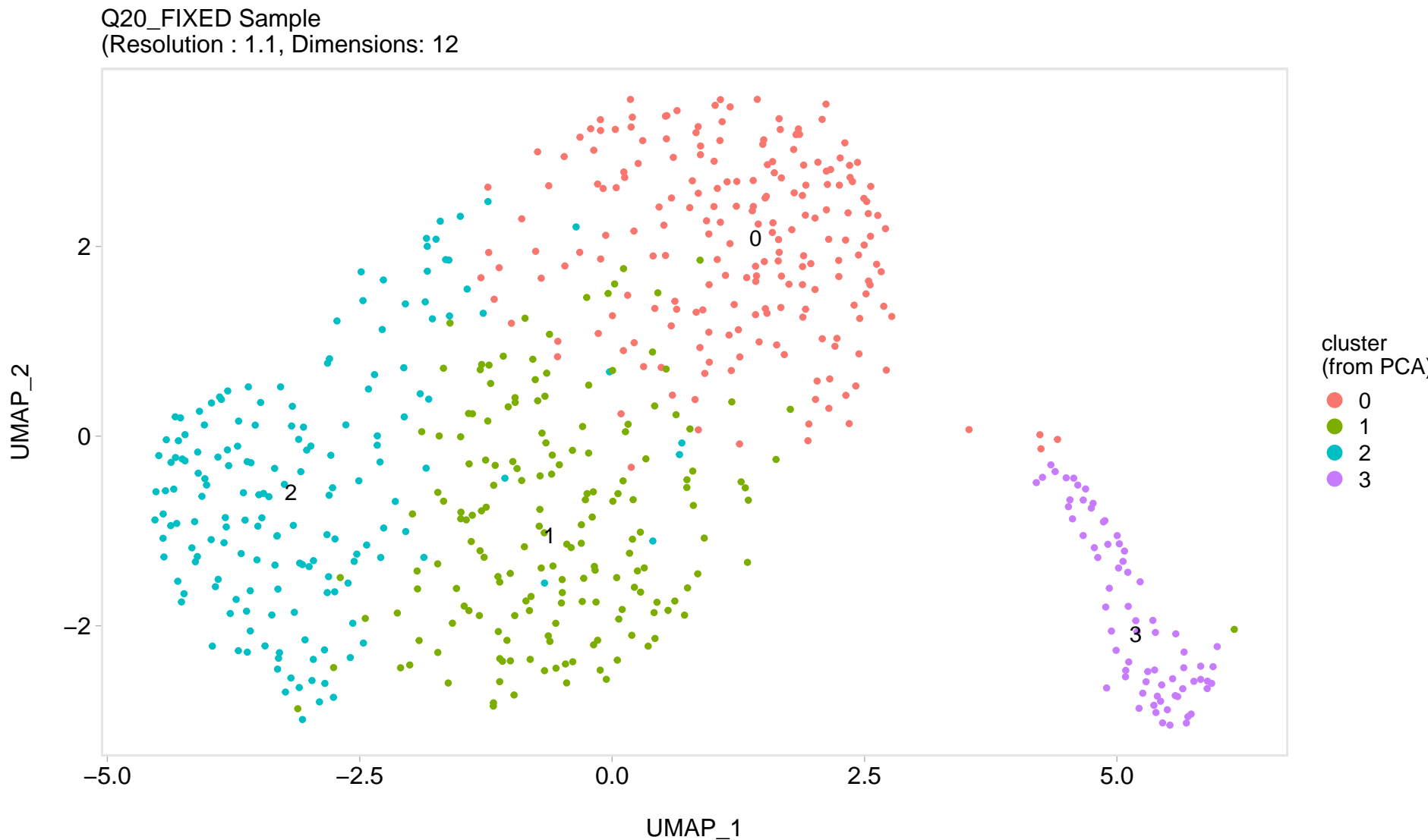
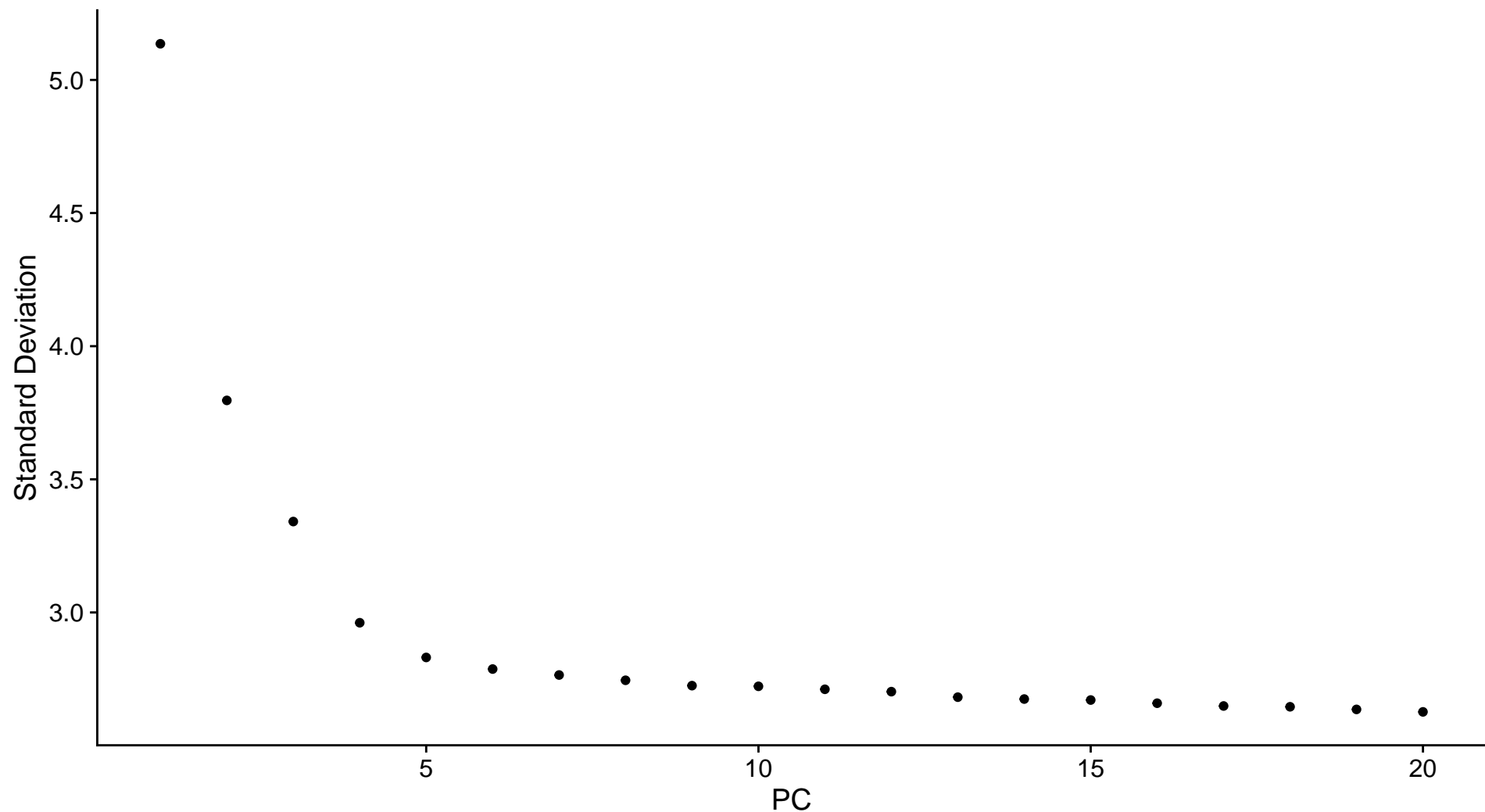


SCRIPT 2 (QC) SUMMARY (Q20_FIXED)

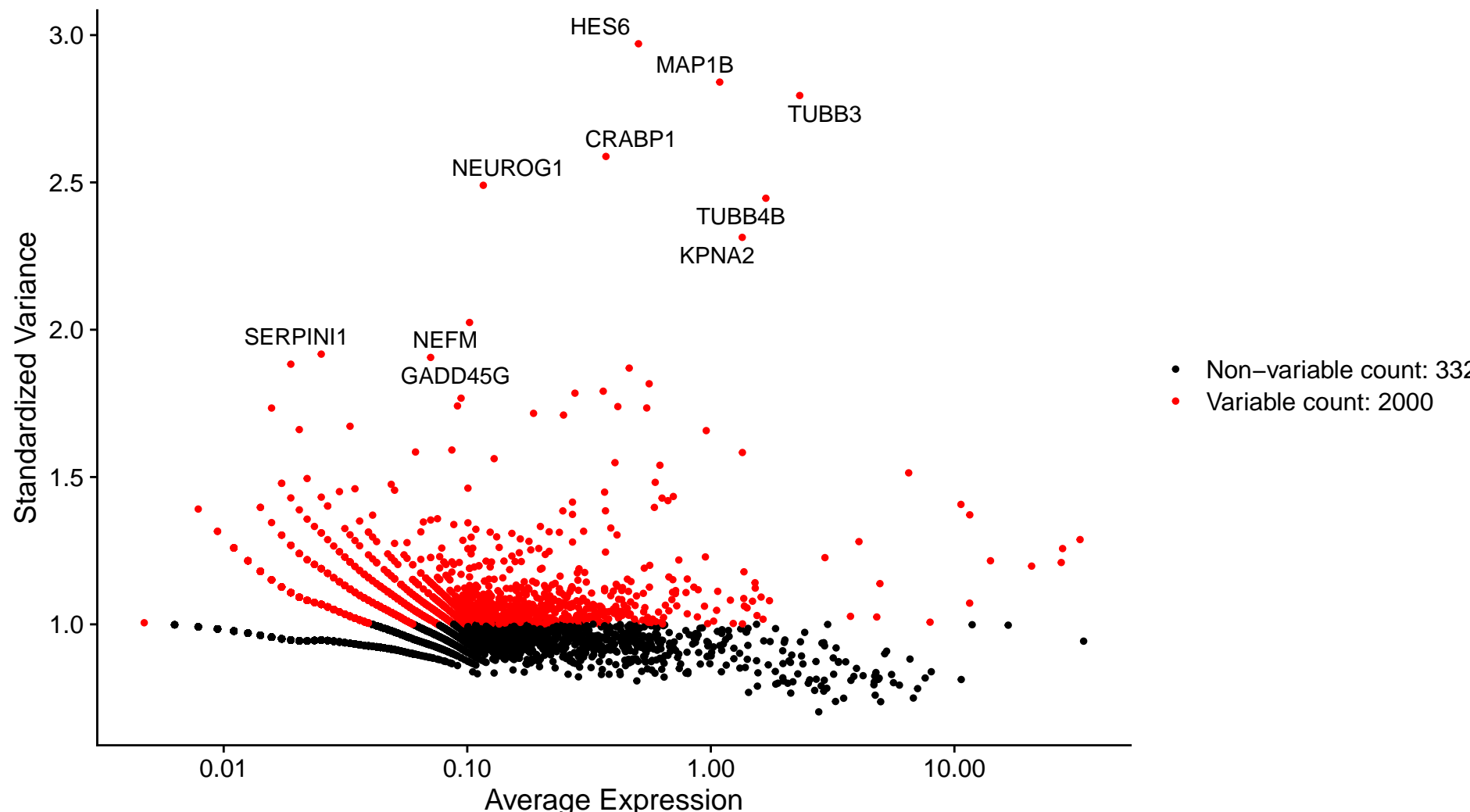
