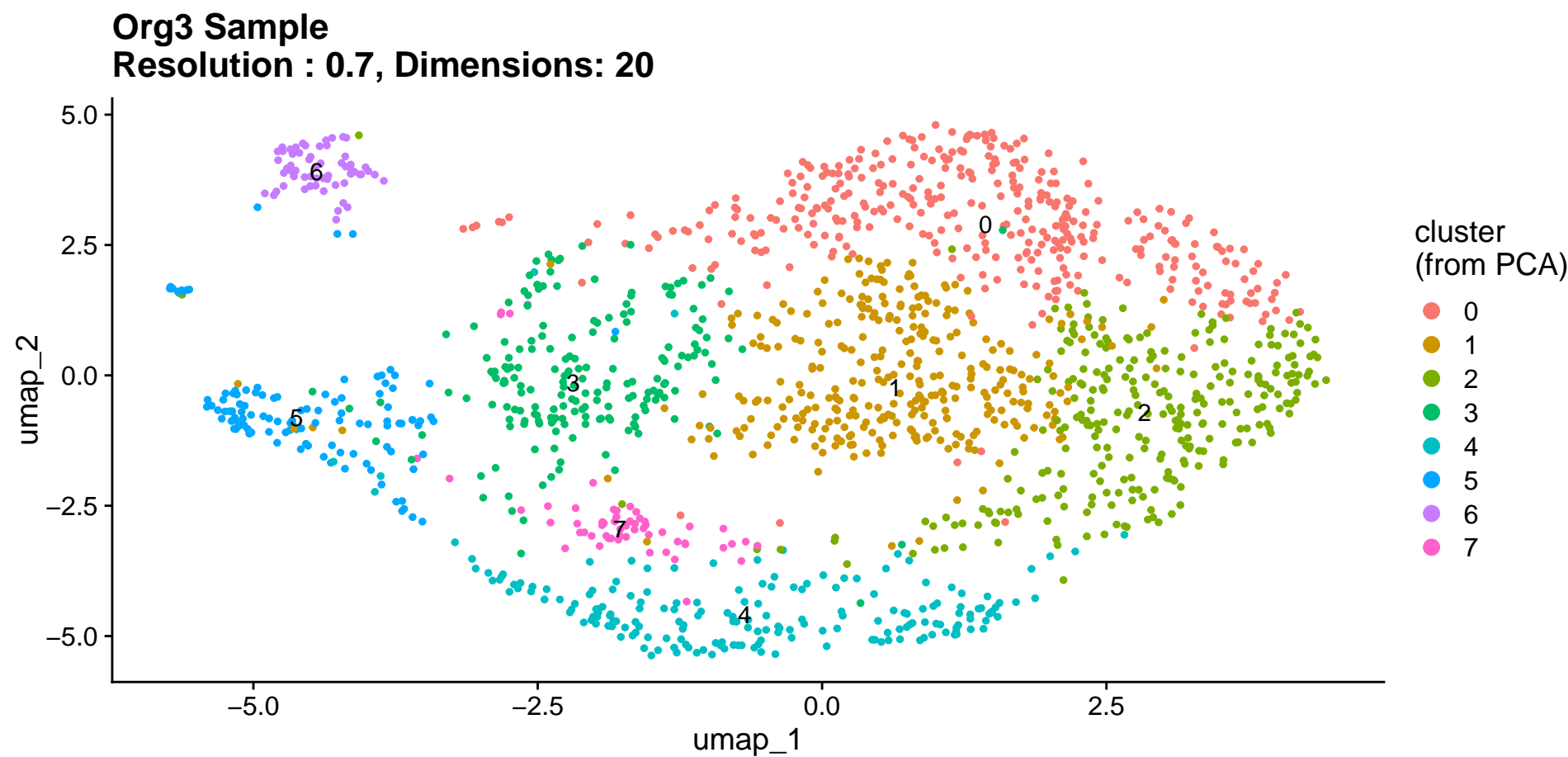
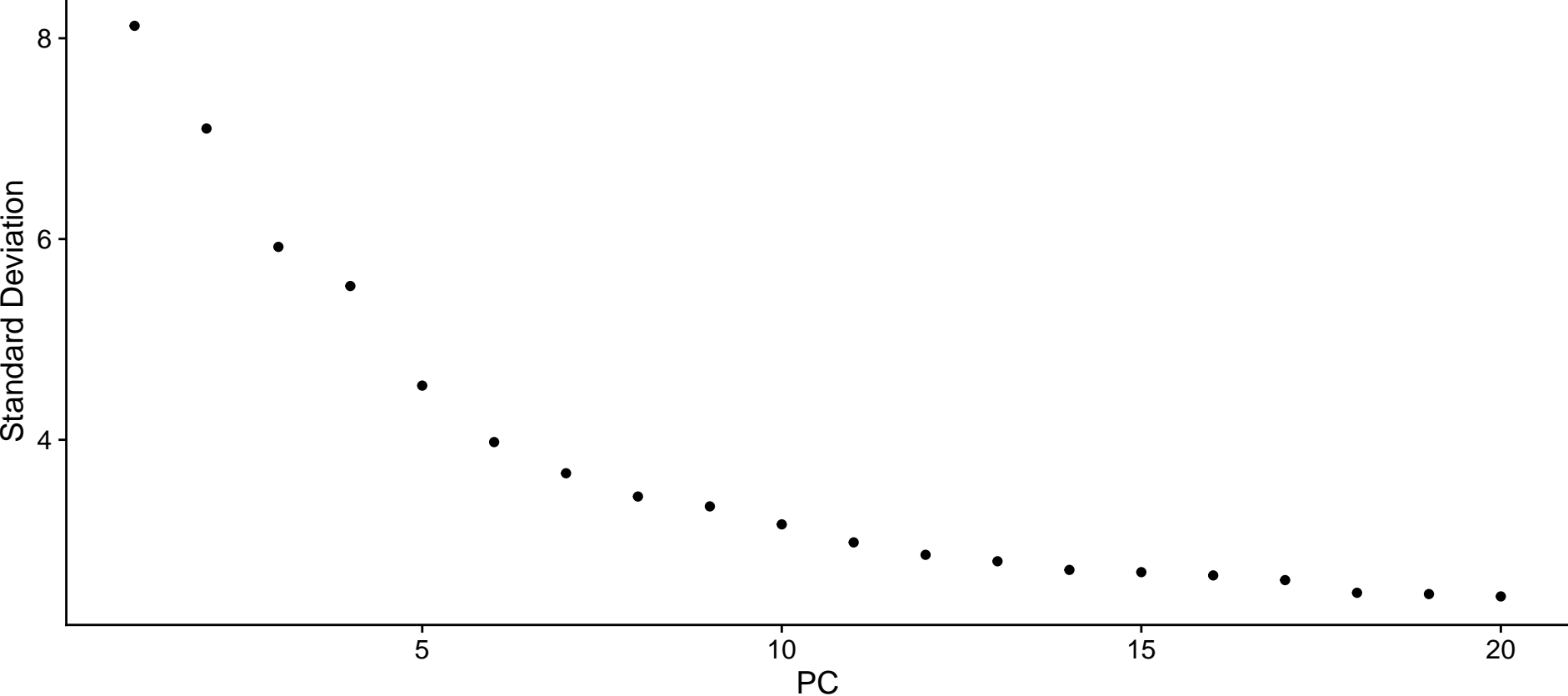


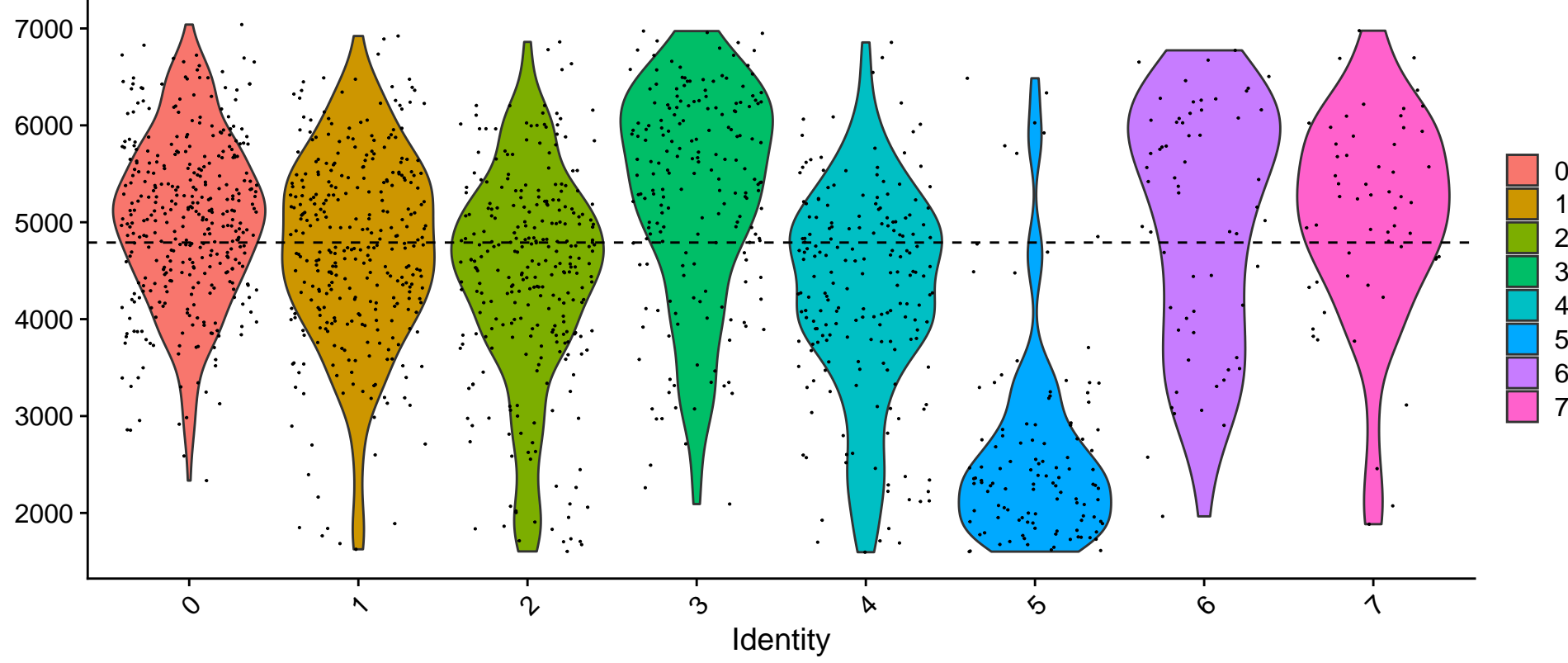
Sample ID	6M_org2
Initial Cell Num	1583
Cells after QC	1562
