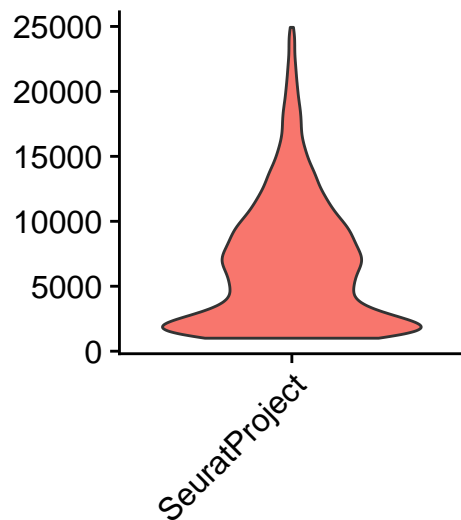
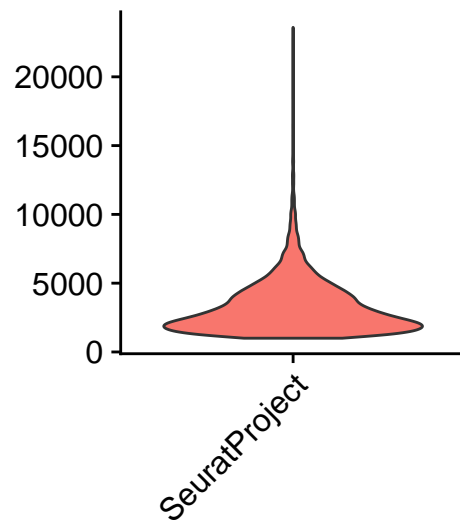
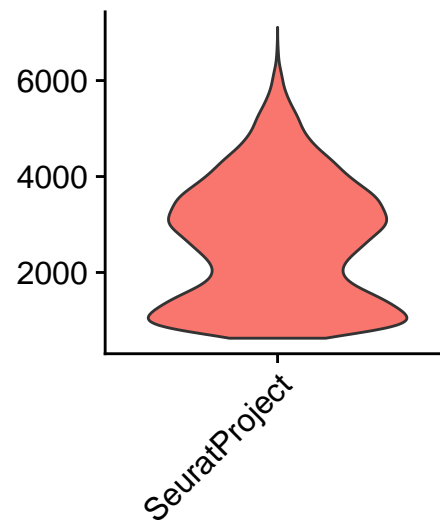


