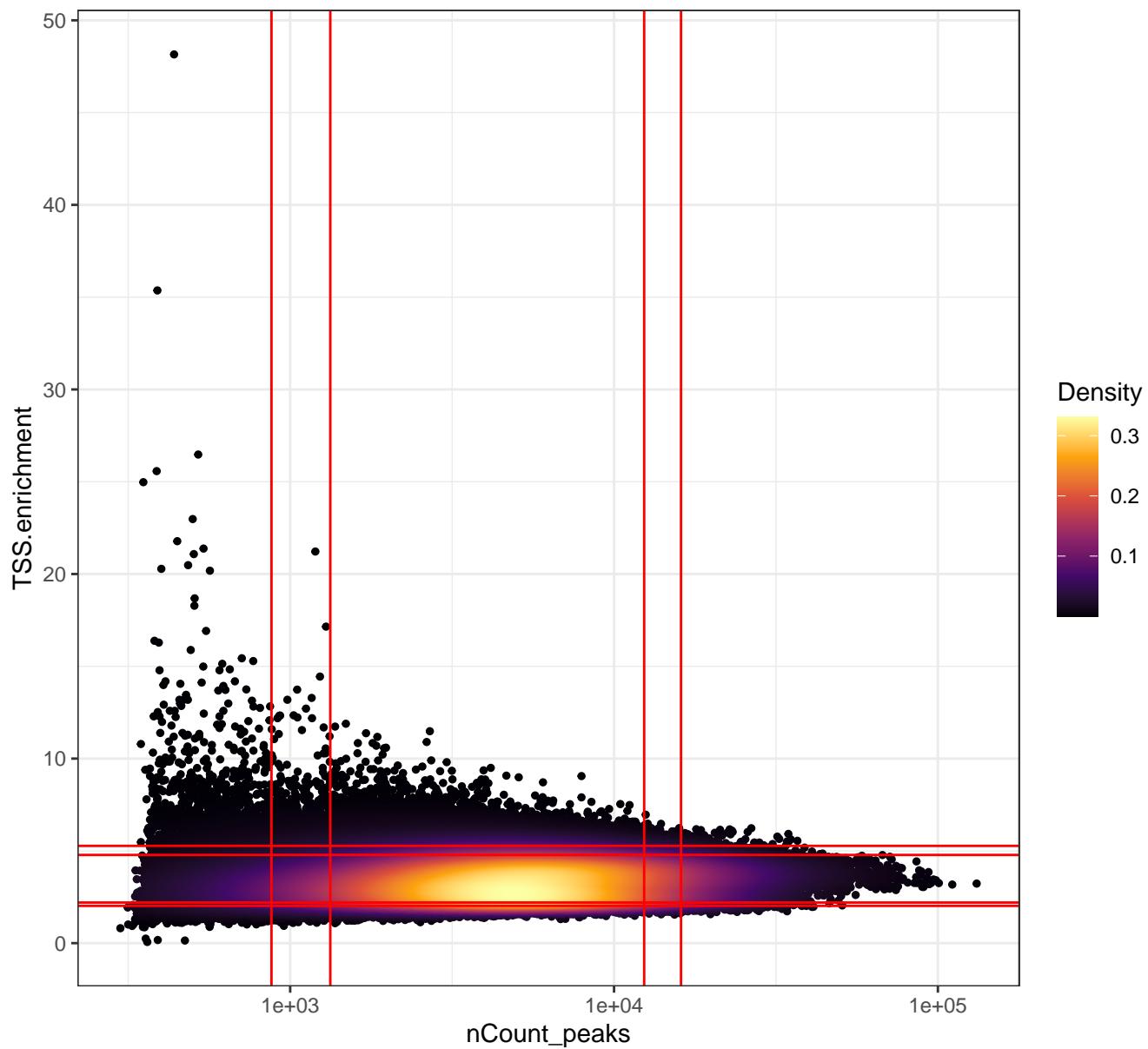
