

Supplemental Figure 1. LPS induces TLR4 dependent HSPC expansion. A)

Representative flow plots of BM LSK 24 hours after PBS or LPS injection. % of **B)** LSK and **C)** LT-HSC (CD150⁺CD135⁻LSK) and ST-HSC (CD150⁻CD135⁺ LSK) expansion in C3H/HeOuJ and C3H/HeJ mice following intraperitoneal injection of 5 µg of LPS; **D)** % LSK expansion in B6, MyD88^{-/-}, and TRIF^{-/-} mice; or **E)** SEV129 mice and IFNAR^{-/-} mice 24 hours following intraperitoneal injection of 5 µg of LPS. *p<0.05 for PBS vs LPS. ** p<0.05 for wild type LPS treated versus knockout LPS treated mouse. Experiments were performed at least 3 separate times with an n=3-4 per group.

Supplemental Figure 2. LT-HSCs from septic mice are capable of long term

reconstitution. A) Schema for identification of Lineage⁻c-kit⁺sca-1⁺ LSK and CD150⁺CD135⁻ LT-HSCs and CD150⁻CD135⁺ activated ST-HSCs 36 hours after sham or CLP surgery. **B)** Gating schema for sorting of LT-HSCs 36 hours after sham or CLP for long term reconstitution experiment with FACS Aria following magnetic sorting for lineage-depletion showing pre-sort (top panels) and **C)** post-sort purities. **D)** Survival of lethally irradiated mice following transfer of ten CD150⁺CD135⁻LSK cells from sham (n=7) or CLP (n=7) mice with 5x10⁵ Lineage⁺ cells for radioprotection, or 5x10⁵ Lineage⁺ cells without CD150⁺CD135⁻LSK cells (n=10).

Supplemental Figure 3. CLP-induced bone marrow reprogramming of

hematopoiesis involves depletion of B cells and increased myeloid differentiation

of hematopoietic progenitors. A) Time course of bone marrow B cell depletion following CLP. Representative flow plots of AA4.1⁺B220^{low} pre-B cells 36 hours after sham treatment (left panels) and CLP treatment (right panels) in **B)** B6 control and **C)** MyD88^{-/-} mice with graphical representations (bottom panels) of % pre-B cells in bone

marrow at 36 hours or Day 7 following sham or CLP treatment. D) 36 hours following sham or CLP treatment, total Lineage⁻ hematopoietic stem and progenitor cells were isolated by magnetic bead sorting and 1×10^4 Lineage⁻ cells were cultured in methylcellulose with the indicated cytokines. 14 days later, colonies were counted with a light microscope. Each point represents Lineage⁻ cells from two mice of the same treatment group. All experiments were performed at least twice with an n=4 per group.

Supplemental Figure 4. Blockade of inflammatory factors that regulate hematopoiesis have no effect on CLP induced LSK expansion. 36 hours following sham or CLP procedure BM was analyzed for LSK expansion. **A)** C57BL/6, IL1R^{-/-} and IL-6^{-/-} animals. **B)** Sham or CLP mice with or without treatment with pegylated soluble TNFR1 (100 mg/kg). **C)** C57BL/6 sham and CLP mice with or without treatment with apocynin (apo-5mg/kg) and n-acetyl cysteine (NAC-100mg/kg). **D)** C57BL/6 sham and CLP mice treated with or without indomethacin (5mg/kg). All treatments were given i.p. 2 hours prior to surgery. Apo+NAC and TNFR1 were also given 24 hours following CLP.

