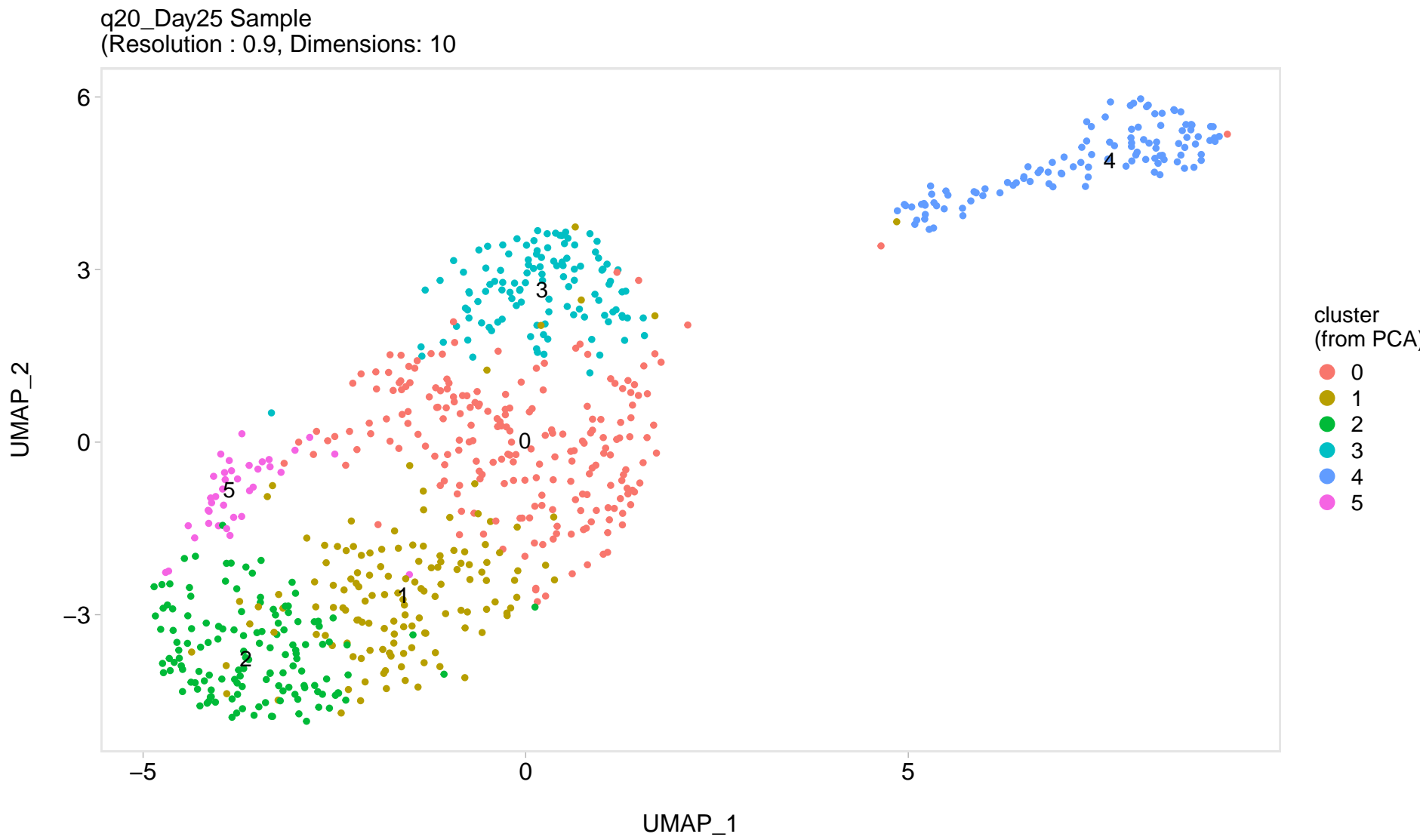
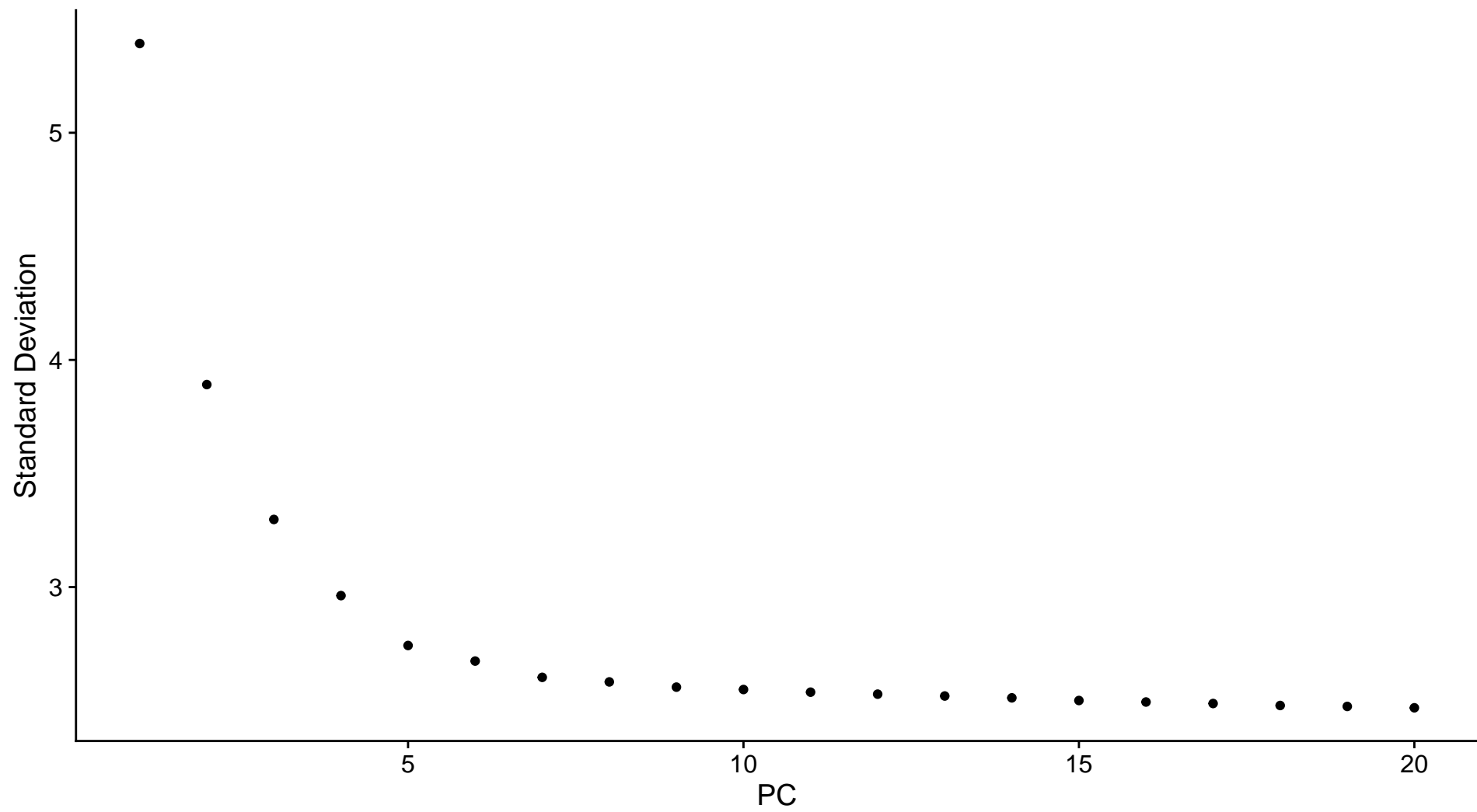


SCRIPT 2 (QC) SUMMARY (q20\_Day25)

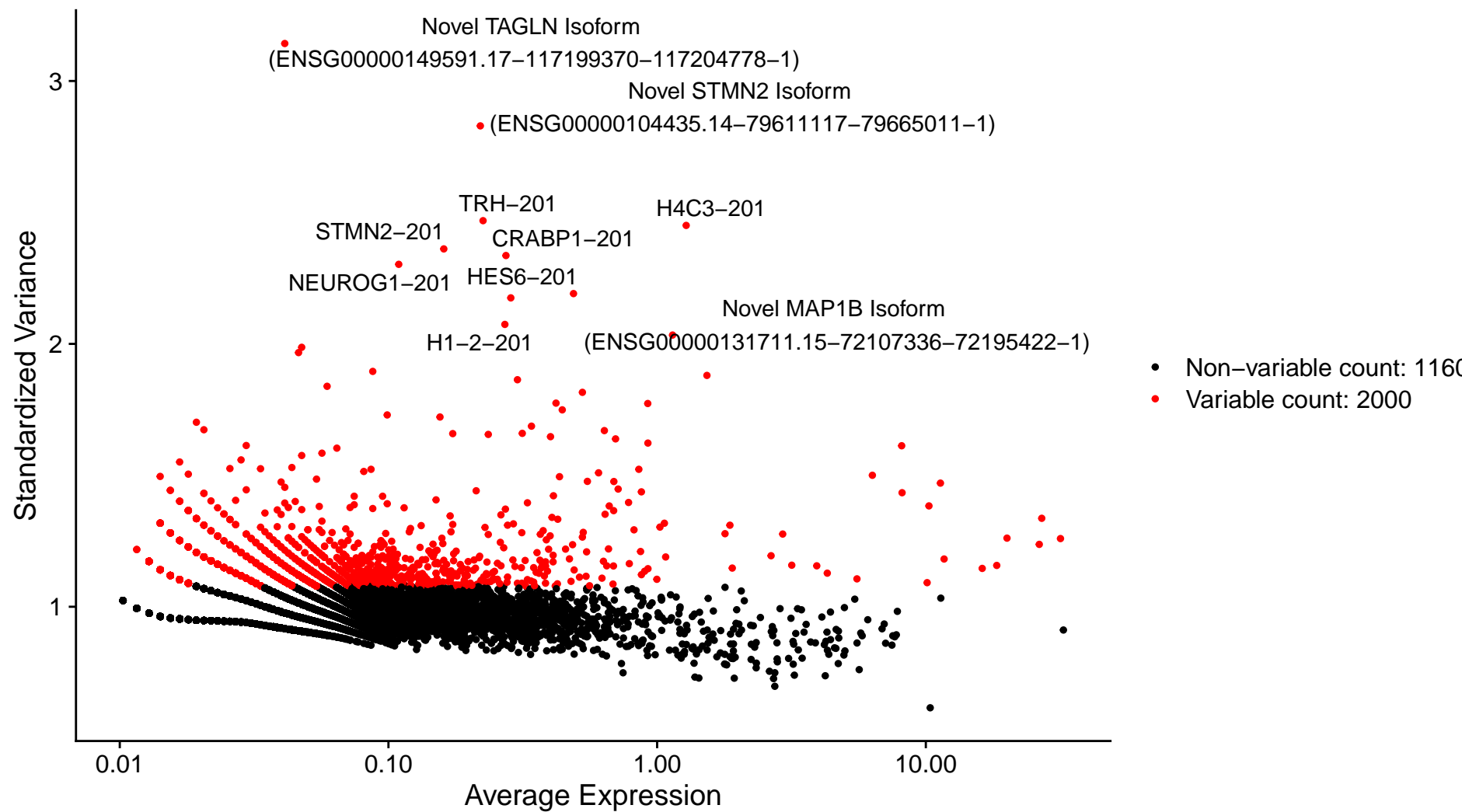
