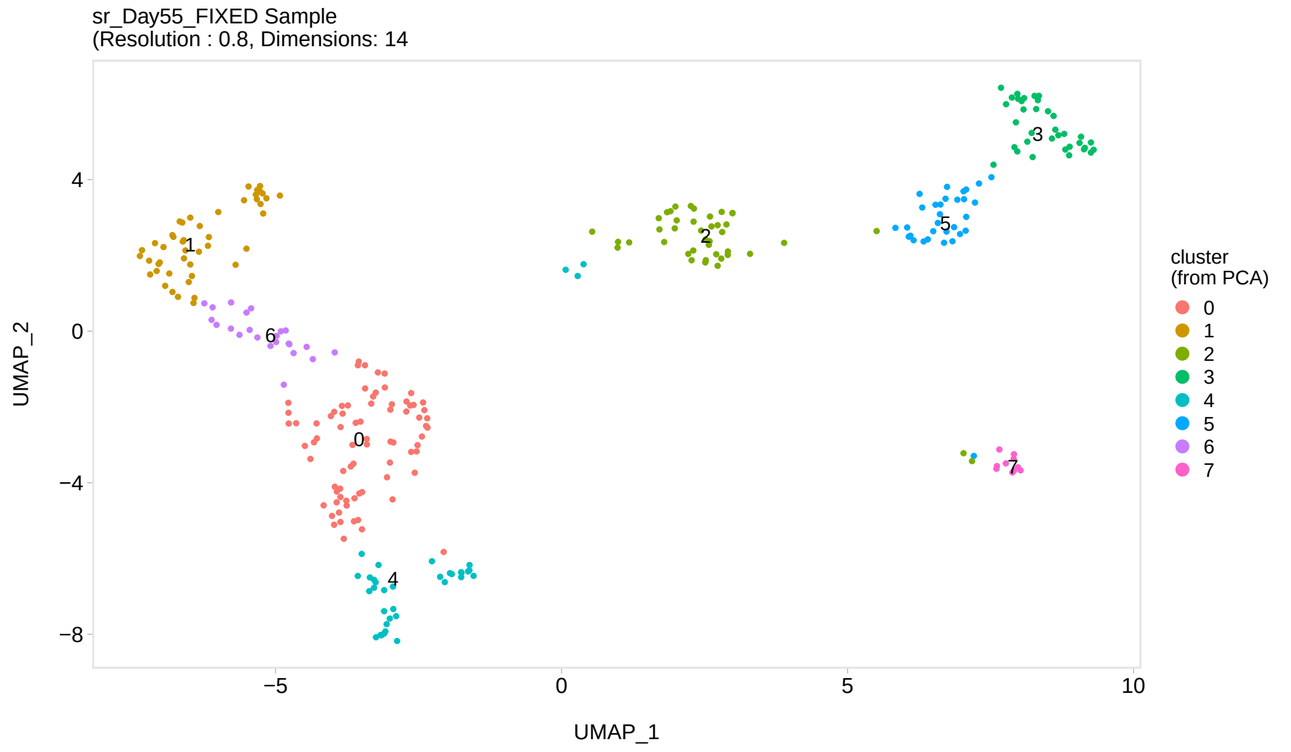
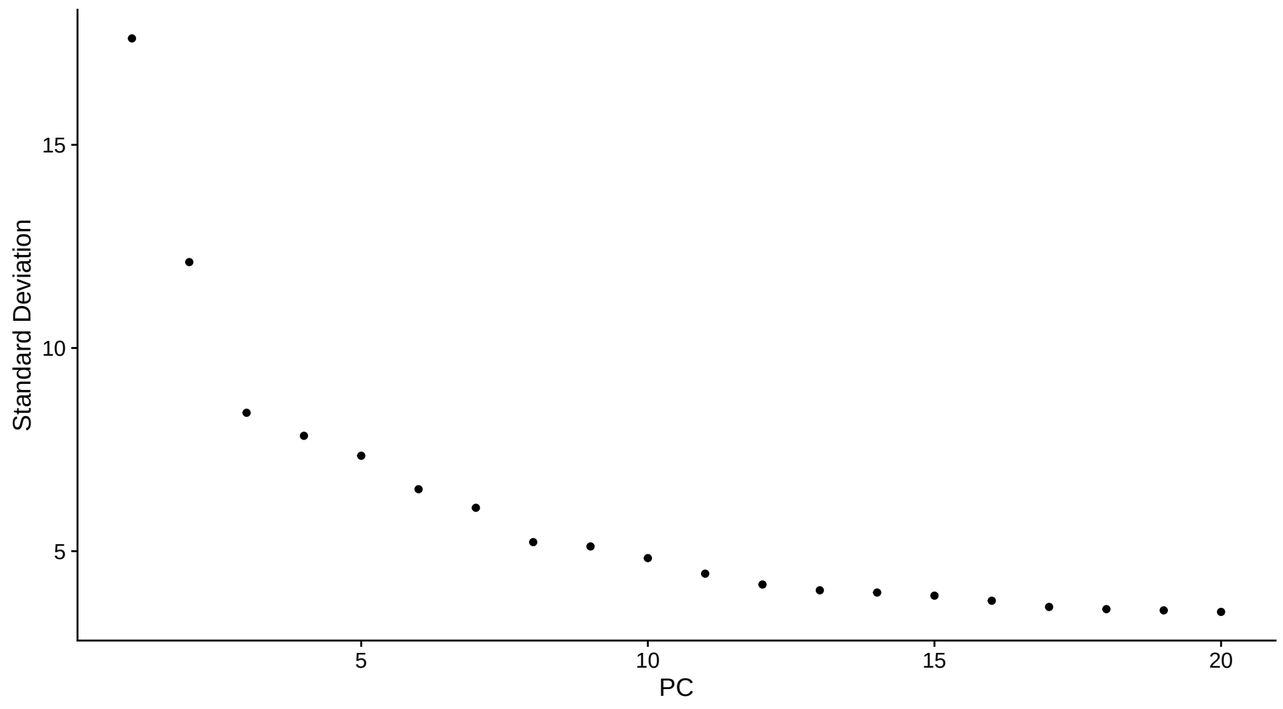
SCRIPT 2 (QC) SUMMARY (sr_Day55_FIXED)

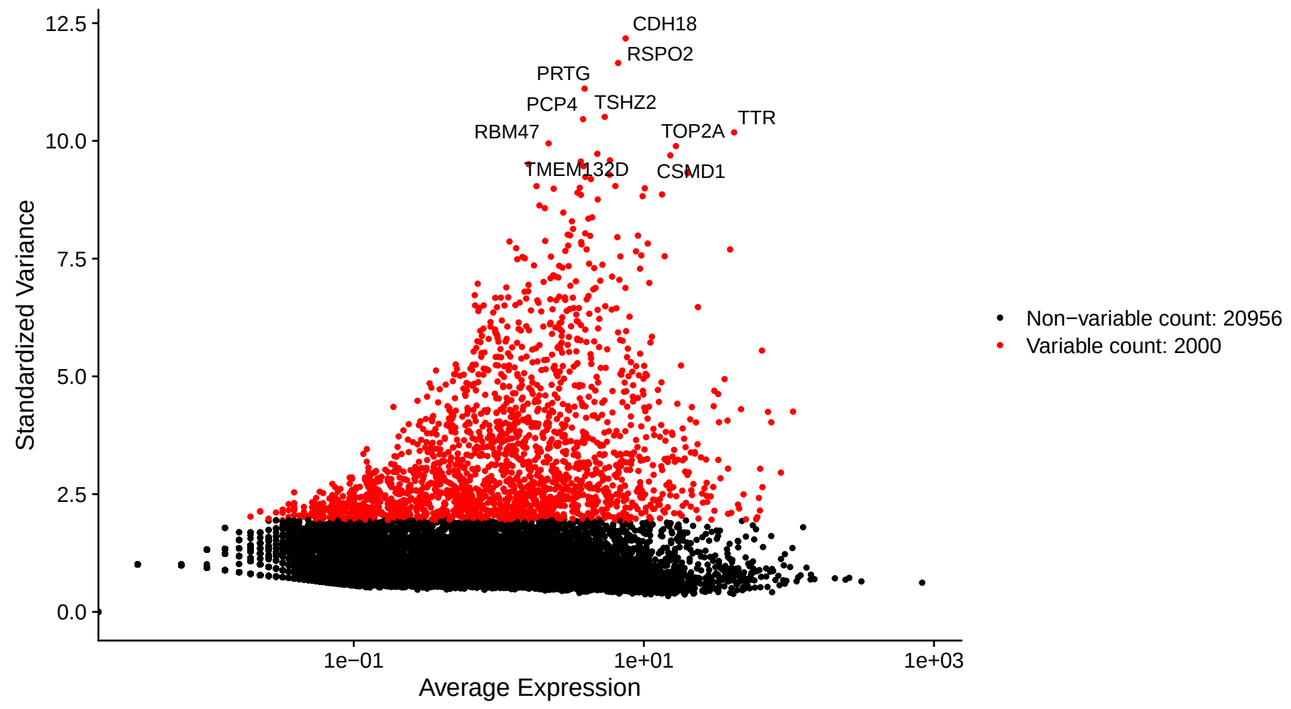
Sample ID	sr_Day55_FIXED