Sample ID	Q20_FIXED
Initial Cell Count	811
Cells after QC	618
Feature Type	Isoform
Min. Feature Threshold	200
Max. Feature Threshold	1000
Min. Counts Threshold	500
Max. Counts Threshold	1e+05
Median Features per Cell	497.5
Median Reads per Cell	949
Mitochondrial Content Threshold (%)	10
Median MT % before filter	0
Median MT % after filter	0
NPCs (UMAP dimensions)	12



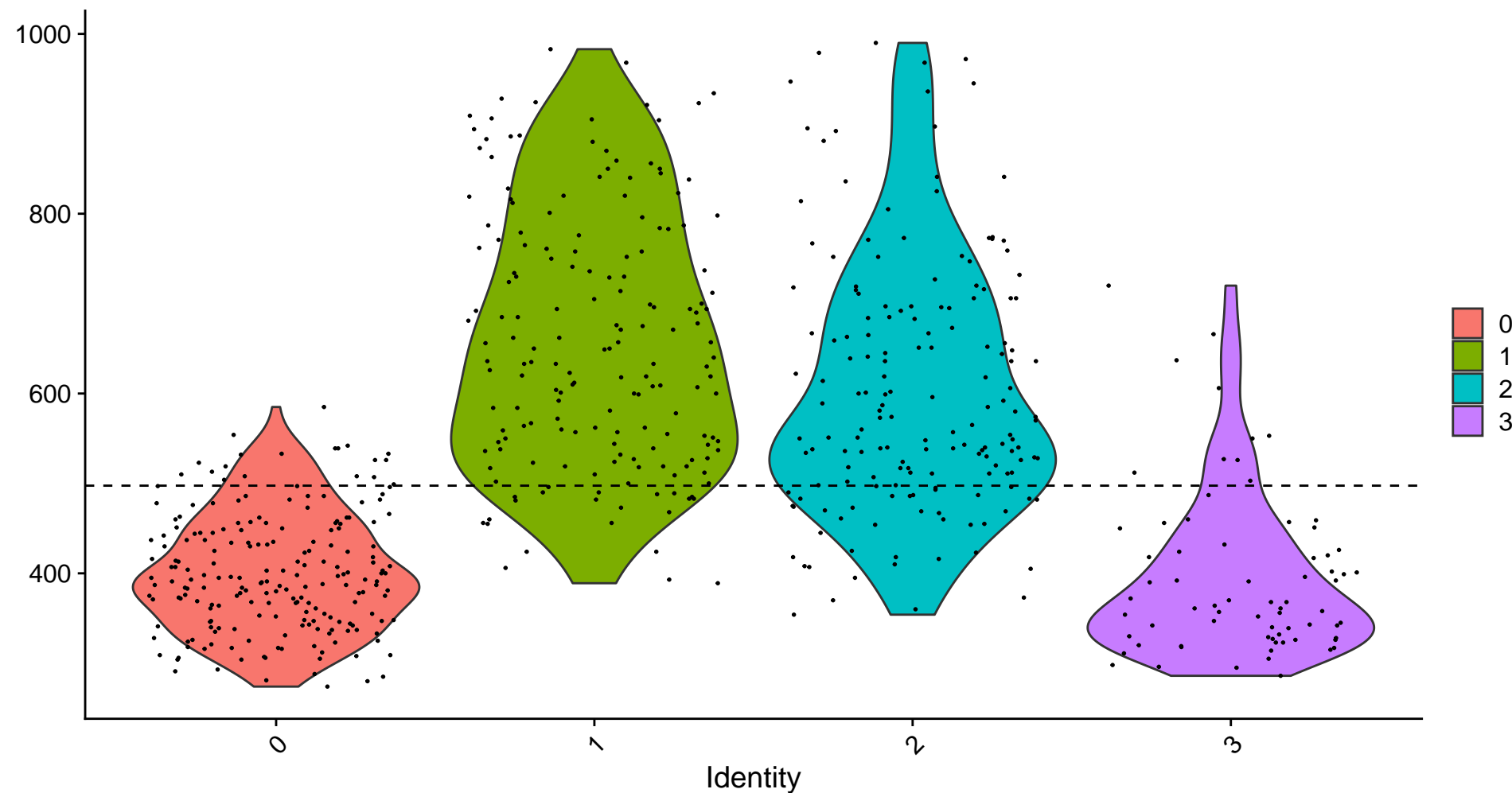
Standard Deviation explained by each Principle Component