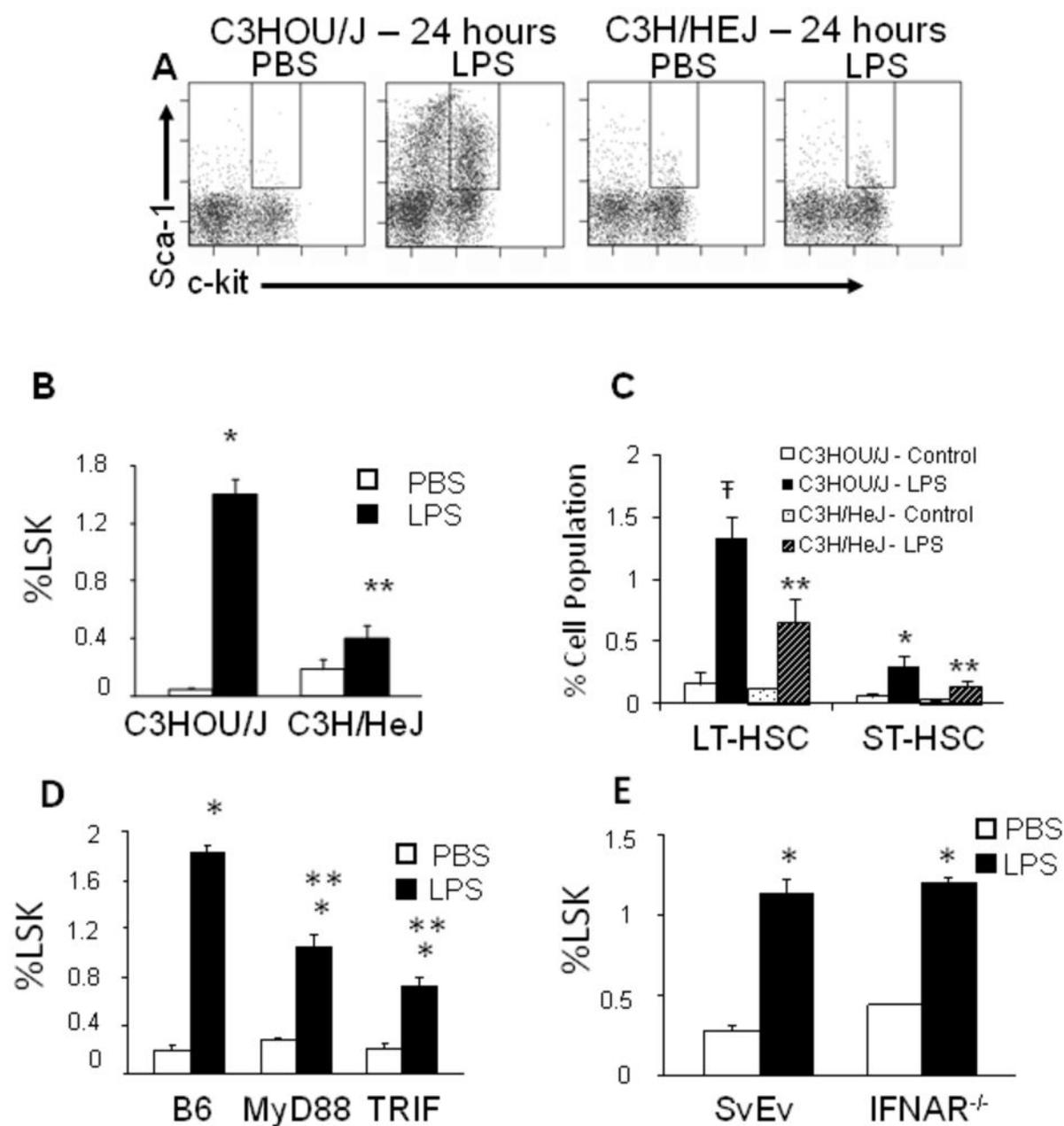
Supplemental Figure 5. Staphylococcal superantigen induces HSPC expansion and S. aureus sepsis induces a temporary leukopenia. **A)** Expansion of LSK, LT-HSCs and ST-HSCs in B6 mice treated IP with 10 μ g of Staphylococcal enterotoxin B (SEB) or same volume of PBS as control. **B)** represents peripheral blood total leukocyte counts.

Supplemental Figure 6. Time course of BM repopulation following CLP. Mice were subjected to Sham or CLP and at different time points hind limb bones were harvested