Sample ID	q20_Day25
Initial Cell Count	811
Cells after QC	753
Feature Type	Isoform
Min. Feature Threshold	200
Max. Feature Threshold	999999999
Min. Counts Threshold	400
Max. Counts Threshold	1e+05
Median Features per Cell	801
Median Reads per Cell	1323
Mitochondrial Content Threshold (%)	10
Median MT % before filter	0
Median MT % after filter	0
NPCs (UMAP dimensions)	10



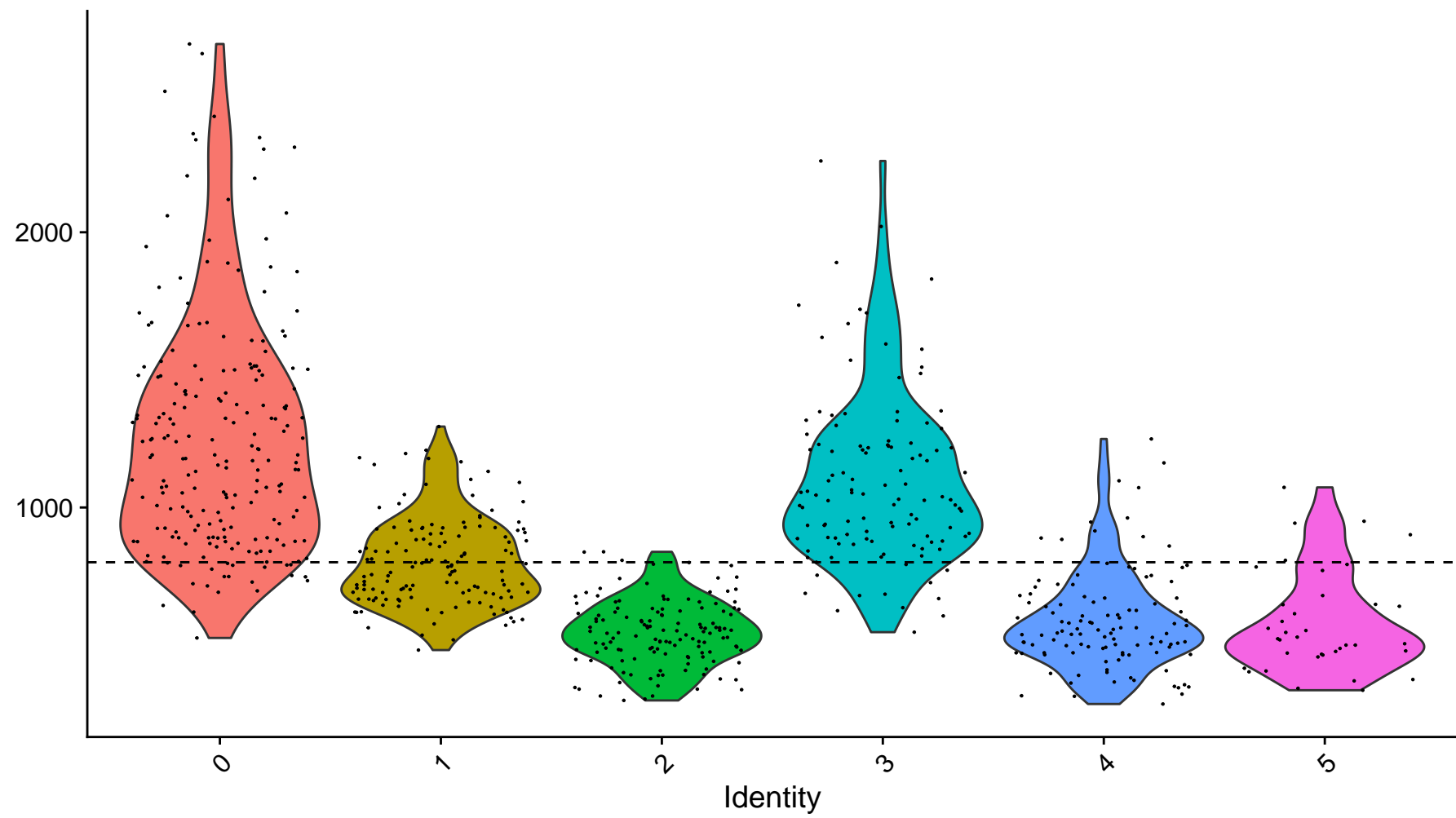
Standard Deviation explained by each Principle Component