**nCount\_RNA**

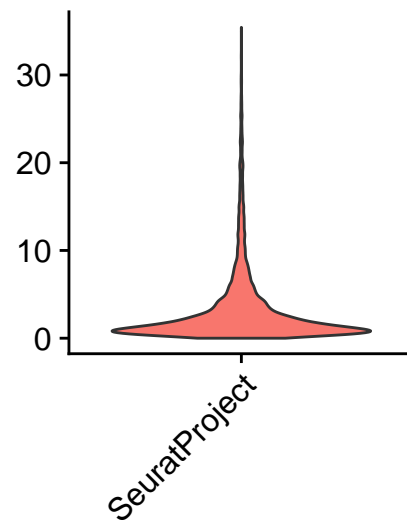
Identity

**nCount\_ATAC**

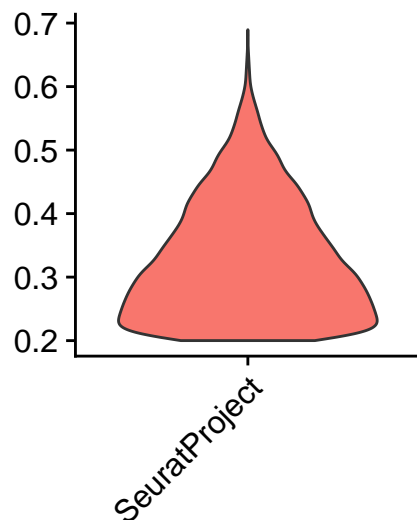
Identity

**nFeature\_RNA**

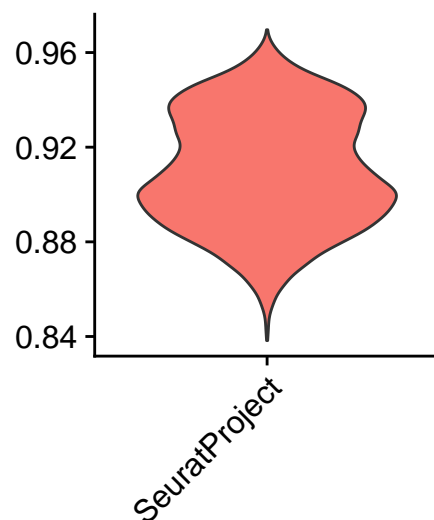
Identity

**mitoPct**

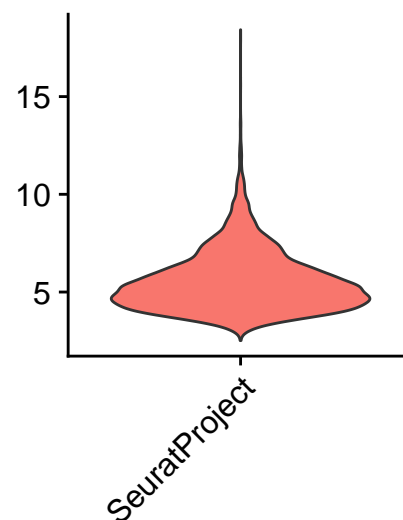
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**FRiP**

