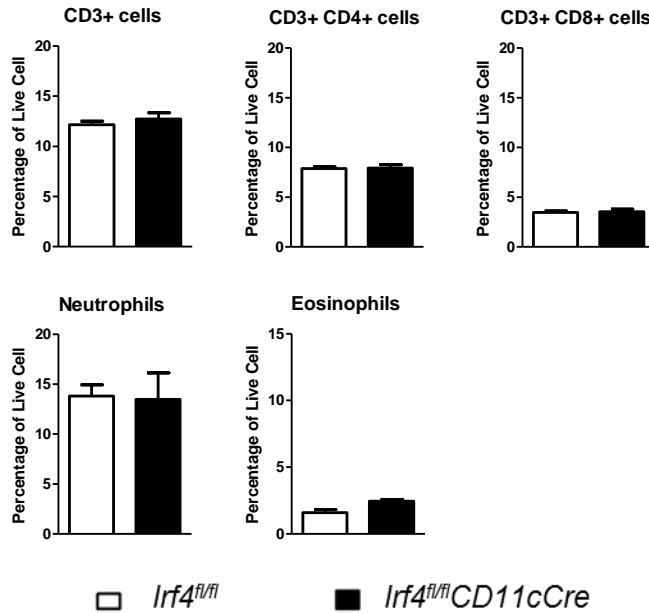
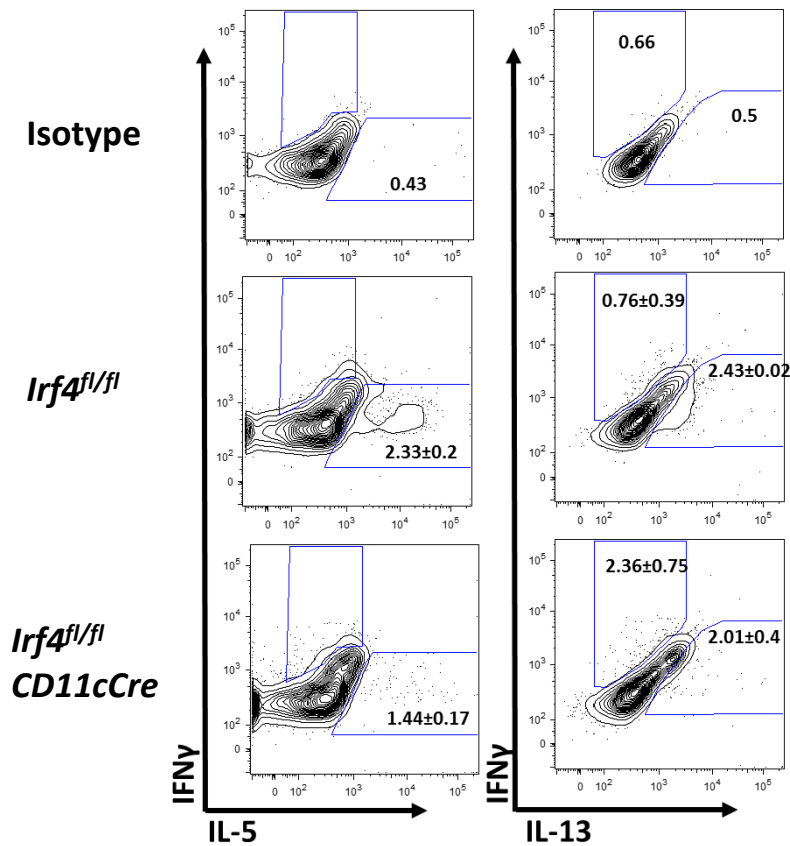


Supplementary Information:

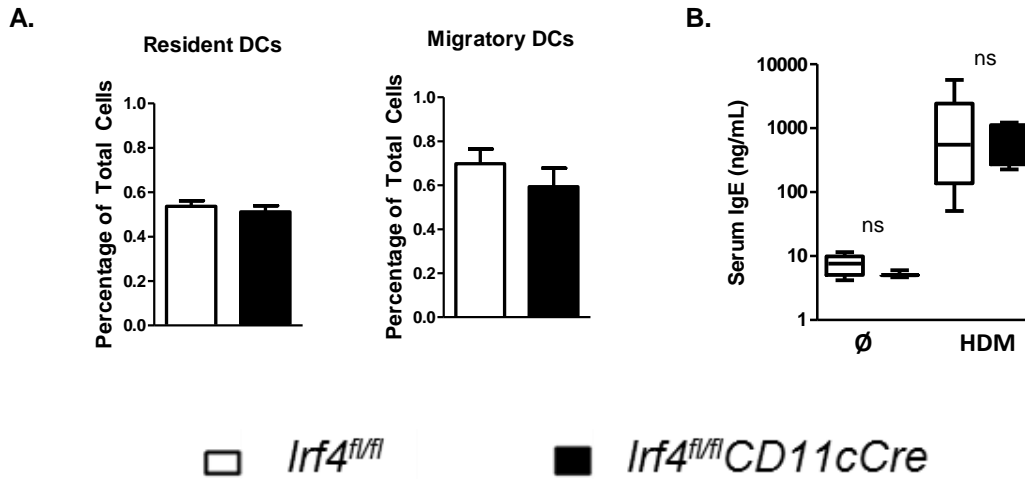


Supplementary Figure S1: *Irf4^{fl/fl}CD11cCre* mice have normal lung cellularity.

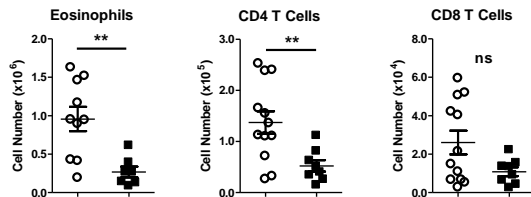
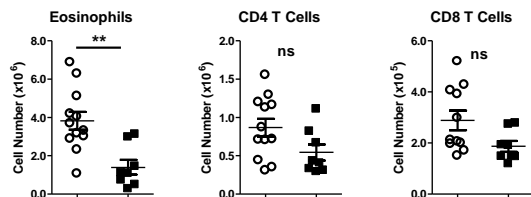
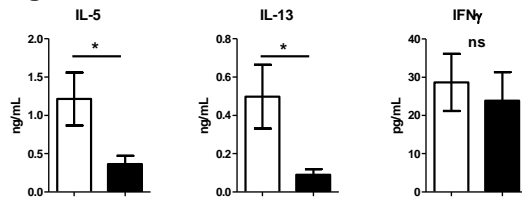
Unsensitized and unchallenged *Irf4^{fl/fl}* or *Irf4^{fl/fl}CD11cCre* mice were sacrificed and assayed for immune cells in the lungs. The percent of lung CD3⁺ cells, CD4⁺ T cells, CD8⁺ T cells, Neutrophils, and Eosinophils were identified by flow cytometry. Data represent the mean \pm SEM (statistical test by student t-test). Data are from three or four animals per group.



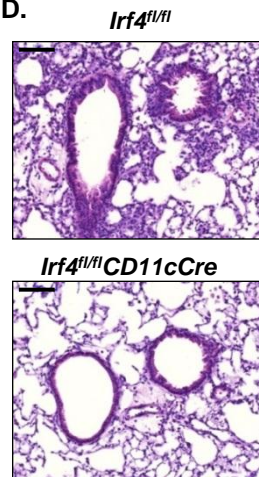
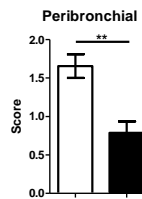
Supplementary Figure S2: Intracellular cytokine staining of T cells from the lungs of HDM-challenged mice. Lung T cells from HDM sensitized and challenged *Irf4^{fl/fl}* or *Irf4^{fl/fl}CD11cCre* mice were stimulated with PMA/Ionomycin, followed by intracellular cytokine staining for IL-5, IL-13, IFN γ , or appropriate isotype controls. Data represent the mean \pm SEM. Data are representative of two independent experiments with three to six mice per group.



