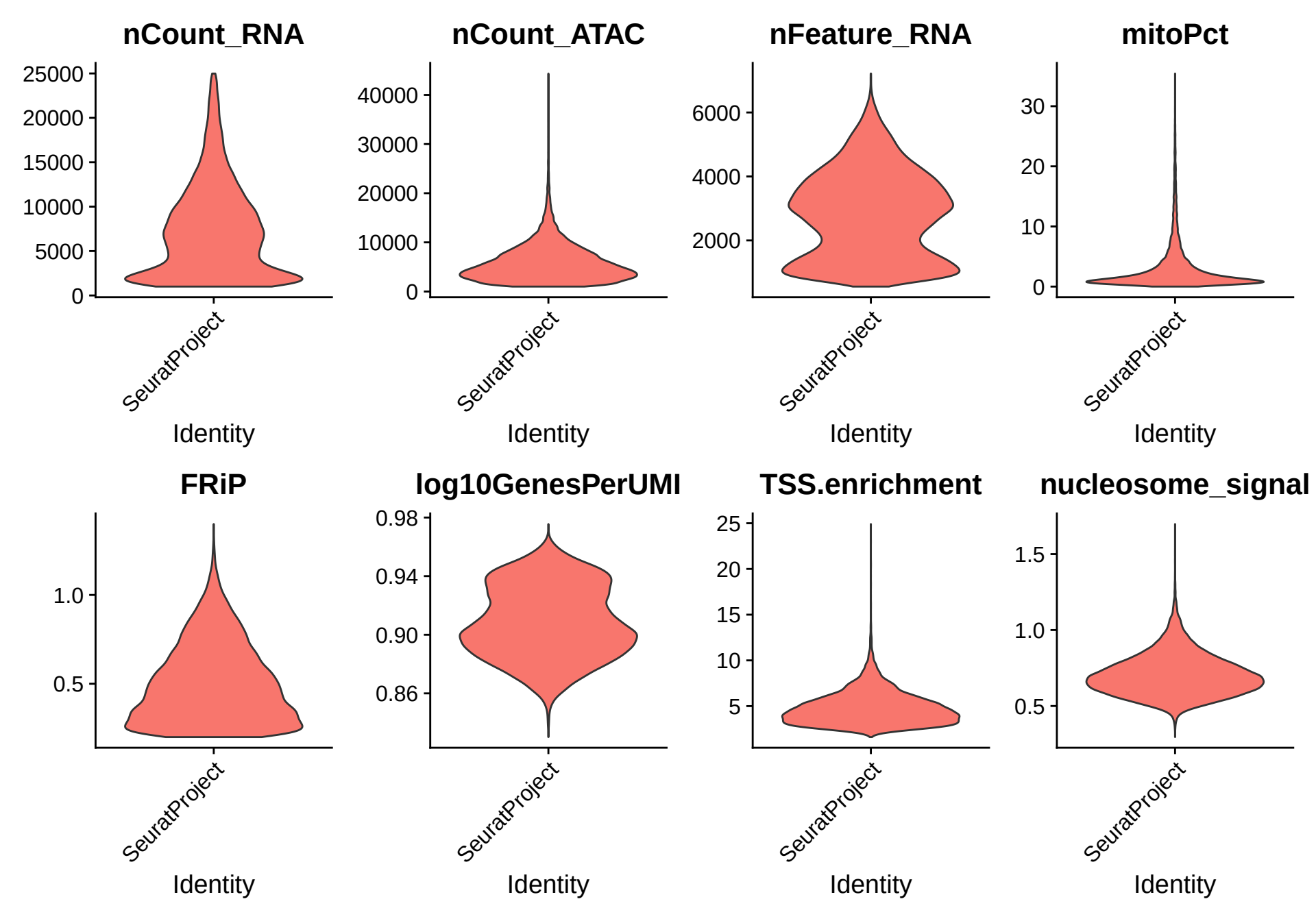
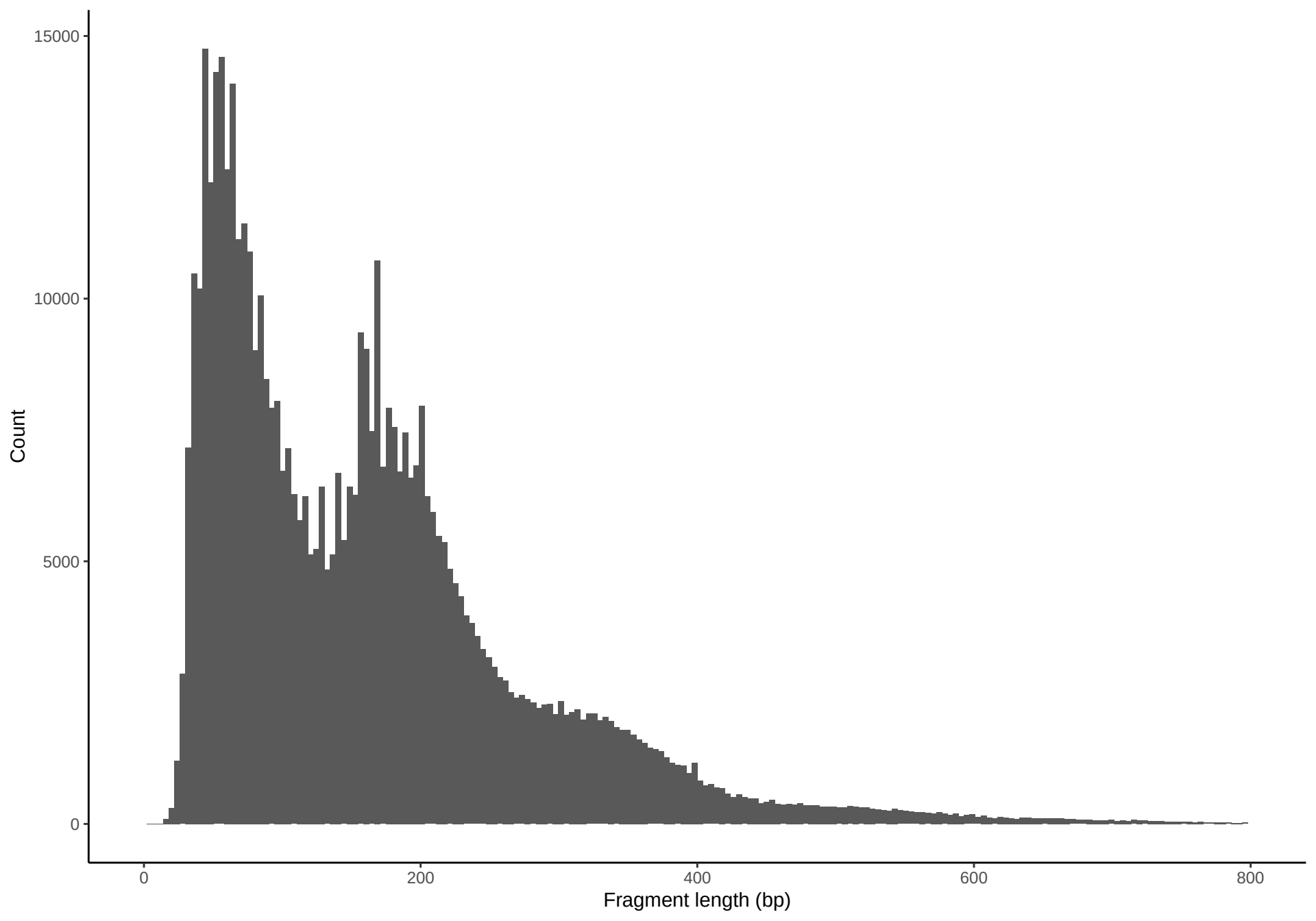
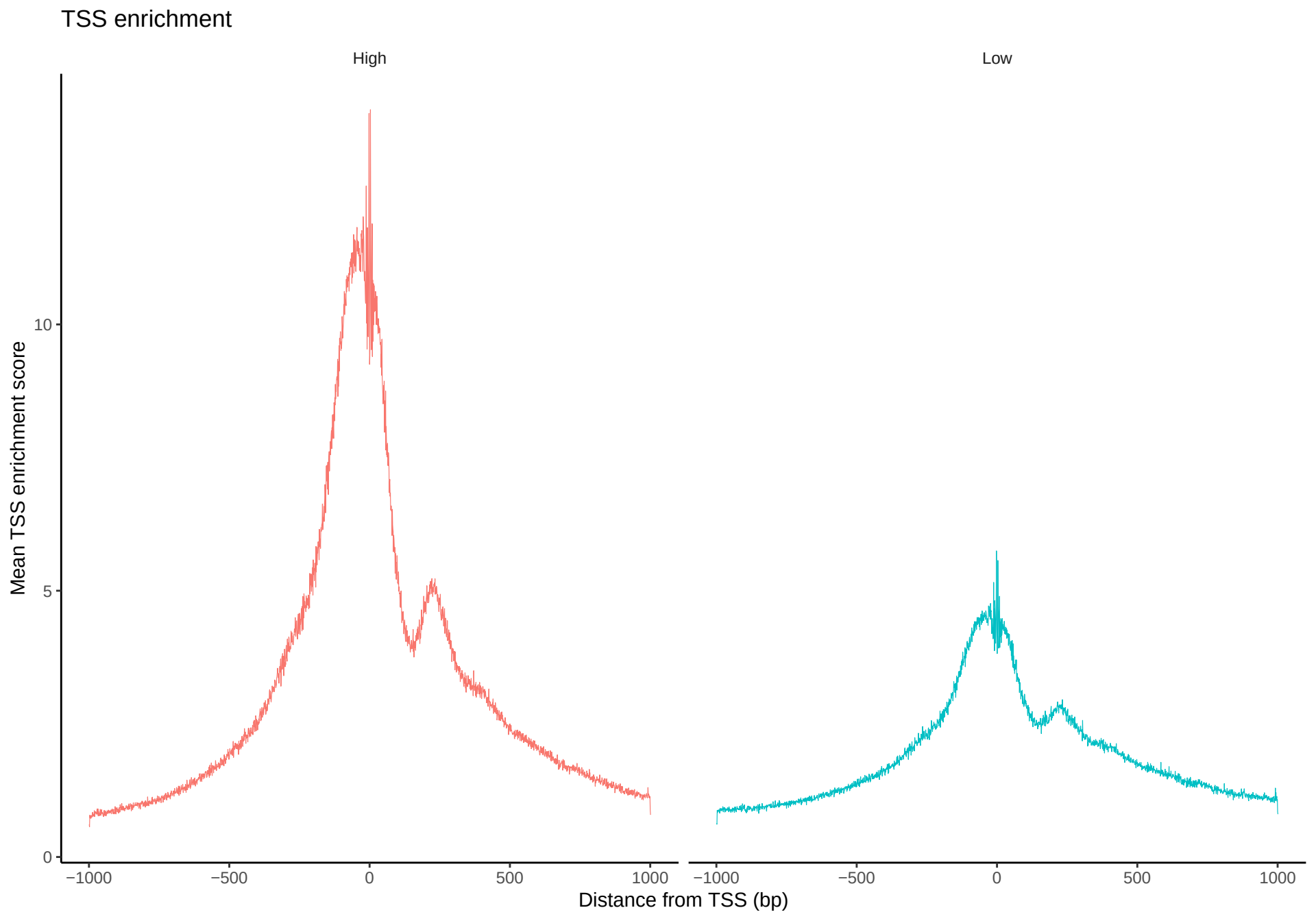
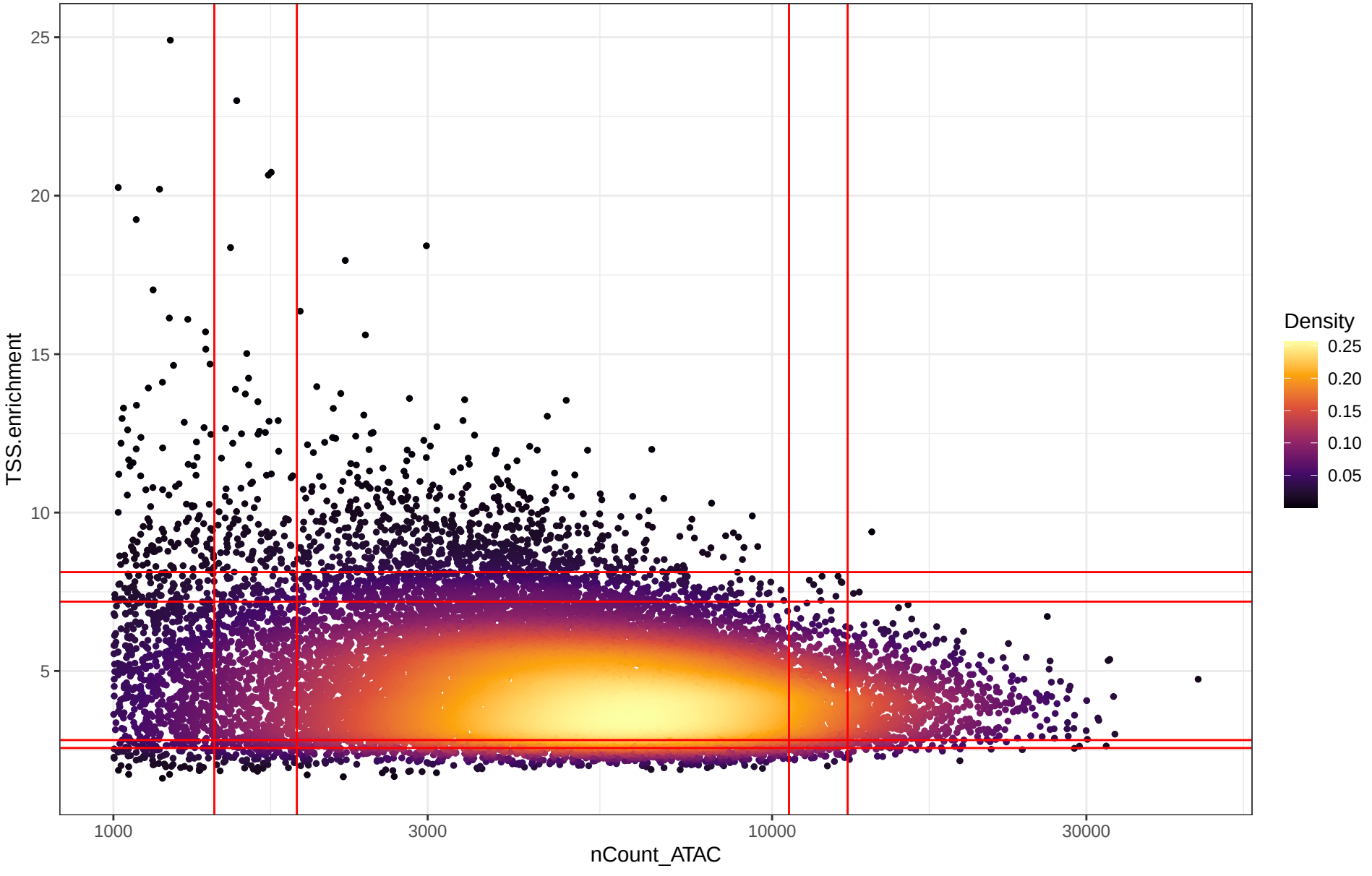

nCount_RNA
25000
20000