Min. Gene Threshold	1574
Max. Gene Threshold	7061
Min. Reads Threshold	500
Max. Reads Threshold	27000
Mitochondrial Content Threshold (%)	15
Median Genes per Cell	4790.5
Median Isoforms per Cell	3241
Median Reads per Cell	13312
Median MT % before filter	10.13
Median MT % after filter	8.71
NPCs (UMAP dimensions)	20
Cluster Resolution	0.7



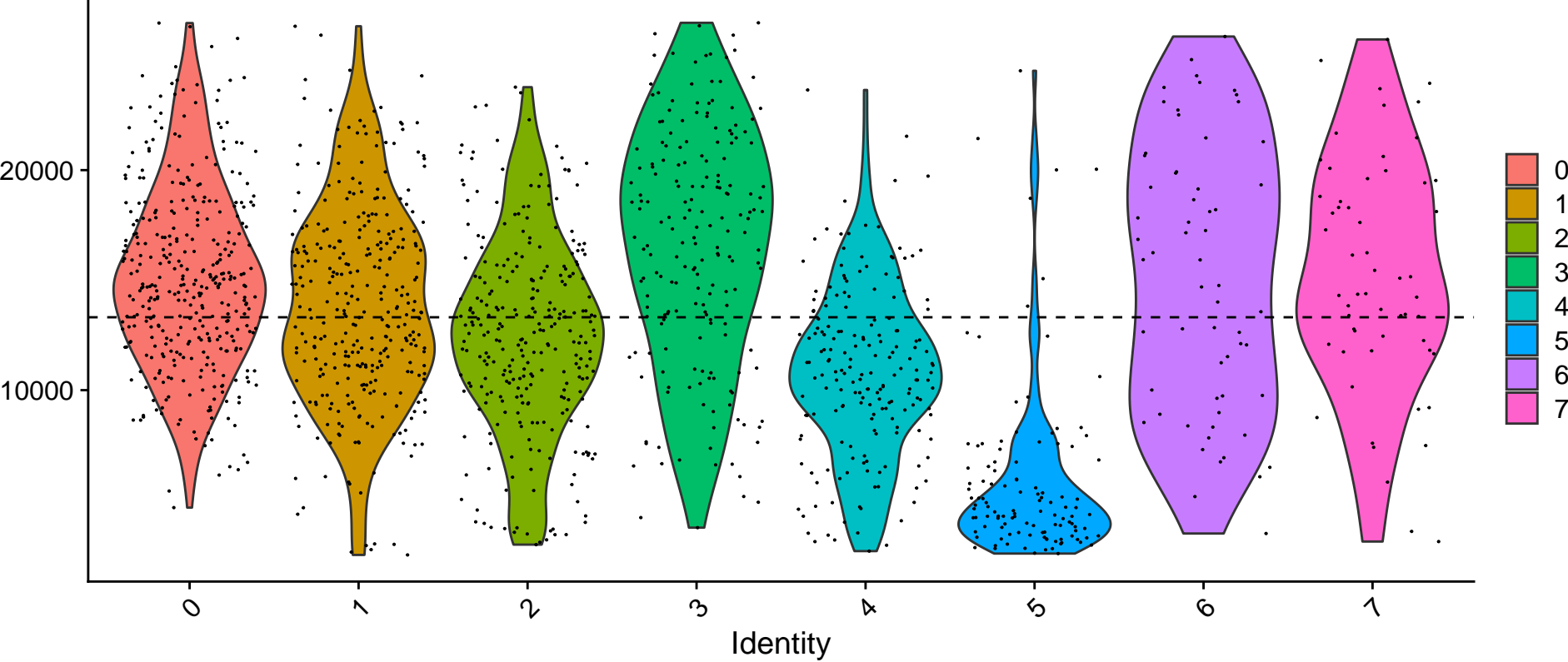
