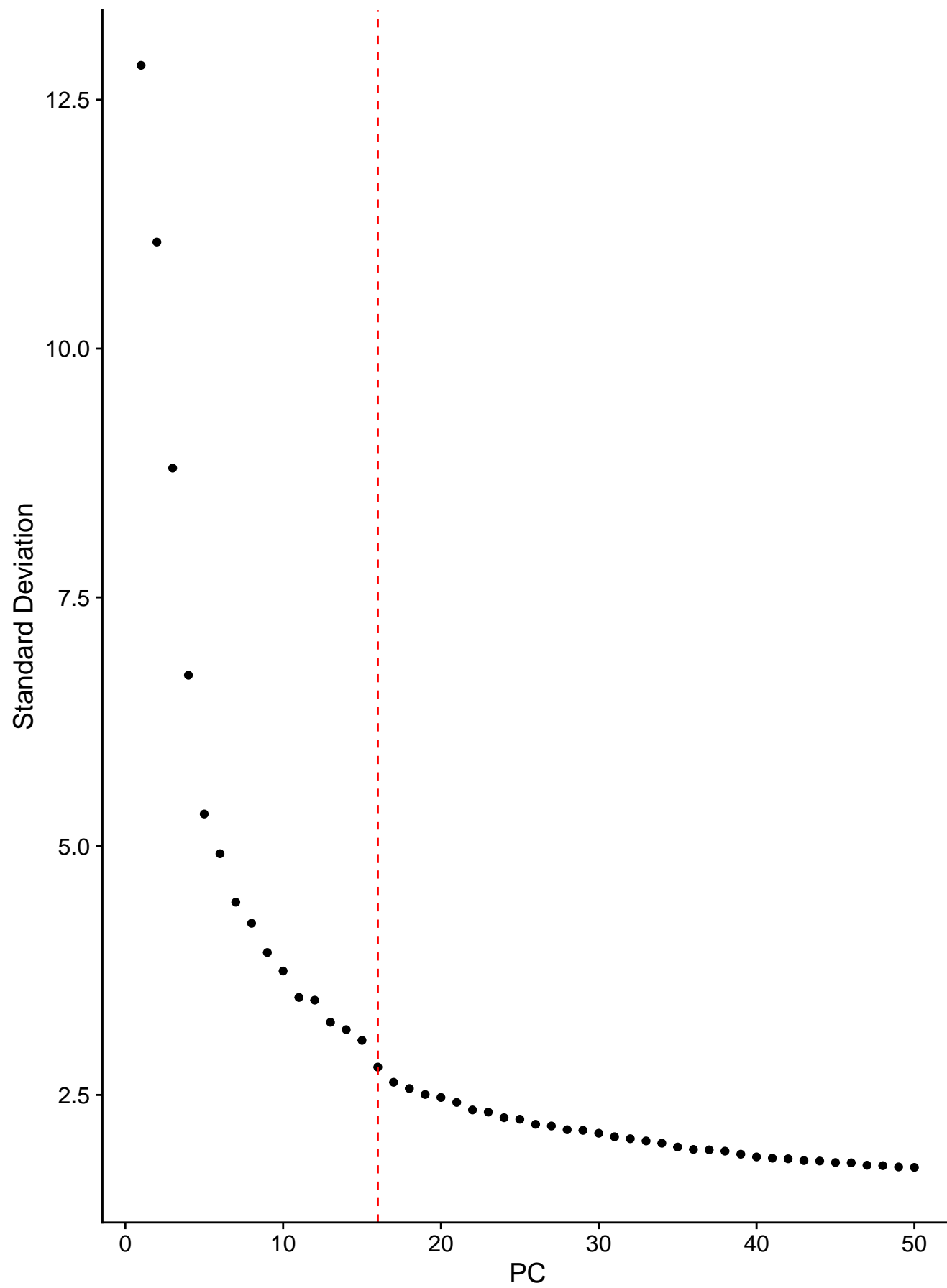
