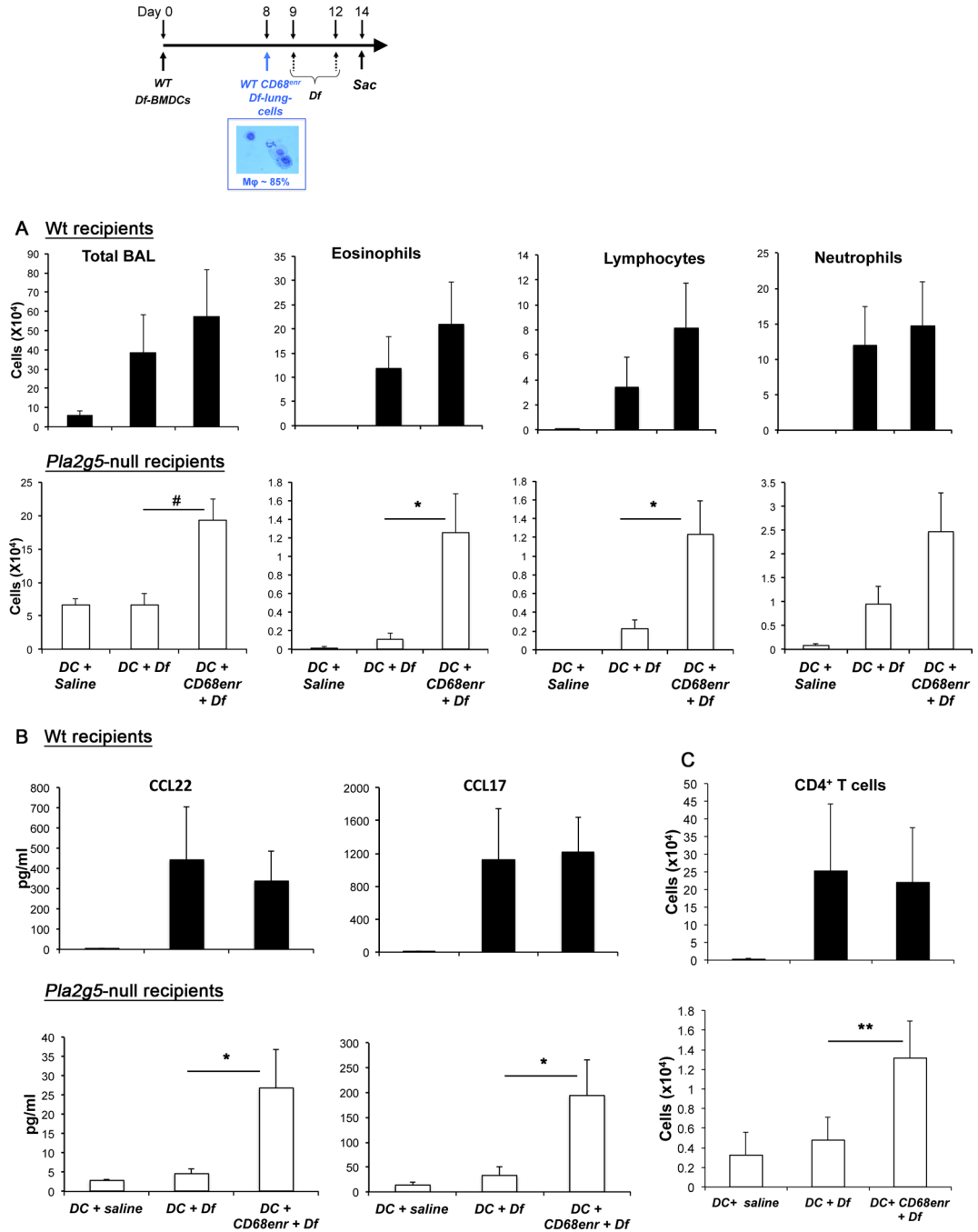


Figure 5. Group V sPLA₂ is required for generation of CCL22 and CCL17 in the lung

(A) Levels of CCL22 and CCL17 measured by respective ELISAs in BAL fluid of WT (filled bars) and *Pla2g5*-null (open bars) mice treated with saline or *Df*. (B) Lung CD4⁺ cell counts and intracellular cytokine expression evaluated by flow cytometry. (C) Total cell and differential counts from BAL fluid of WT and *Pla2g5*-null mice treated with saline, *Df*, *Df* + Isotype control, or *Df* + CCL22 Antibody (Ab); or (D) saline, *Df*, *Df* + Isotype control, or *Df* + CCL17 Ab. Values are mean \pm SEM from three independent experiments with 8–17 mice per group (A, B) or from two independent experiments with 5–9 mice per group (C, D). *, $p < 0.05$; **, $p < 0.03$; #, $p < 0.01$



