



Figure 1—Figure Supplement 2. Analysis of WT clusters.

Expression of marker genes (Y-axis) in each WT cluster (X-axis) in a heatmap. Genes that reach the criteria (fraction ≥ 0.8 , specificity ≥ 0.2 , q -value ≤ 0.01) are defined as marker genes. Expression value is visualized with range between -2 to 2.