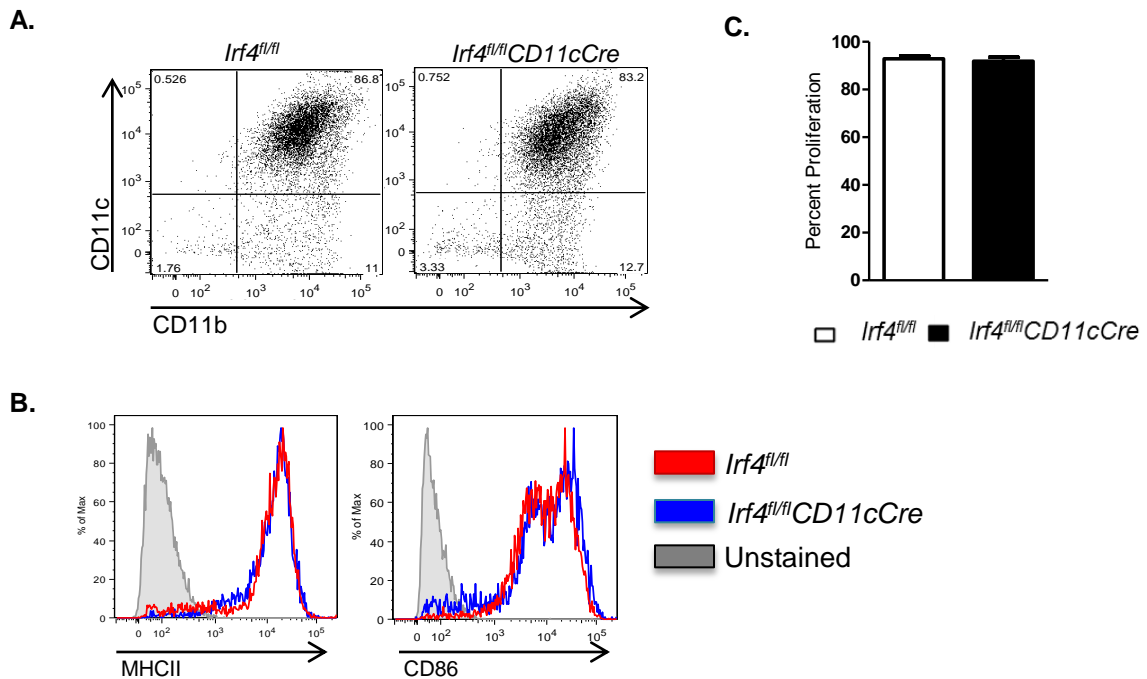
Supplementary Figure S3: *Irf4^{fl/fl}CD11cCre* mice show similar DC migration and IgE responses following HDM sensitization and challenge. (A) Mediastinal lymph nodes from HDM sensitized and challenged *Irf4^{fl/fl}* or *Irf4^{fl/fl}CD11cCre* mice were assayed for infiltration of migratory and resident DC populations. (B) Serum from unchallenged or sensitized and challenged mice were assayed for total IgE levels. Data represent the mean \pm SEM (statistical test by student t-test). Data is representative of two independent experiments with four to six replicates per group.

