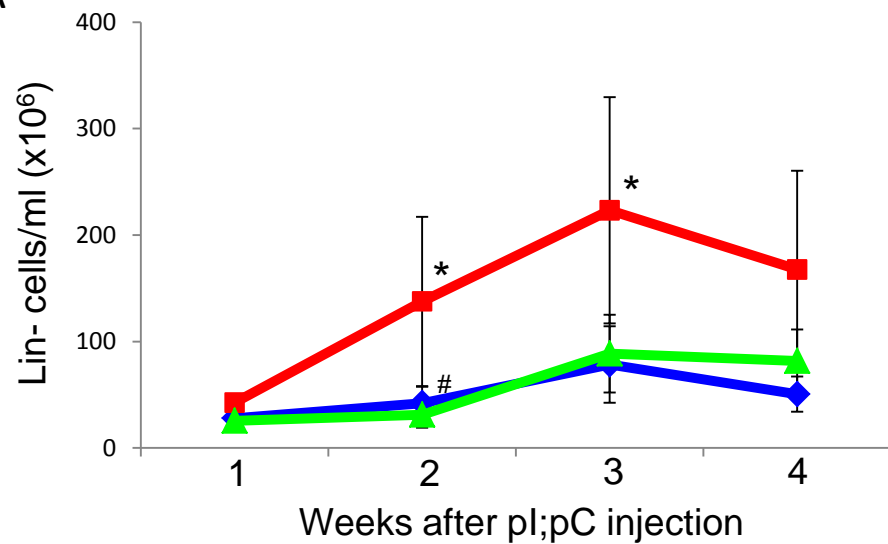
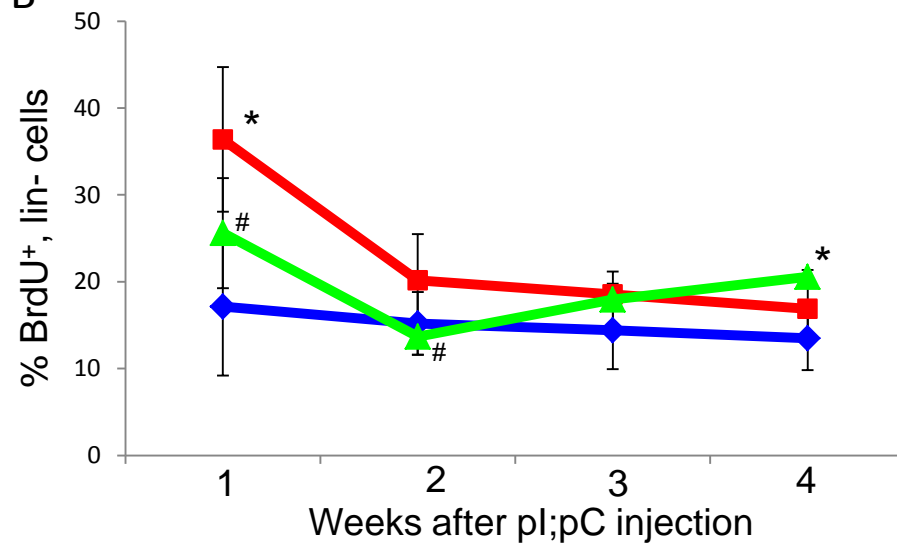


