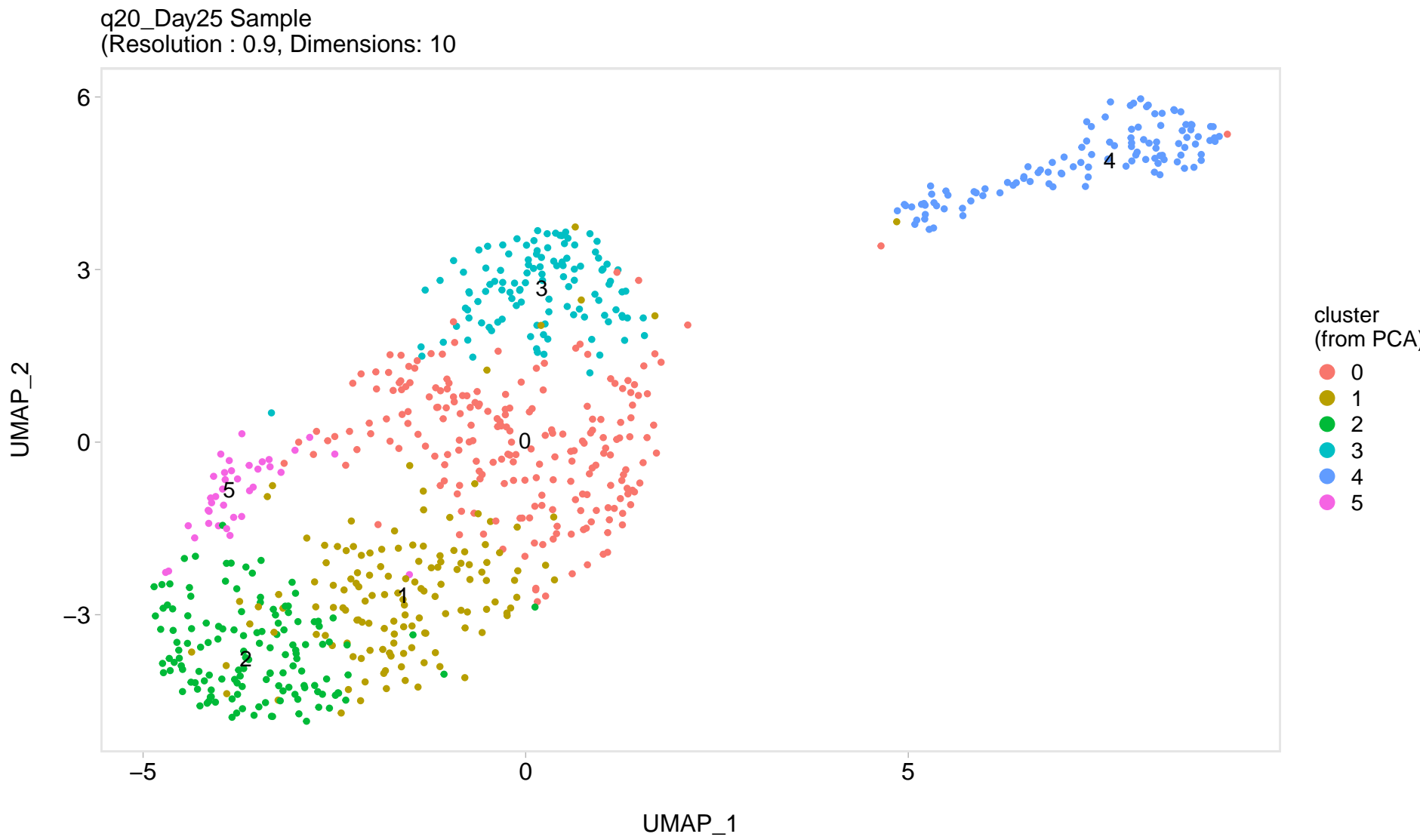
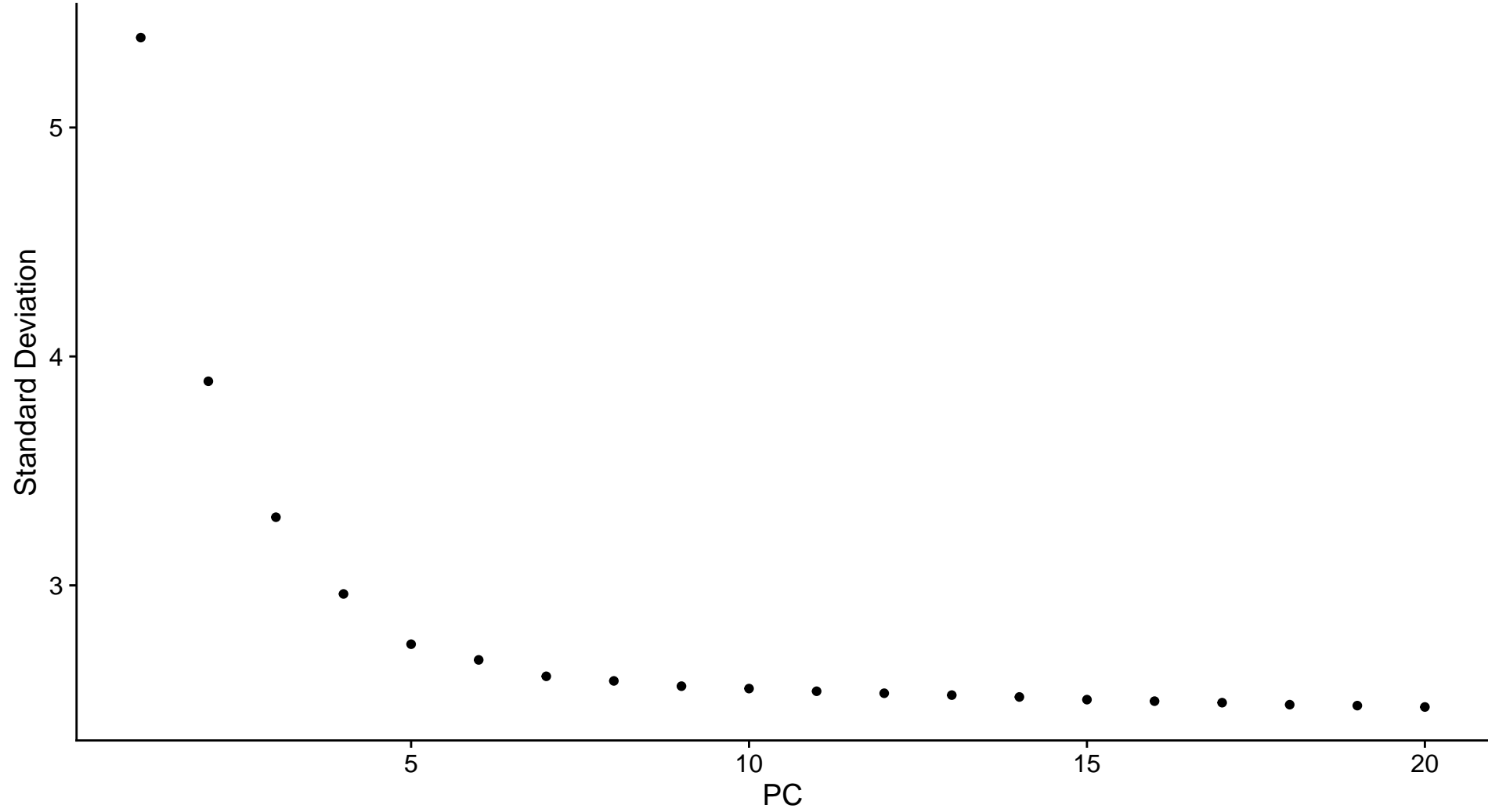


SCRIPT 2 (QC) SUMMARY (q20_Day25)

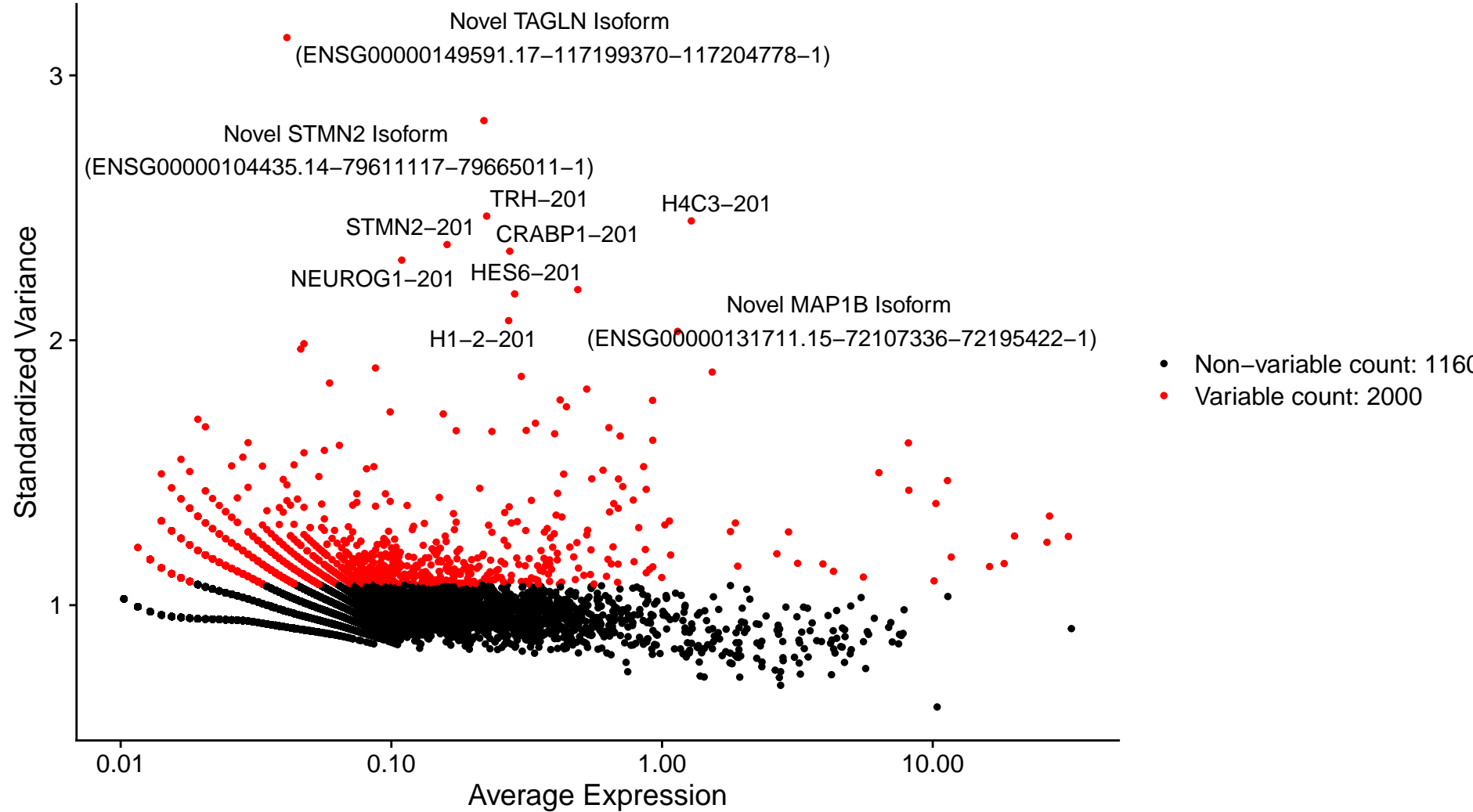