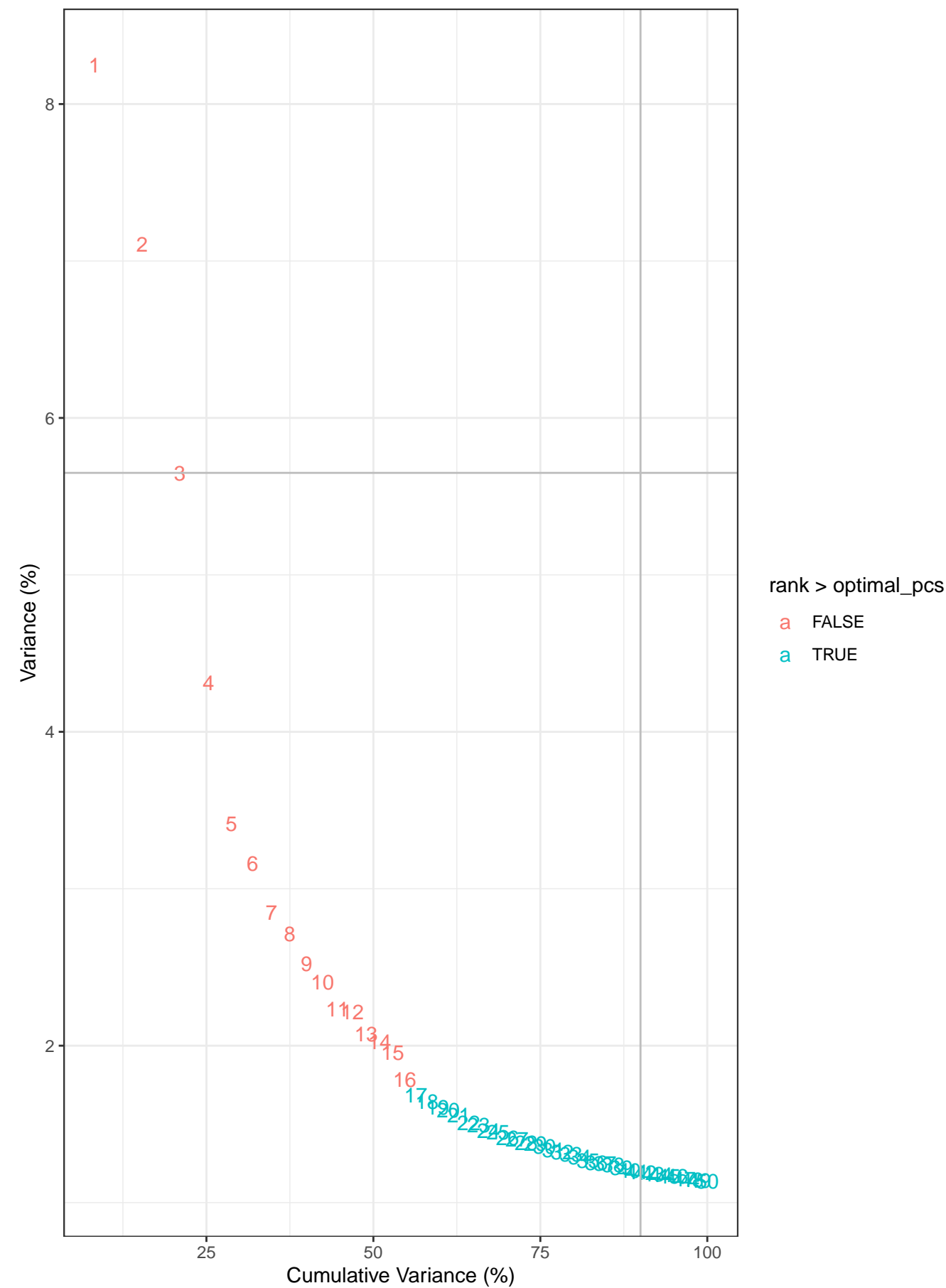