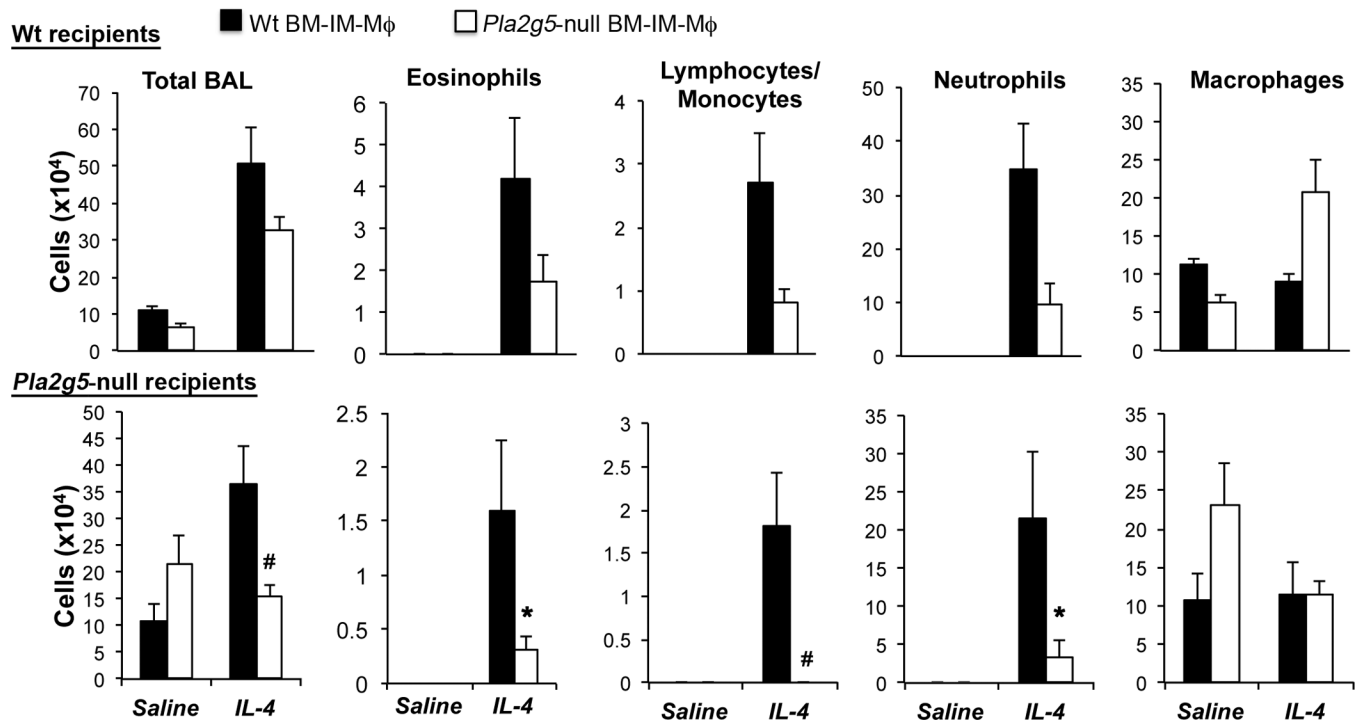


Figure 7. Transfer of WT but not *Pla2g5*-null BM-IM-M ϕ into *Pla2g5*-null recipient mice amplifies IL-4-induced pulmonary inflammation

WT (upper row) and *Pla2g5*-null (lower row) recipient mice received WT (filled bars) and *Pla2g5*-null (open bars) BM-IM-M ϕ intratracheally at day 0, followed by IL-4 (5 μ g/dose) intratracheally at day 1, 2 and 5. Mice were euthanized 36 h after the last dose. Values are mean \pm SEM from one representative of two separate experiments with 5–10 mice per group. *, $p < 0.03$; #, $p < 0.01$

Table I

List of mouse and human primers used

	Product	5'-3' Forward primer	5'-3' Reverse primer
Mouse	CCL22	CATCATGGCTACCCTGCGTGTC	CCTCCTCCCTAGGACAGTTTATGGA
	CCL17	TTGTGTTTCGCTGTAGTGATA	CAGGAAGTTGGTGAGCTGGTATA
	Arg-1	CAGAAGAATGGAAGAGTCAG	CAGATATGCAGGGAGTCACC
	Ym1	TCACAGGTCTGGCAATTCTTCTG	TTTGTCTTAGGAGGGCTTCCTC
	MMP-12	TCAATTGGAATATGACCCCTG	ACCAGCAAGCACCTTCACTAC
	GV-sPLA ₂	TGGTTCCTGGCTTGAGTGAG	TTCGCAGATGACTAGGCCATT
	CXCL10	ATGGCTGTCCTAGCTCTGTACT	ACTTAGAACTGACGAGCCTGAG
	iNOS	CCCTTCCGAAGTTTCTGGCAGCAGC	GGCTGTCAGAGCCTCGTGGCTTTGG
	GAPDH	TCAACAGCAACTCCCACTTTCCA	ACCTGTGCTGTAGCCGTATCA
Human	GV-sPLA ₂	GGCTTGAGTGTGCCTGCTG	CGCAGGTGACCACGCCCCAC
	GAPDH	GATGACATCAAGAAGGTGGTGAA	GTCTTACTCCTGGAGGCCATGT

The primers for mouse CCL11 were from SABiosciences.