Sample ID	q20_Day25
Initial Cell Count	811
Cells after QC	752
Feature Type	Isoform
Min. Feature Threshold	200
Max. Feature Threshold	999999999
Min. Counts Threshold	400
Max. Counts Threshold	1e+05
Median Features per Cell	800.5
Median Reads per Cell	1322.5
Mitochondrial Content Threshold (%)	10
Median MT % before filter	0
Median MT % after filter	0
NPCs (UMAP dimensions)	10



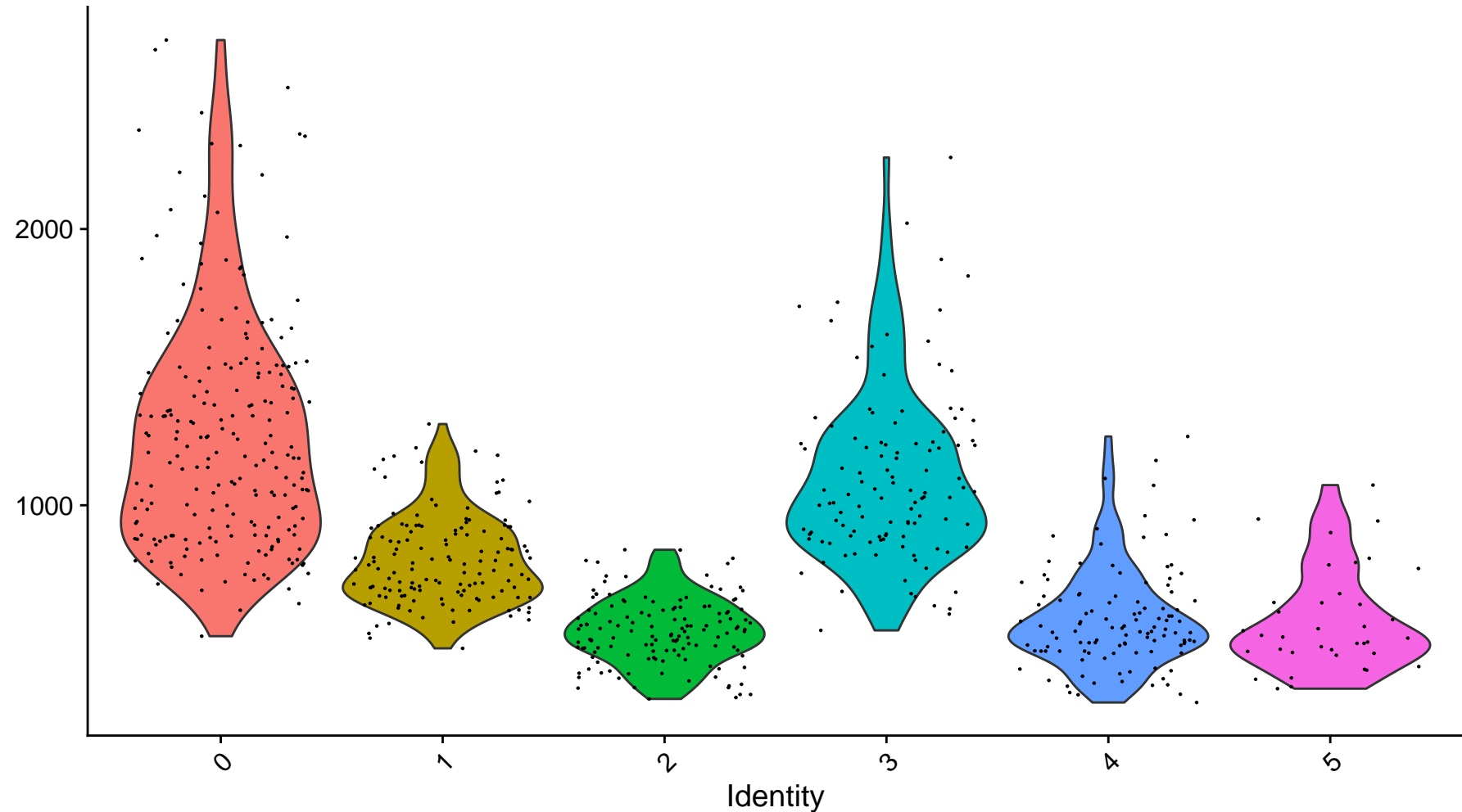
