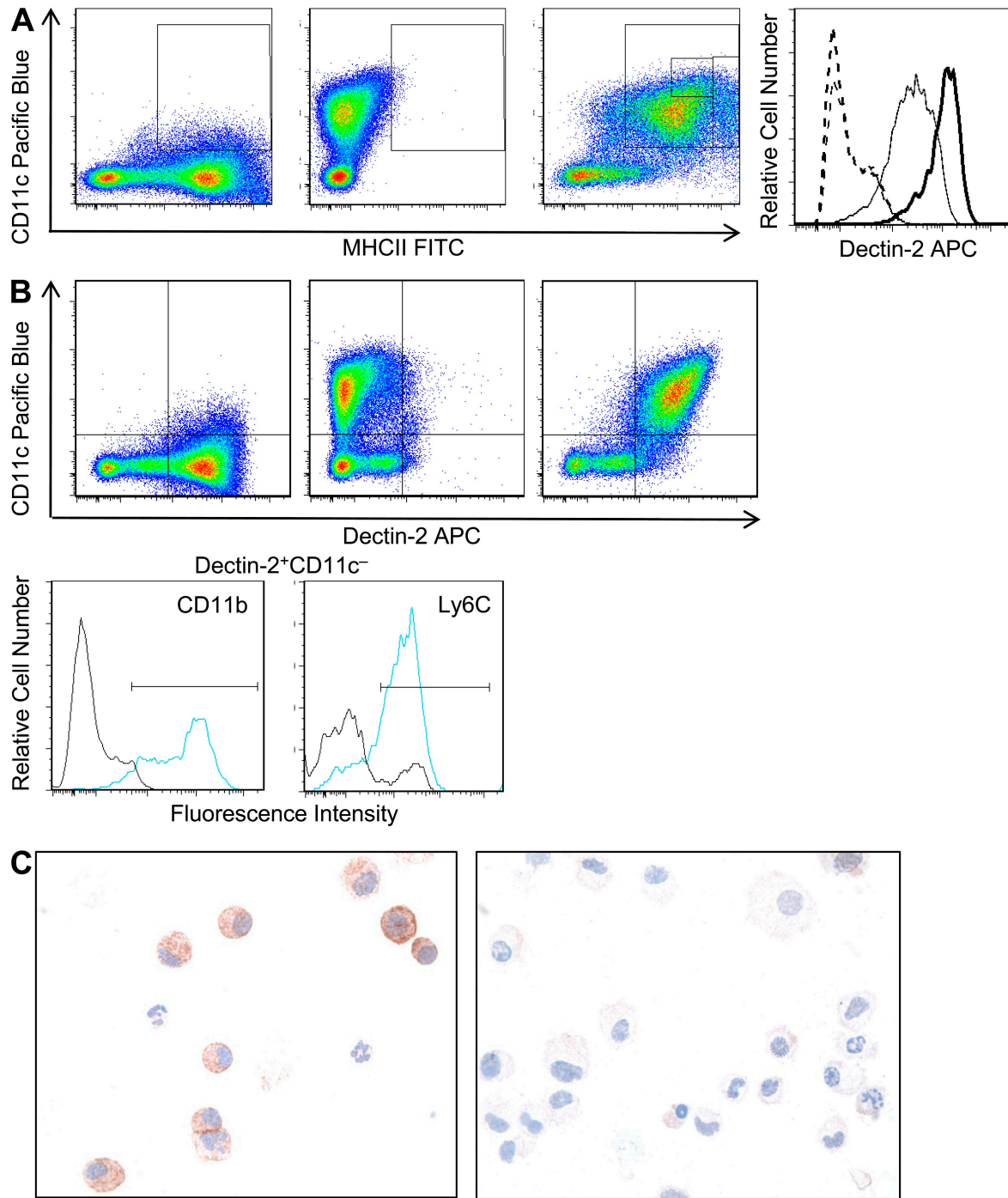
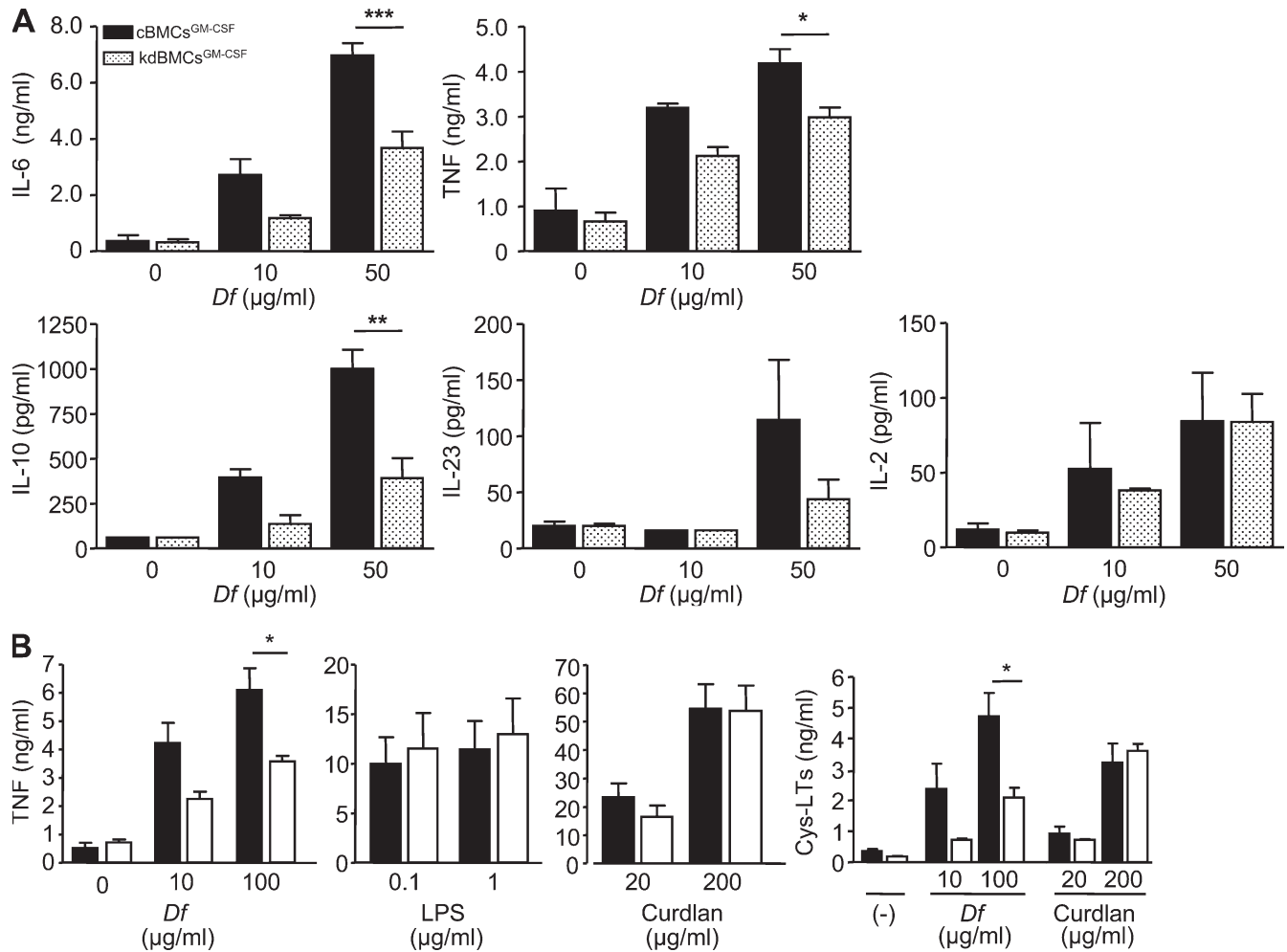


SUPPLEMENTAL MATERIAL

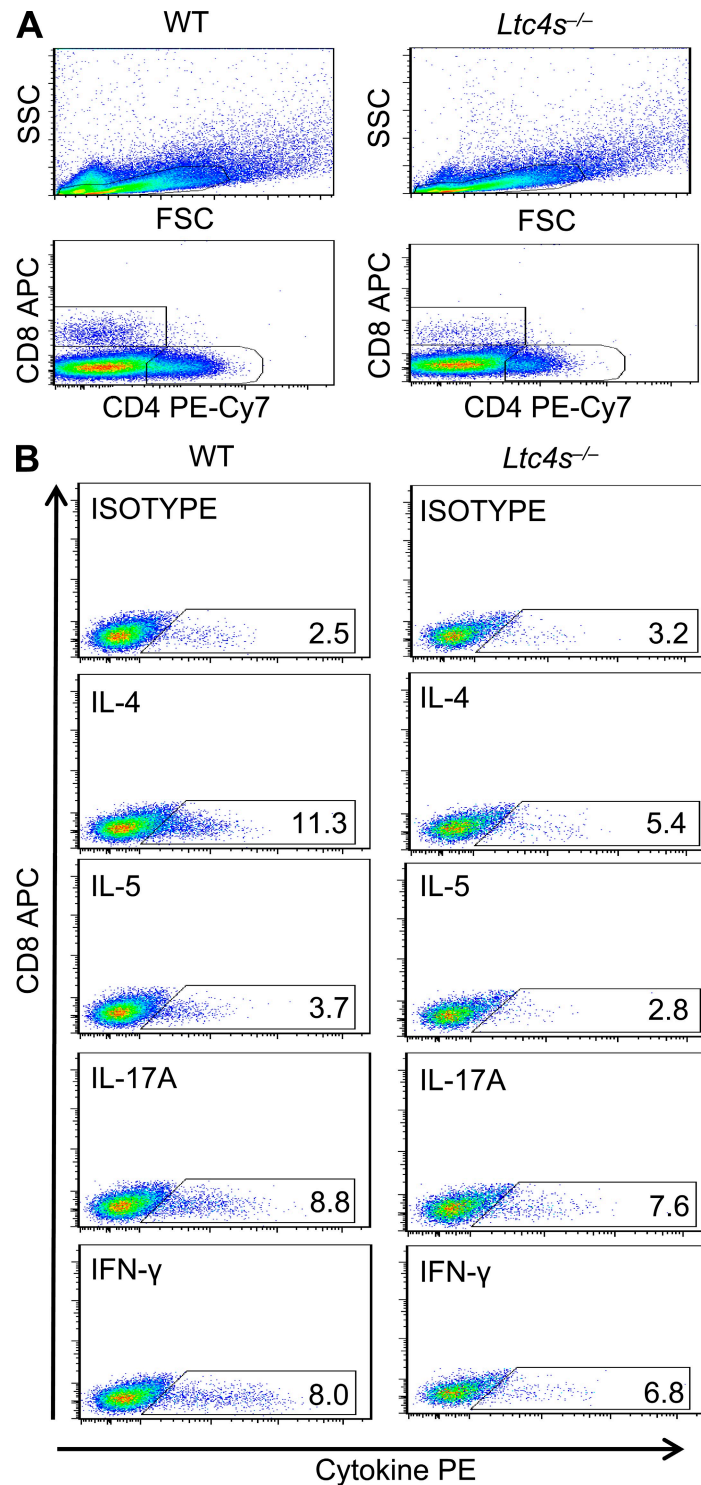
Barrett et al., <http://www.jem.org/cgi/content/full/jem.20100793/DC1>



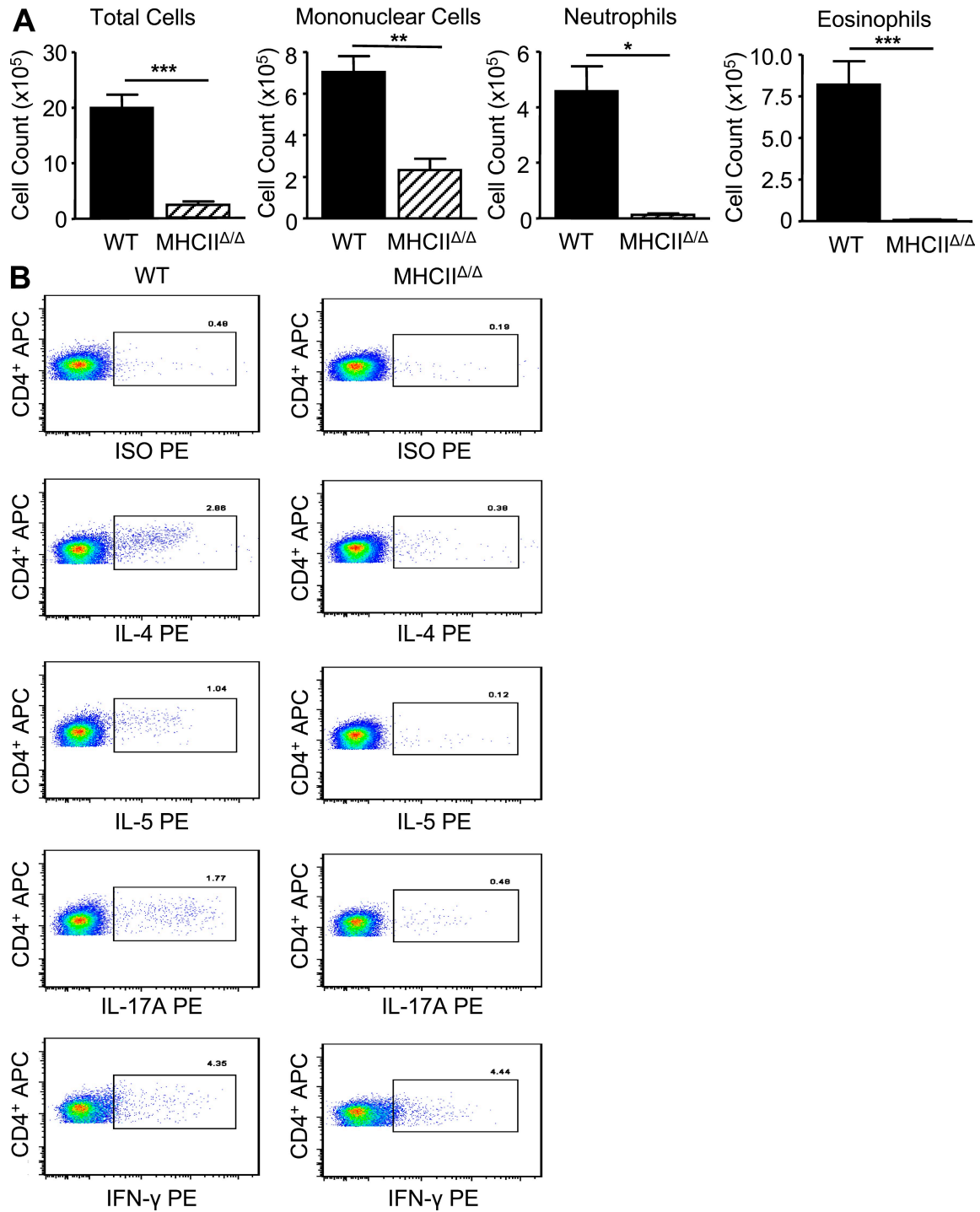
**Figure S1. Dectin-2 expression in GM-CSF-expanded BM cultures.** Day-7 BMCs<sup>GM-CSF</sup> were harvested and stained for the cell surface expression of CD11c, MHCII, and Dectin-2 and analyzed by flow cytometry. (A) Left, a representative histogram is shown of CD11c and MHCII expression, as gated by the isotype controls. Gating on a CD11c<sup>hi</sup>MHCII<sup>low</sup> and a CD11c<sup>+</sup>MHCII<sup>hi</sup> subset is also indicated (small box and small rectangle, respectively). MHCII staining was performed with saturating antibody concentrations. Right, Dectin-2 expression on the CD11c<sup>hi</sup>MHCII<sup>low</sup> subset (thick black line) and the CD11c<sup>+</sup>MHCII<sup>hi</sup> subset (thin black line) is shown, as well as the isotype control staining for each population (thick dashed and thin dashed lines, respectively). (B) Top, a representative histogram is shown of CD11c and Dectin-2 expression, as gated by the isotype controls. Bottom, CD11b and Ly6C expression on the Dectin-2<sup>+</sup>CD11c<sup>-</sup> population. (C) Dectin-2 expression on cytospin preparations from day-7 BMCs<sup>GM-CSF</sup> detected by immunocytochemistry. Positive staining is indicated in brown. Left, anti-Dectin-2. Right, rat IgG2a control. Results are representative of four cultures from two independent experiments.