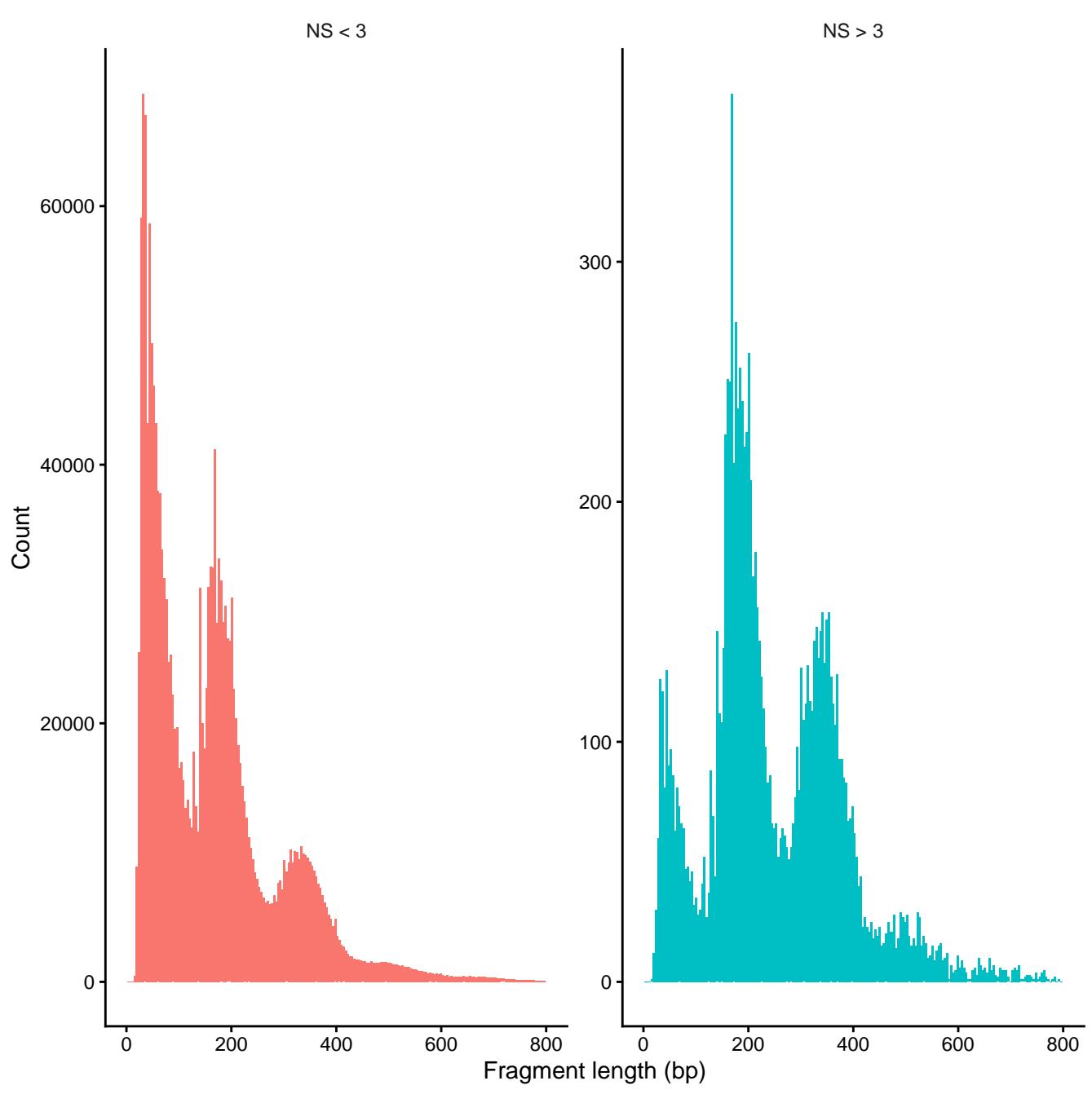