Initial Cell Count	505
Cells after QC	299
Feature Type	Gene
Min. Feature Threshold	3500
Max. Feature Threshold	999999999
Min. Counts Threshold	500
Max. Counts Threshold	1e+05
Median Features per Cell	7780
Median Reads per Cell	40910
Mitochondrial Content Threshold (%)	10
Median MT % before filter	3.73376623376623
Median MT % after filter	3.52587744591822
NPCs (UMAP dimensions)	14



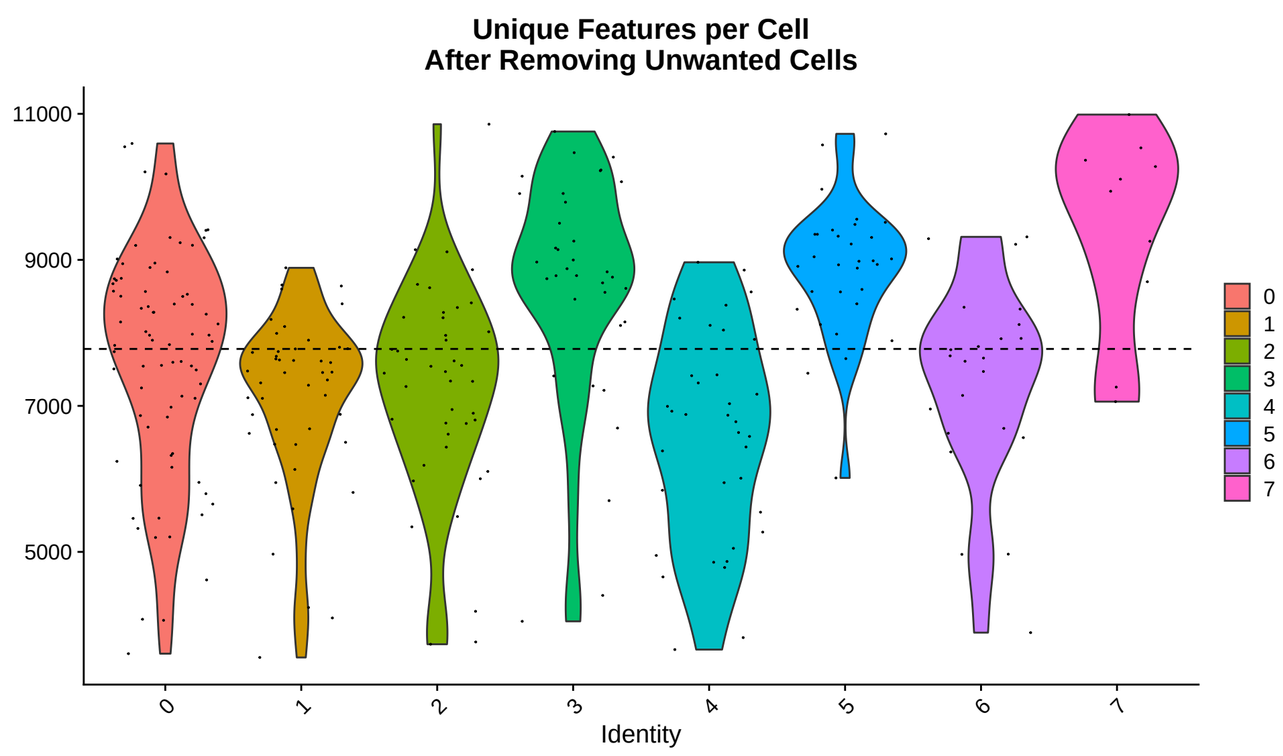
Standard Deviation explained by each Principle Component



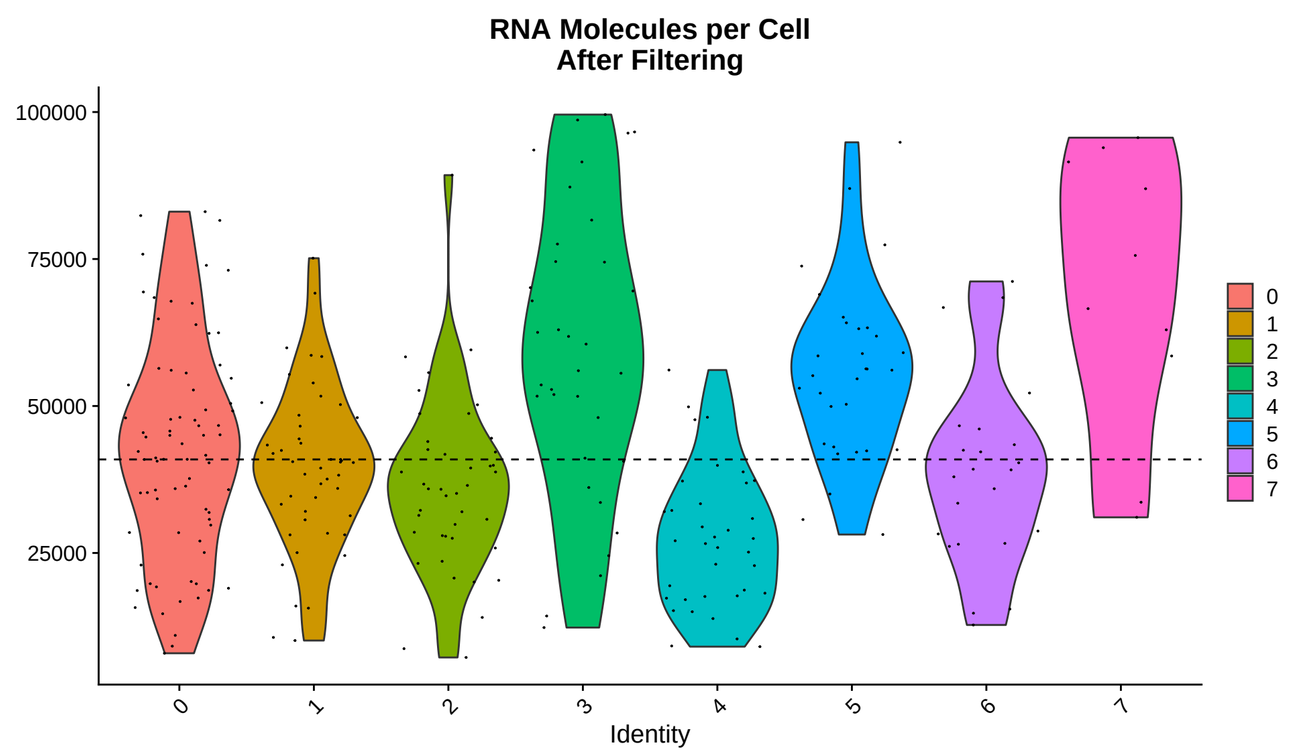
Top 2000 Variable Features across Cells



