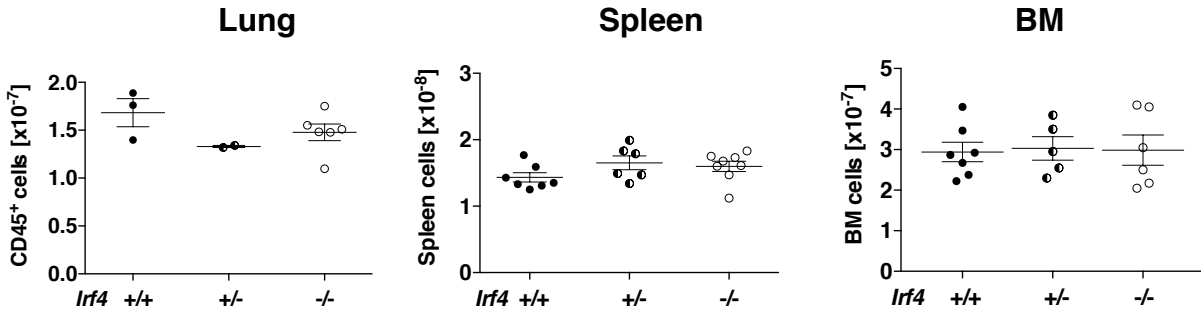


Fig. S1

A *CD11c-cre Irf4* mice



B *CD11c-cre Irf8* mice

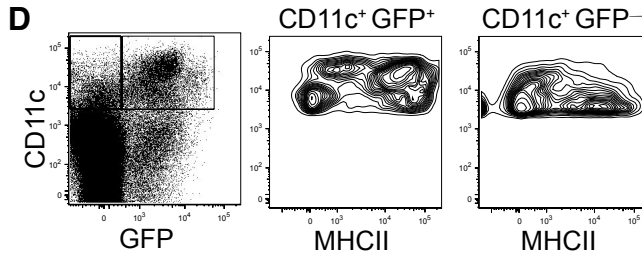
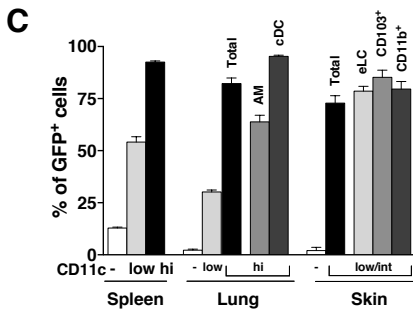
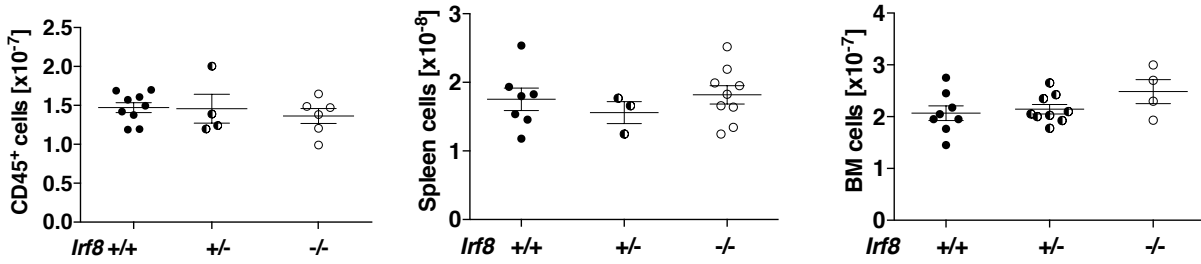