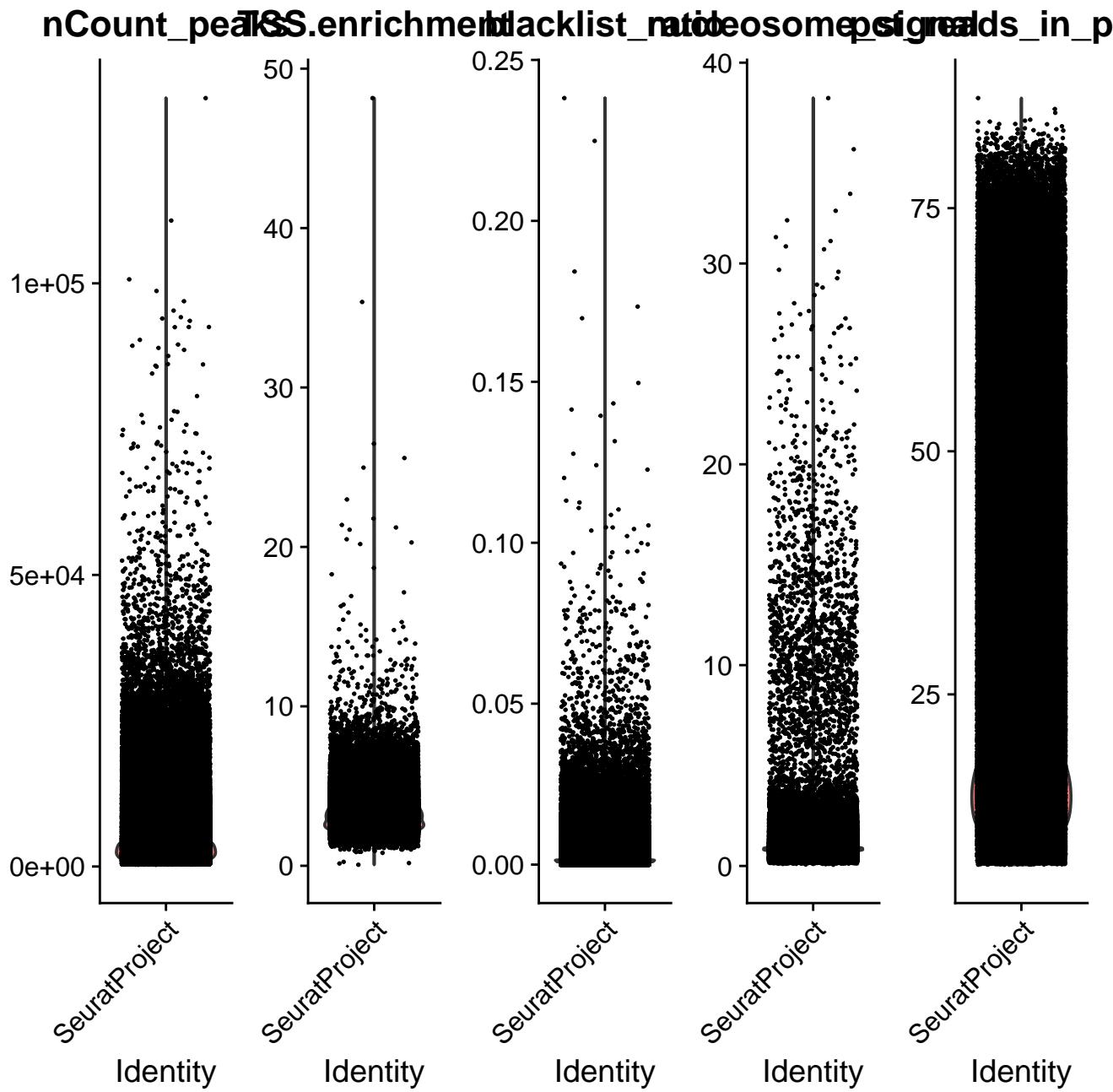
Quantiles

nCount_peaks: 5%:875.4 10%:1330 90%:12389.2 95%:16098.6