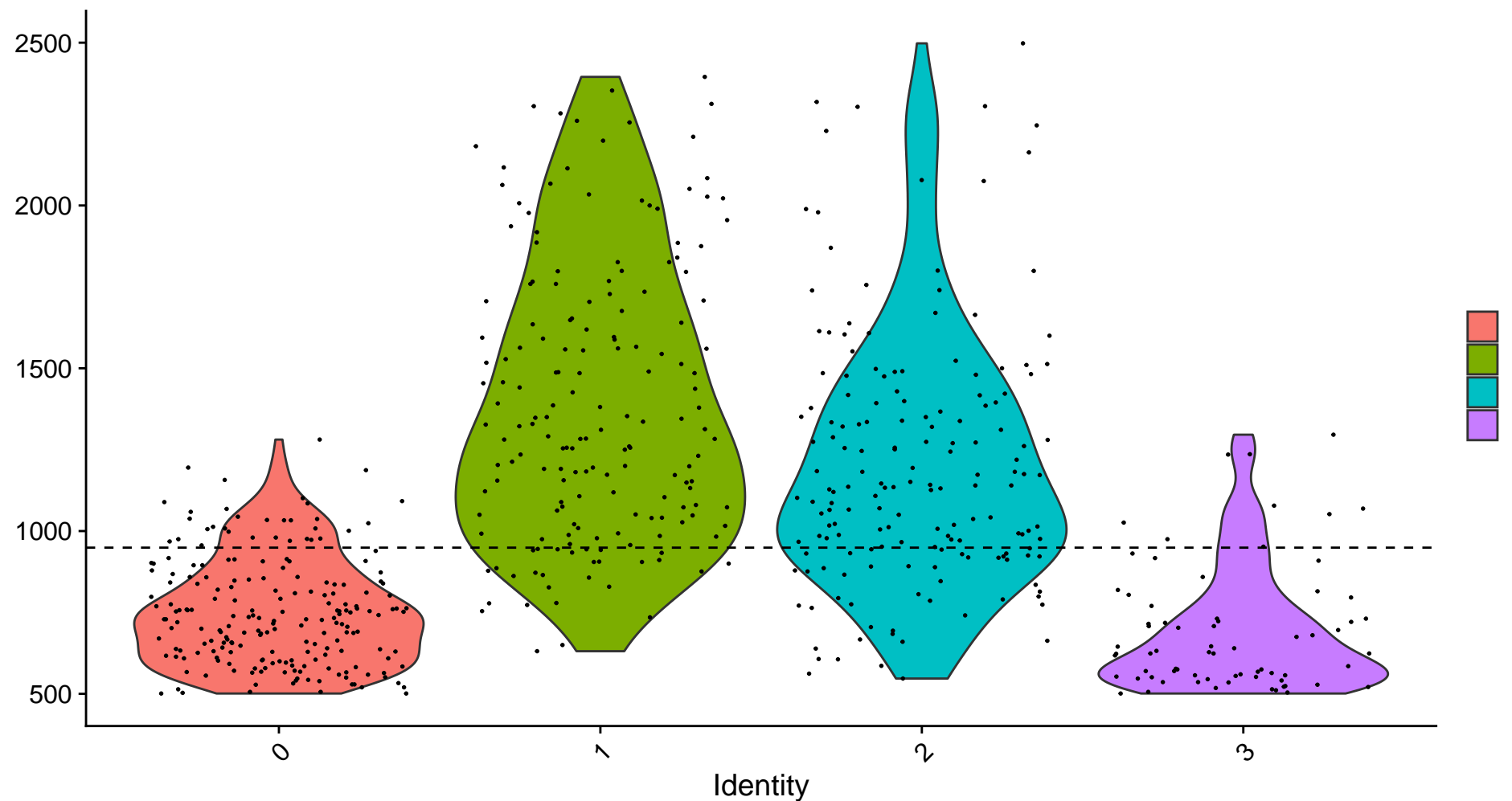
Top 2000 Variable Genes across Cells



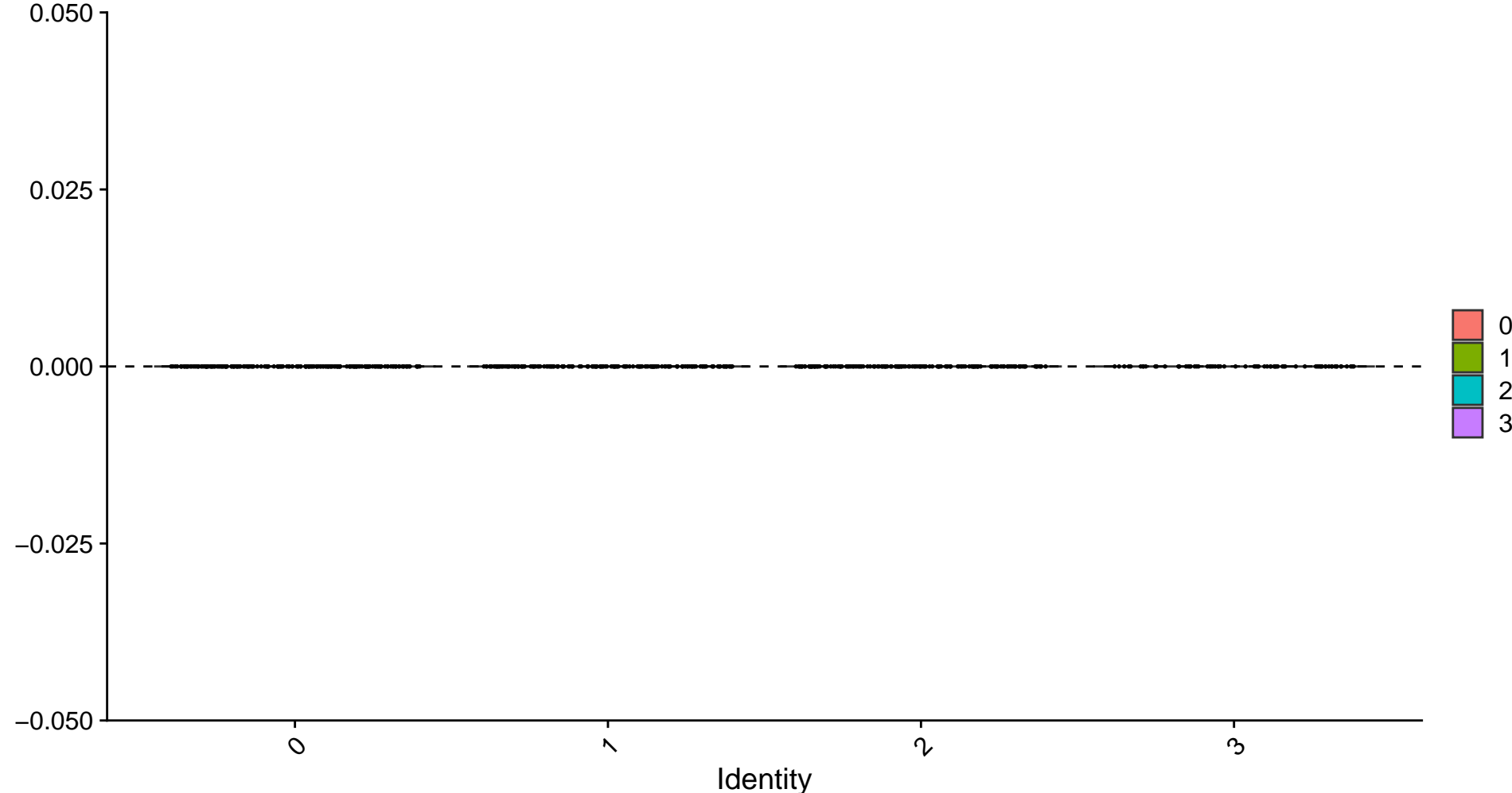
