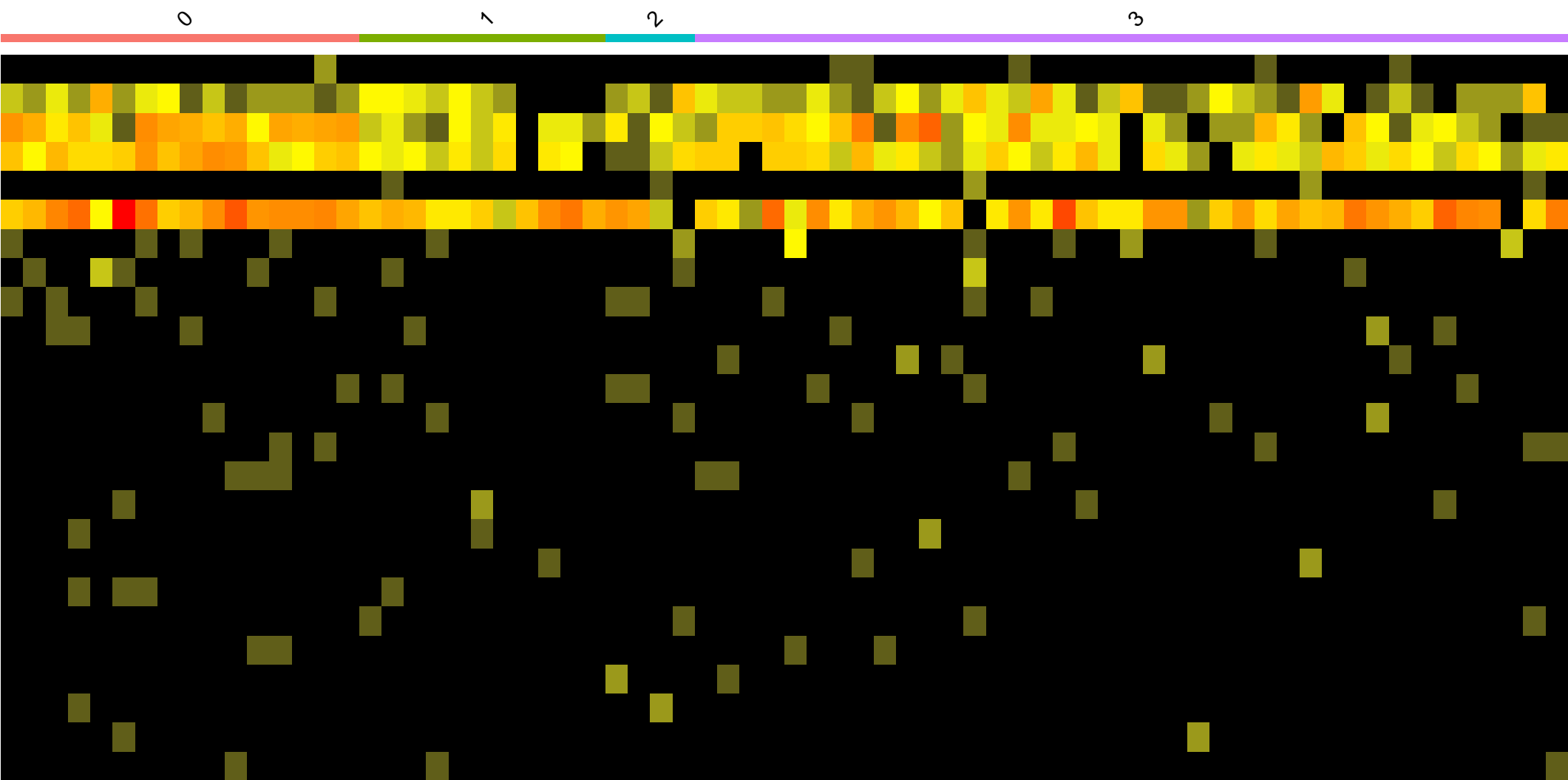**Cluster 30, Batch Correction: harmony_rmclusts_res4**
70 out of 5175 total neurons, reclustered (UFC)
Gene List: Total Average Cluster Expression, Top 20

Cluster 30, Batch Correction: harmony_rmclusts_res4
70 out of 5175 total neurons, reclustered (UFC)
Gene List: Sub Cluster Average Expression, Top 5