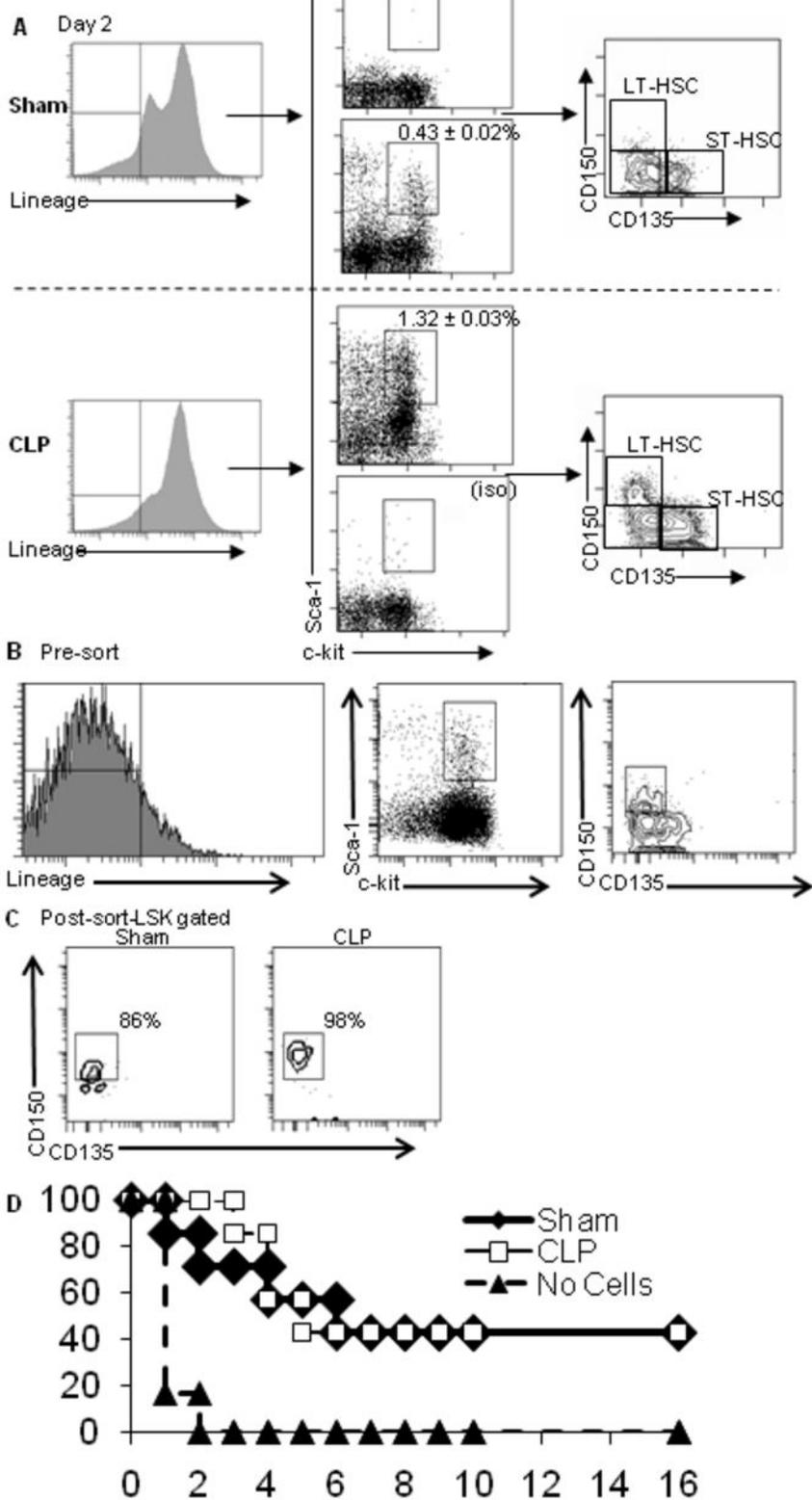
for H&E staining. All micrographs are 100x magnification. **A)** At 12 hours, there is increased dilatation of BM vasculature and increased space from cell loss is beginning. **B)** By 36 hours after CLP, BM space is maximal and there is continued dilatation of the BM vasculature. **C)** 96 hours after CLP, there is continued BM space but the BM vasculature has returned to normal. **D)** By Day 7, the BM has completely regained cellularity with dramatically increased myeloid elements. **E)** Time course of the total BM cellularity following sham or CLP * $p < 0.05$ control vs treatment by Student's t-test. Experiments were performed three independent times with at least $n=3$ per group. Figure represents data from one experiment.