Standard Deviation explained by each Principle Component



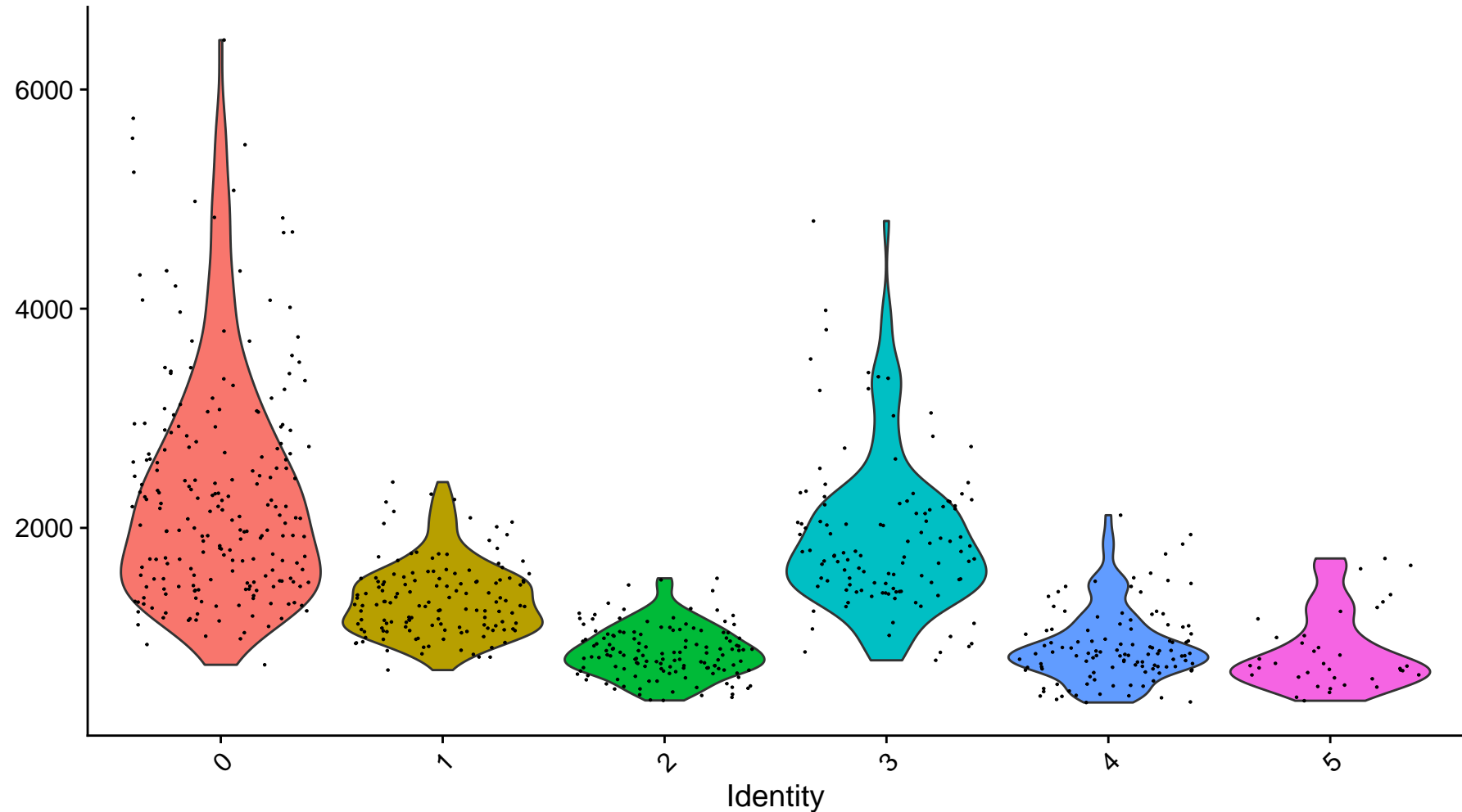
Top 2000 Variable Features across Cells



