

Yang FF, et al. Figure S2

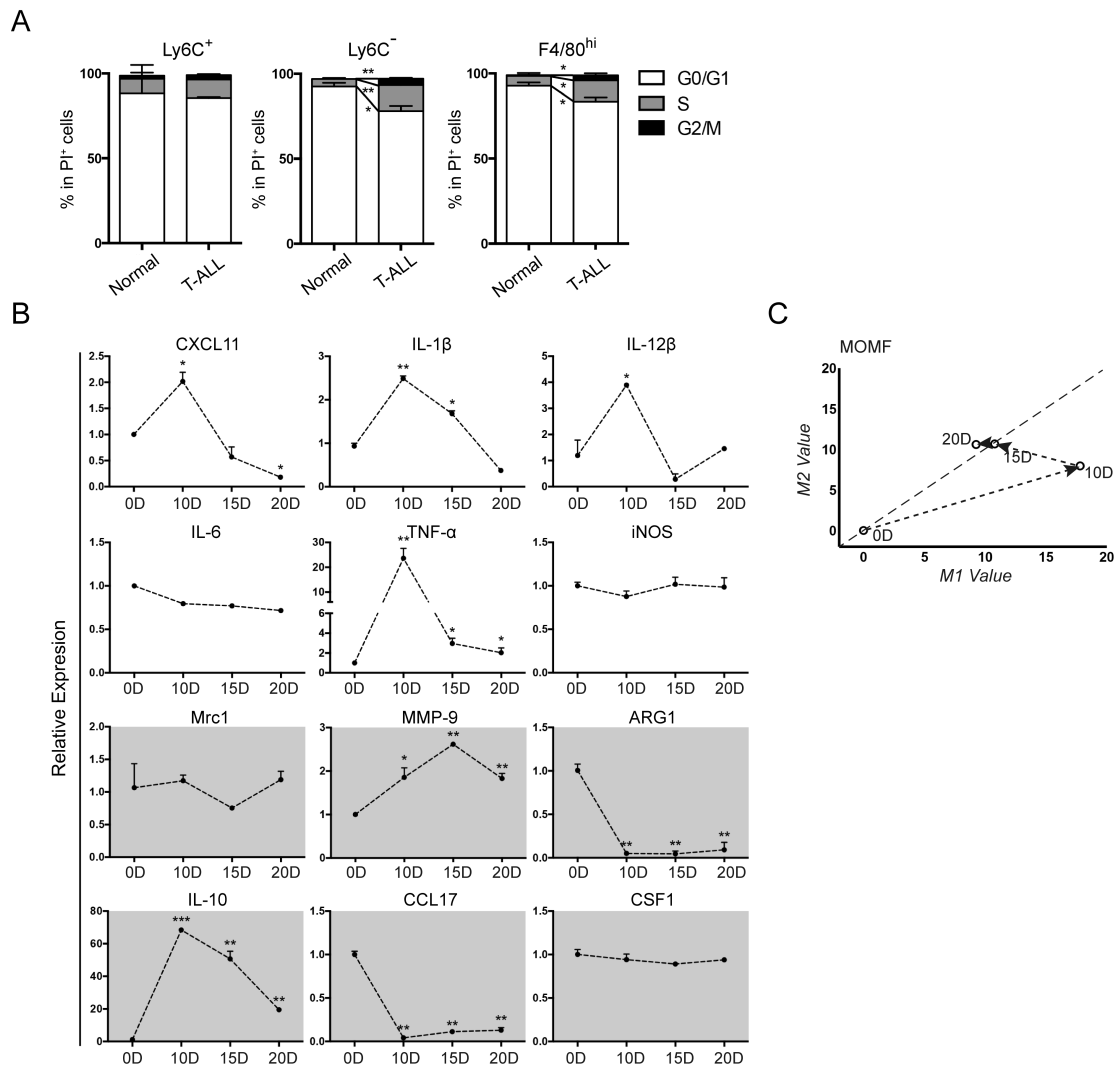


Figure S2. Cell cycle analysis of SP LAM subsets and phenotypic evolution of MOMF in the liver of T-ALL mice.

(A) Normal macrophage subsets and LAM subsets (day 15) were sorted from the SP. Cell cycle was analyzed by propidium iodide staining (n=5). Data are representative of three independent experiments. Student's t test was performed. (B) T-ALL cells were transplanted into WT mice. Mice were sacrificed at the indicated time points, and the expression of phenotype-associated genes in MOMF was monitored by RT-PCR. For each gene, the RQ value of normal MOMF (day 0) was designated as 1.000. (C) A two-dimensional illustration of MOMF phenotype is shown. The phenotypic evolution of MOMF is indicated by arrows. Data are representative of three independent experiments. One-way ANOVA was performed (*: p<0.05; **: p<0.01; ***: p<0.001).