6M_Org – PCA Elbow Analysis (Pre-Integration)

6M_Org: Elbow Plot (Optimal PCs = 16)

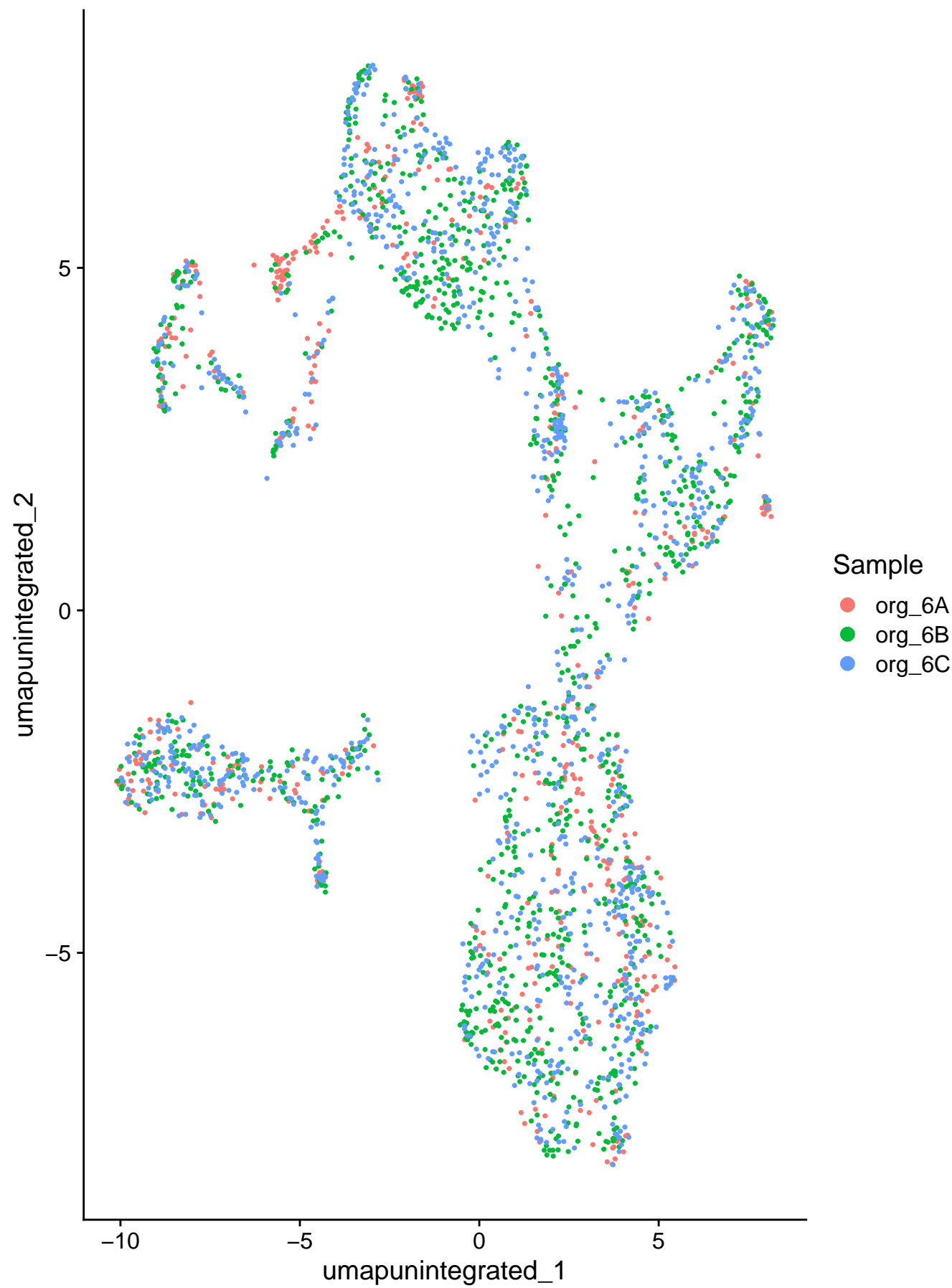


6M_Org: Quantitative PC Selection

