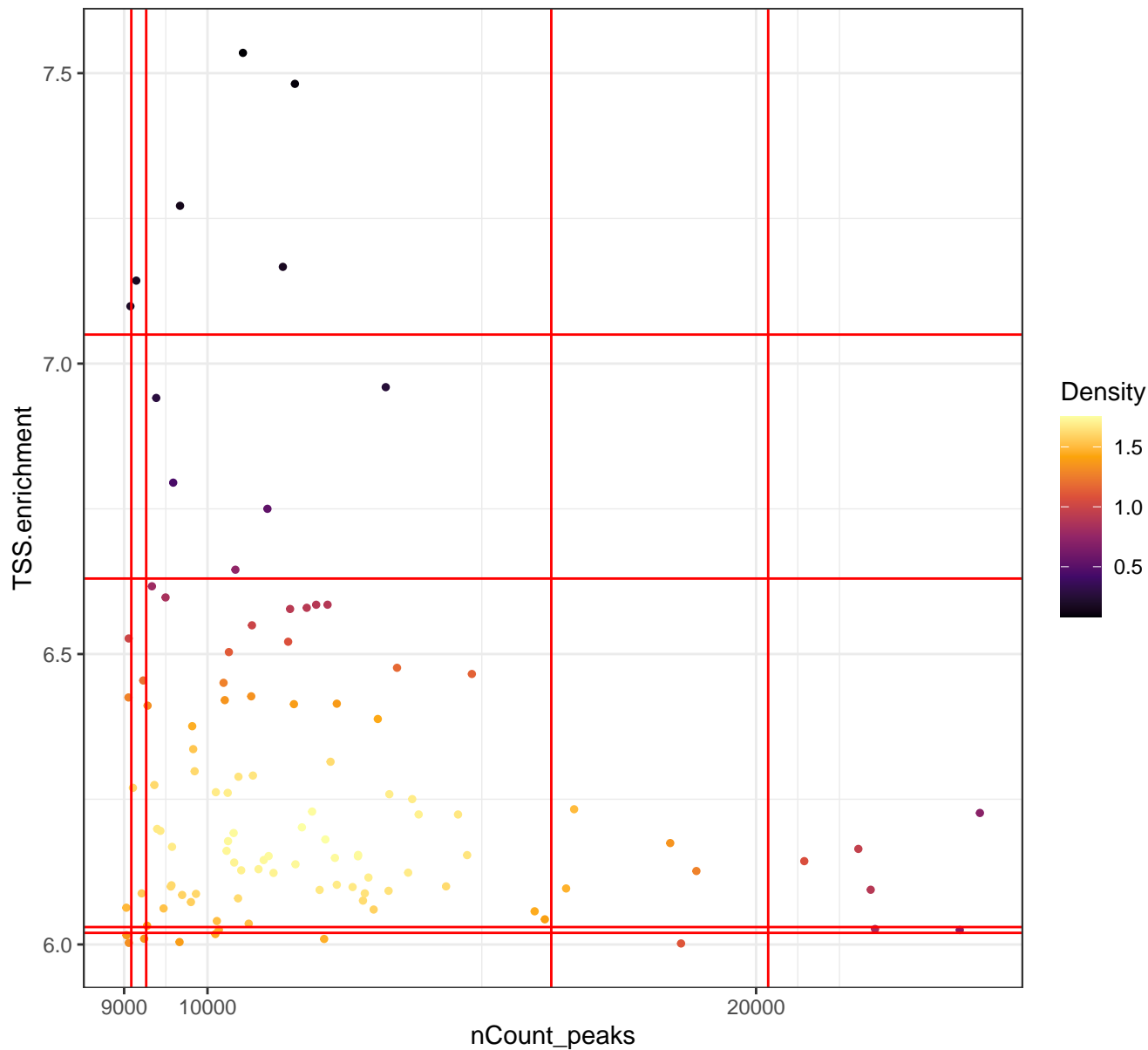


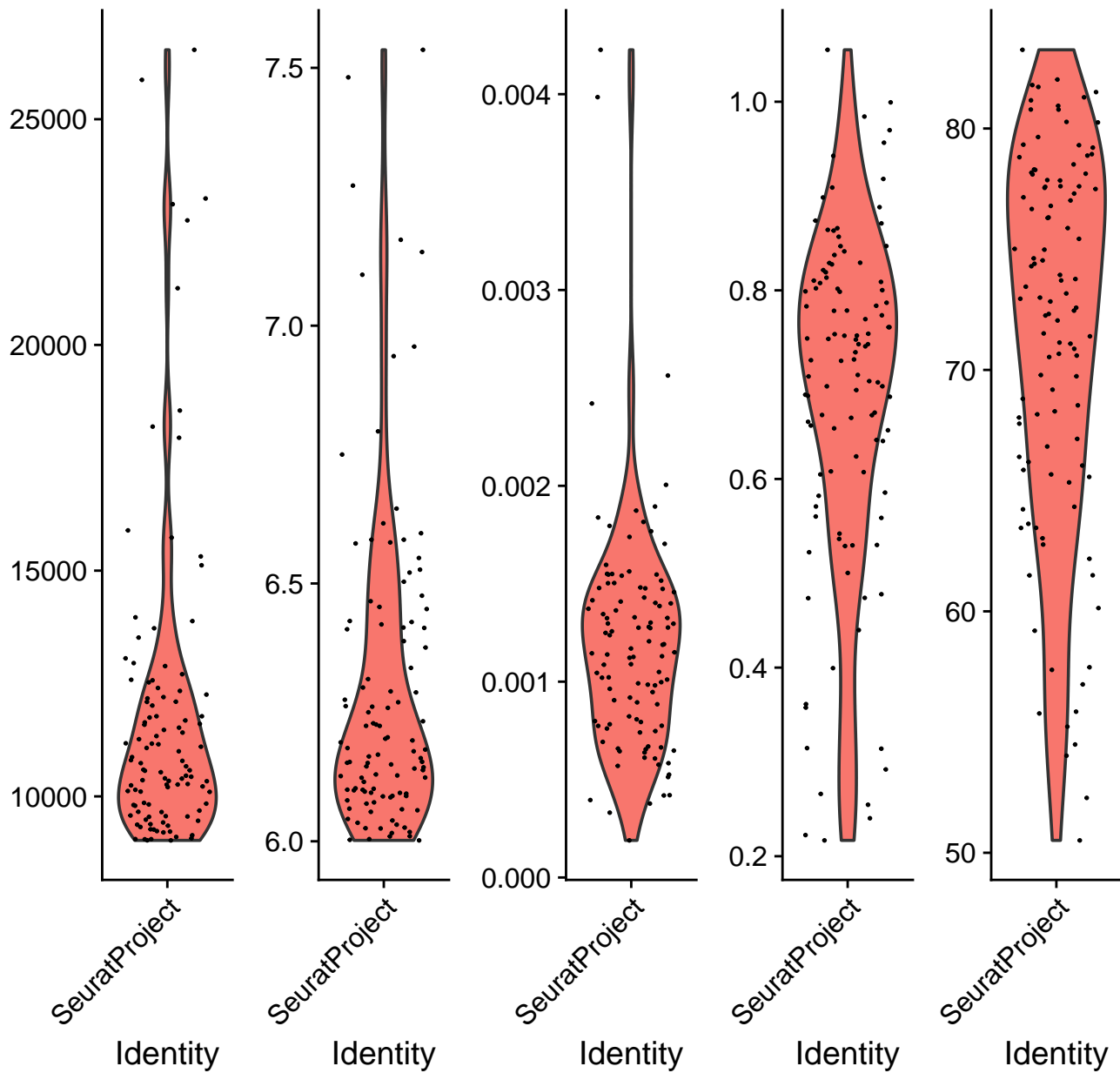
Quantiles