Fig. S1. Total numbers of cells in the lungs, spleen and BM of *CD11c-cre-Irf4* and *CD11c-cre-Irf8* +/+, +/- and -/- mice. (A) The total numbers of lung CD45⁺ cells (n=3-6), splenocytes (n=6-8) and bone marrow cells (n=5-7) in multiple *CD11c-cre-Irf4* +/+, +/- and -/- mice are compiled. (B) The total numbers of lung CD45⁺ cells (n=4-9), splenocytes (n=5-6) and bone marrow cells (n=4-9) in multiple *CD11c-cre-Irf8* +/+, +/- and -/- mice are compiled. (C) To illustrate the penetrance of the *Irf4* deletion in *CD11c-cre-Irf4* -/- mice, shown is the percentage of GFP⁺ cells within the CD11c⁻, CD11c^{low} and CD11c^{hi} fractions of the spleen, lung and skin. Within the lung CD11c^{hi} fraction, the percentage of GFP⁺ alveolar macrophages (AM) and cDCs is also reported. Within the skin CD11c^{low/int} fraction, the percentage of GFP⁺ epidermal Langerhans cells (eLCs), dermal CD103⁺ DCs and CD11b⁺ dermal DCs is also reported. (D) Shown is the expression of CD11c and GFP in the lung of *CD11c-cre-Irf4* +/- mice. (E) Shown is the expression of MHCII and GFP in the dermis of *CD11c-cre-Irf4* -/- mice