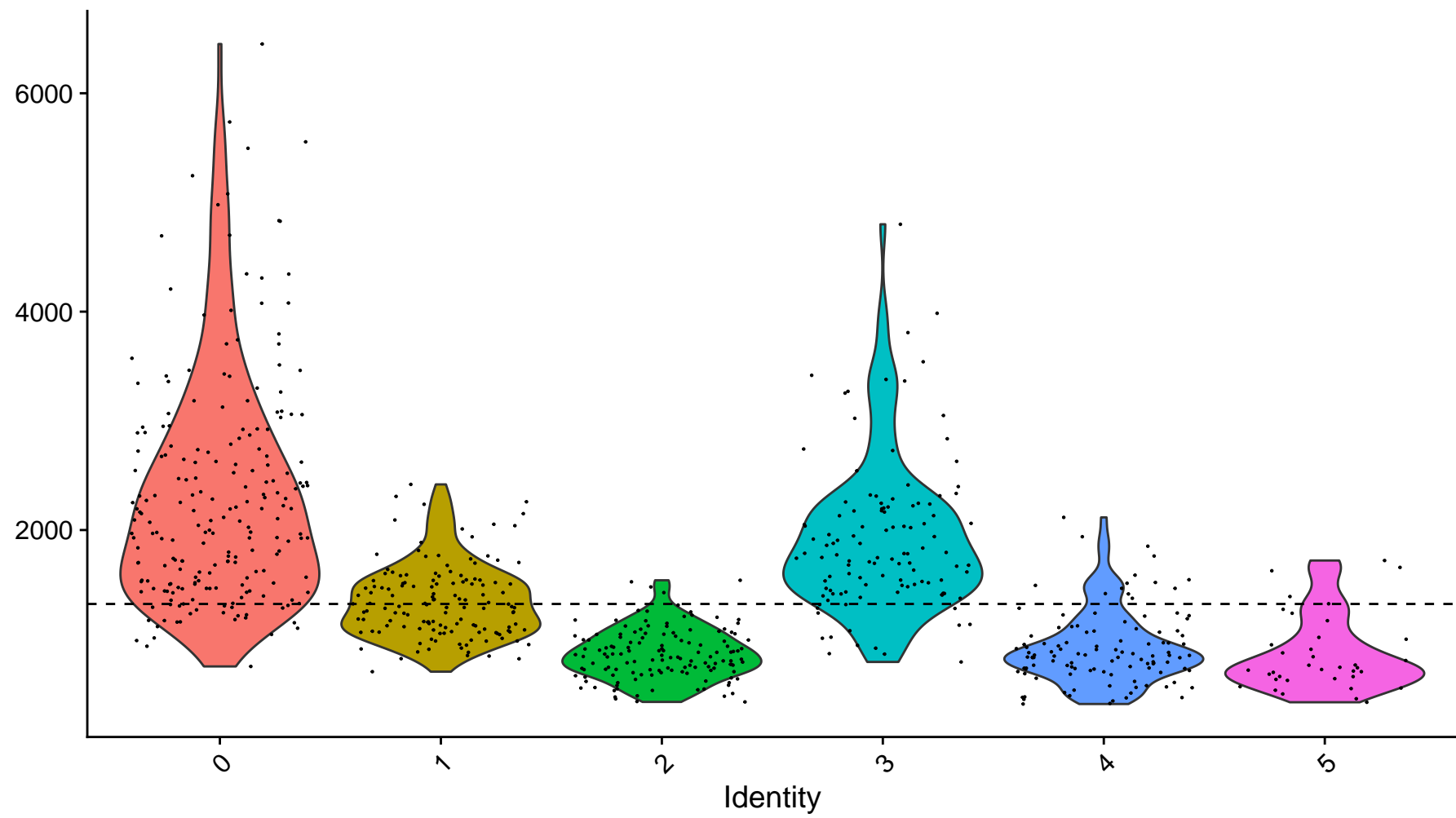
Top 2000 Variable Features across Cells



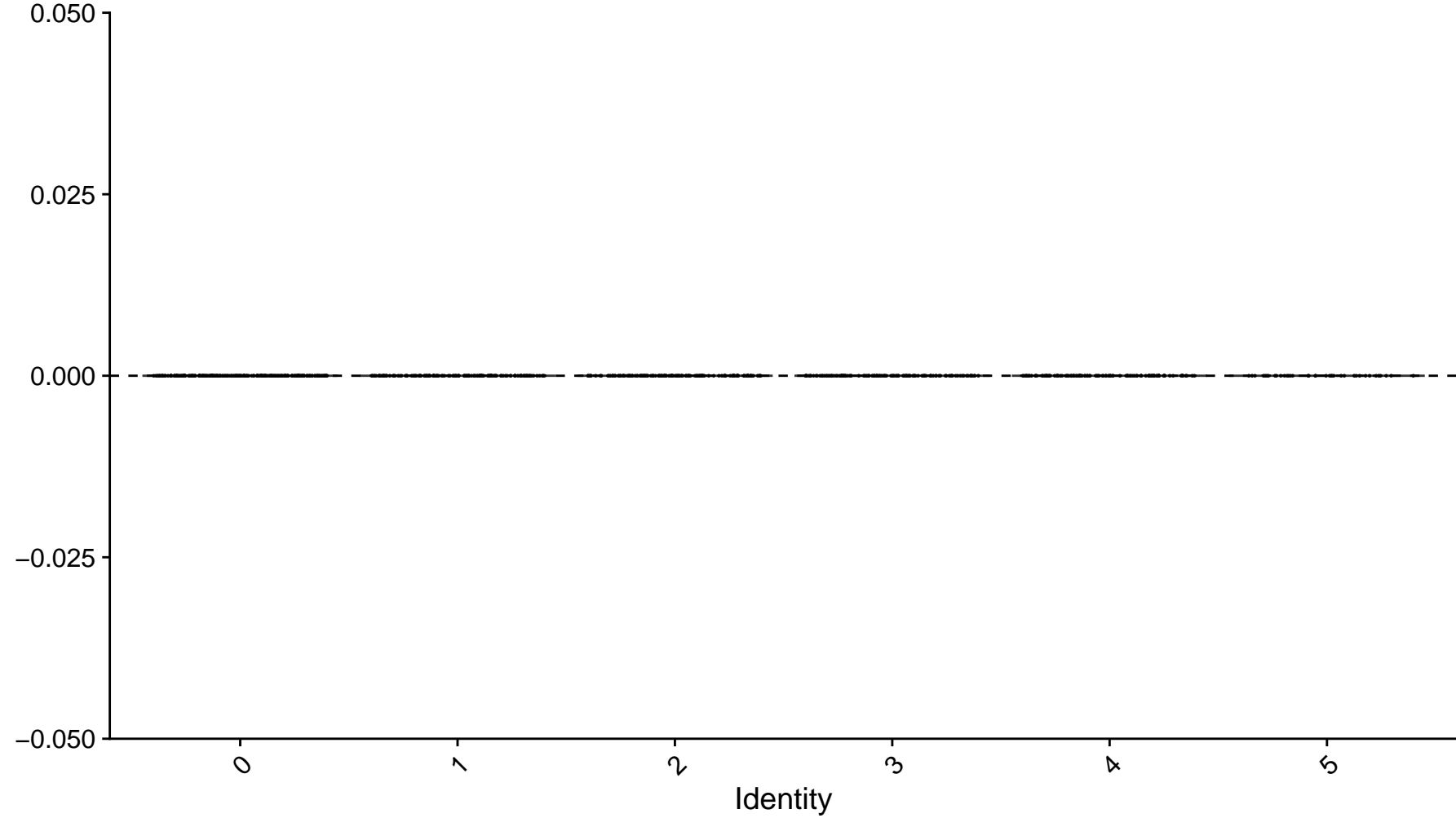