nCount_peaks: 5%:9082.15 10%:9254.8 90%:15440.1 95%:20307.85

TSS.enrichment: 5%:6.02 10%:6.03 90%:6.63 95%:7.05

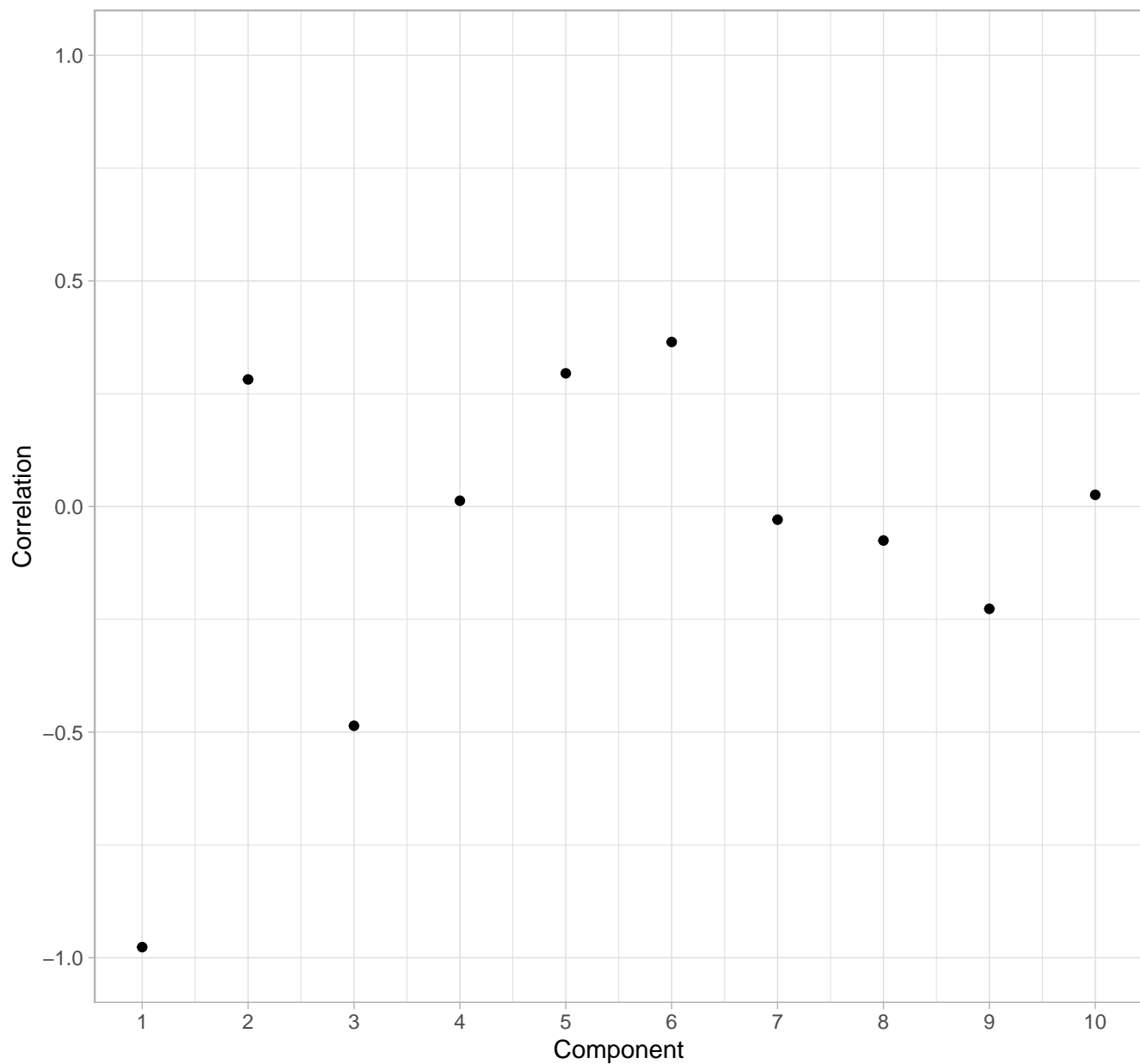