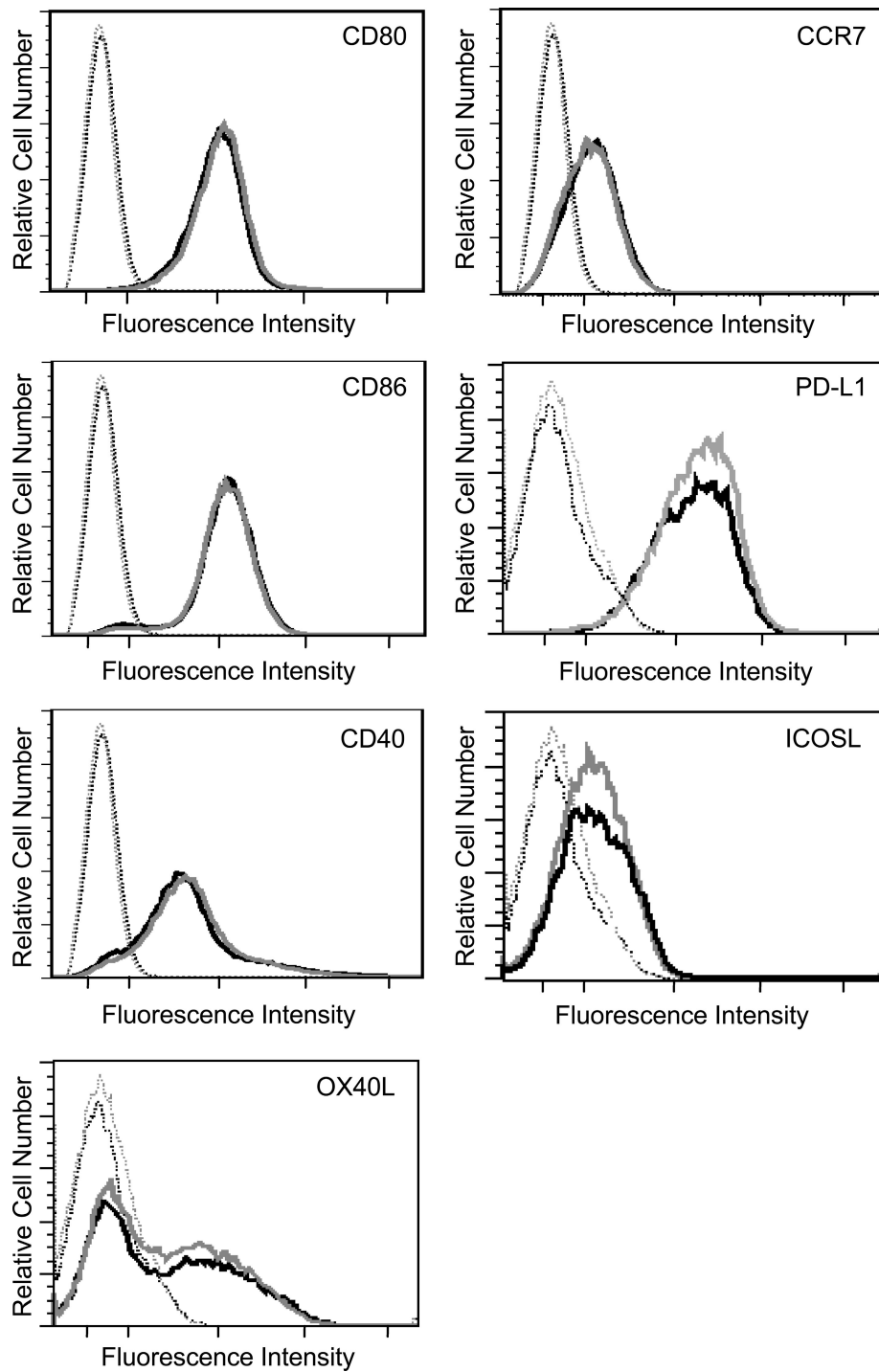
**Figure S2. Cys-LT and cytokine production by BMCs<sup>GM-CSF</sup> after lentiviral knockdown.** BMCs<sup>GM-CSF</sup> were infected with viral particles containing Dectin-2 shRNA (kdBMCs<sup>GM-CSF</sup>), an empty vector control (cBMCs<sup>GM-CSF</sup>), or a nontargeting vector control (ntcBMC<sup>GM-CSF</sup>), selected with puromycin, and harvested at day 7. (A) BMCs<sup>GM-CSF</sup> were stimulated for 30 h with *Df* at the indicated concentrations, and cytokines in the supernatant were measured by ELISA. Results are means  $\pm$  SEM ( $n = 5-6$  mice per group) from three independent experiments. \*,  $P = 0.01$ ; \*\*,  $P = 0.004$ ; \*\*\*,  $P = 0.002$ . Significance was determined with an unpaired Student's *t* test. (B) Left, TNF production from