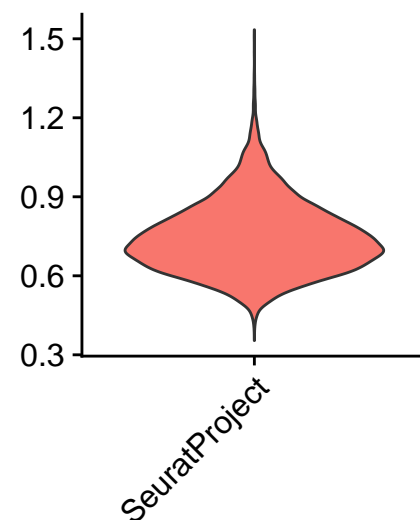
Identity

**log10GenesPerUMI**

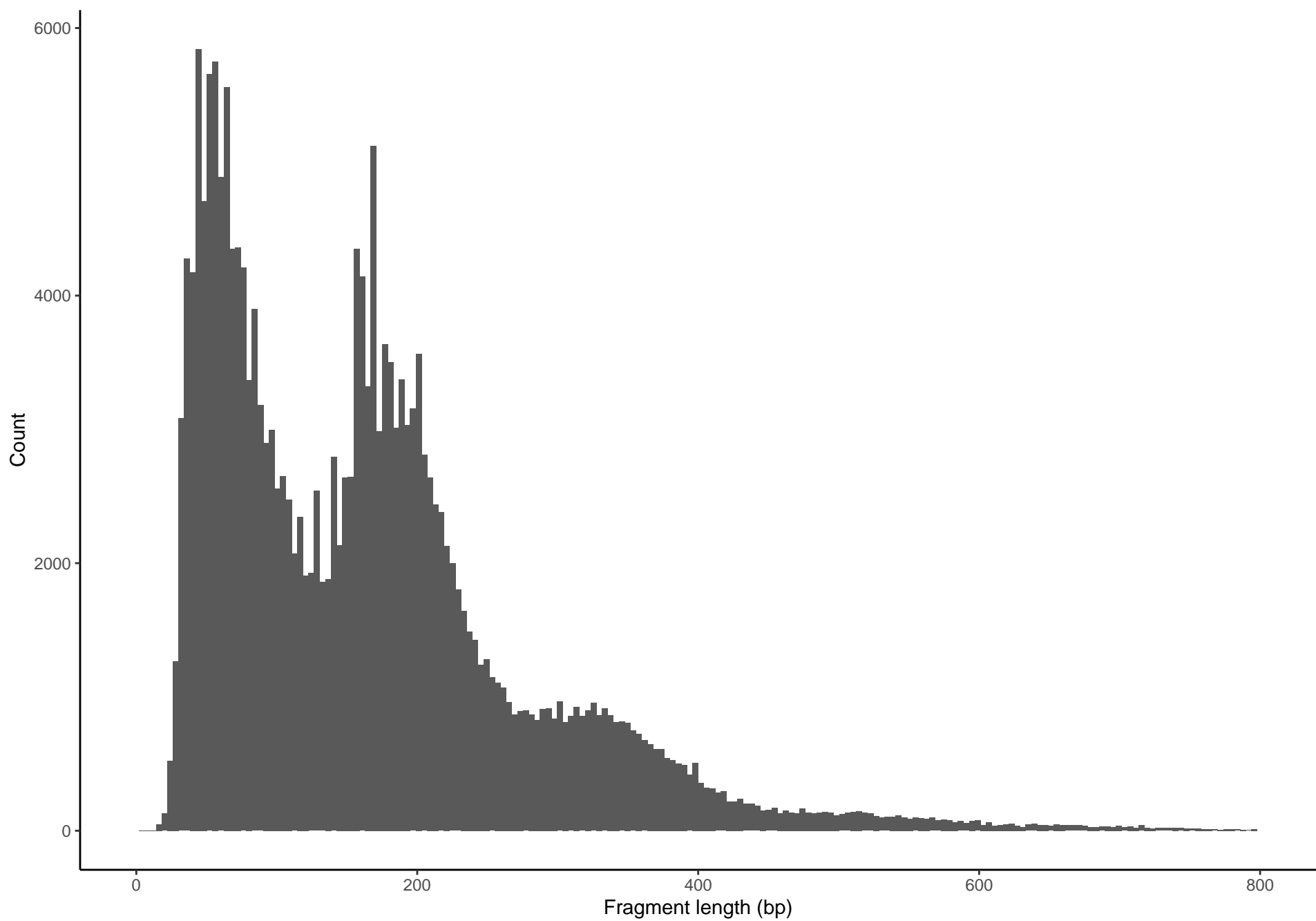
Identity

**TSS.enrichment**