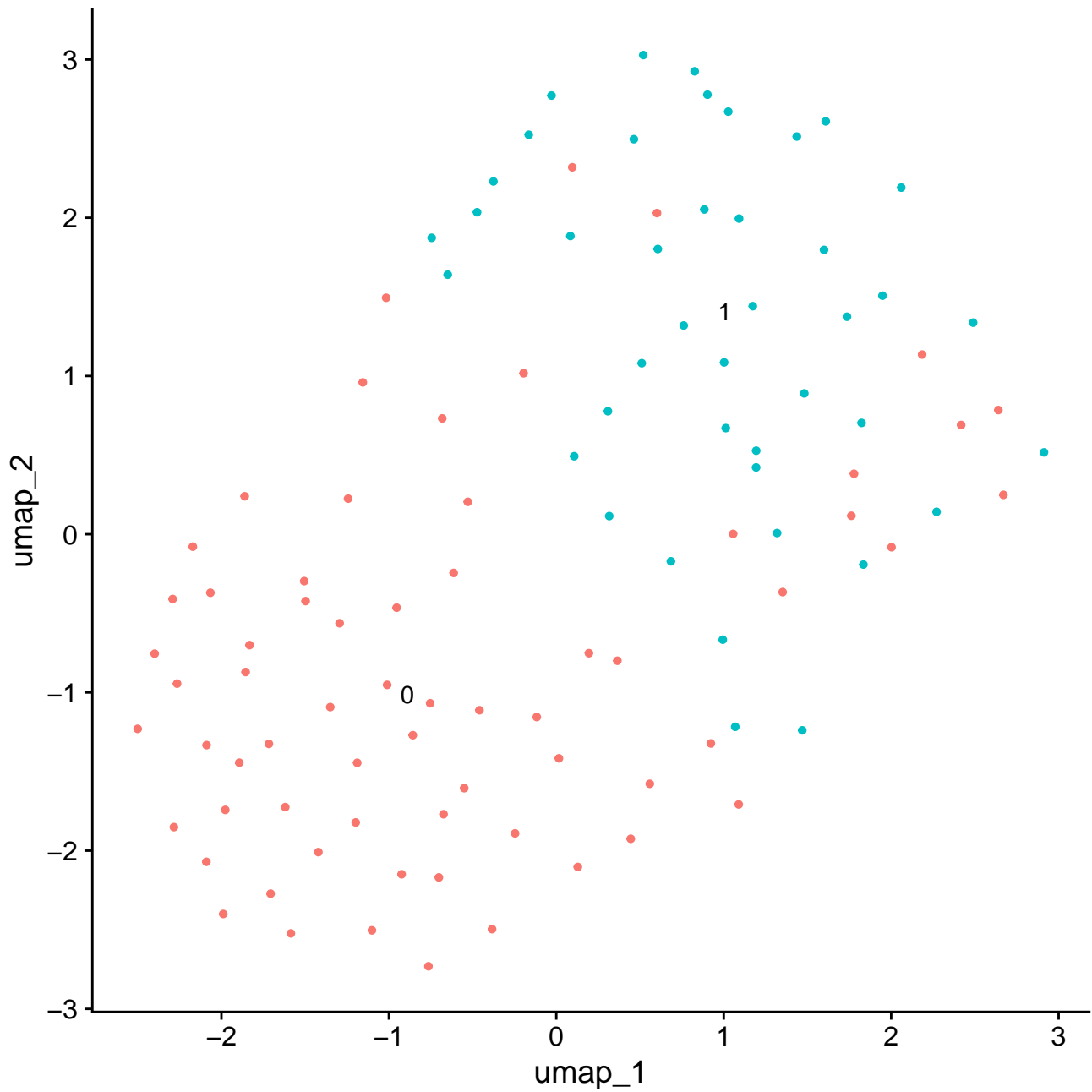
nCount_peaks TSS.enrichment blacklist_ratioosome_posreads_in_p