cBMCs<sup>GM-CSF</sup> (black) or kdBMCs<sup>GM-CSF</sup> (white) in response to *Df*, curdlan, or LPS at 4h. Results are means  $\pm$  SEM ( $n = 5-6$ ) from three independent experiments. \*,  $P = 0.04$ . Significance was determined with an unpaired Student's *t* test. Right, Cys-LT production from cBMCs<sup>GM-CSF</sup> (black) or kdBMCs<sup>GM-CSF</sup> (white) in response to 10 and 100  $\mu$ g/ml *Df* or 20 and 200  $\mu$ g/ml curdlan at 60 min. Results are means  $\pm$  SEM ( $n = 4-5$ ) from two independent experiments. \*,  $P = 0.02$ . Significance was determined with an unpaired Student's *t* test.



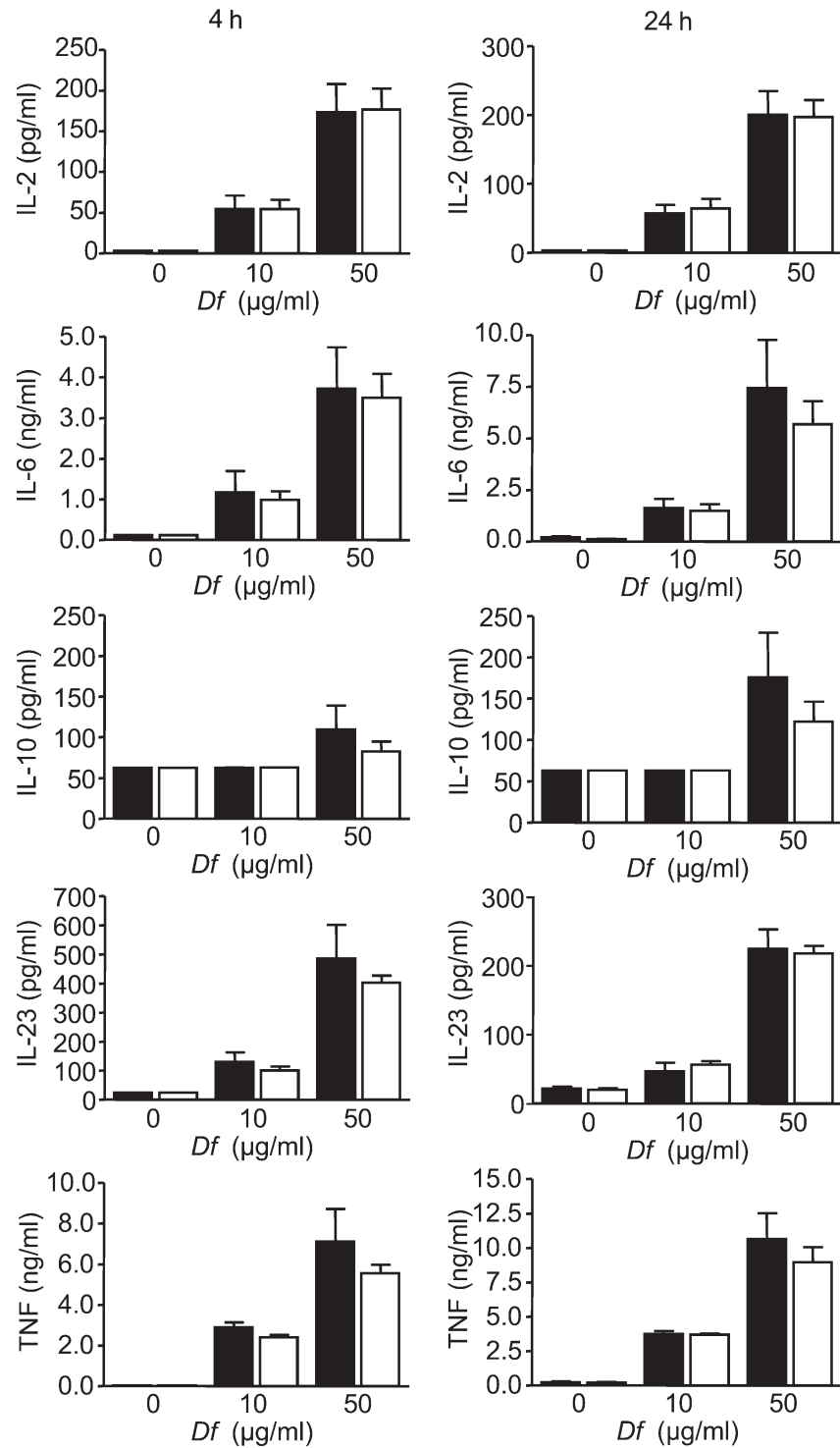
**Figure S3. *Df*-elicited CD4<sup>+</sup> Th2 cell recruitment to the lung.** WT mice were sensitized with saline-pulsed or *Df*-pulsed WT or *Ltc4s*<sup>-/-</sup> BMCs<sup>GM-CSF</sup> and challenged with *Df* as described in the Fig. 4 legend. Pulmonary mononuclear cells were isolated and stimulated with 50 ng/ml PMA and 1  $\mu$ M ionomycin for 2 h. 2.5  $\mu$ M monensin was added and cells were stimulated for an additional 8 h, fixed, permeabilized, stained for cell surface expression of CD4 and CD8 and for intracellular expression of IL-4, IL-5, IL-17A, and IFN- $\gamma$ , and analyzed by flow cytometry. (A) Cell size (top) and CD4 and CD8 expression (bottom) from mice sensitized with *Df*-pulsed WT (left) or *Ltc4s*<sup>-/-</sup> (right) BMCs<sup>GM-CSF</sup>. (B) Intracellular cytokine staining in CD4<sup>+</sup> cells in the lymphocyte gate from mice sensitized with *Df*-pulsed WT (left) or *Ltc4s*<sup>-/-</sup> (right) BMCs<sup>GM-CSF</sup> with the percentage of cytokine-positive cells noted.