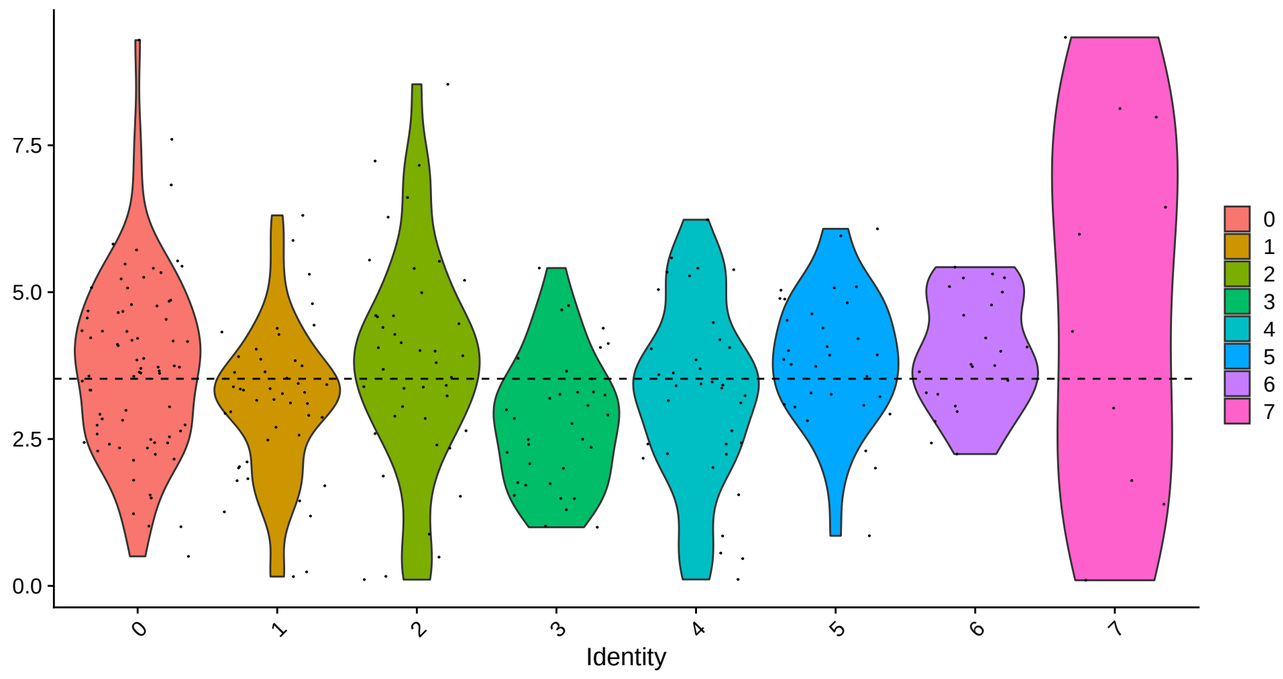
Unique Features per Cell After Removing Unwanted Cells



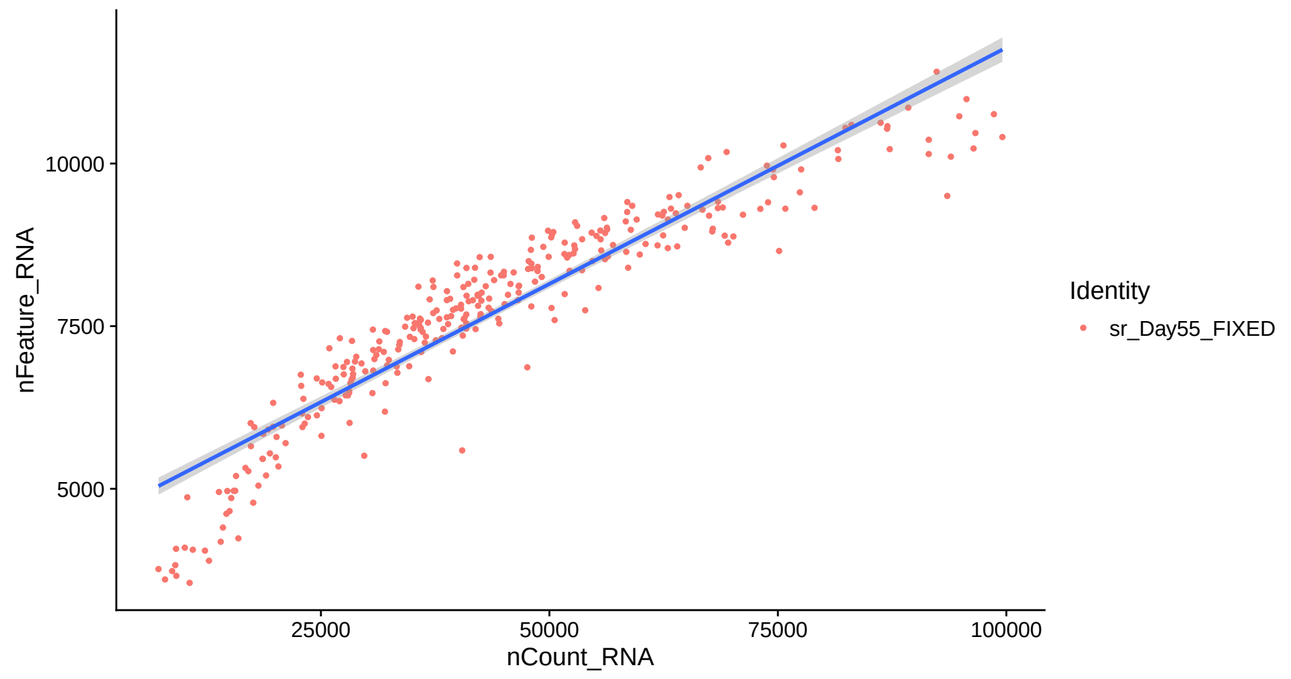
RNA Molecules per Cell After Filtering