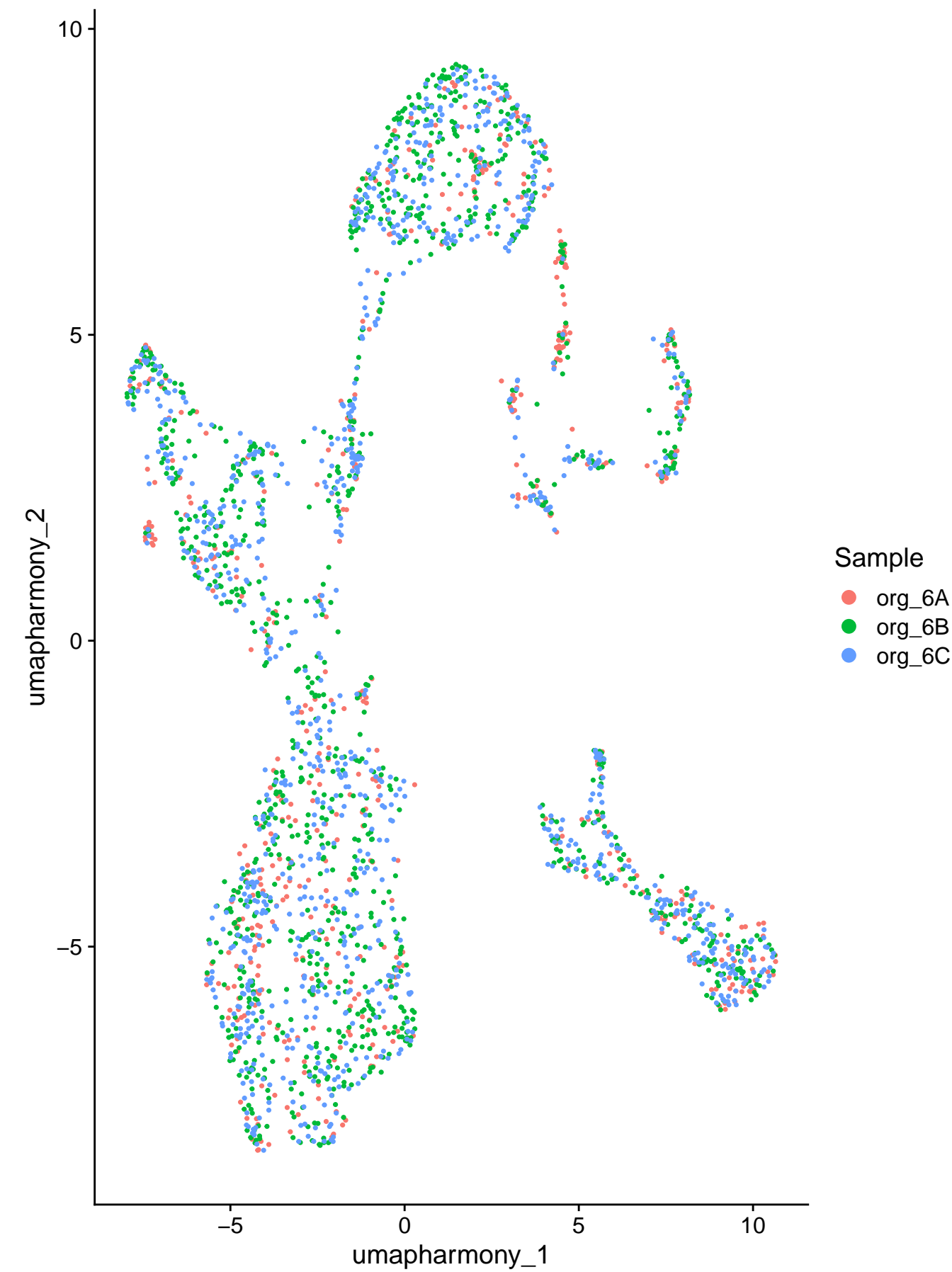


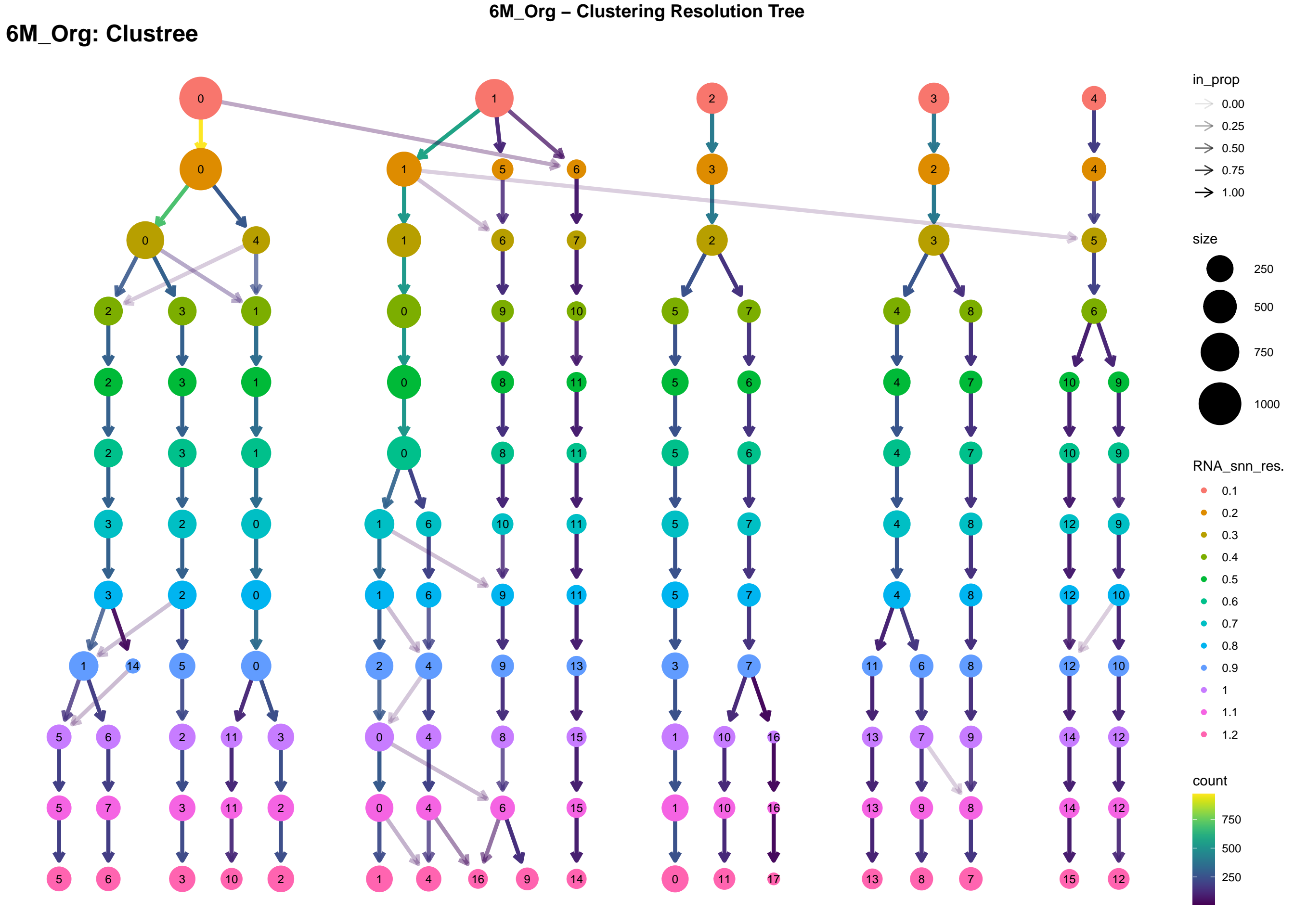
6M_Org – Harmony Integration Comparison

6M_Org: Before Integration