Supplemental Figure 4

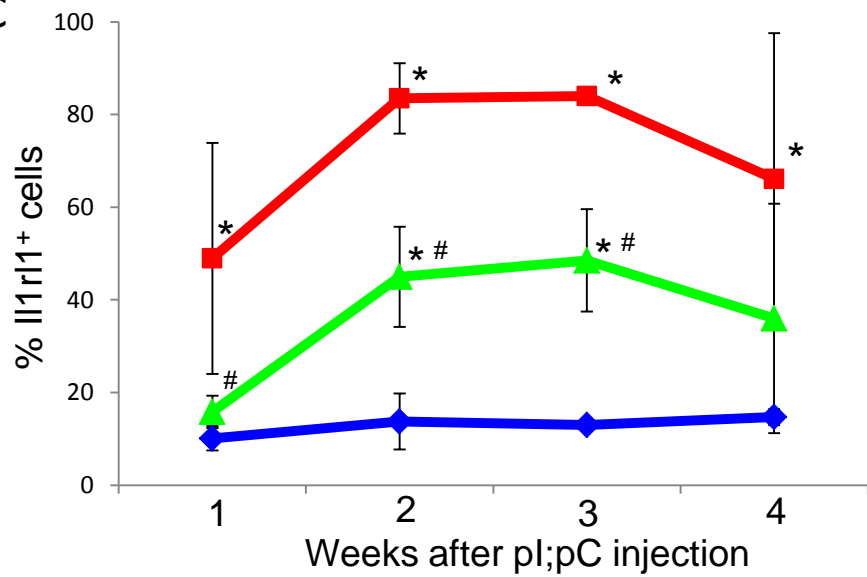
A



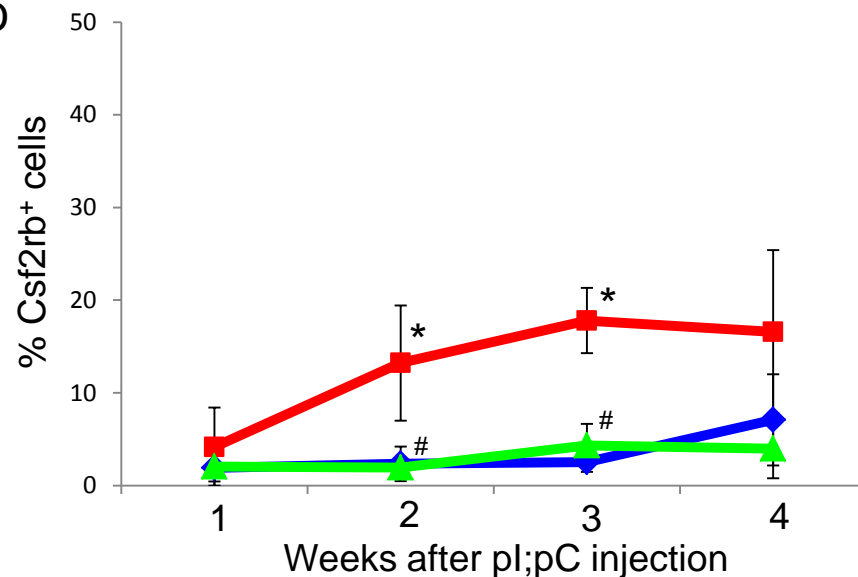
B



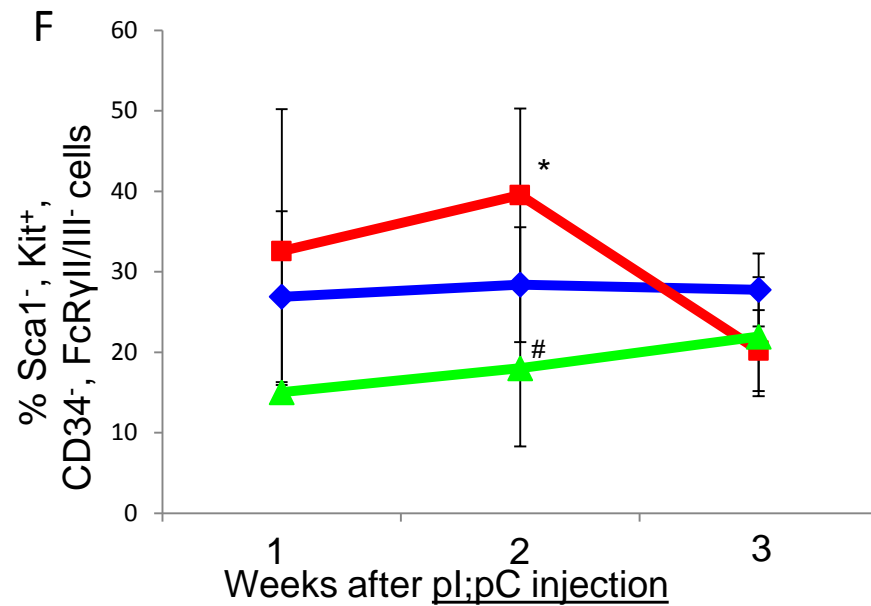
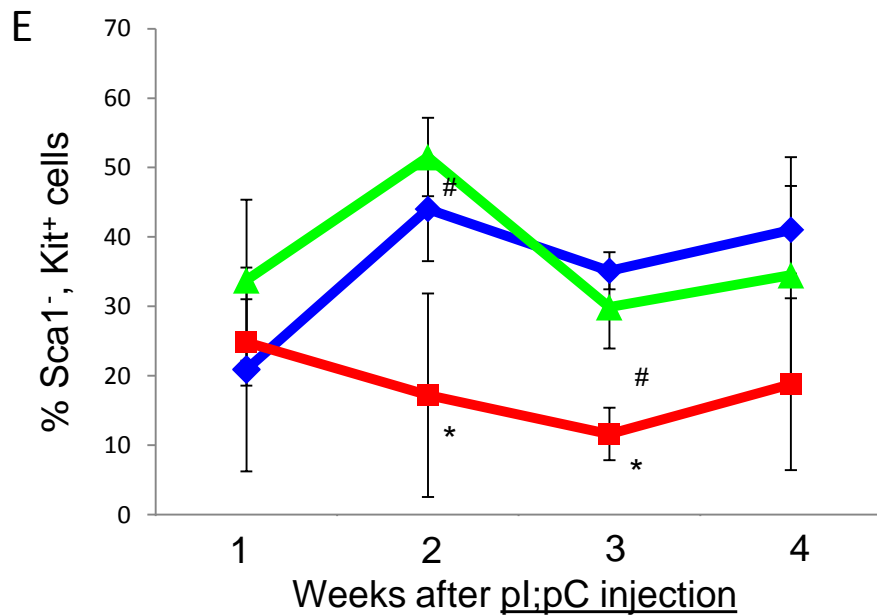
C



D



◆ *Runx1*^{+/+}
 ■ *Mx1-Cre*⁺, *Cbfb*^{+I56M}, *Runx1*^{+/+}
 ▲ *Mx1-Cre*⁺, *Cbfb*^{+I56M}, *Runx1*^{+/-}



◆ *Runx1*^{+/+} ■ *Mx1-Cre*⁺, *Cbfb*^{+l56M}, *Runx1*^{+/+} ▲ *Mx1-Cre*⁺, *Cbfb*^{+l56M}, *Runx1*^{l/z}

Supplemental Figure 4. *Runx1* activity is required for adult hematopoietic defects induced by *Cbfb-MYH11*. Line graphs of (A) the number of *lin*⁻ cells per ml ($\times 10^6$) in the bone marrow, (B) percentage (%) of BrdU⁺ cells in the *lin*⁻ bone marrow cell population, (C) the percentage of *Il1rl1*⁺ cells in the *lin*⁻ bone marrow cell population, (D) the percentage of *Csf2rb*⁺ cells in the *lin*⁻ bone marrow cell population, (E) the percentage of *Sca1*⁻, *Kit*⁺ cells in the *lin*⁻ bone marrow cell population, and (E) the percentage of *CD34*⁻, *FcRγII/III*⁻ cells in the *Sca1*⁻, *Kit*⁺, *lin*⁻ bone marrow cell population, in mice of the indicated genotypes, at the indicated time points after *Cbfb-MYH11* induction. $N \geq 3$ for each genotype. * indicates $p \leq 0.05$ as compared to *Runx1*^{+/+} mice. # indicates $p \leq 0.05$ as compared to *Mx1-Cre*⁺, *Cbfb*^{+l56M}, *Runx1*^{+/+} mice.