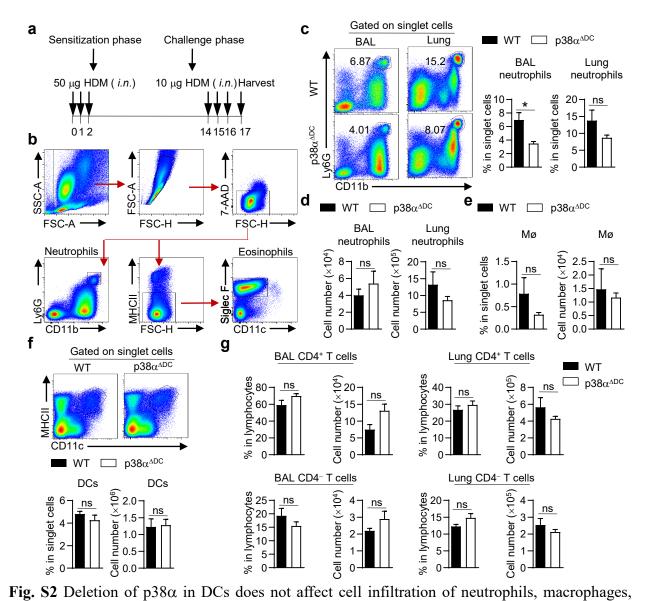


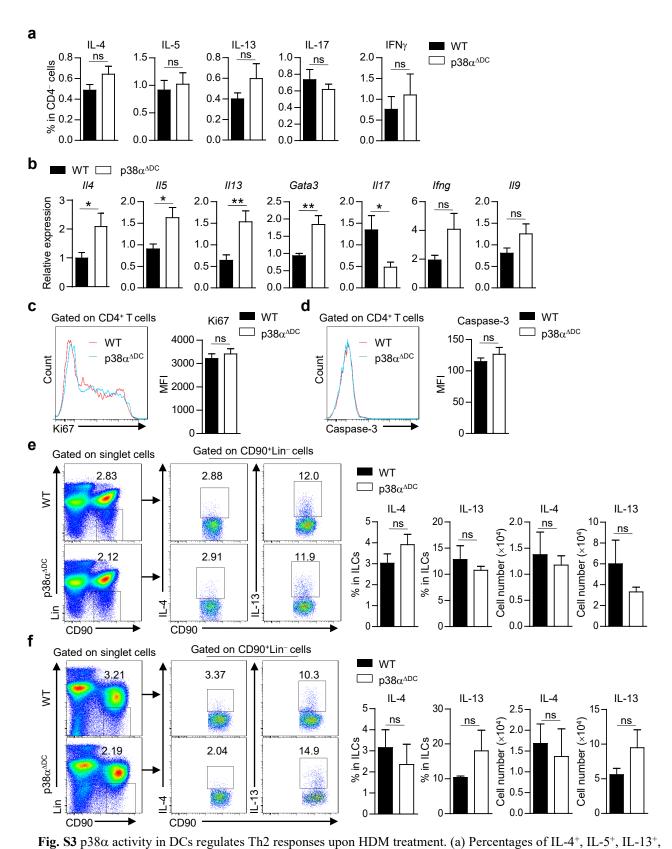
Fig. S1 HDM treatment increases DC infiltration in the lung and BAL and deletion of p38 α in DCs does not affect lung immune homeostasis. (a, b) WT mice were *i.n.* sensitized with 50 μ g HDM for 3 consecutive days (n = 4-5). Percentage and cell number of DCs (CD11c+MHCIIhigh) in the lung (a)

and BAL (b) were examined 24 h after the last challenge. (c) *Mapk14* expression was examined in flow cytometry-sorted naïve WT and p38 α -deficient lung DCs (CD11c+MHCIIhigh) by qPCR (n = 3).(d) p38 expression was measured in WT and p38 α -deficient splenic DCs by Western blot.(e) Flow cytometry and proportions of total DCs, cDC1 and cDC2 subsets in the lung of naïve WT and p38 α -deficient (n = 4). (f) Mean of fluorescence intensity (MFI) of CD40, CD80, CD86 and OX40L expression in naïve WT and p38 α -deficient lung DCs (n = 3). (g) Total cell number isolated from lung tissues of naïve WT and p38 α -deficient (n = 3). (h) Flow cytometry (left) and proportions (right) of eosinophils, neutrophils, CD4+TCR β + T cells and CD4-TCR β + T cells in lung tissues of WT and p38 α -deficient (n = 3). *P < 0.05, **P < 0.01. ns, not significant. Data are representative of three (a, b and d-h) independent experiments or pooled of three independent experiments with consistent results (c). Student's t-test (a-c, e-h) was performed and data are presented as mean \pm SEM.