Supplemental Figure 7. Expansion of total cell numbers in response to CLP or *S. aureus* from selected experiments. Total cell counts for LSK, LT-HSC, and ST-HSCs were calculated. **A)** Total cell number expansion of LT-HSC and ST-HSC in Sham and CLP treated WT (C57BL/6) mice, 36 hours after surgery (experiment from Figure 1 B and C). **B)** Total cell number expansion of LSK, LT-HSC and ST-HSC in WT, MyD88^{-/-} and TRIF^{-/-} mice 36 hours after Sham or CLP (experiment from Figure 2B). **C)** Total cell number expansion of LSK, LT-HSC and ST-HSC in WT versus MyD88^{-/-}TRIF^{-/-} (DKO) mice 36 hours after IV infection with 5×10^6 SA (experiment from Figure 2D). **D)** Total cell number expansion of LSK, LT-HSC and ST-HSC in Rag1^{-/-} mice 36 hours after IV infection with 5×10^6 SA (experiment from Figure 2E). * $p < 0.05$, $\mp p < 0.001$ by Student's t test or one way ANOVA.

Supplemental Figure 1

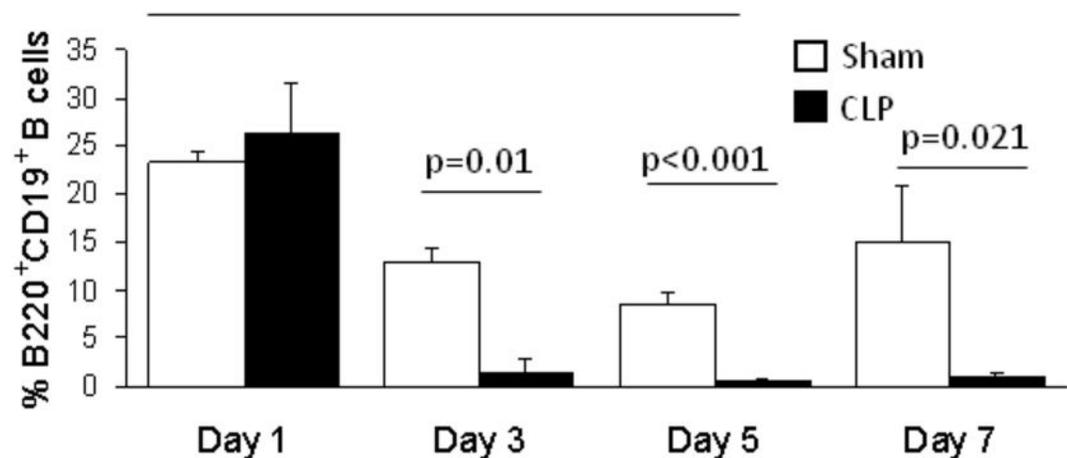


Supplemental Figure 2



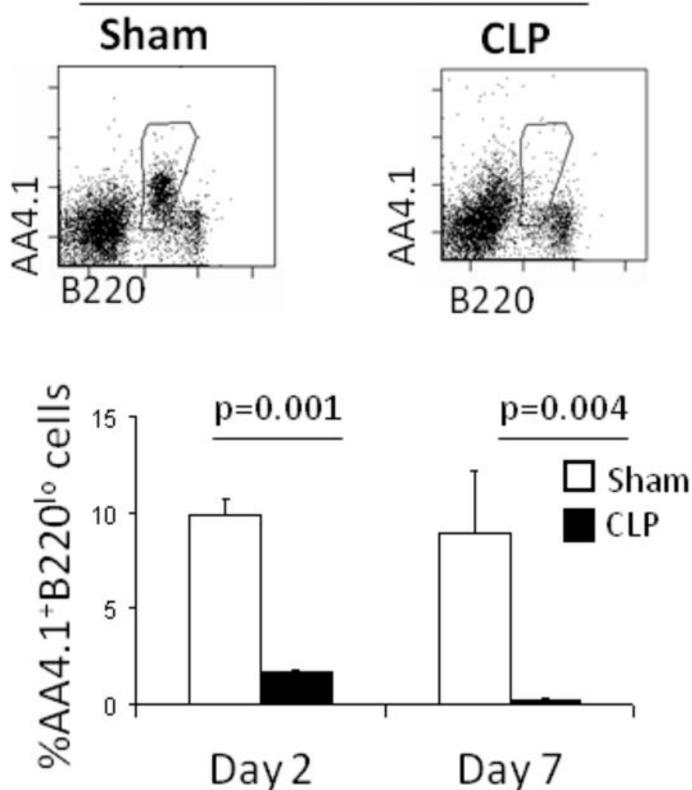
A

C57Bl/6 Total BM B cells

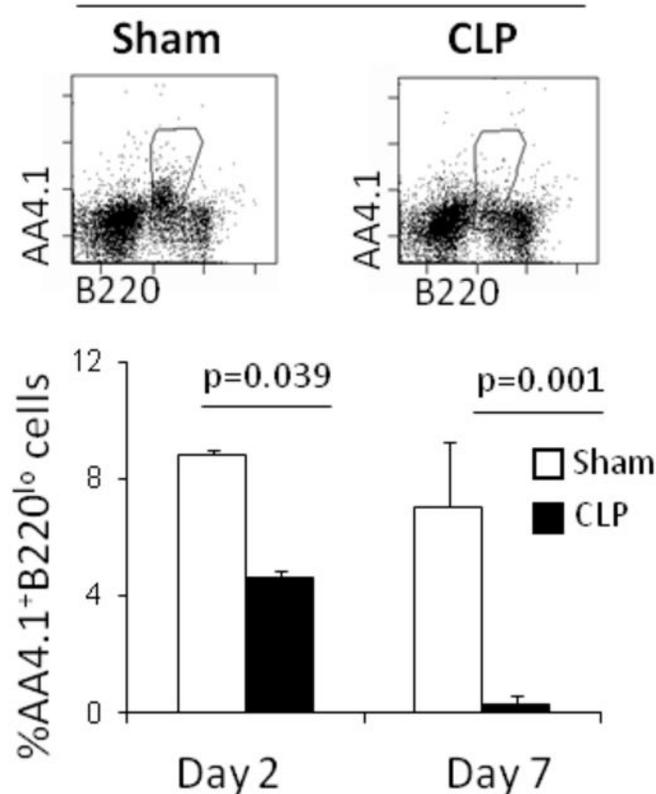


B

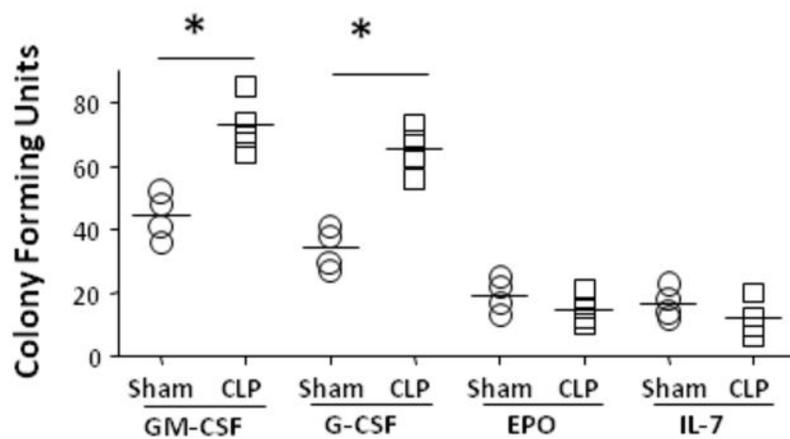
C57Bl/6 Day 2