TSS.enrichment: 5%:2.02 10%:2.2 90%:4.78 95%:5.27

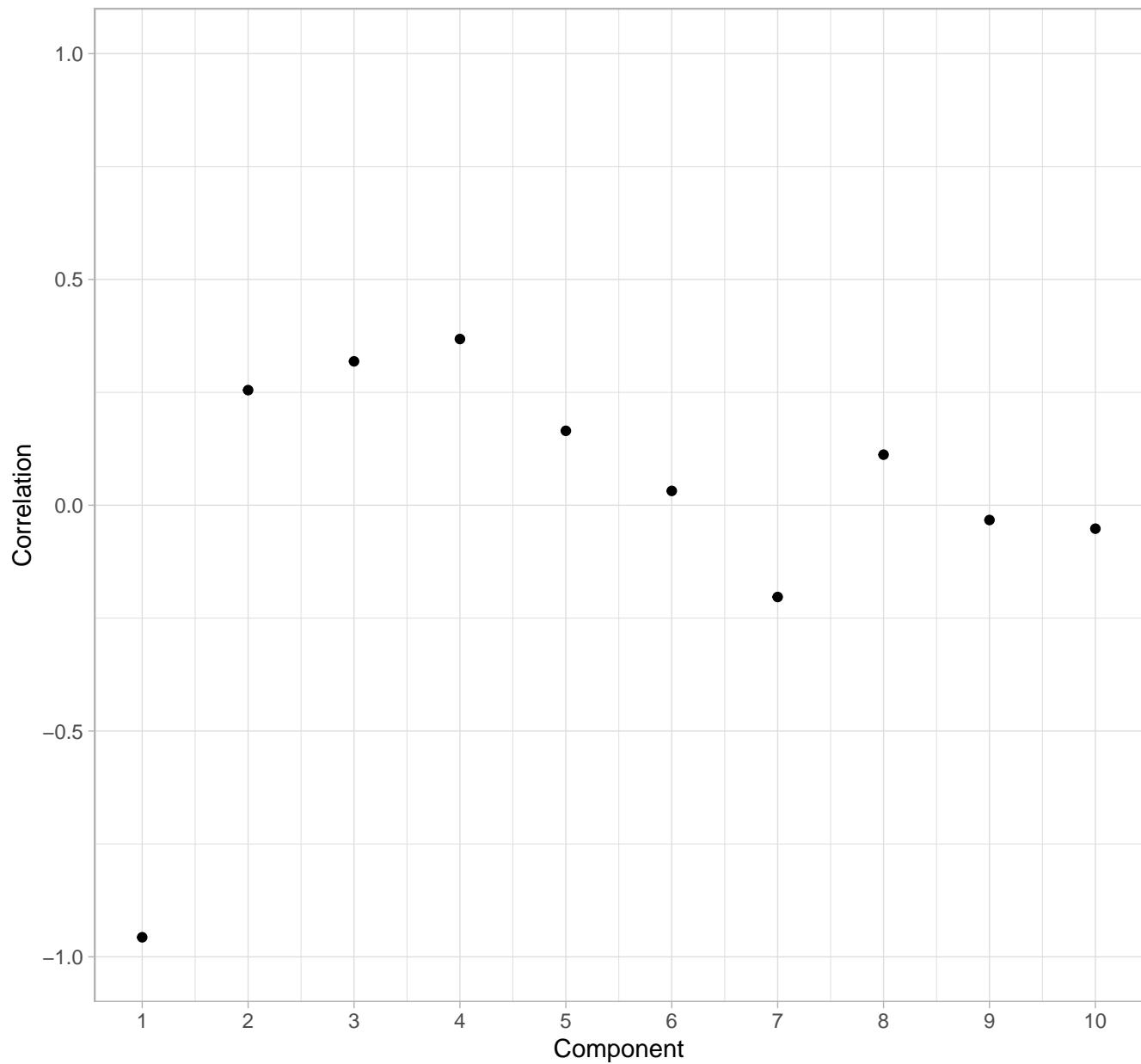


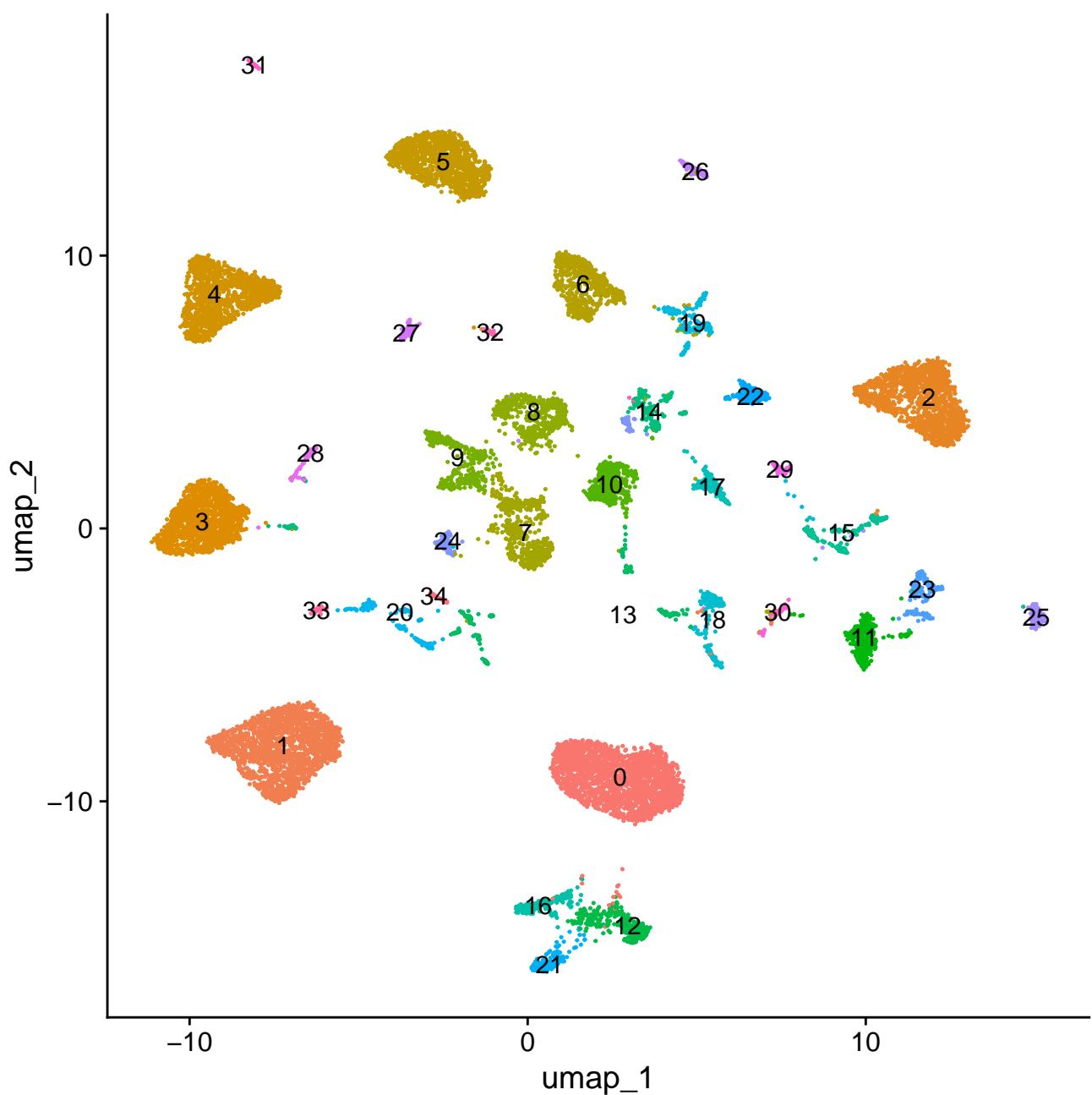