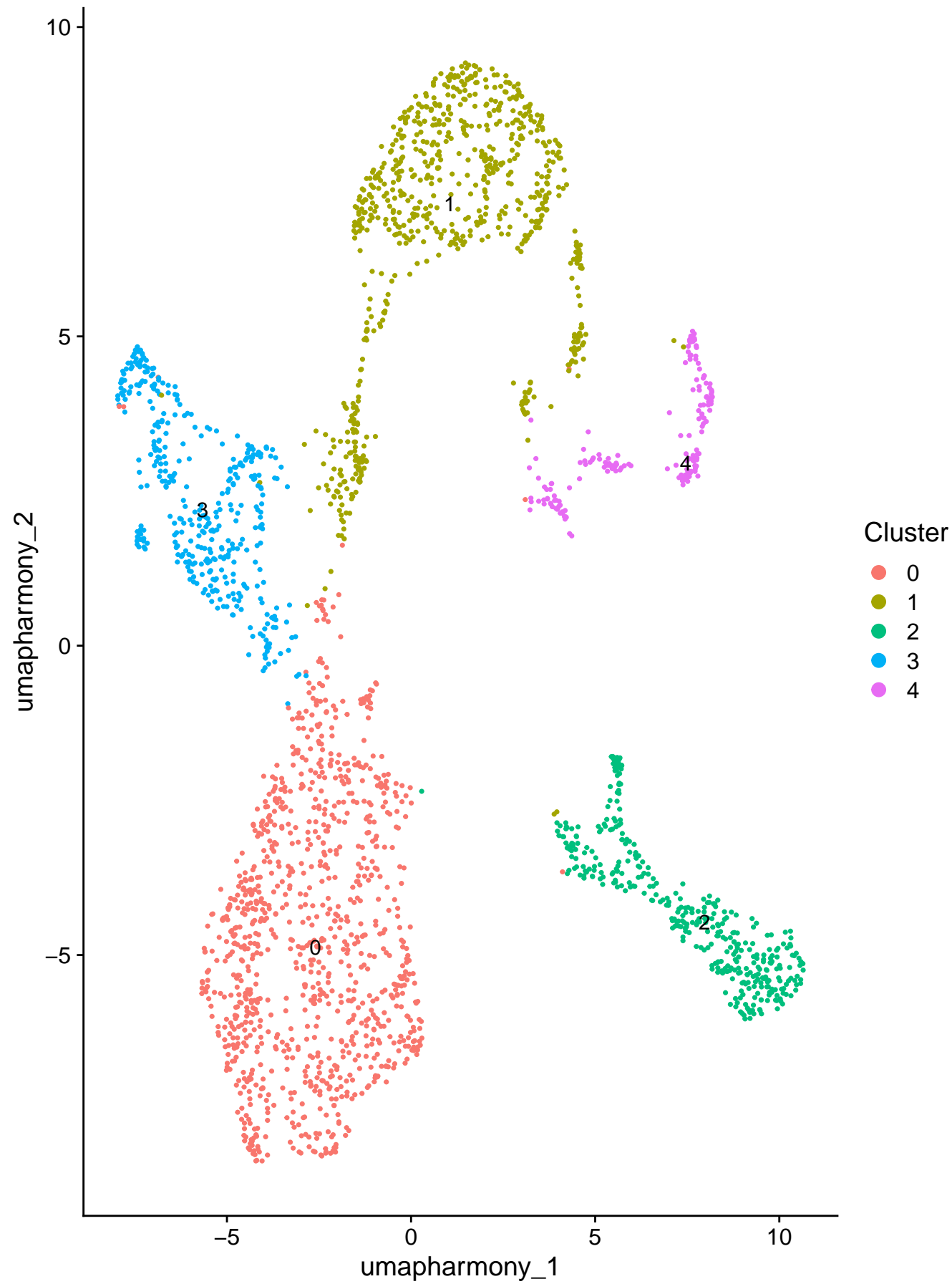
6M_Org: After Harmony Integration





6M_Org – Final Clustering Results

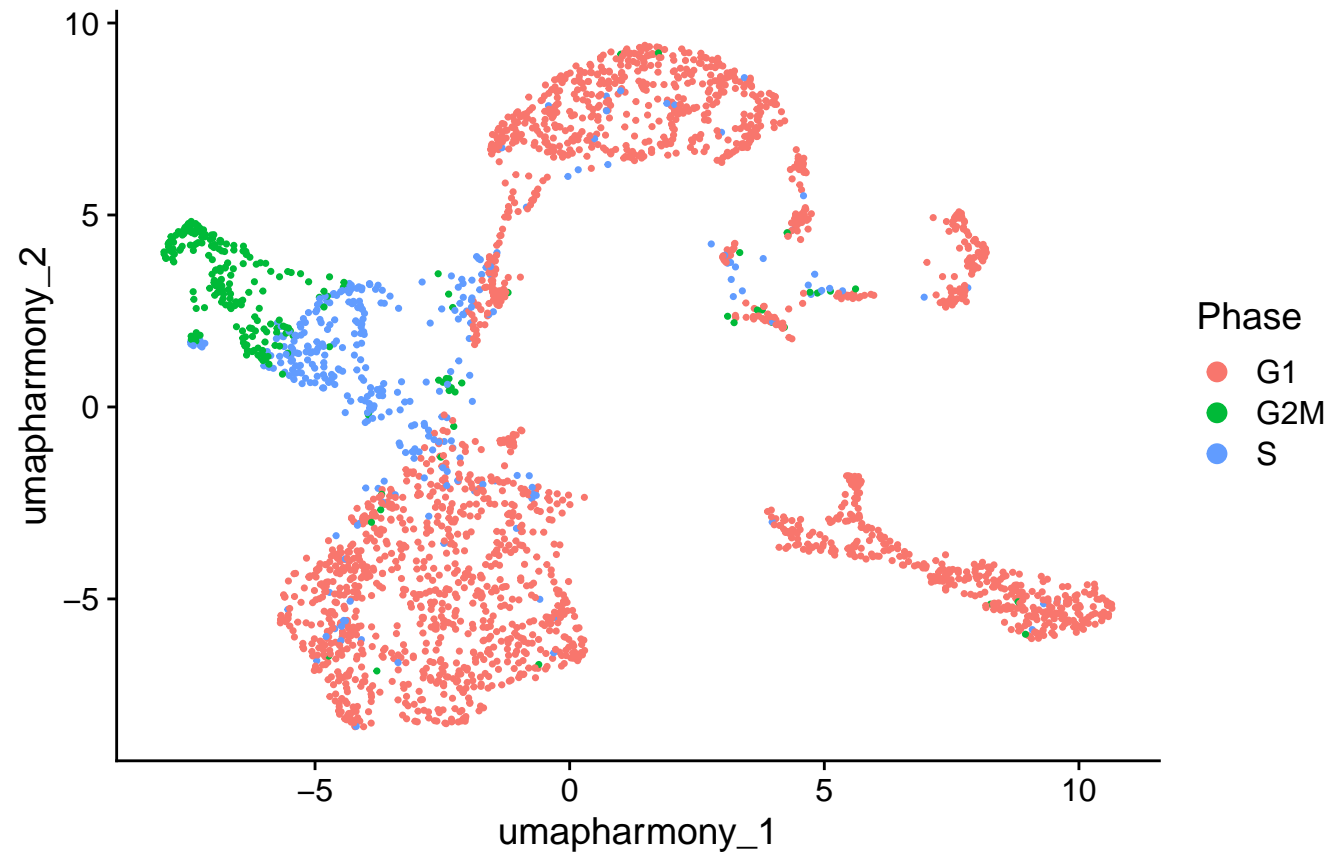
6M_Org: Final Clusters