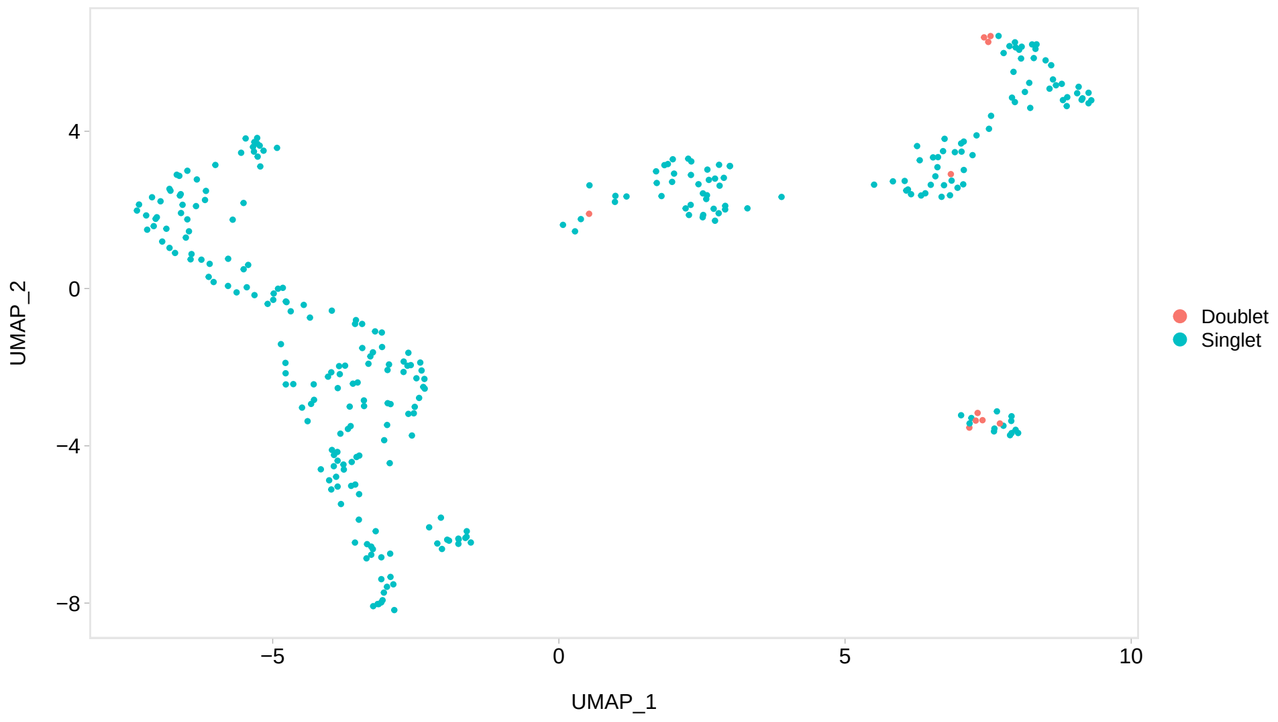
Mitochondrial Content (%) After Removing Unwanted Cells



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	58397.5	8911	10
2	Singlet	40910.0	7780	299