Standard Deviation explained by each Principle Component



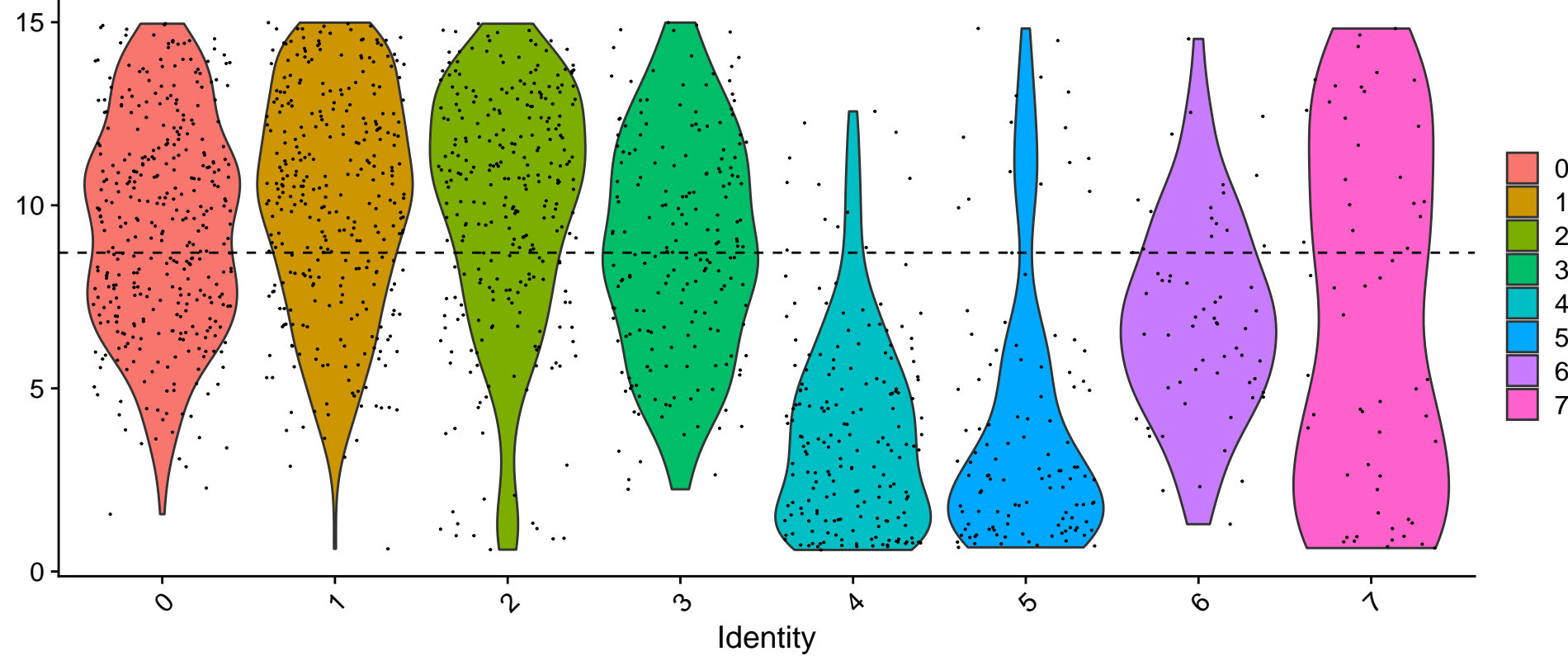
Unique Genes per Cell
After Removing Unwanted Cells



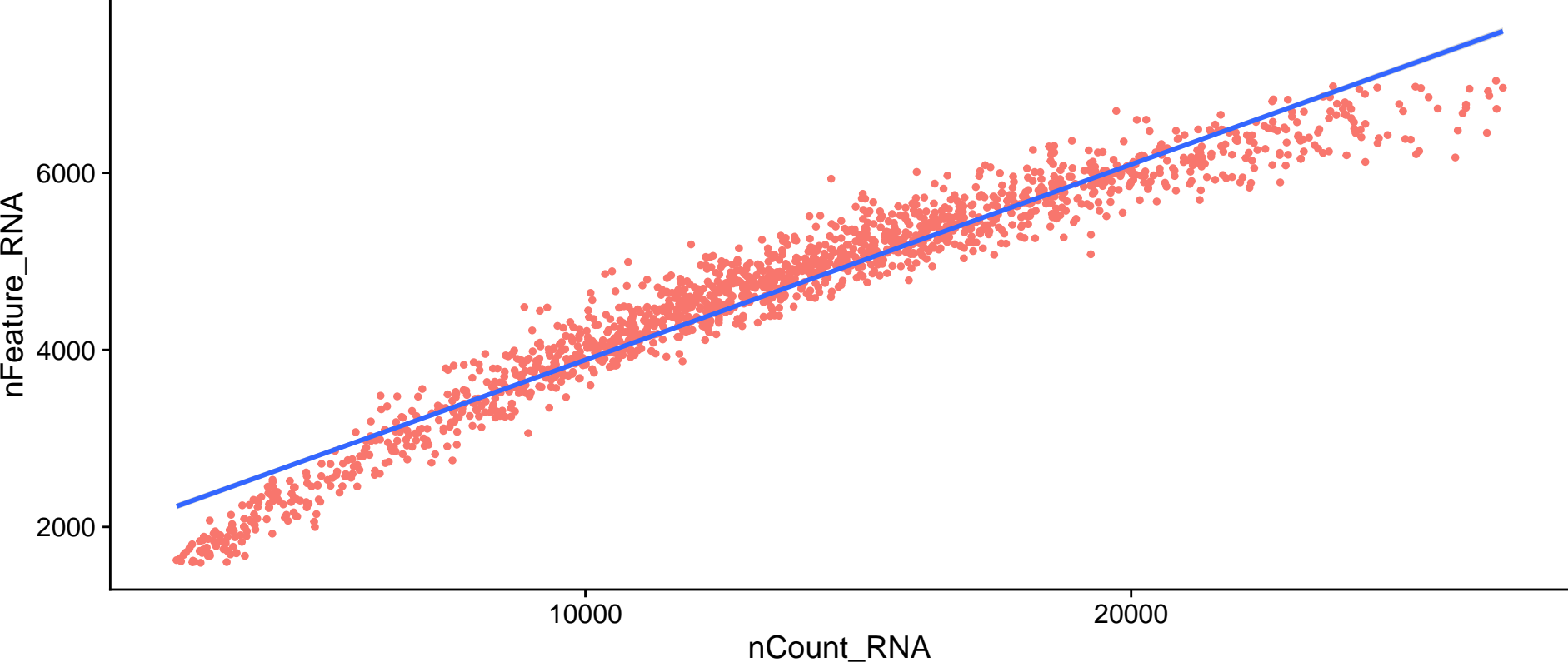