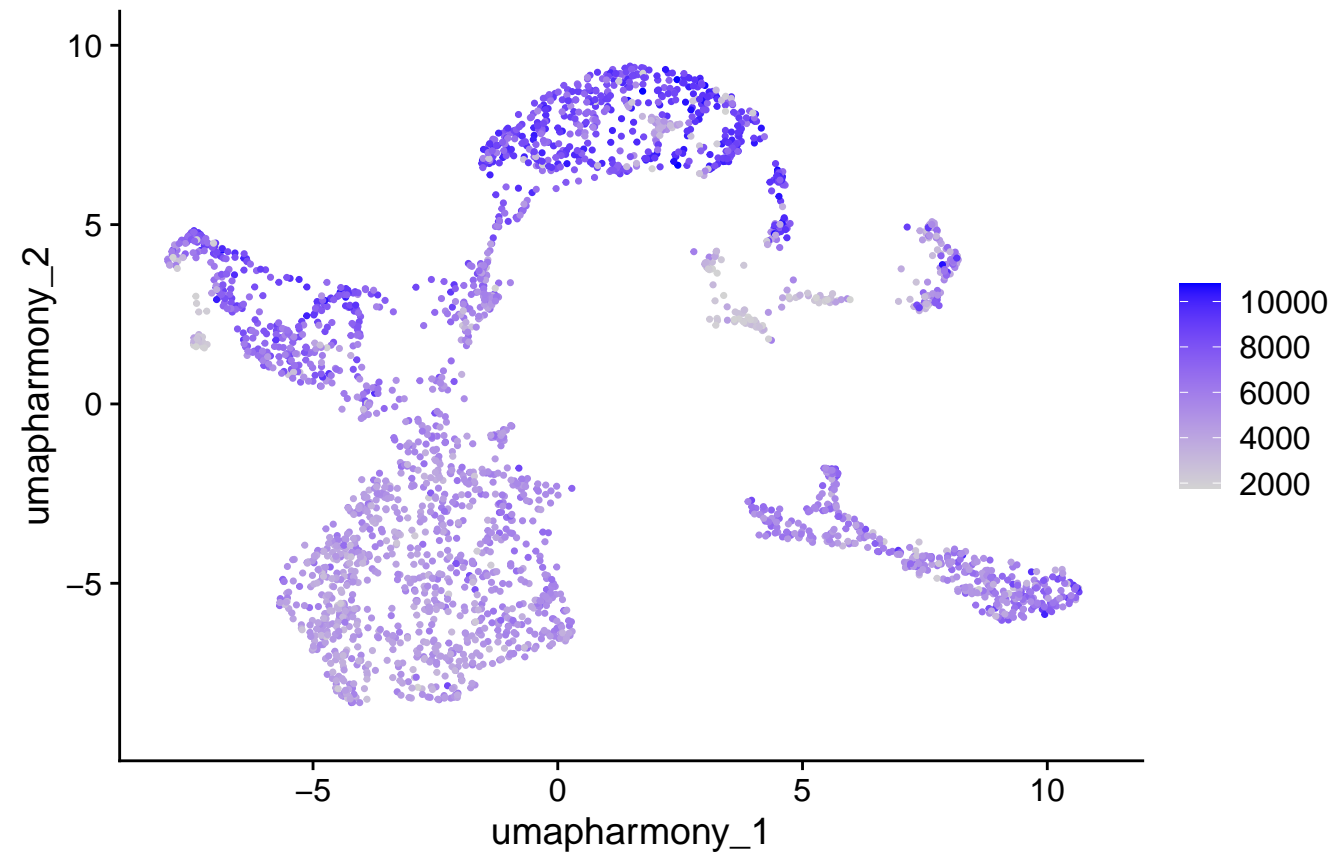
	res_vals	num_clusters	avg_sil_vals
1	0.1	5	0.384888775406927
2	0.2	7	0.375614626505589
3	0.6	12	0.260421633650227
4	0.3	8	0.253590903352673
5	0.4	11	0.230426171738615
6	0.8	13	0.226061621629569
7	0.9	15	0.218182293815375
8	1.2	18	0.192964408929519
9	1	17	0.192537432156566