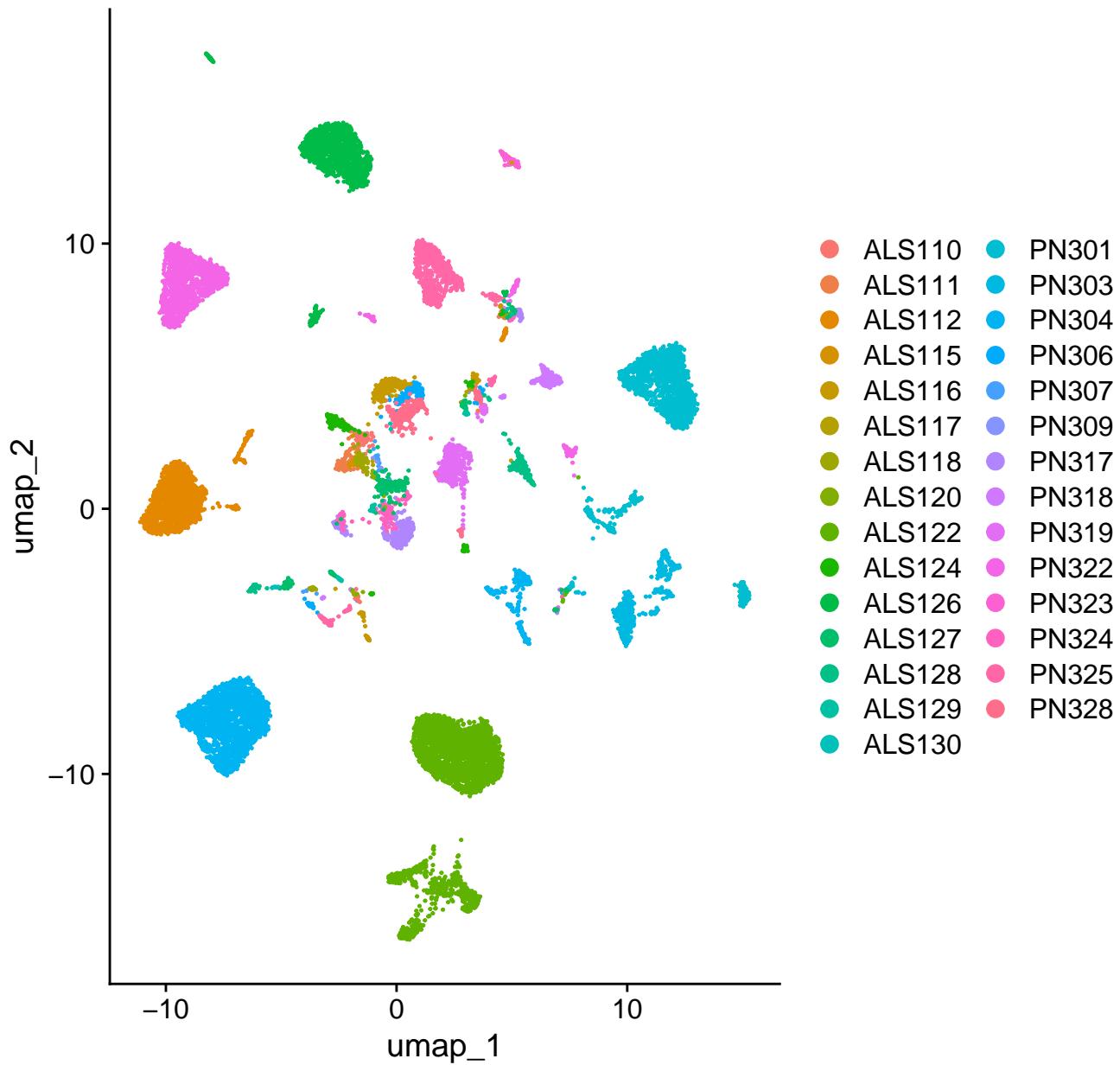
Correlation between depth and reduced dimension components