15000
10000
5000
0
SeuratProject
Identity
nCount_ATAC
40000
30000
20000
10000
0
SeuratProject
Identity
nFeature_RNA
6000
4000
2000
SeuratProject
Identity
mitoPct
30
20
10
0
SeuratProject
Identity
FRiP
1.0
0.5
SeuratProject
Identity
log10GenesPerUMI
0.98
0.94
0.90
0.86
SeuratProject
Identity
TSS.enrichment
25
20
15
10
5
SeuratProject
Identity
nucleosome_signal
1.5
1.0
0.5
SeuratProject
Identity

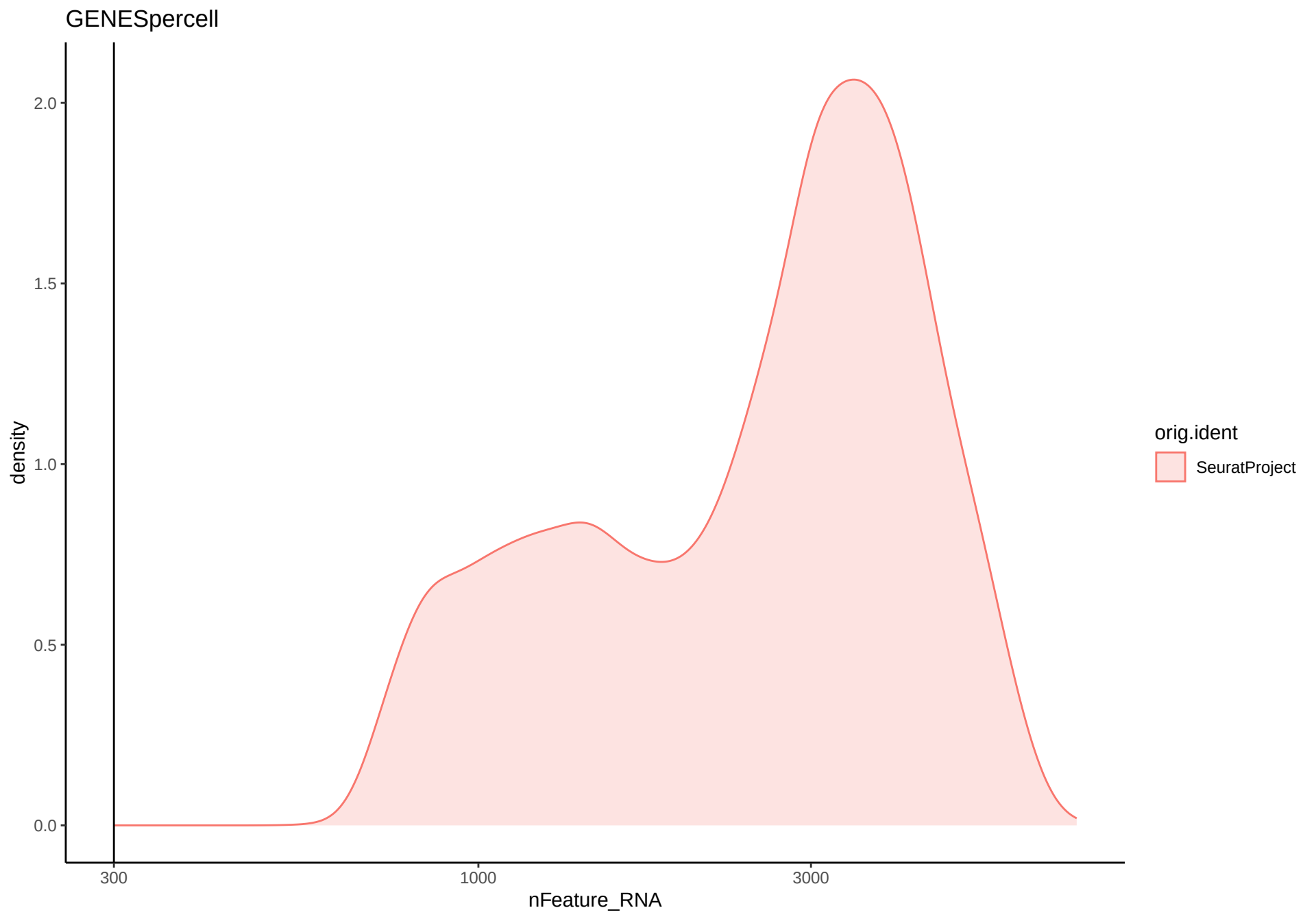

15000
10000
5000
0
Count
0
200
400
600
800
Fragment length (bp)

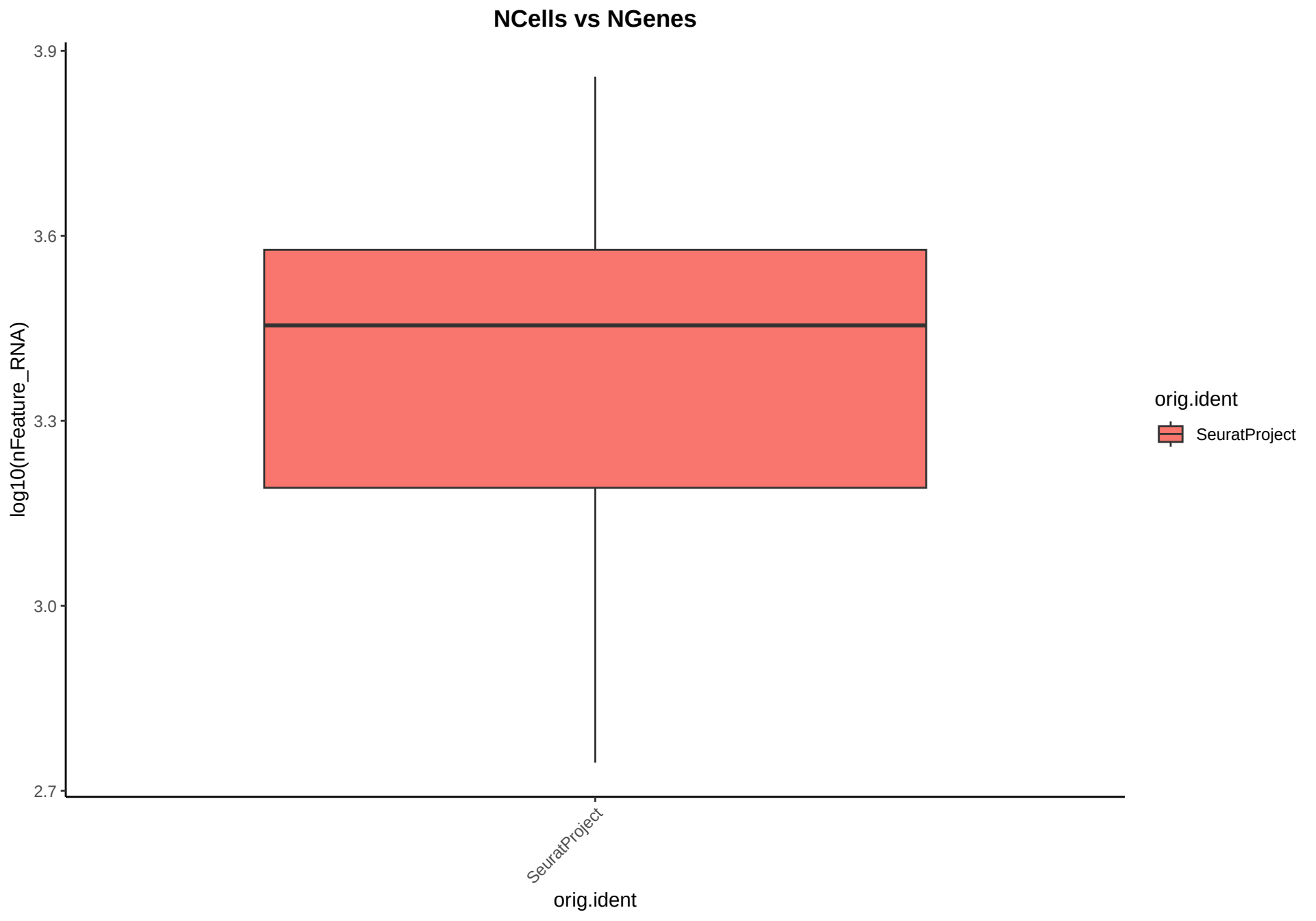

TSS enrichment
High