6M_Org – QC Metrics on Integrated Data

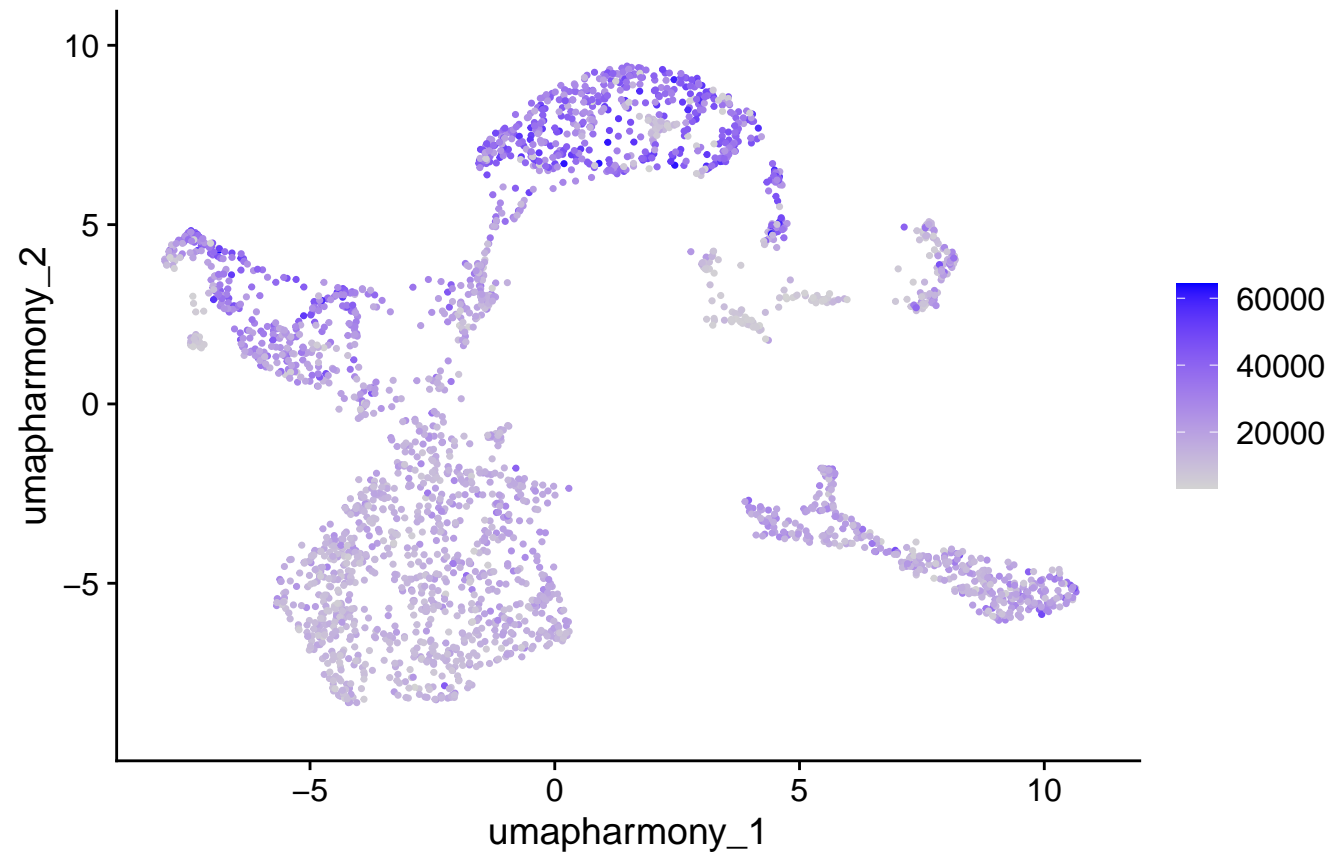
6M_Org: Cell Cycle Phase



6M_Org: nFeature_RNA



6M_Org: nCount_RNA



6M_Org: Final Clusters

