

**Figure S1**. **Transcriptional profiles of selected cytokines and chemokines in ear dermal cells during the course of infection.** C57BL/6 mice were infected in the ear dermis with 1000 *Lm*Sd metacyclic promastigotes and total mRNA was isolated from the ear tissue at the indicated times during infection, reverse transcribed and analyzed by real-time PCR. Target genes were normalized to endogenous control, and the values shown are the fold increase relative to expression in naïve ears. Data are represented as means ± SD of 8 samples (4 mice, 8 ears) at each time point. The results are representative of 2 independent experiments.







**Figure S3. Neutrophils are still recruited to the chronic lesions in** *LmSd* **infected** *CXCL1-/- mice***.** Mice were infected in the ear dermis with 1000 *Lm*Fn or *Lm*Sd metacyclic promastigotes and ear tissues were processed at 20 wks post-infection. The total numbers of CD11b+ and CD11b+Ly6C<sup>int</sup> Ly6G<sup>+</sup> (PMN) per ear are shown with means ± SD.



Fig S4. Treatment with RB6 antibody fails to moderate pathology or to reduce neutrophil recruitment to the site. C57BL/6 mice were infected in the ear dermis with 1000 *Lm*Sd metacyclic promastigotes. At 10 wks post-infection, mice were injected i.p. with 0.5 mg RB6-8C5 or GL113 every 3-4 days for a total of 5 injections. Lesion scores were measured at 12.5 wks, and ear tissue was processed to determine the total numbers of CD11b+ and CD11b+Ly6C<sup>int</sup> Ly6G<sup>+</sup> (PMN) per ear. Values shown are means  $\pm$  SD. This experiment was repeated a total of three times with similar results.