DCs and T cells during asthma pathogenesis. (a) Experimental protocol for HDM-induced asthma. (b) Representative gating strategy for the evaluation of eosinophils and neutrophils. (c, d) Flow cytometry, proportion and cell number of neutrophils in the BAL and lungs (n = 4). (e) The frequency and cell number of macrophages (Mø) in the BAL (n = 5). (f) Flow cytometry, proportion and cell number of DCs in the lungs (n = 5). (g) The frequencies and cell numbers of

CD4⁺ and CD4⁻ T cells in the BAL and lungs of WT and p38 $\alpha^{\Delta DC}$ mice (n = 5). *P < 0.05. ns, not significant. Data are representative of three (c-g) independent experiments. Student's t-test (c-g) was performed and data are presented as mean \pm SEM.



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IL-17⁺ and IFN γ ⁺CD4⁻ cells in lung tissues determined by intracellular staining (n = 13). (b) mRNA expression of T

cell-related genes in HDM-sensitized lung tissues measured by qPCR and normalized to Hprt (n = 9-10). (c, d) Cells were isolated from lung tissues of HDM-sensitized WT and p38 $\alpha^{\Delta DC}$ mice, and then analyzed by Ki67 and active Caspase-3 staining of CD4⁺ T cells (n = 5). (e) Flow cytometry (left), proportions (middle) and cell numbers (right) of IL-4⁺ and IL-13⁺ cells in ILCs in the lung of HDM-sensitized WT and p38 $\alpha^{\Delta DC}$ mice (n = 6). (f) Flow cytometry (left), proportions (middle) and cell numbers (right) of IL-4⁺ and IL-13⁺ cells in ILCs in the lung of papain-sensitized WT and p38 $\alpha^{\Delta DC}$ mice (n = 3). *P < 0.05, **P < 0.01. ns, not significant. Data are pooled of three (a) or two (b) independent experiments or representative of two (c-f) independent experiments. Student's t-test (a-f) was performed and data are presented as mean \pm SEM.

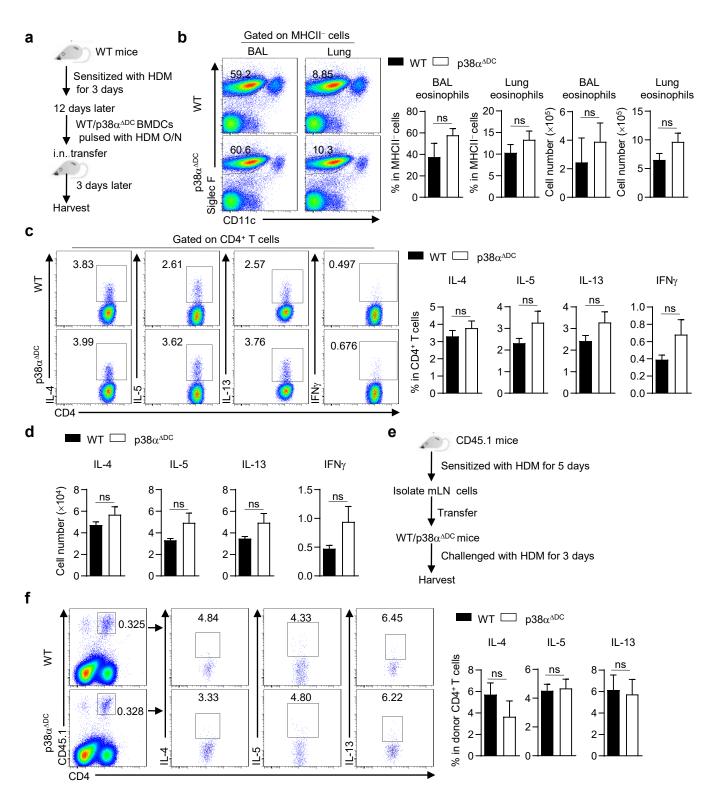


Fig. S4 p38 α activity in DCs is not required for HDM-induced effector allergic inflammation. (a-d) WT mice were sensitized with HDM for 3 consecutive days. The mice were rested for 12 days and *i.n.* transferred with 2×10^6 HDM-pulsed WT or p38-deficient BMDCs, and then analyzed 72 h after BMDC transfer (n = 3-4).