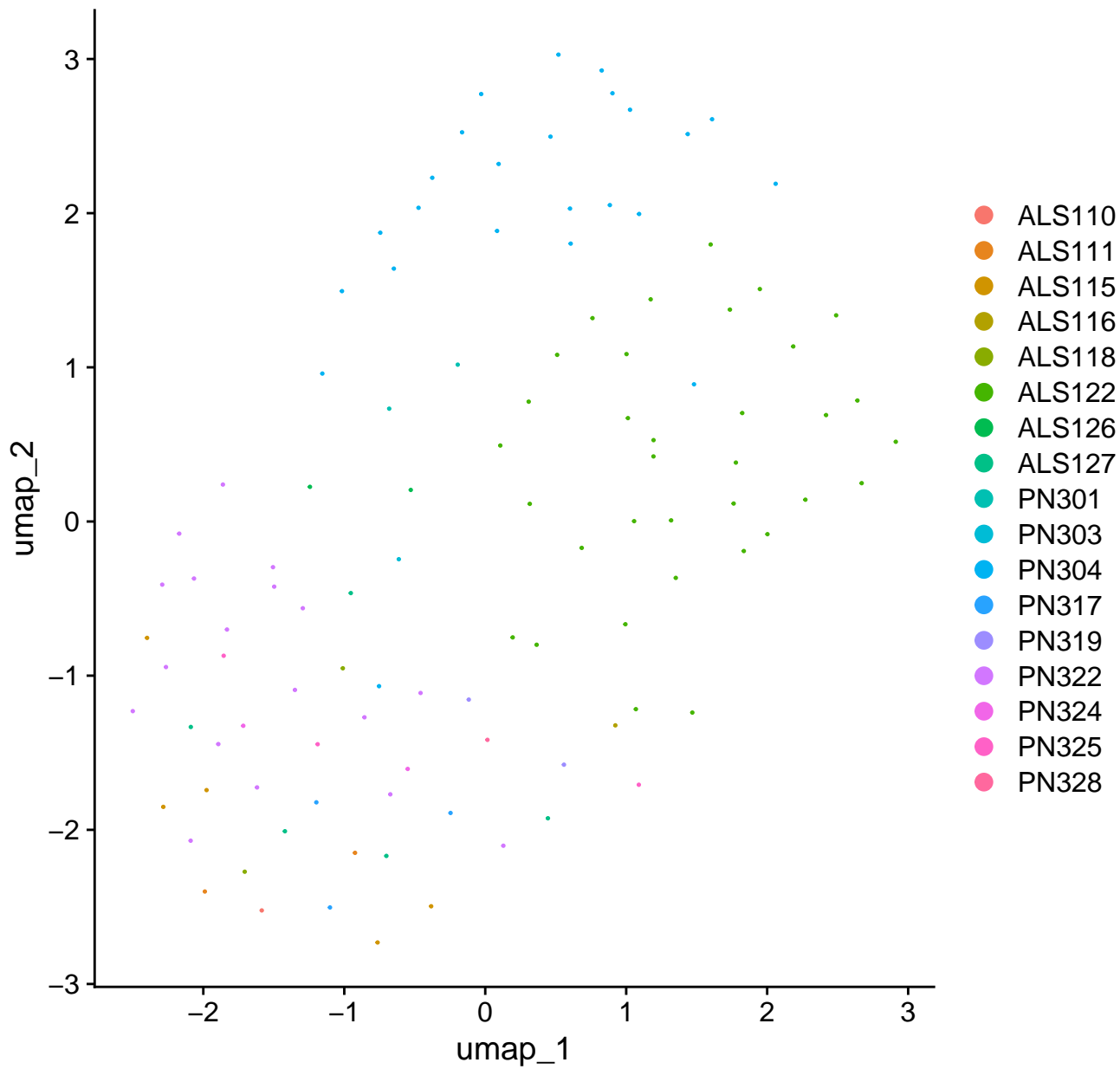
Correlation between depth and reduced dimension components