A. BAL**B. Lung****C. Lung**

□ *Irf4^{fl/fl}* ■ *Irf4^{fl/fl}CD11cCre*

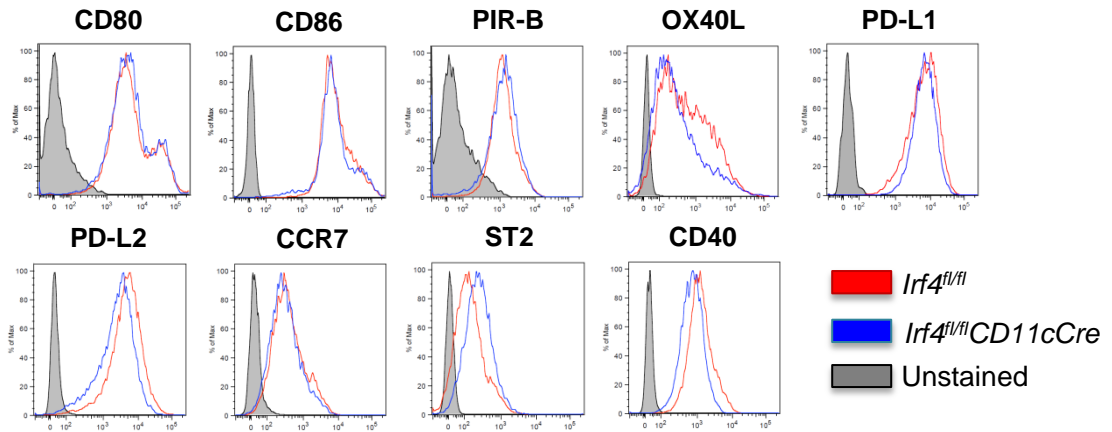
D.**E.**

Supplementary Figure S4: IRF4 in DCs regulates the development of Th2 responses toward HDM. *Irf4^{fl/fl}* or *Irf4^{fl/fl}CD11cCre* mice were sensitized i.p. with HDM/Alum and challenged i.t. with HDM, then sacrificed at the peak of inflammation on day 4 after challenge. (A) BAL eosinophil, CD4⁺, and CD8⁺ T cells numbers recovered from sensitized and challenged mice. (B) Lung eosinophil, CD4⁺, and CD8⁺ T cell numbers recovered from sensitized and challenged mice. (C) Lung T cells were restimulated and measured for secreted cytokines. (D) H&E stains of lung from sensitized/challenged (top) *Irf4^{fl/fl}* and (bottom) *Irf4^{fl/fl}CD11cCre* mice. (E) Peribronchovascular scoring of lung sections from sensitized and challenged mice for cellular infiltration. Histology bar = 100 μ m. Data represent the mean \pm SEM (statistical test by student t-test; *, $P < 0.05$; **, $P < 0.01$; ***, $P < 0.001$). Data are representative of three independent experiments, with three to six replicates per experiment.

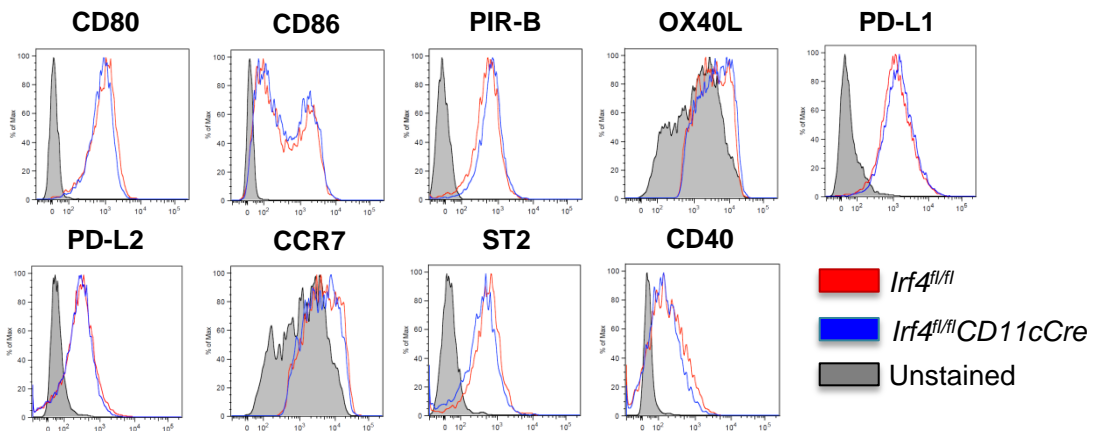


Supplementary Figure S5: *Irf4^{fl/fl}CD11cCre* BMDCs develop and are activated similar to *Irf4^{fl/fl}* BMDCs. (A) BMDCs from *Irf4^{fl/fl}* or *Irf4^{fl/fl}CD11cCre* mice were assayed for CD11c⁺ CD11b⁺ cells. (B) BMDCs were activated with LPS for eight hours and assayed for expression of MHCII and CD86. (C) *Irf4^{fl/fl}* or *Irf4^{fl/fl}CD11cCre* BMDCs were activated with LPS and used to present OVA to CFSE-labeled OTII cells, and T cell proliferation was measured five days later. Data represent the mean \pm SEM (statistical test by student t-test). Data are representative of three or more independent experiments.

A. GMCSF BMDCs



B. Flt3L BMDCs



Supplementary Figure S6: Phenotypic analysis of GM-CSF and Flt3L-derived *Irf4*^{fl/fl}CD11cCre BMDCs following HDM stimulation. (A) GM-CSF-derived and (B) Flt3L-derived BMDCs from *Irf4*^{fl/fl} or *Irf4*^{fl/fl}CD11cCre mice were assayed for expression of activation molecules following HDM-treatment. Data are representative of three unique experiments.