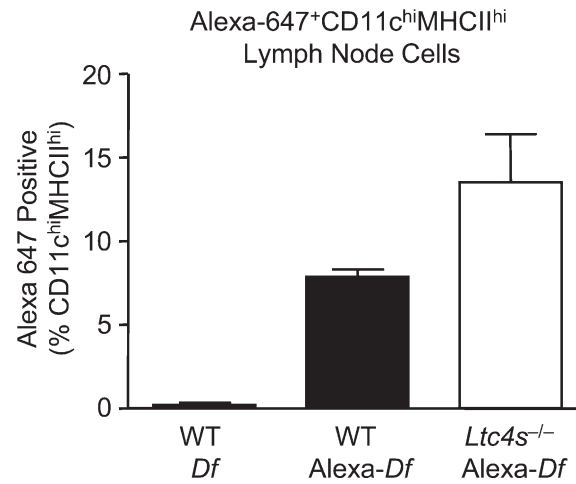
**Figure S4. BMCs<sup>GM-CSF</sup> MHCII expression is required for *Df*-elicited pulmonary inflammation and CD4<sup>+</sup> T cell cytokine generation.** WT C57BL/6 mice were sensitized with intranasal administration of 10<sup>4</sup> *Df*-pulsed WT or MHCII<sup>Δ/Δ</sup> BMCs<sup>GM-CSF</sup>, challenged with 10 μg *Df* on day 22, day 24, and day 26, and analyzed on day 27. (A) Cells from the BAL fluid were counted and 400 cells/slide in cytospins were counted for specific cell types. Results are means ± SEM (*n* = 8–10 mice per group) from two experiments. \*, *P* = 0.0005; \*\*, *P* = 0.0002; \*\*\*, *P* = 0.0001. Significance was determined with an unpaired Student's *t* test. (B) Parabronchial lymph node cells were isolated, stimulated as in Fig. S3 for a total of 6 h, stained for cell surface expression of CD4 and CD8 and for intracellular expression of IL-4, IL-5, IL-17A, and IFN-γ, and analyzed by flow cytometry. Representative histograms show intracellular cytokine staining on CD4<sup>+</sup> cells in the lymphocyte gate from WT recipients of *Df*-pulsed WT (left) or MHCII<sup>Δ/Δ</sup> (right) BMCs<sup>GM-CSF</sup>.



**Figure S5. WT and *Ltc4s*<sup>-/-</sup> BMCs<sup>GM-CSF</sup> have comparable cell surface expression of costimulatory markers.** WT and *Ltc4s*<sup>-/-</sup> BMCs<sup>GM-CSF</sup> were grown to day 7, pulsed with 100 µg/ml *Df* for 24 h, harvested, and stained for the cell surface expression of CD11c, MHCII, CD80, CD86, CD40, OX40L, CCR7, PD-L1, and ICOSL, and analyzed by flow cytometry. Representative histograms are shown of costimulatory marker expression on CD11c<sup>+</sup>MHCII<sup>+</sup> BMCs<sup>GM-CSF</sup>. Specific stainings are shown for WT (thick black lines) and *Ltc4s*<sup>-/-</sup> (thick gray lines) BMCs<sup>GM-CSF</sup>, and isotype control stainings are shown for WT (dotted black lines) and *Ltc4s*<sup>-/-</sup> (dotted gray lines) BMCs<sup>GM-CSF</sup>.



**Figure S6.** WT and *Ltc4s*<sup>-/-</sup> BMCS<sup>GM-CSF</sup> generate comparable concentrations of *Df*-elicited cytokines at 4 and 24 h. WT (black bars) and *Ltc4s*<sup>-/-</sup> (white bars) BMCS<sup>GM-CSF</sup> were grown to day 7 and stimulated with *Df* at the indicated concentrations for either 4 or 24 h, and cytokines in the supernatants were measured by ELISA. Results are means ± SEM (*n* = 4 mice per group) from two independent experiments.



**Figure S7. WT and *Ltc4s*<sup>-/-</sup> BMCs<sup>GM-CSF</sup> show comparable migration to the lung-draining lymph nodes.** Day-7 BMCs<sup>GM-CSF</sup> were stimulated with 100 µg/ml of unlabeled *Df* or Alexa Fluor 647-*Df* for 18 h, and 10<sup>6</sup> cells were transferred intranasally into WT recipients. 24 h later, the draining lymph nodes were harvested and digested, and single cell suspensions were stained for CD11c-PE-Cy7 and MHCII-PE and analyzed by flow cytometry. The percentage of CD11c<sup>hi</sup>MHCII<sup>hi</sup> cells that were Alexa Fluor 647 positive is shown. Results are means ± SEM (*n* = 4–5 mice per group) from two independent experiments.