Assay: peaks Reduction: Isi

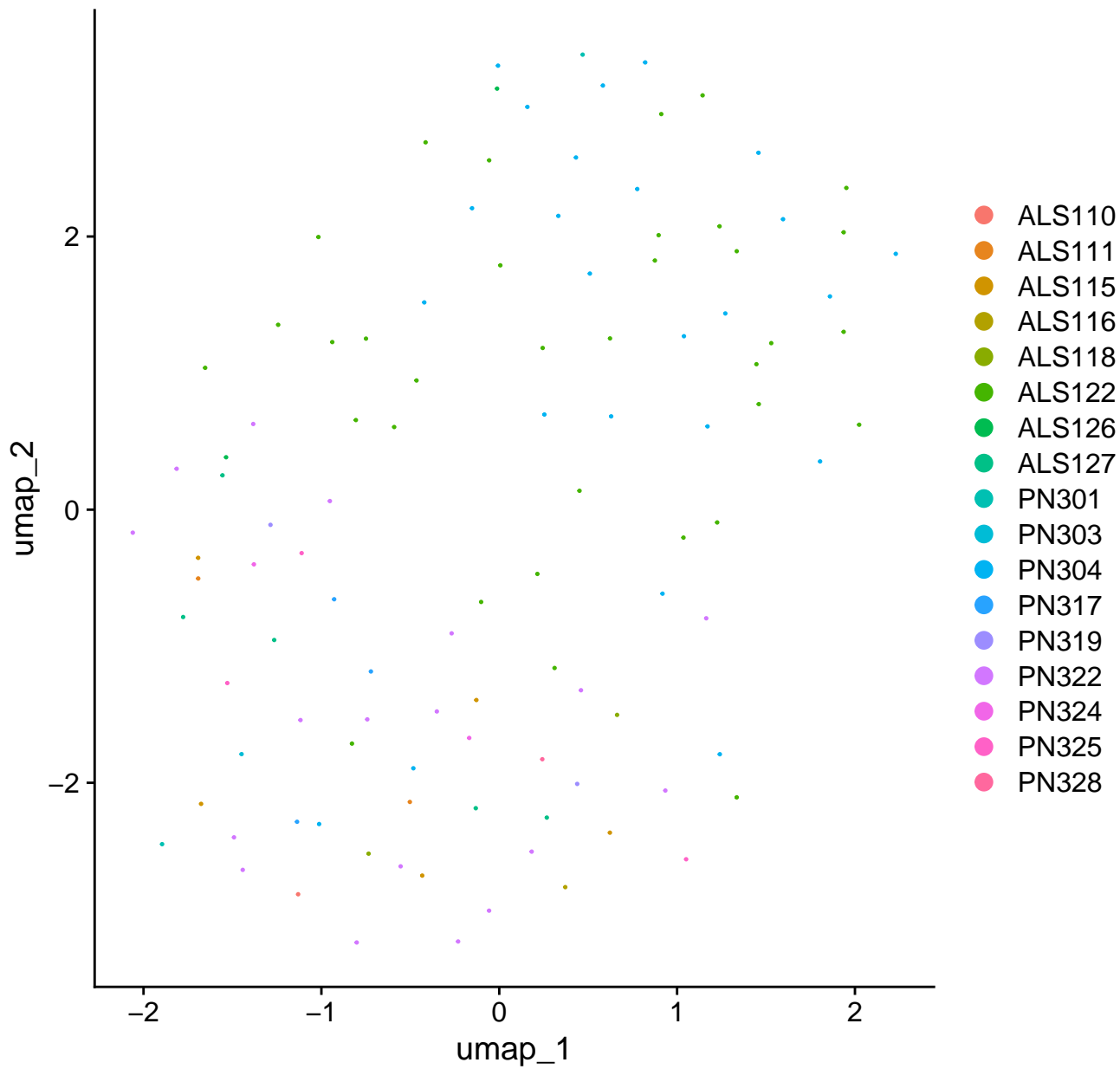


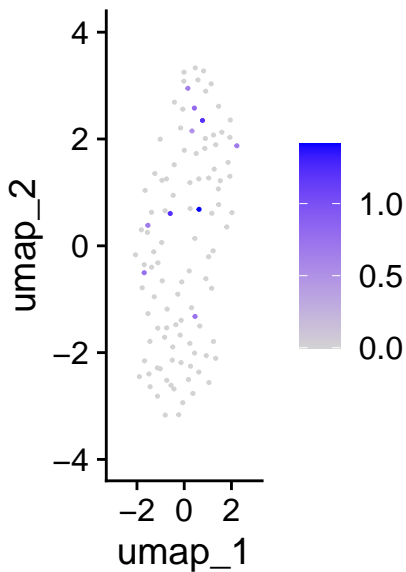
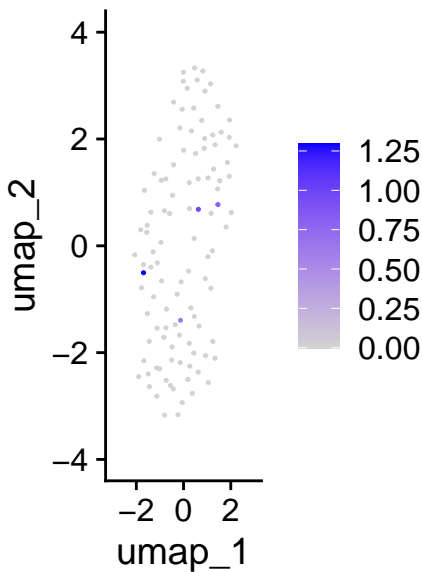