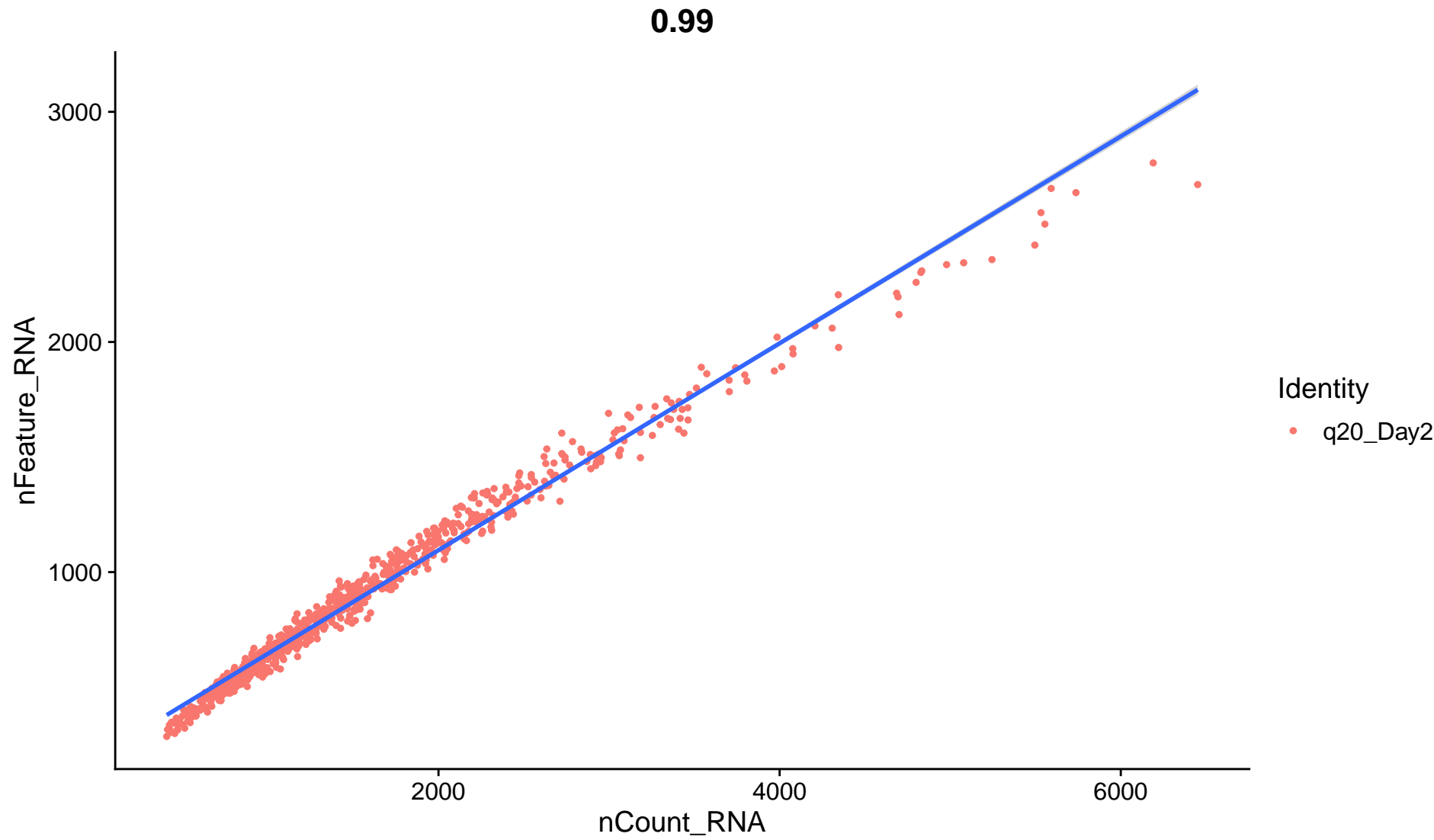
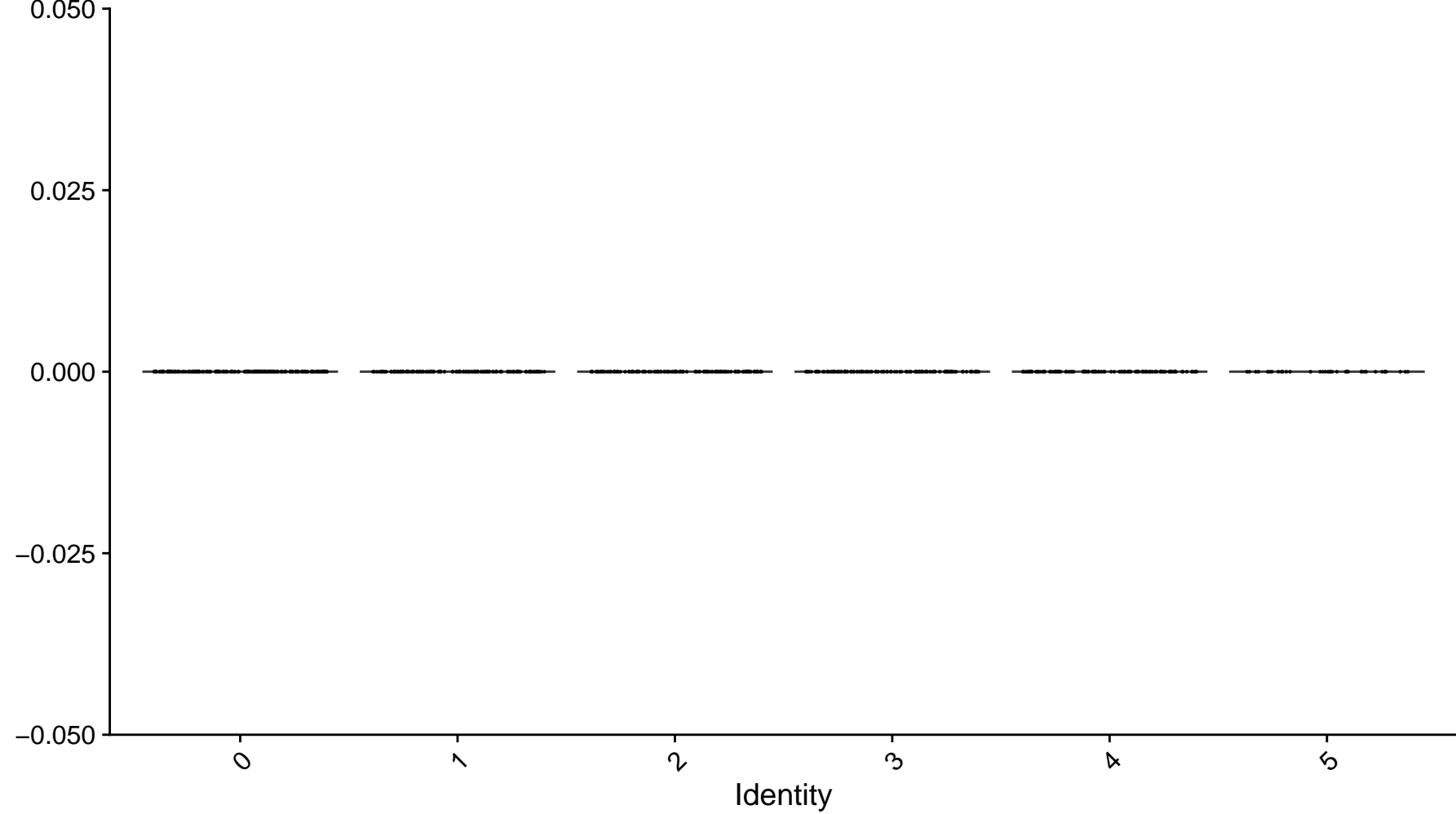
Unique Features per Cell
After Removing Unwanted Cells



