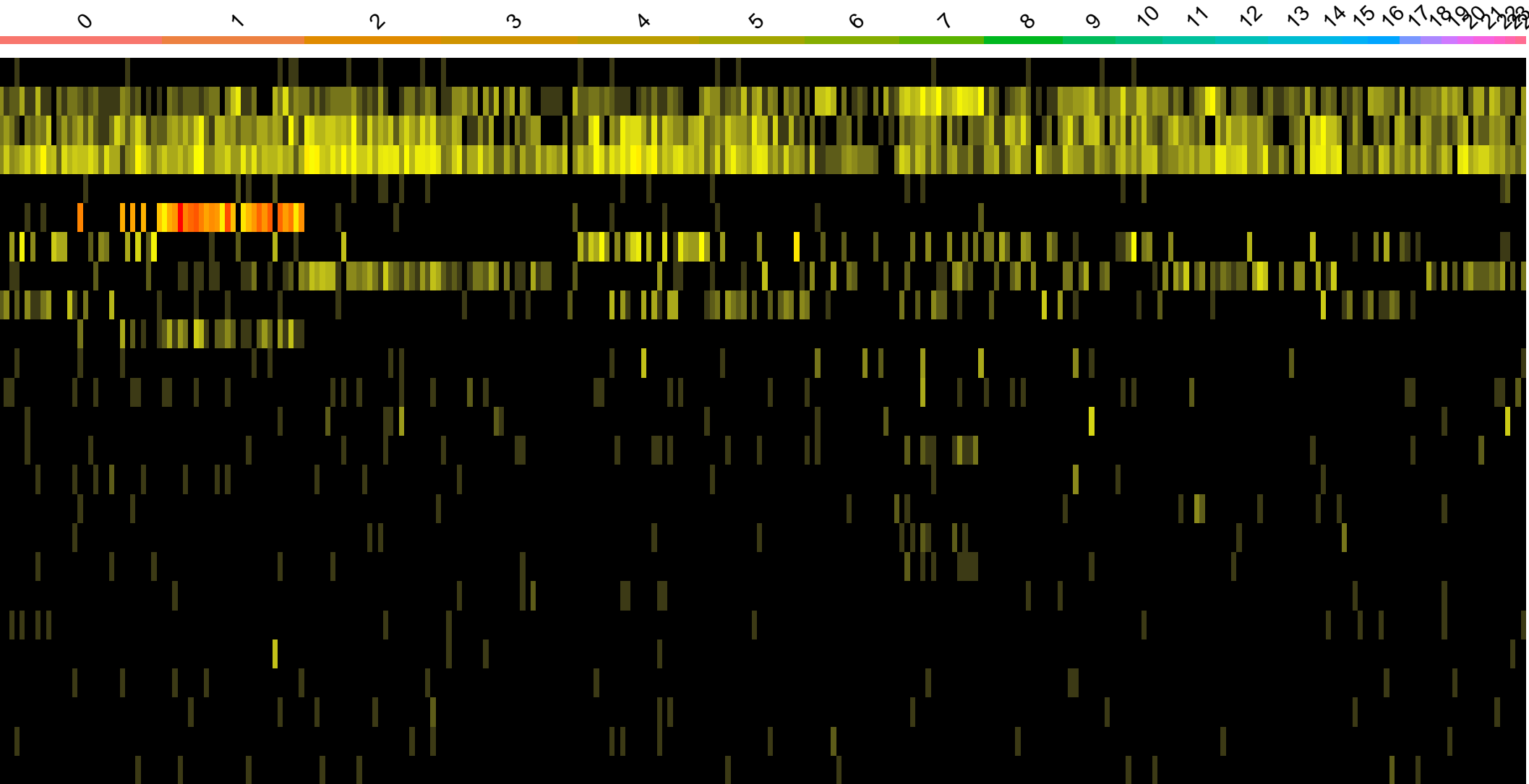


Cluster 0, Batch Correction: harmony_rmclusts_res4
290 out of 5175 total neurons, reclustered (USFC)
Gene List: Total Average Cluster Expression, Top 20



Cluster 0, Batch Correction: harmony_rmclusts_res4
290 out of 5175 total neurons, reclustered (USFC)
Gene List: Sub Cluster Average Expression, Top 5