Reads per Cell
After Filtering



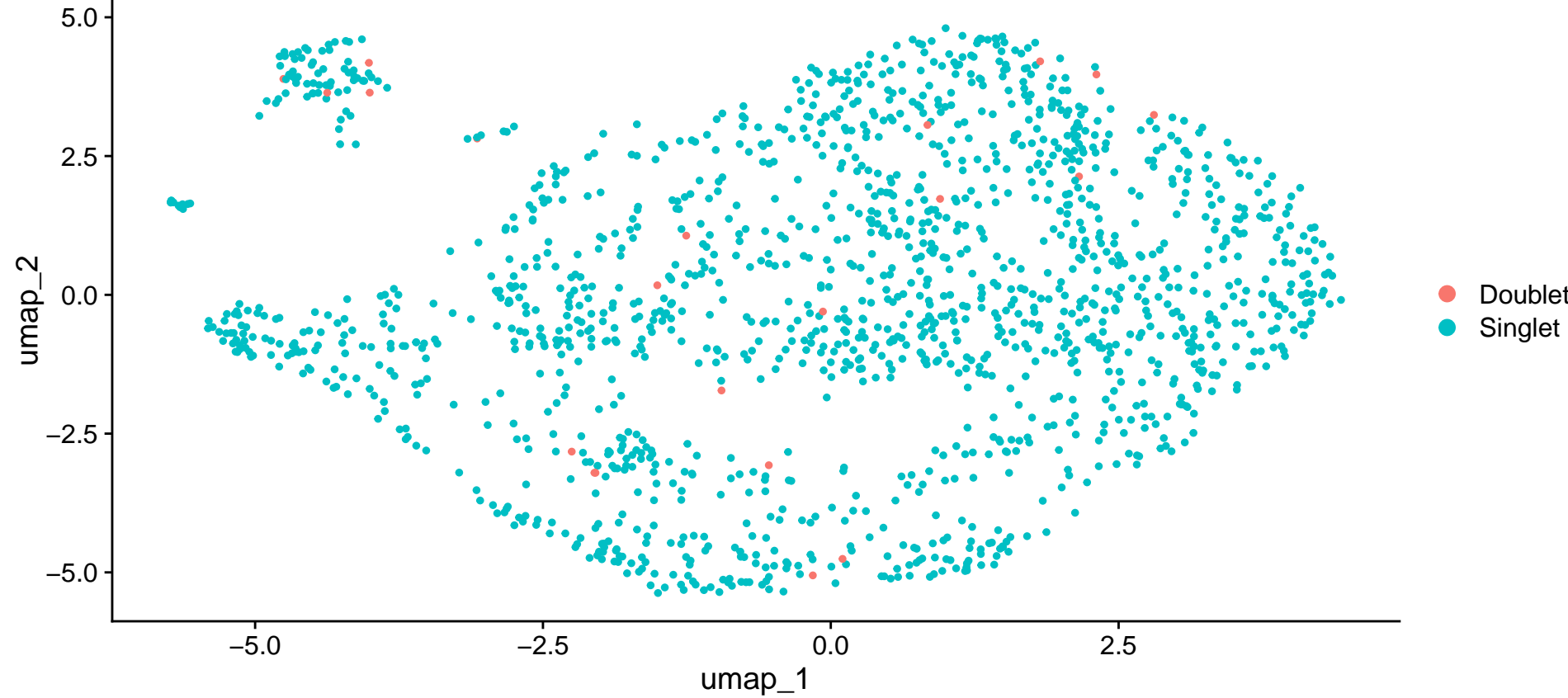
Mitochondrial Content (%)
After Removing Unwanted Cells



Association between reads
and unique genes per cell AFTER filering



Doublets that were detected and removed



	DF.classifications	median_nCount_RNA	median_nFeature_RNA	count
1	Doublet	18756	6020.0	21
2	Singlet	13312	4790.5	1562