Assay: peaks Reduction: lsi

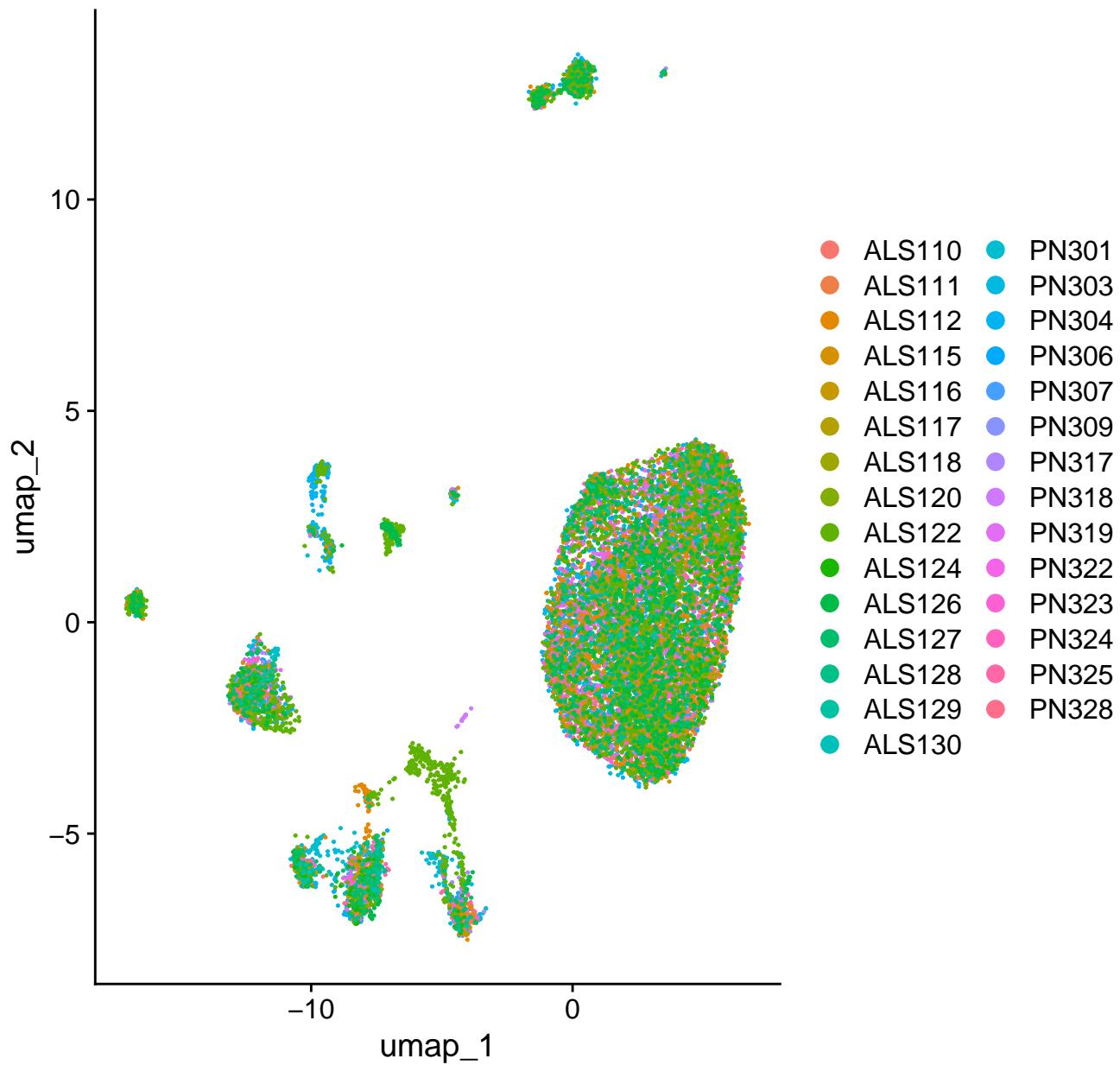


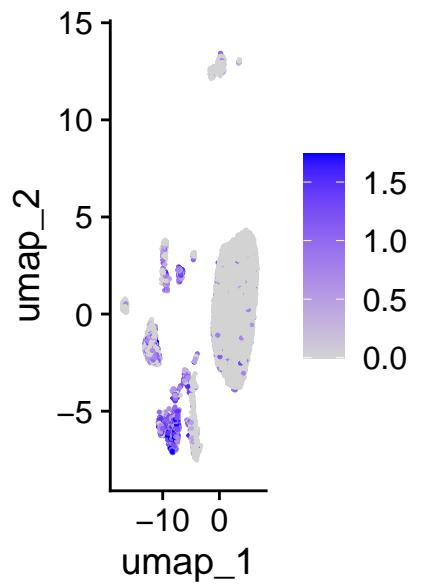
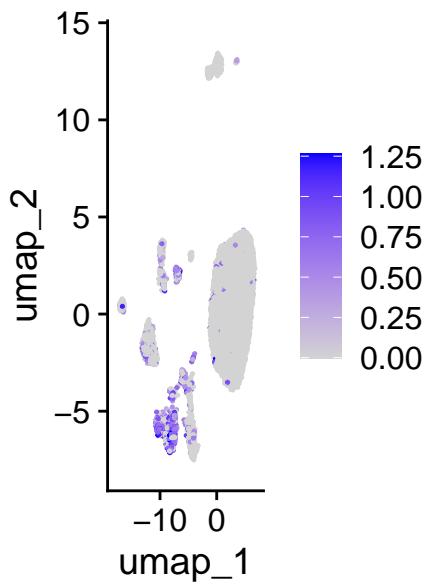
