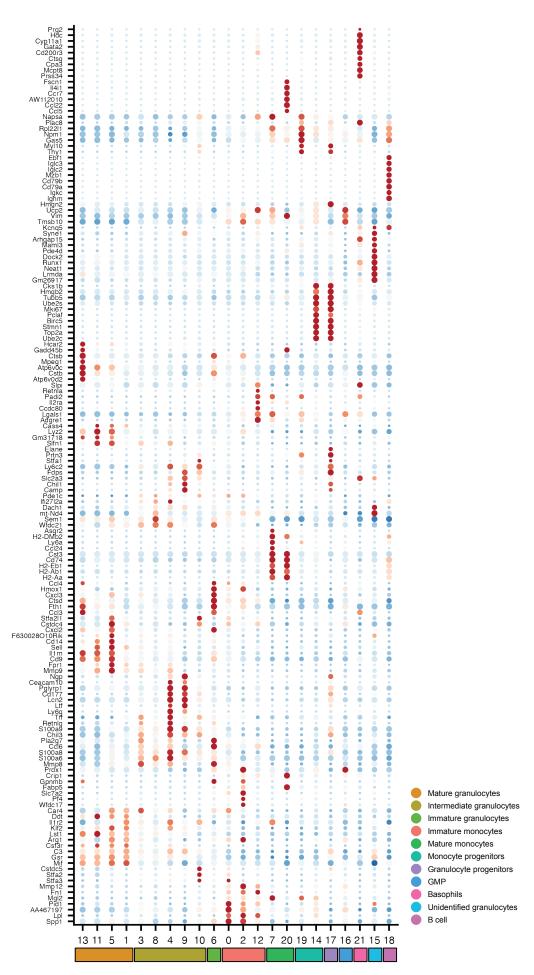
Percent Expressed • 0 • 25 • 50 • 75 • 100



Supplementary Figure s3. Top marker genes used for cluster identification. Normalized expression and per cluster percentage expression of top 10 marker genes for each cell cluster for the scRNA sequencing analysis on the integrated sample including MDSCs generated \pm 1.5 μ M DAB.