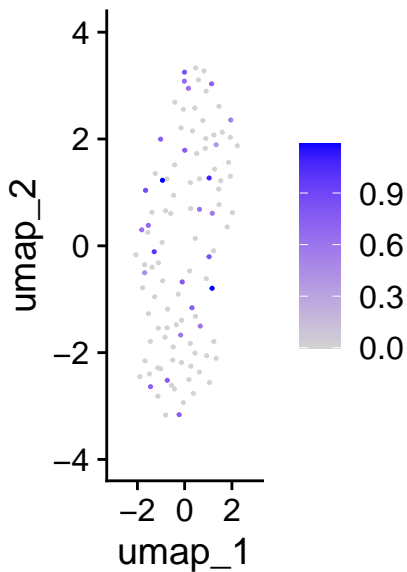
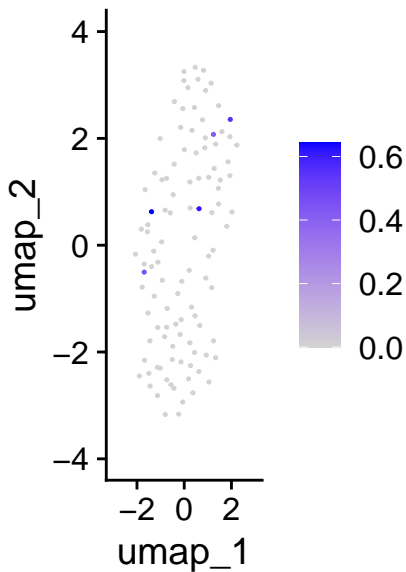
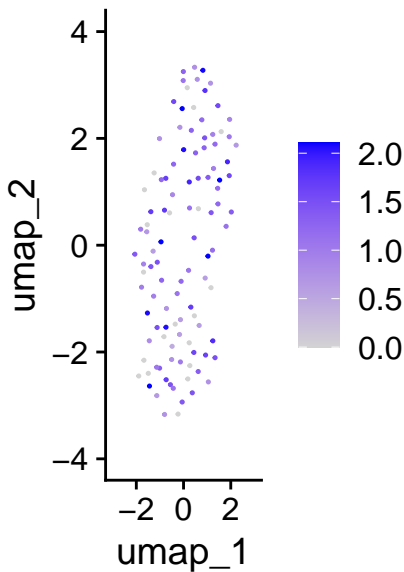