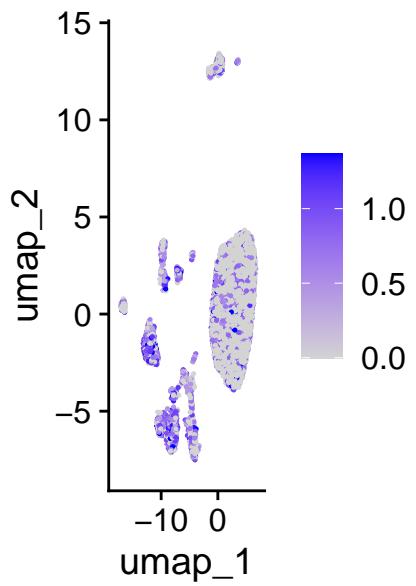
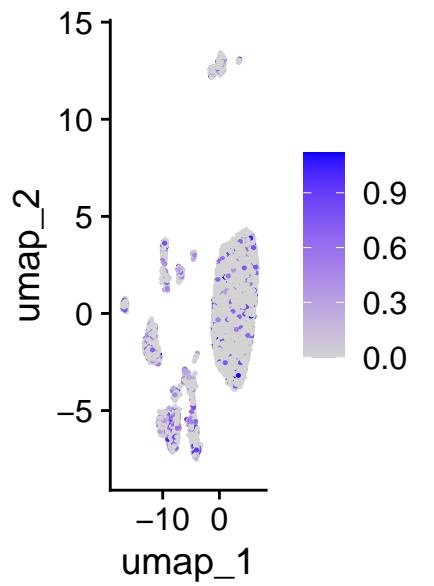
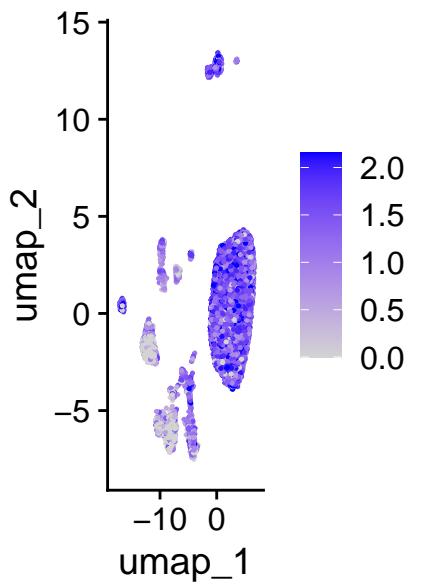


