



Figure S6: Gating scheme for other immune cell subsets

All tumors were analyzed for the percentage of CD45⁺ leukocytes that were macrophages, monocytes, neutrophils, DCs or NK cells. Lineage-negative cells (Thy1.2-CD4-CD8-B220-CD138-Sca-1⁻) were analyzed for expression of CD11c, CD11b, Ly6C, Ly6G, MHC II as indicated (middle rows). Thy1.2-B220⁻ cells were analyzed for CD49 β expression as indicated to identify NK cells (bottom row). All data shown are from a normal spleen.