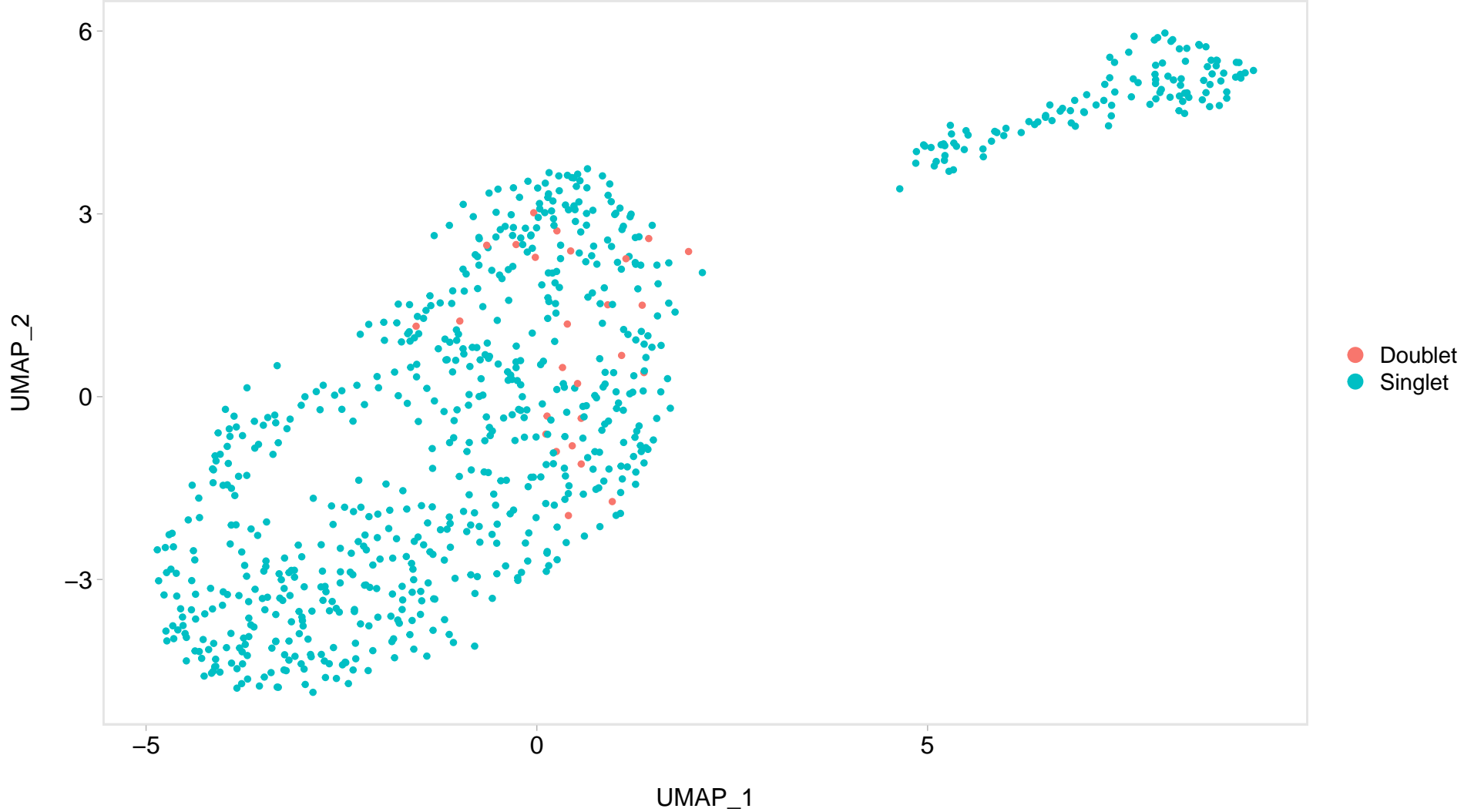
RNA Molecules per Cell
After Filtering



Mitochondrial Content (%)
After Removing Unwanted Cells



Doublets that were detected and removed



	DF.classifications	median_nCount_RNA	median_nFeature_RNA	count
1	Doublet	2599.5	1426.5	26
2	Singlet	1322.5	800.5	752