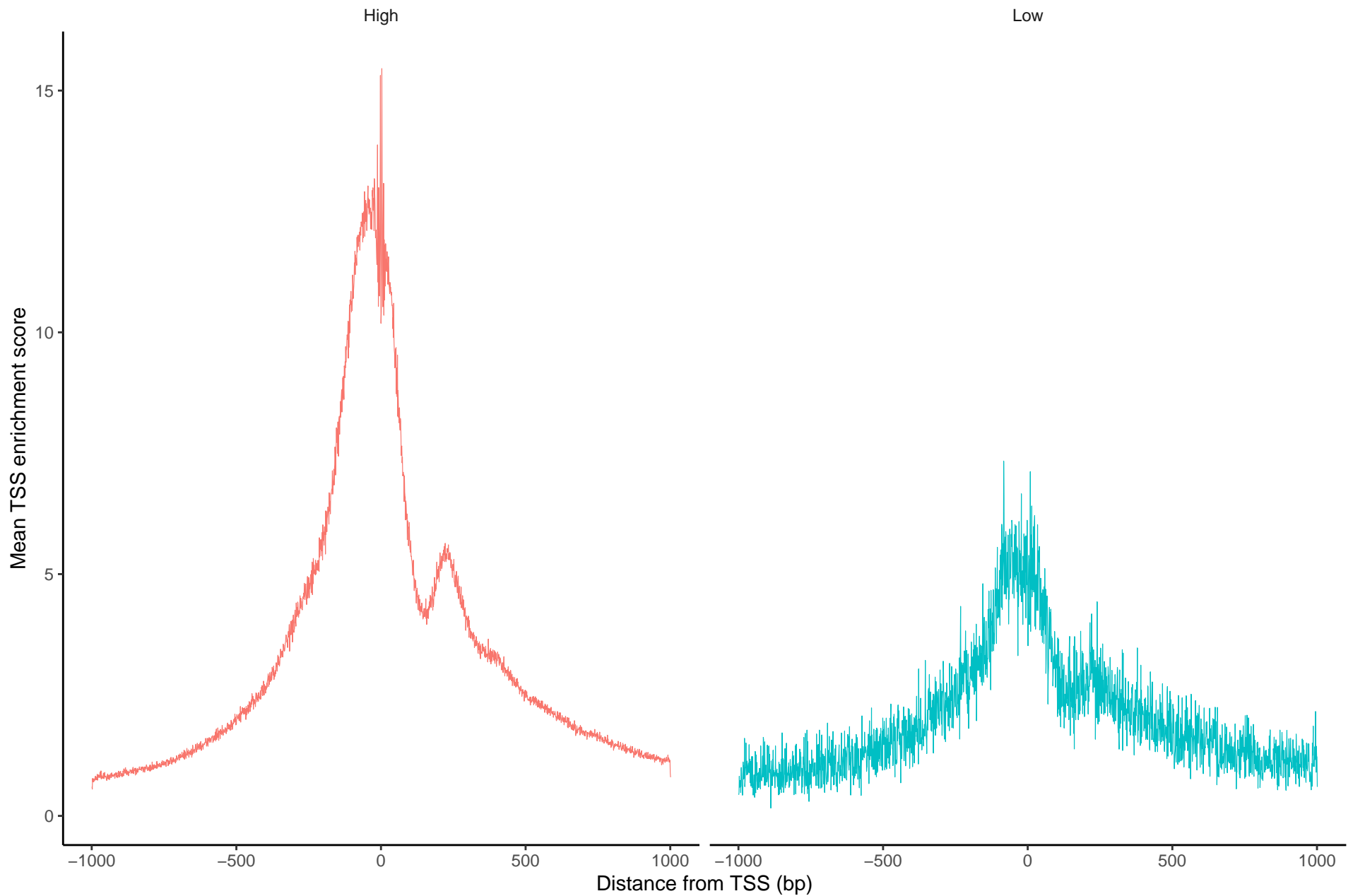
Identity

**nucleosome\_signal**

Identity



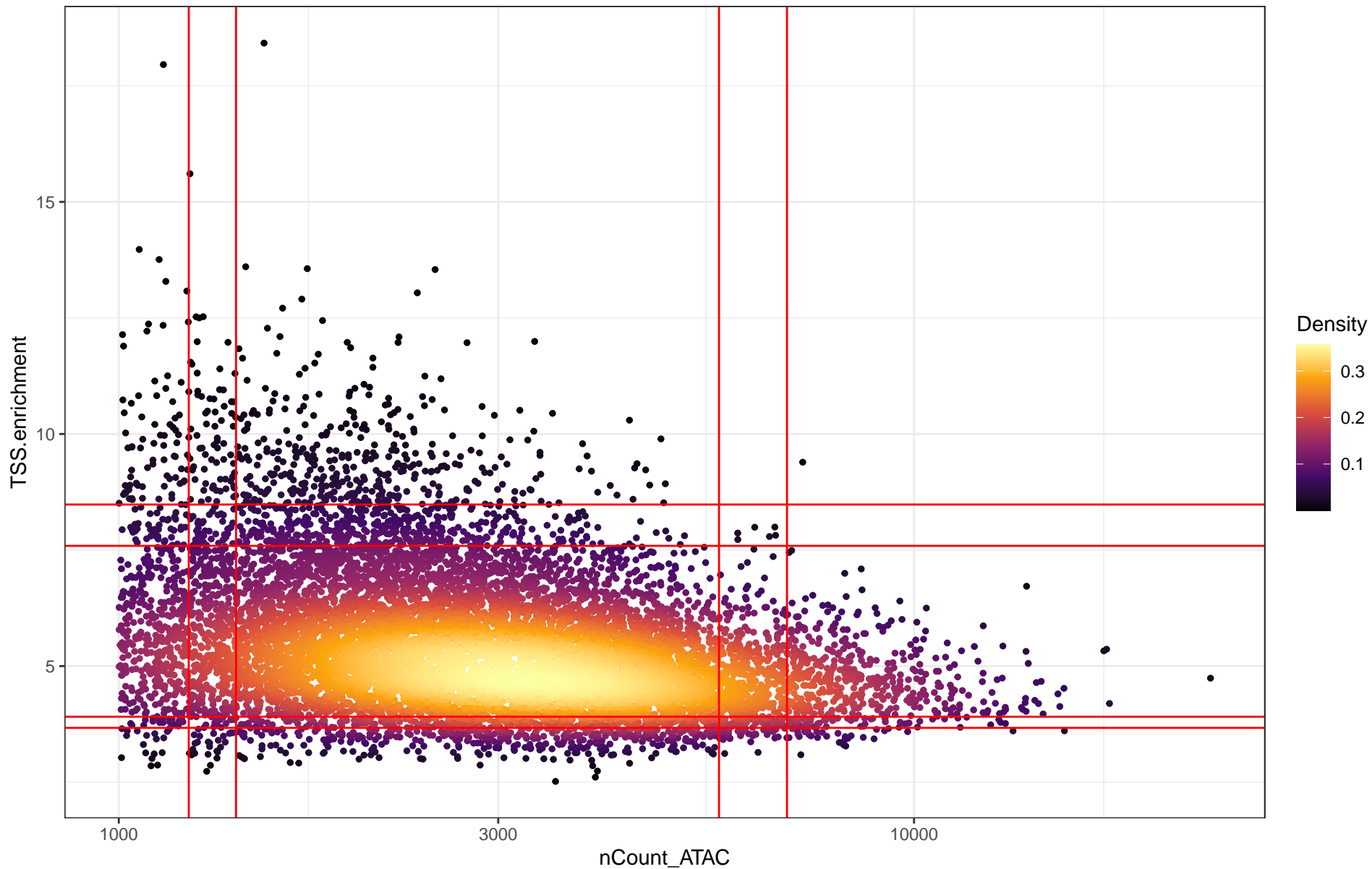
# TSS enrichment



# Quantiles