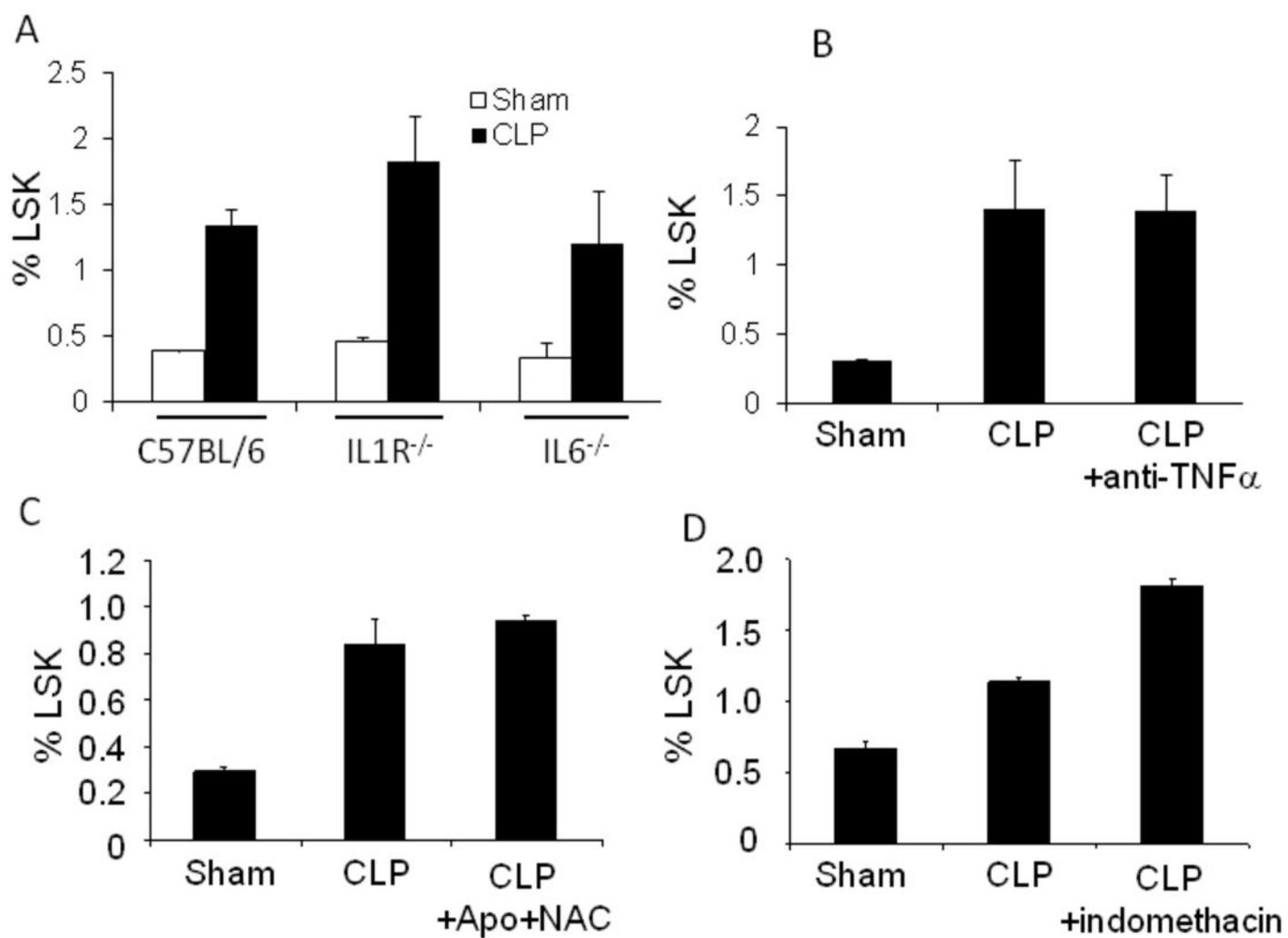
C

MyD88^{-/-} Day 2

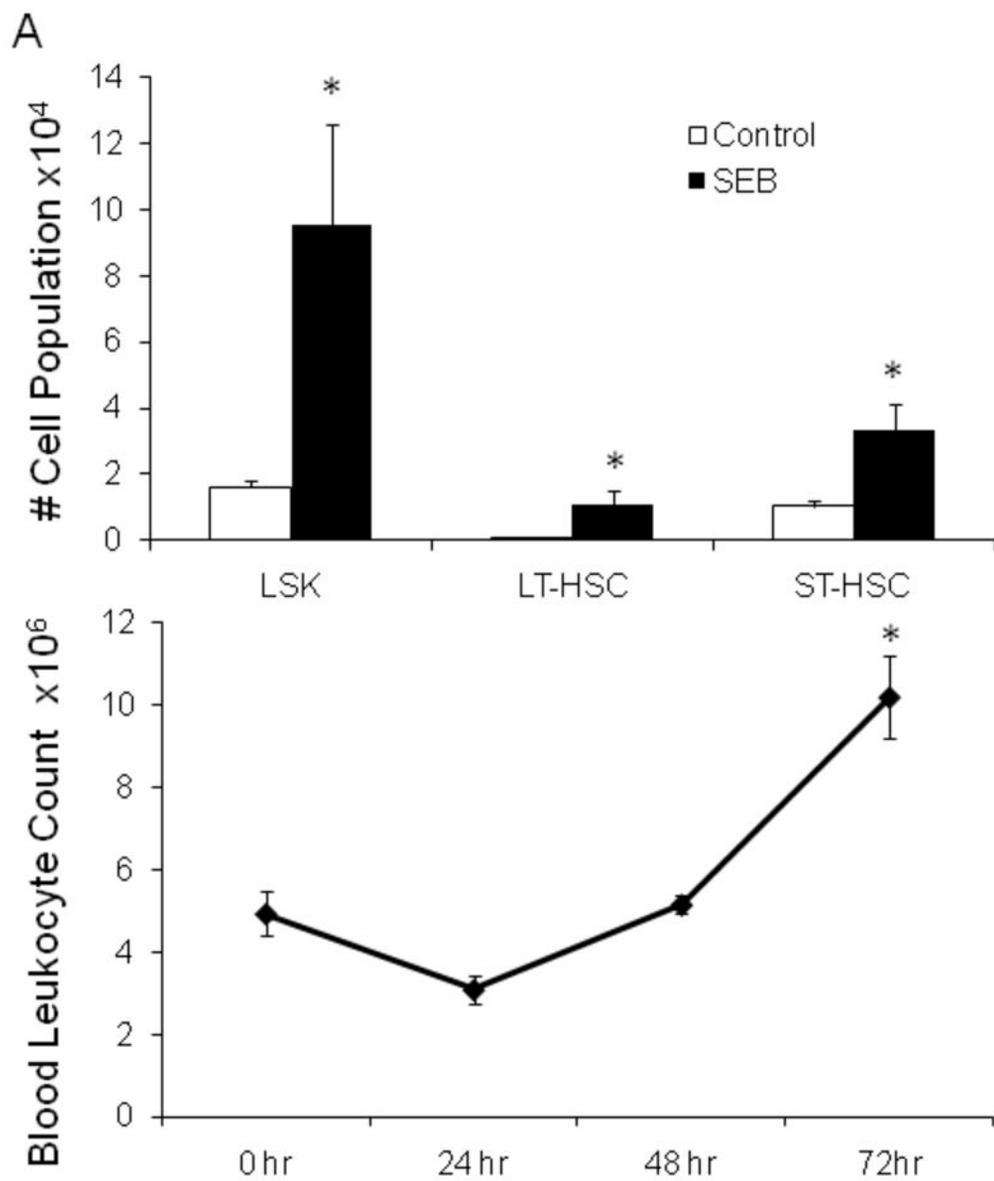
D



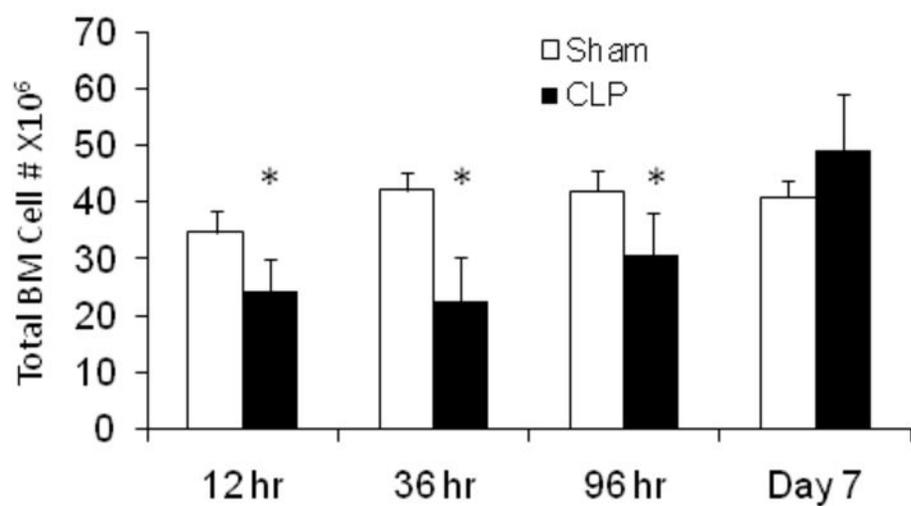
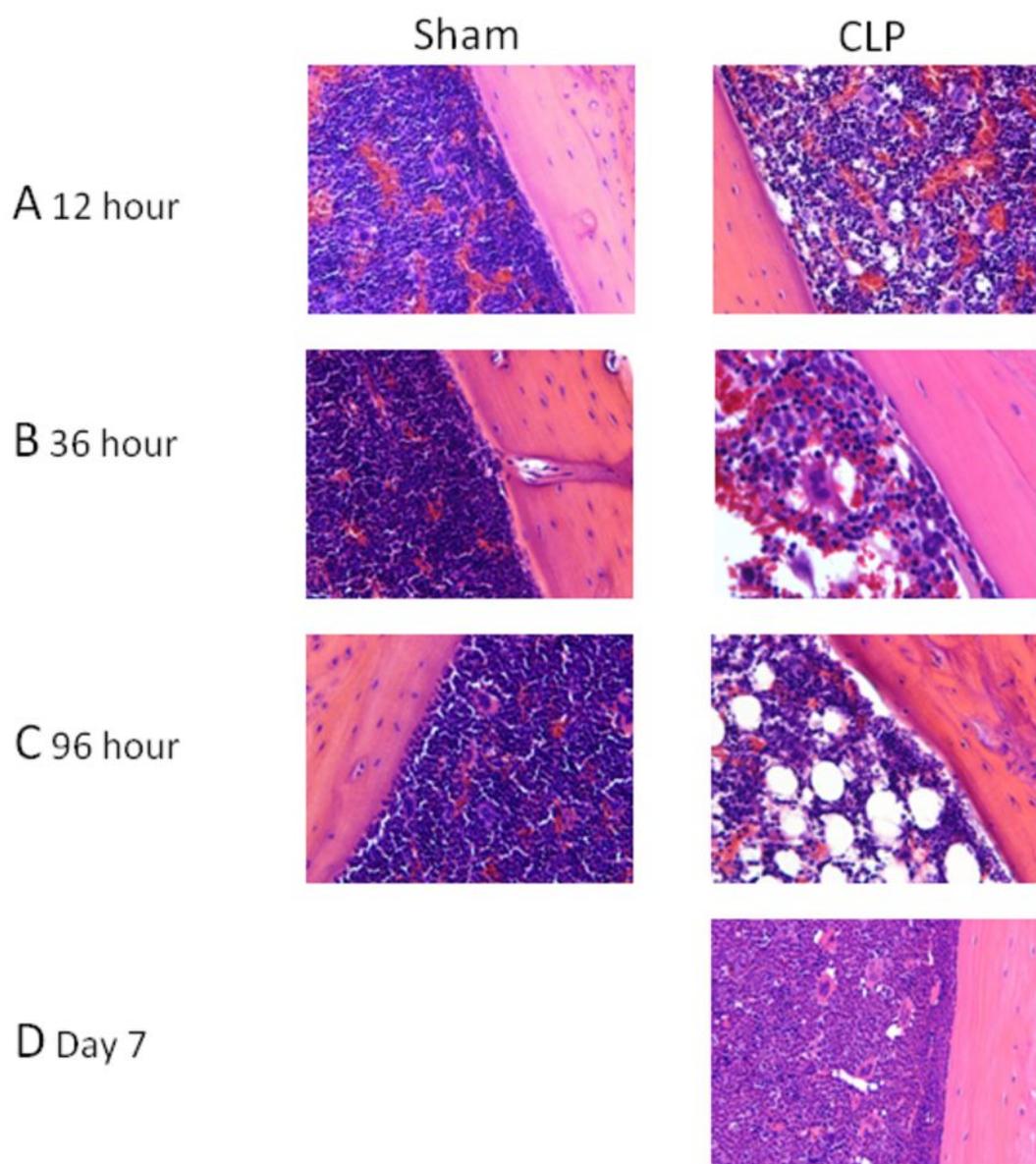
Supplemental Figure 4



Supplemental Figure 5



Supplemental Figure 6



Supplemental Figure 7