dataset

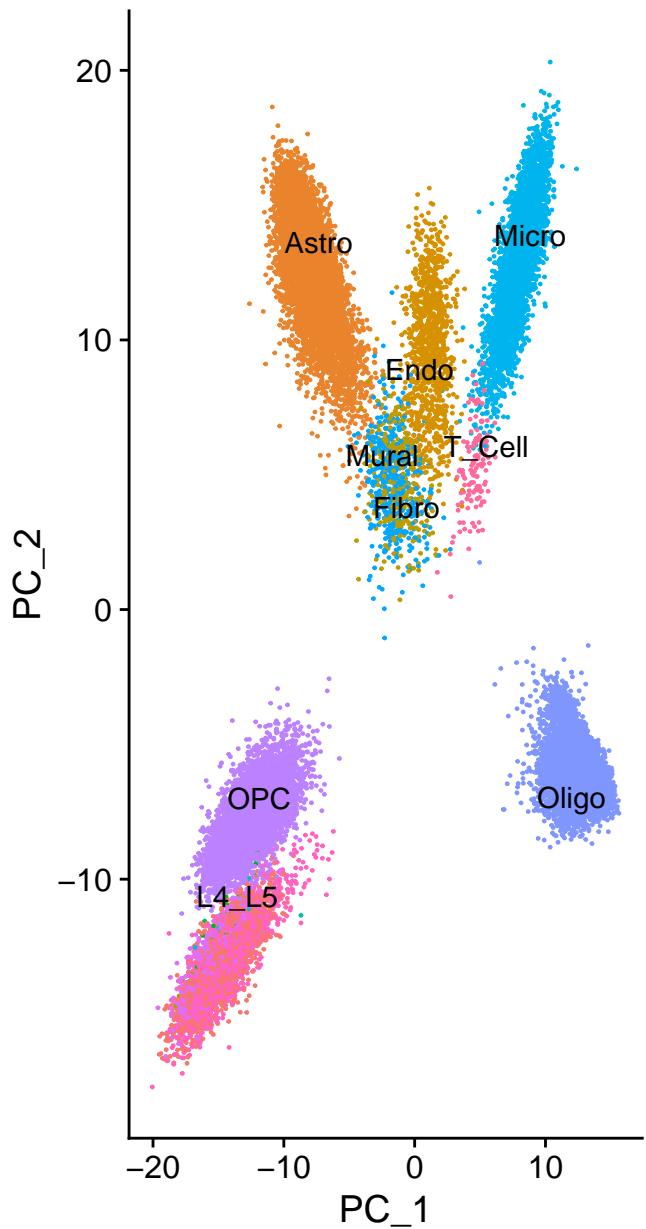


dataset

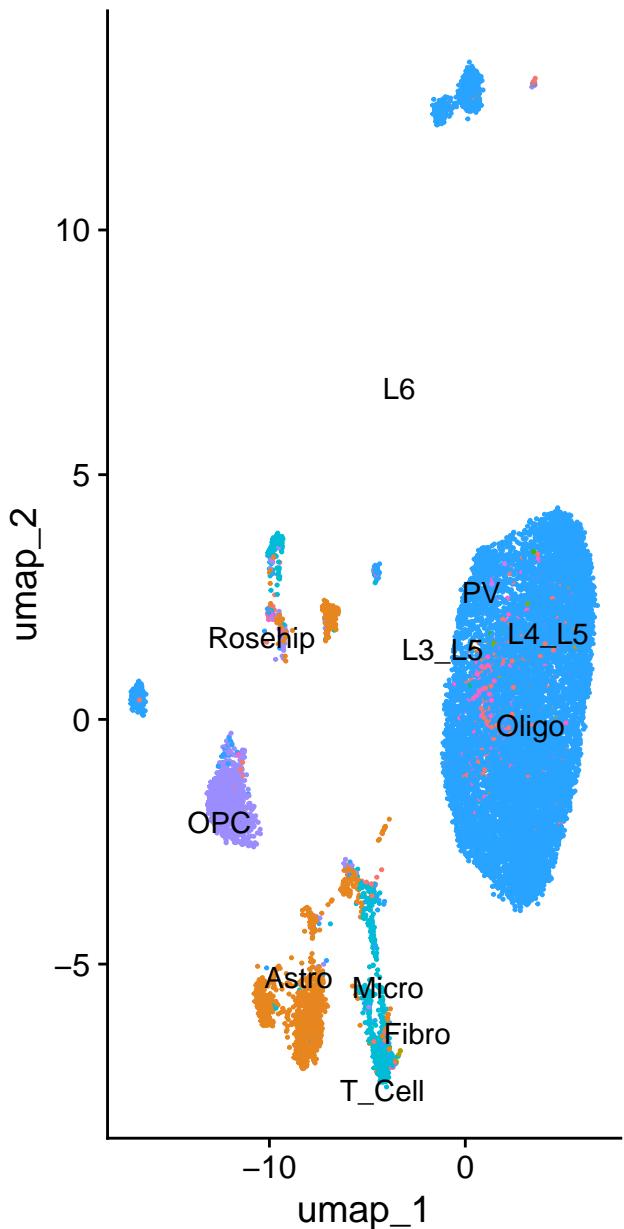


GFAP**AQP4****VEGFA****VEGFB****SHTN1**

scRNA-seq



scATAC-seq



prediction.score.max