Unique Features per Cell
After Removing Unwanted Cells



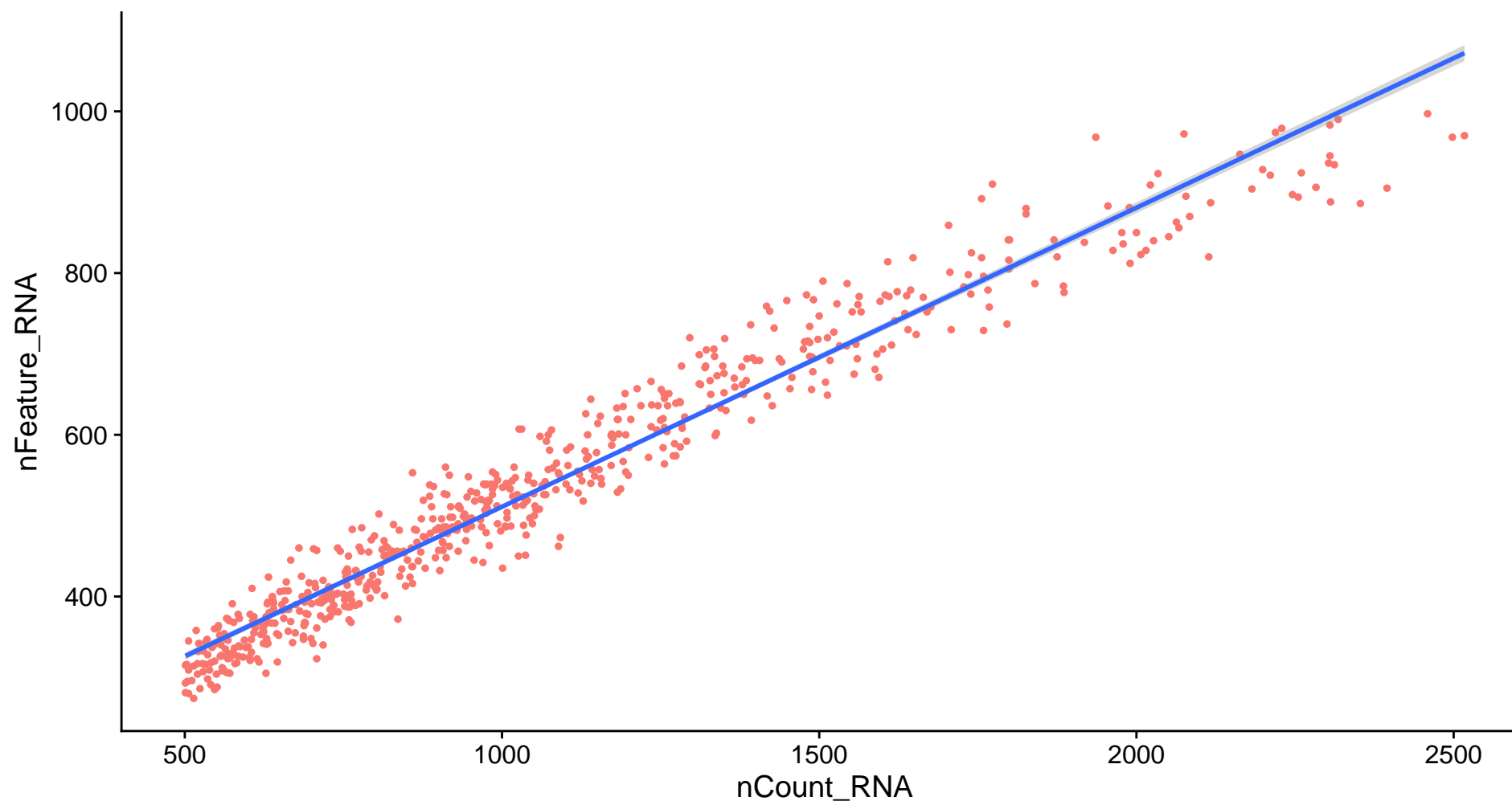
RNA Molecules per Cell
After Filtering



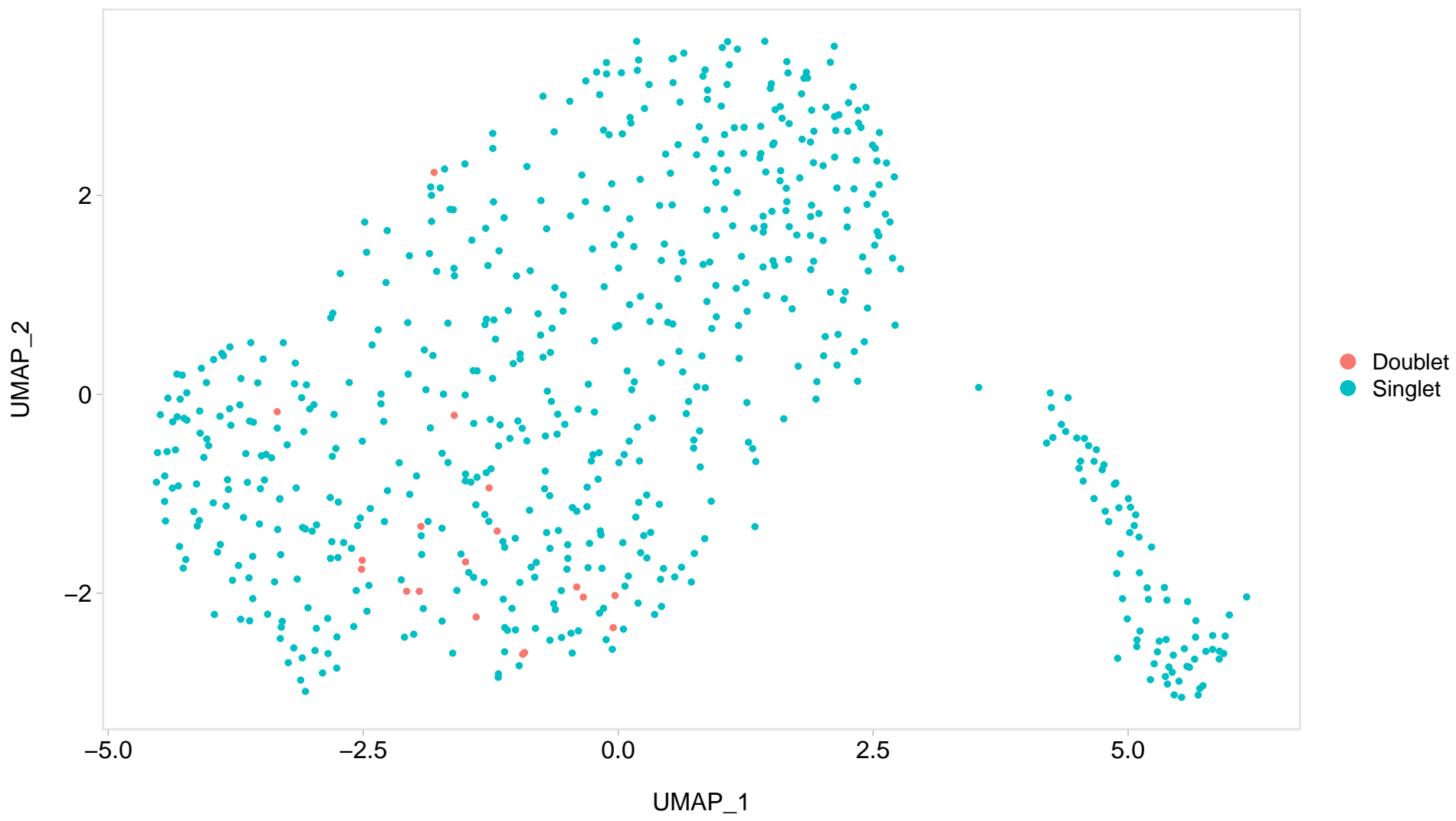
Mitochondrial Content (%)
After Removing Unwanted Cells



Association between reads
and unique features per cell AFTER filering



Doublets that were detected and removed



	DF.classifications	median_nCount_RNA	median_nFeature_RNA	count
1	Doublet	1583	778.0	18
2	Singlet	949	497.5	618