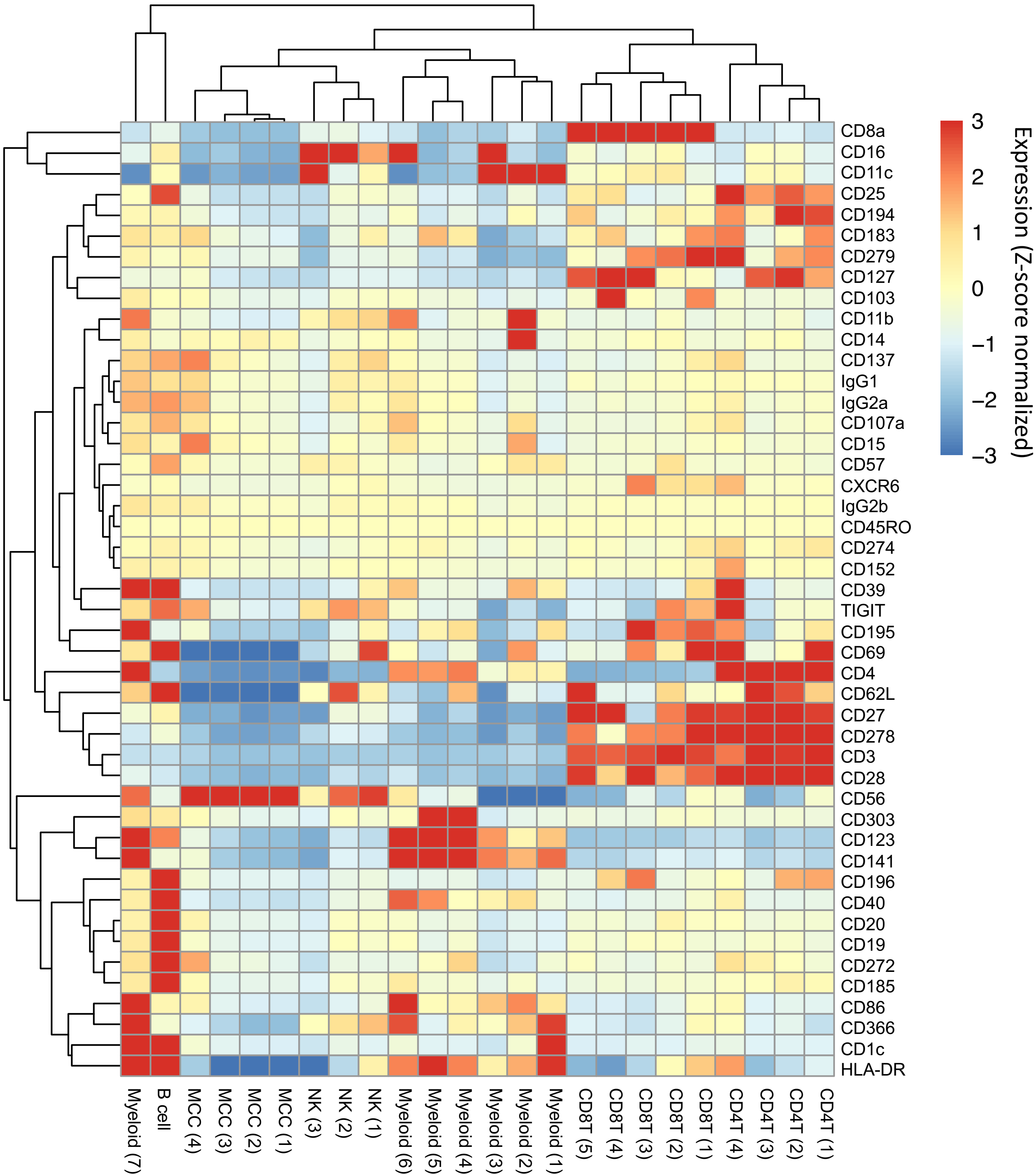




Supplemental Figure 1. Clustering justification of single cell RNAseq. Rows represent proteins measured via CITEseq. Rows represent clusters. Cell color average protein expression for a given protein in each cluster normalized to a Z statistic for each protein.