nCount\_ATAC: 5%:1224.65 10%:1404.3 90%:5685.4 95%:6922.4

TSS.enrichment: 5%:3.67 10%:3.91 90%:7.59 95%:8.48



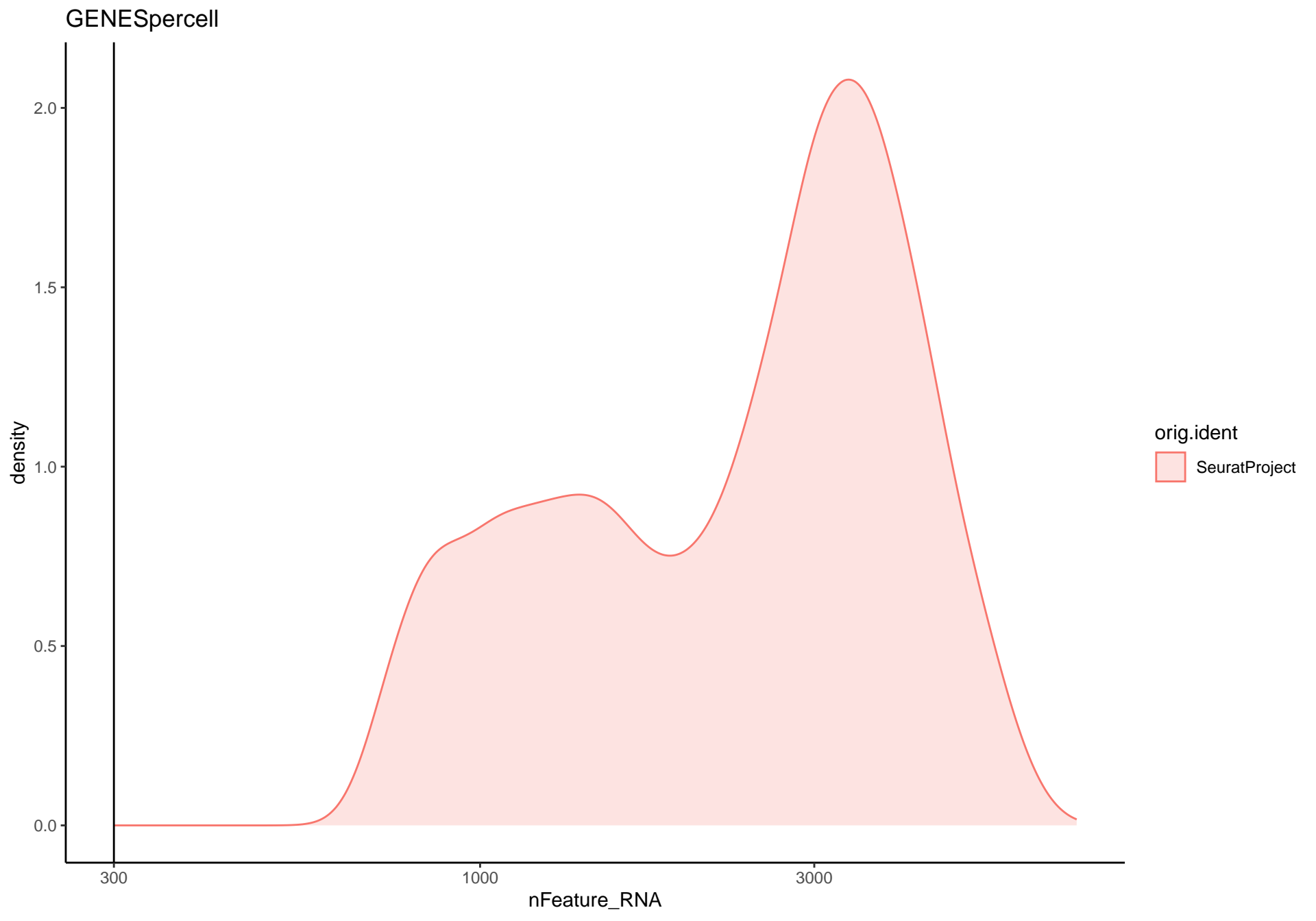
GENESpercell