dataset

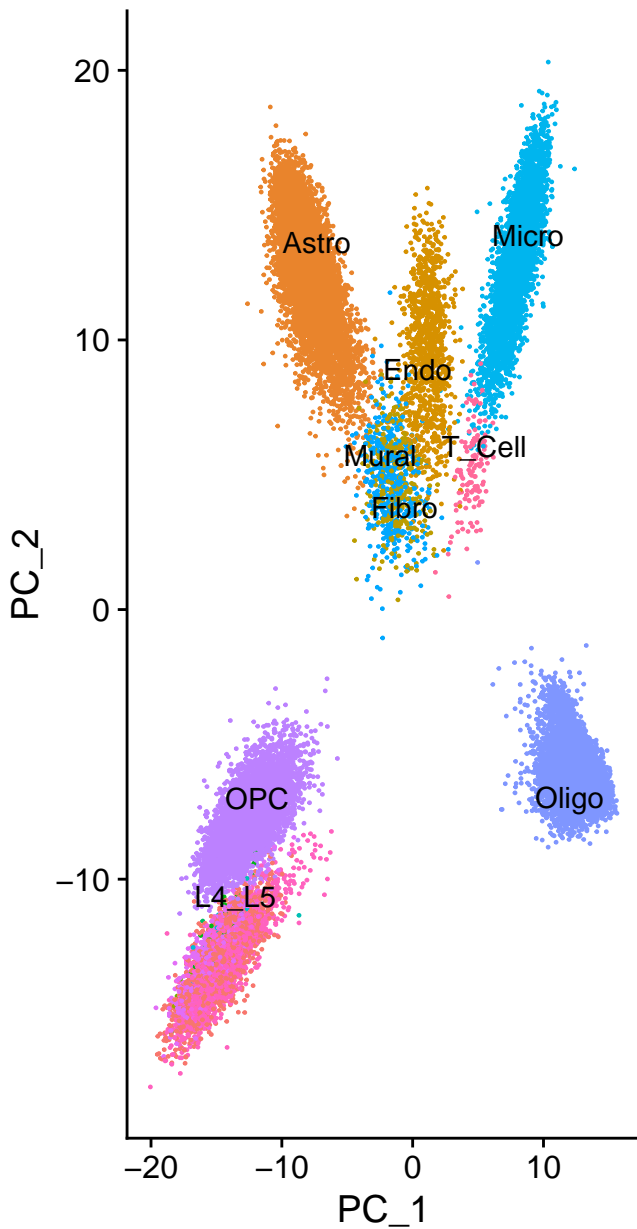


dataset

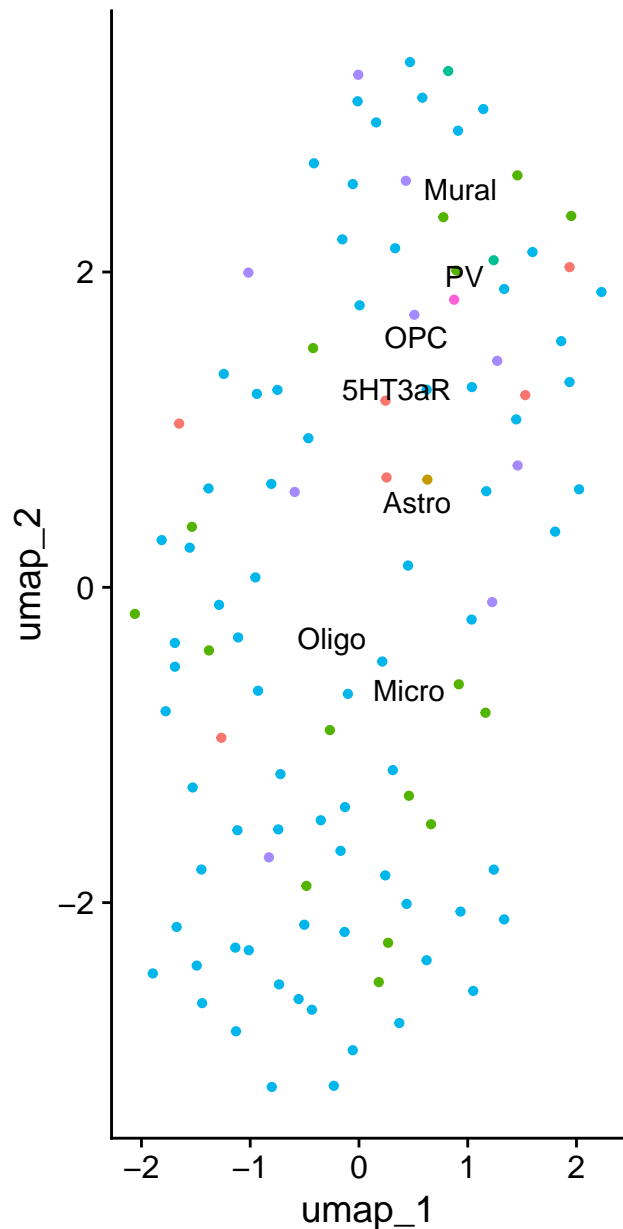


GFAP**AQP4****VEGFA****VEGFB****SHTN1**

scRNA-seq



scATAC-seq



prediction.score.max