Experimental schema (a). Flow cytometry, proportions and cell number of eosinophils in the BAL and Lung (b). Flow cytometry, proportions (c) and cell numbers (d) of IL-4⁺, IL-5⁺, IL-13⁺ and IFN γ ⁺ CD4⁺ T cells in the lung. (e and f) 10 × 10⁶ mLN cells isolated from HDM-sensitized CD45.1 mice were *i.n.* injected into WT and p38 α ^{Δ DC} recipient CD45.2 mice. The recipients were then *i.n.* challenged with 10 μ g HDM for 3 consecutive days and analyzed 24 h after the last challenge. Experimental schema (e). Flow cytometry and proportions of donor IL-4⁺, IL-5⁺ and IL-13⁺ CD4⁺ T cells (f) (n = 3-4). ns, not significant. Data are representative of two (b-d, f) independent experiments. Student's *t*-test was performed and data are presented as mean \pm SEM.

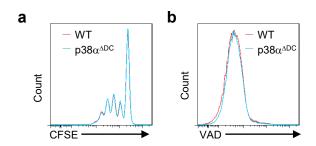


Fig S5 p38α deletion in DCs does not affect T cell proliferation or survival in vitro. Naïve OT-II CD4⁺ T cells (CD4⁺CD25⁻CD62L⁺CD44⁻) co-cultured with WT and p38α-deficient lung DCs. Proliferation (a) and survival (b) of OT-II CD4⁺ T cells were detected 72 h and 96 h later, respectively. Data are representative of two independent experiments (a, b). Every DC sample was pooled from at least three WT or p38α^{ΔDC} mice.

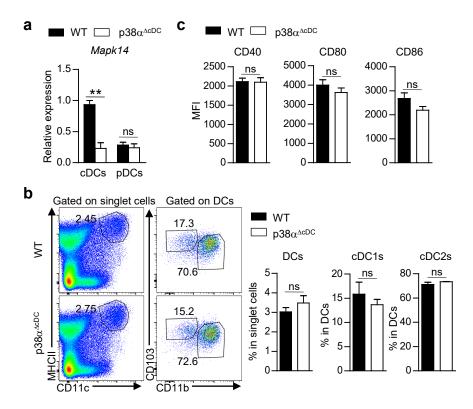


Fig. S6 p38α deletion in cDCs does not affect DC subsets and activation upon HDM treatment. (a) Mapk14 expression in naïve WT and p38α^{ΔcDC} cDCs (CD11c^{hi}MHCII^{hi}) and pDCs (CD11c^{dim}PDCA-1⁺) (n = 3). (b) Flow cytometry (left) and proportions (right) of total DCs, cDC1 and cDC2 subsets in the lung of HDM-treated WT and p38α^{ΔcDC} mice (n = 4). (c) MFI of CD40, CD80, CD86 expression in HDM-treated WT and p38α-deficient lung DCs (n = 4). **P < 0.01. ns, not significant. Data are pooled from two independent experiments (a) or representative of two independent experiments with consistent results (b, c). Two-way ANOVA (a) and student's *t*-test (b, c) was performed and data are presented as mean ± SEM.

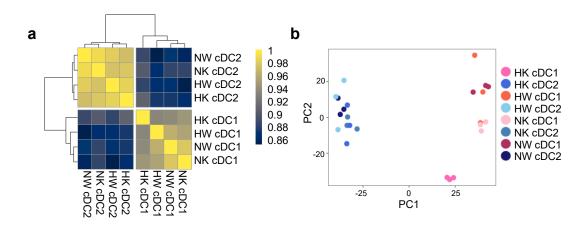


Fig. S7 The effect of p38 α deficiency is more pronounced under HDM-stress in the cDC1 population. (a) Pearson correlation and clustering of cDC1s and cDC2s. (b) PCA plot of the RNAseq results. NW, NK, HW and HK are short for naïve WT, naïve p38 α knock-out, HDM-treated WT and HDM-treated p38 α knock-out, respectively.

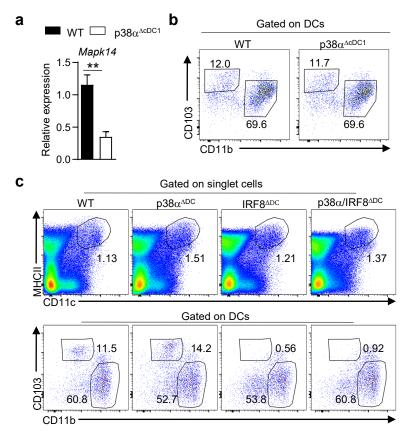


Fig. S8 p38α deletion in cDC1s does not affect DC subsets and percentage. (a) *Mapk14* expression was examined in flow cytometry-sorted naïve WT and p38α-deficient splenic cDC1s by qPCR (n = 4). (b) Flow cytometry of cDC1s and cDC2s in the lung of naïve WT and p38α $^{\Delta cDC1}$ mice. (c)The percentages of DCs, cDC1s and cDC2s were detected by flow cytometry. Data are pooled of two (a) or representative of two (b and c) independent experiments with consistent results.