density

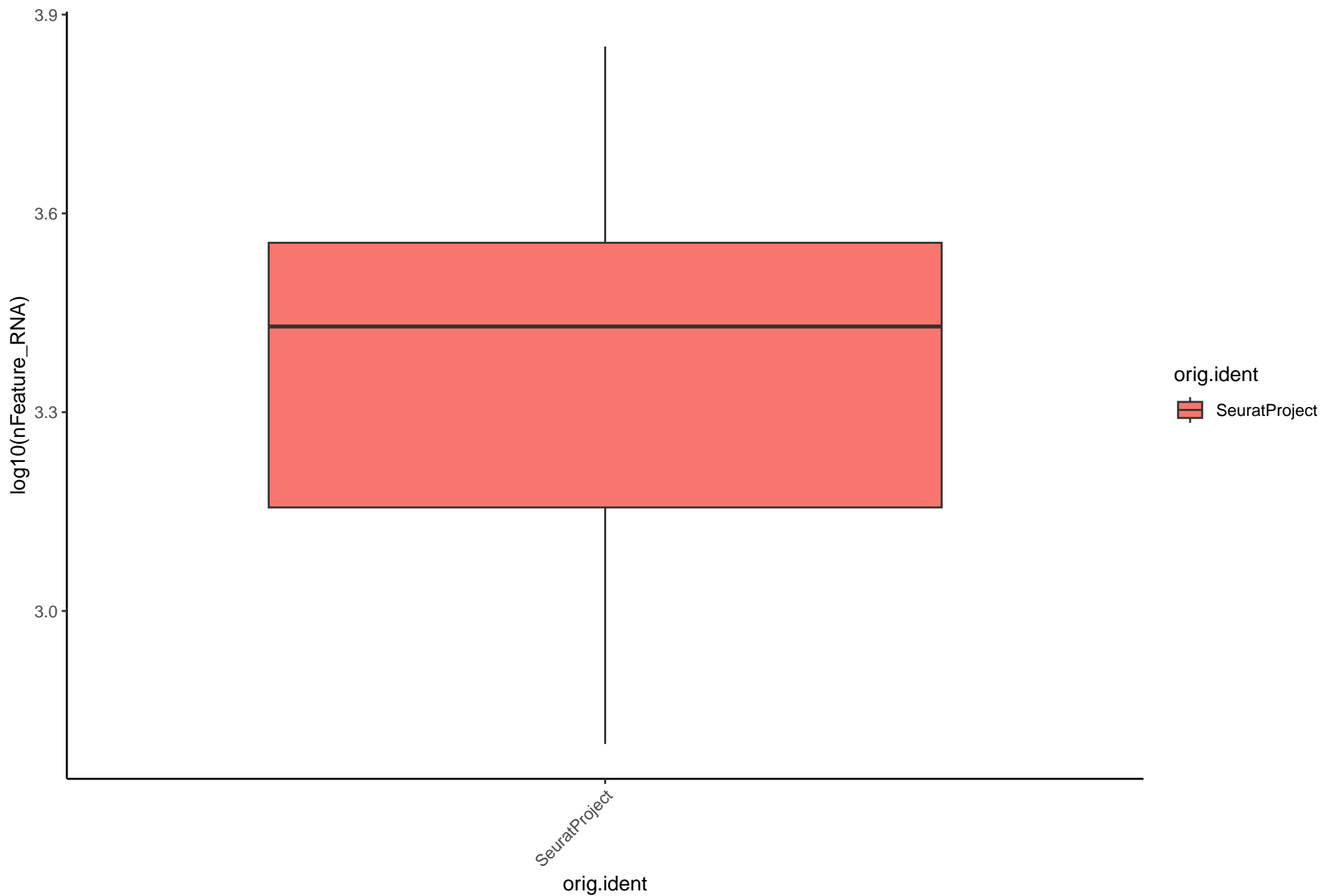
orig.ident

SeuratProject

nFeature\_RNA



# NCells vs NGenes



UMIvsGENESpercell Ncell:: 10534