Fig. S2

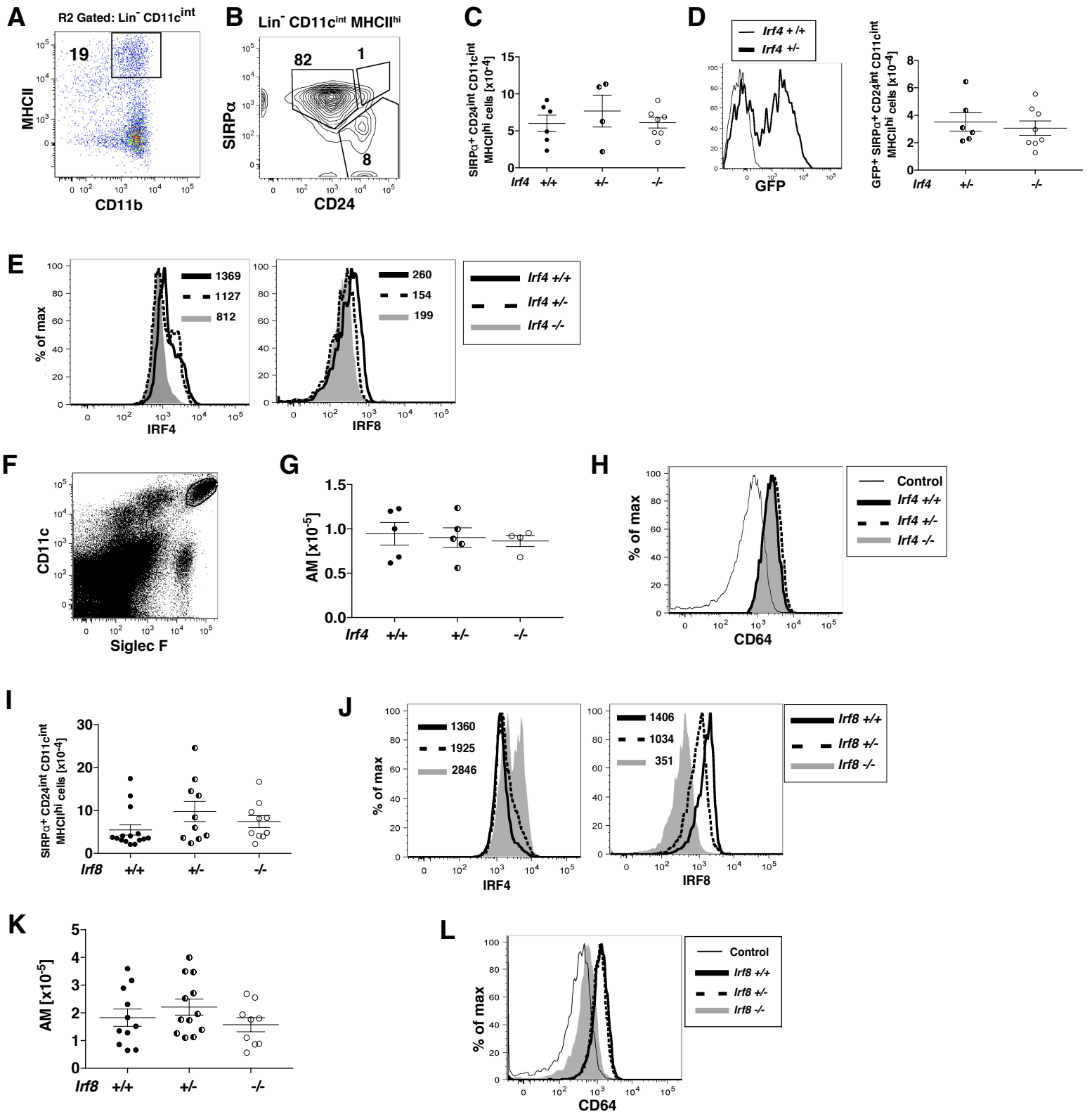


Fig. S2. Multiple DC and macrophage populations are not affected by *Irf4* or *Irf8* deficiency in CD11c⁺ cells. (A-H) Data are from *CD11c-cre-Irf4* *+/+*, *+/-* and *-/-* mice. (A) In the lungs of *CD11c-cre-Irf4* *+/+* mice, definition of CD11b⁺MHCII^{hi} cells in the CD11c^{int}SiglecF⁻ R2 gate (shown in Fig. 1A). (B) DCs are SIRPα⁺CD24^{int} (in P2 gate). Shown are the gates for the P1, P2 and P3 subsets of CD11c^{hi} cDCs in gate R1. (C) Total numbers of CD11c^{int}MHCII^{hi}CD11b⁺CD24^{int} cells in the lungs of *CD11c-cre-Irf4* *+/+*, *+/-* and *-/-* mice, n=4-6. (D) Only a subset of CD11c^{int}MHCII^{hi} DCs (gated in panel A) is GFP⁺ in *+/-* mice. Numbers of GFP⁺CD11c^{int}MHCII^{hi} DCs in multiple *+/-* and *-/-* mice are compiled (n=6-7). (E) IRF4 and IRF8 protein levels were determined in *+/+*, *+/-* and *-/-* CD11c^{int}MHCII^{hi}CD11b⁺CD24^{int} DCs by intracellular staining. The MFI of each histogram is indicated. (F) Definition of CD11c^{hi}SiglecF^{hi} alveolar macrophages in the lungs of *+/+* mice. (G) Numbers of alveolar macrophages in the lungs are compiled from multiple *+/+*, *+/-* and *-/-* mice, n= 4-5. (H) The expression of CD64 on alveolar macrophages in *+/+*, *+/-* and *-/-* mice is shown relative to an isotype control. (I-L) Data are from *CD11c-cre-Irf8* *+/+*, *+/-* and *-/-* mice. (I) Total numbers of CD11c^{int}MHCII^{hi}CD11b⁺CD24^{int} cells in the lungs of *CD11c-cre-Irf8* *+/+*, *+/-* and *-/-* mice, n=10-15. (J) IRF4 and IRF8 protein levels were determined in *+/+*, *+/-* and *-/-* CD11c^{int}MHCII^{hi}CD11b⁺CD24^{int} DCs by intracellular staining. The MFI of each histogram is indicated. (K) Numbers of alveolar macrophages in the lungs are compiled from multiple *+/+*, *+/-* and *-/-* mice, n= 9-11. (L) The expression of CD64 on alveolar macrophages in *+/+*, *+/-* and *-/-* mice is shown. The data were evaluated using a one-way ANOVA (panels C, G, I and K) or an unpaired t test (panel D); no significant differences between genotypes were identified.