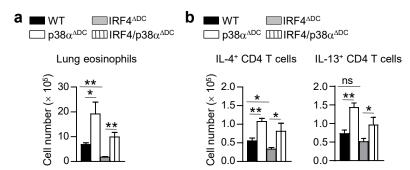


Fig. S9 p38α-deletion in DCs promotes HDM-induced lung inflammation in a cDC2-independent manner. WT, p38α^{ΔDC}, IRF4^{ΔDC} and IRF4/p38α^{ΔDC} mice were sensitized and challenged with HDM for asthma induction (n = 3-5). (a) Cell number of eosinophils in lung tissues. (b) Cell numbers of IL-4⁺CD4⁺ T cells and IL-13⁺CD4⁺ T cells in lung tissues. *P < 0.05; **P < 0.01. ns, not significant. Data are pooled from two experiments (a, b). Student's *t*-test (a, b) was performed and data are presented as mean ± SEM.

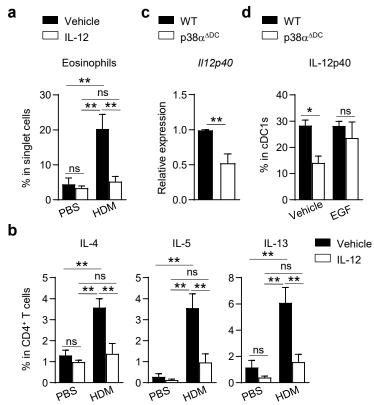


Fig. S10 p38α regulates IL-12 expression through c-FOS signaling pathway in DCs. (a and b) WT mice were induced asthma with HDM and PBS used as control. IL-12p70 was *i.n.* administrated after PBS or HDM treatment (n = 4-5). Eosinophil infiltration in lung tissues was detected by flow cytometry and quantified (a). The percentages of IL-4+, IL-5+ and IL-13+ CD4+ T cells in lung tissues were detected by flow cytometry and quantified (b). (c) mRNA expression of II12p40 in WT and p38-deficient Flt3L-DCs stimulated with HDM for 5 h (n = 5). (d) Percentages of IL-12p40 in HDM-stimulated WT and p38α-deficient lung cDC1s pretreated with vehicle or EGF (n = 4-5). **P < 0.01. ns, not significant. Data are pooled of four (c) or two (d) independent experiments or representative of two (a, b) independent experiments with consistent results. Student's *t*-test was performed and data are presented as mean ± SEM.

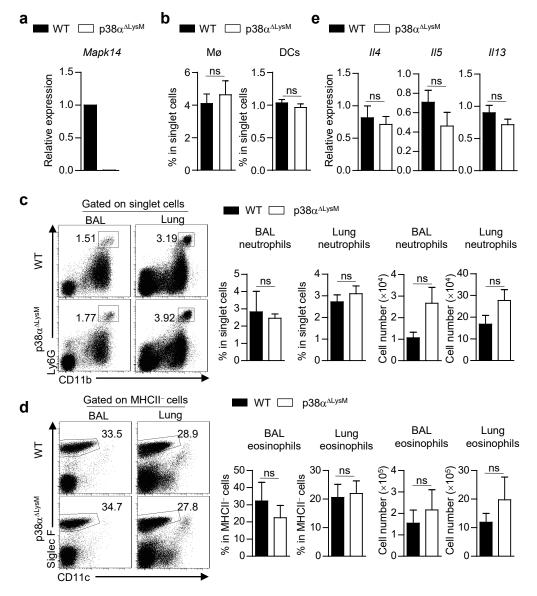


Fig. S11 p38α activity in AMs is not required for Th2-mediated allergic lung inflammation. (a) Mapk14 expression in naïve WT and p38α^{ΔLysM} AMs (n = 1, 1 sample pooled from 3 mice). Percentages of Macrophages (Mφ) and DCs in naïve WT and p38α^{ΔLysM} lung tissues (b) (n = 3). (c-e) WT and p38α^{ΔLysM} mice were sensitized and challenged with HDM to induce asthma. Frequencies and cell numbers of neutrophils and eosinophils in the BAL and lung tissues of HDM-treated WT and p38α^{ΔLysM} mice (c, d) (n = 4). RNA analysis of II4, II5 and II13 in lung tissues of WT and p38α^{ΔLysM} mice (e) (n = 5-6). ns, not significant. Data are representative of three (a-e) independent experiments. Student's t-test was performed and data are presented as mean \pm SEM.