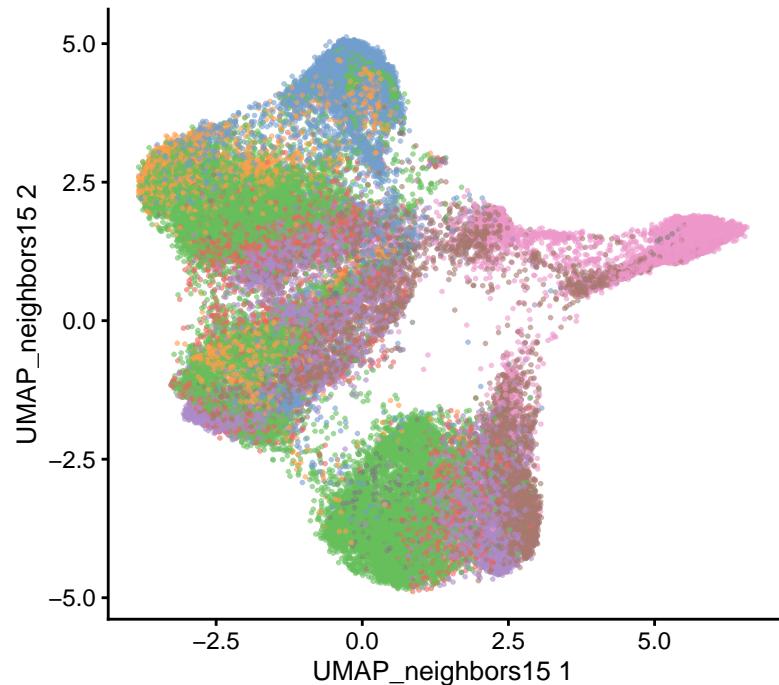
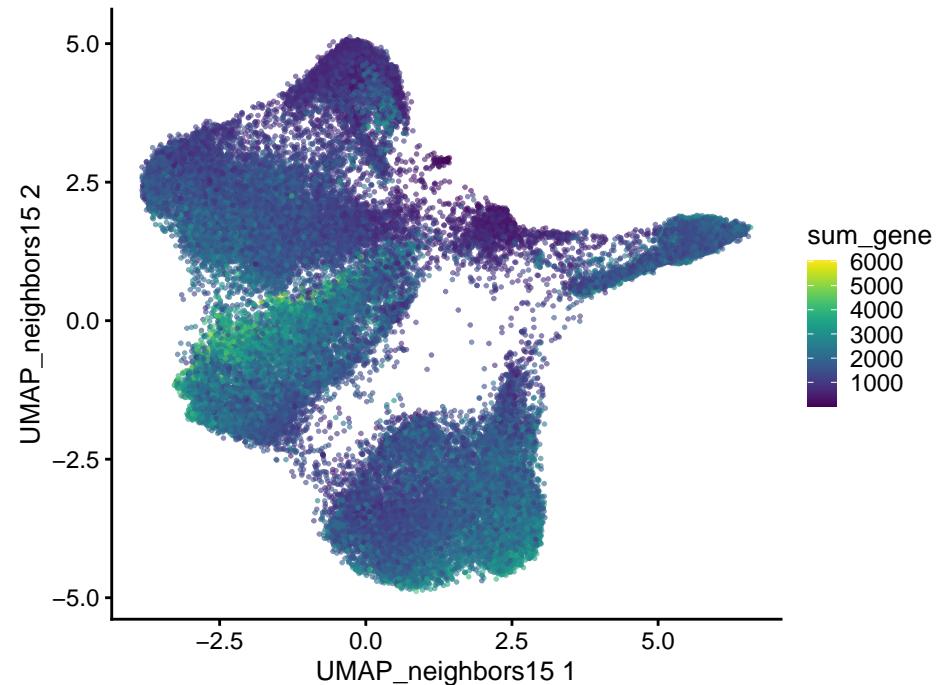
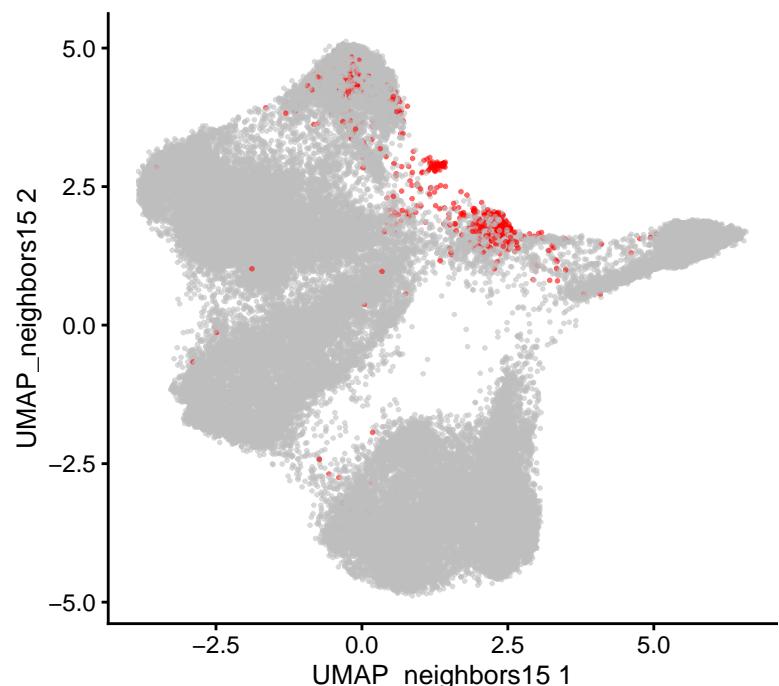


Manually annotated cluster assignment**Unique genes per cell****Cells excluded by <500 threshold****Cells excluded by detecting local outliers (SpotSweeper)**