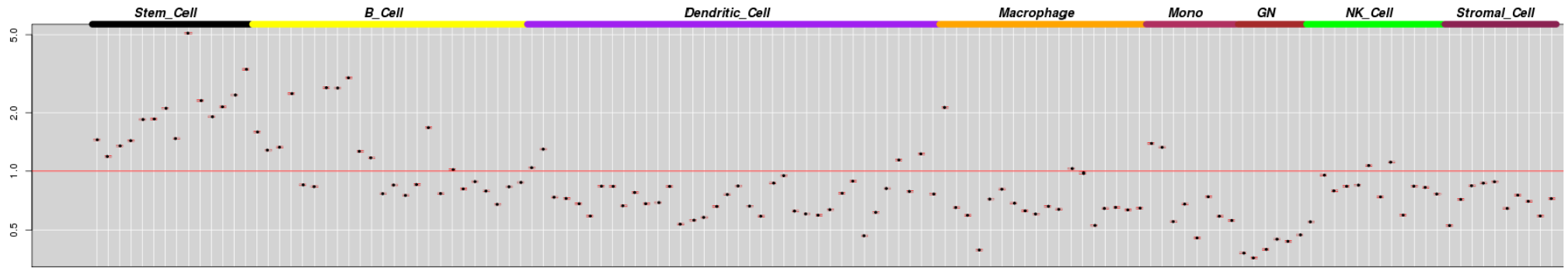
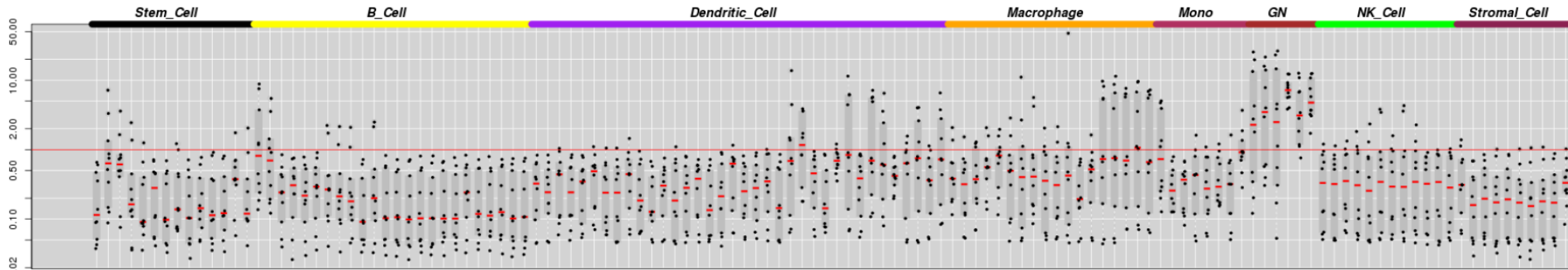


A**B**

Supplementary Figure 3: Gene mapping of the mouse microenvironment and immune cells. Wplots of mouse genes that were differentially expressed in the microenvironment of Gal-7+ tumors. (A) Wplot representing the expression of Zmynd19 gene. Its expression is highlighted in stem cells and multi-lineage progenitor cells. (B) Wplot representing the expression of Stfa1, Stfa2, Retnlg, S100a8, S100a9, IL1 β , CXCL2, CCL3, CCL4, and Arg1 genes in granulocytes, dendritic cells and macrophages, moderately enriched in stem cells, β cells and NK cells. The red line shows the basal expression; signals above this threshold can be considered as enriched.