Unique Features per Cell  
After Removing Unwanted Cells



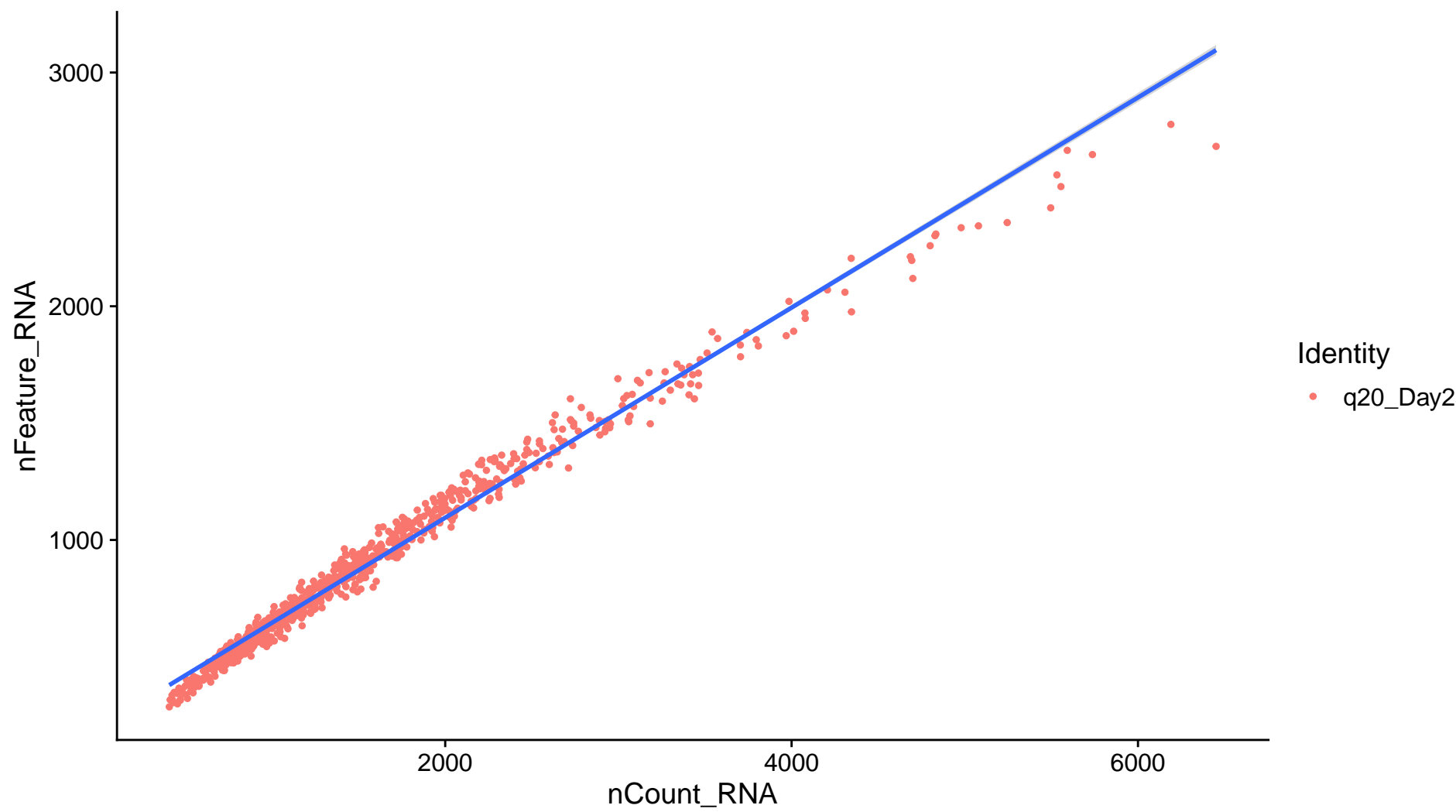
RNA Molecules per Cell  
After Filtering