Low
Mean TSS enrichment score
10
5
0
−1000
−500
0
500
1000
Distance from TSS (bp)

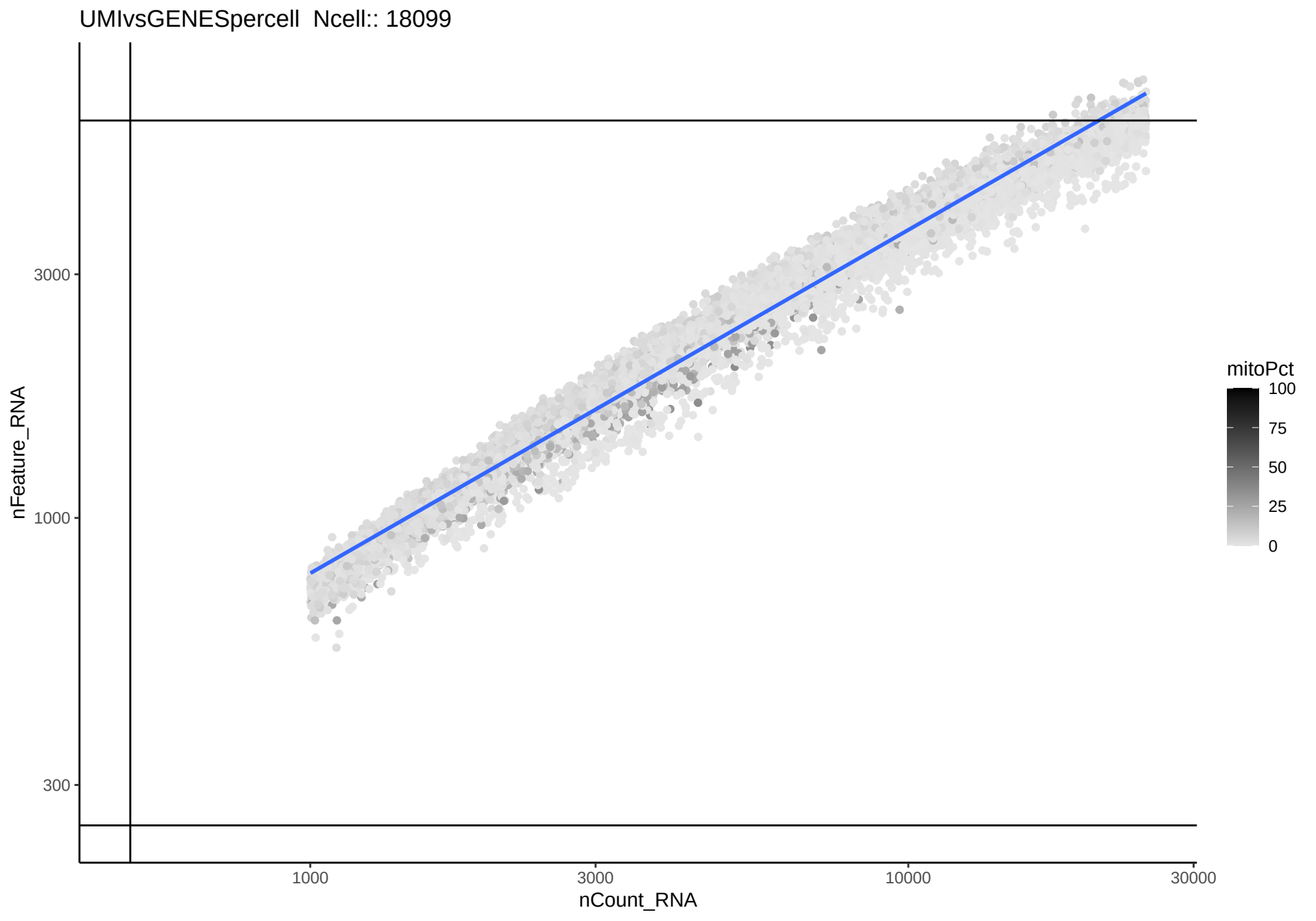
Quantiles

nCount_ATAC: 5%:1423 10%:1898.8 90%:10608.4 95%:13021.1
TSS.enrichment: 5%:2.57 10%:2.82 90%:7.19 95%:8.12




GENESpercell
density
2.0
1.5
1.0
0.5
0.0
300
1000
3000
nFeature_RNA
orig.ident
SeuratProject


NCells vs NGenes
3.9
3.6
3.3
3.0
2.7
log10(nFeature_RNA)
SeuratProject
orig.ident
orig.ident
SeuratProject


UMIvsGENESpercell Ncell:: 18099
nFeature_RNA
3000
1000
300
1000
3000
10000
30000
nCount_RNA
mitoPct
100
75
50
25
0