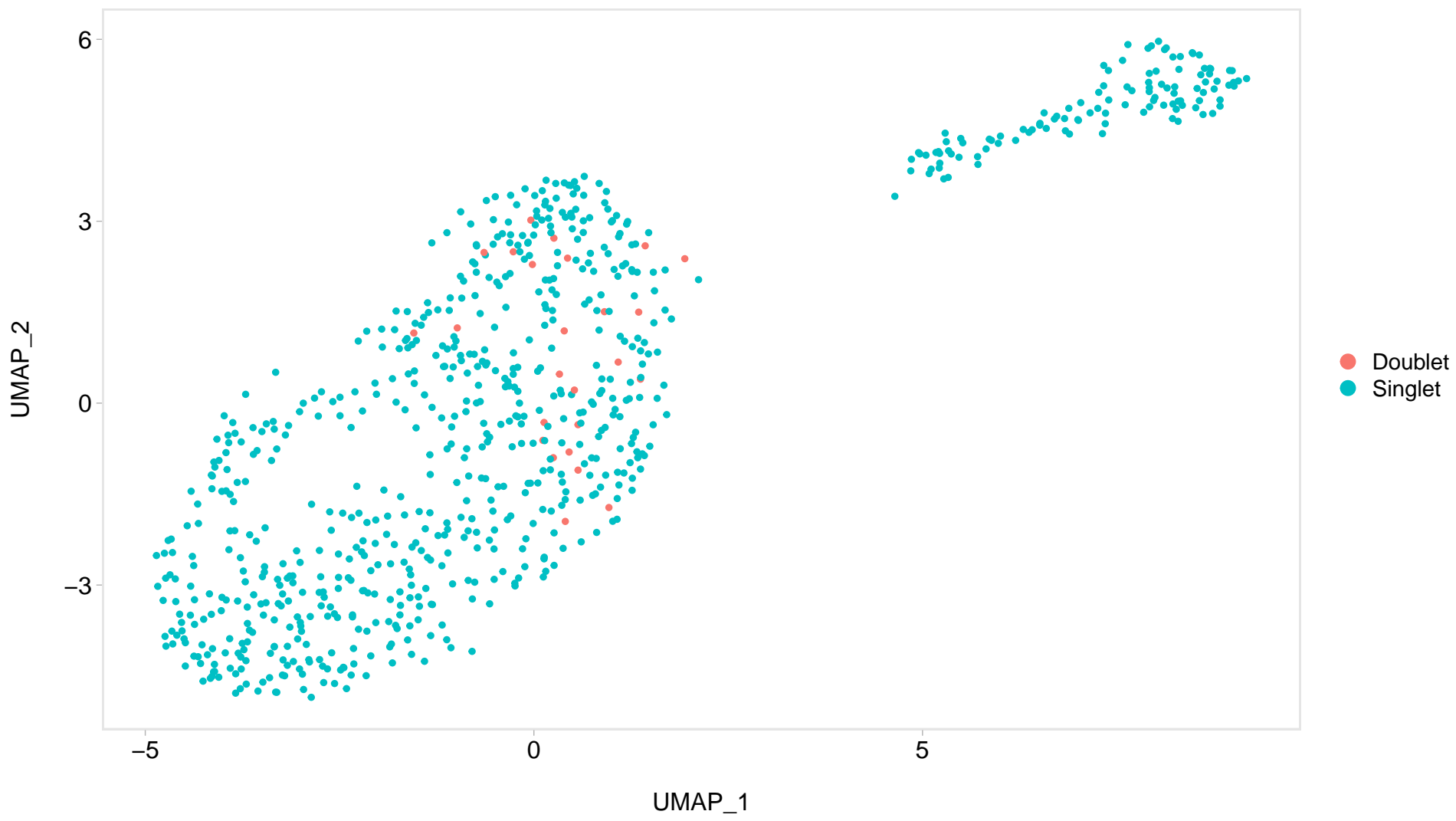
Mitochondrial Content (%)  
After Removing Unwanted Cells



Association between unique features and number of reads per cell



Doublets that were detected and removed



	DF.classifications	median_nCount_RNA	median_nFeature_RNA	count
1	Doublet	2635	1434	25
2	Singlet	1323	801	753