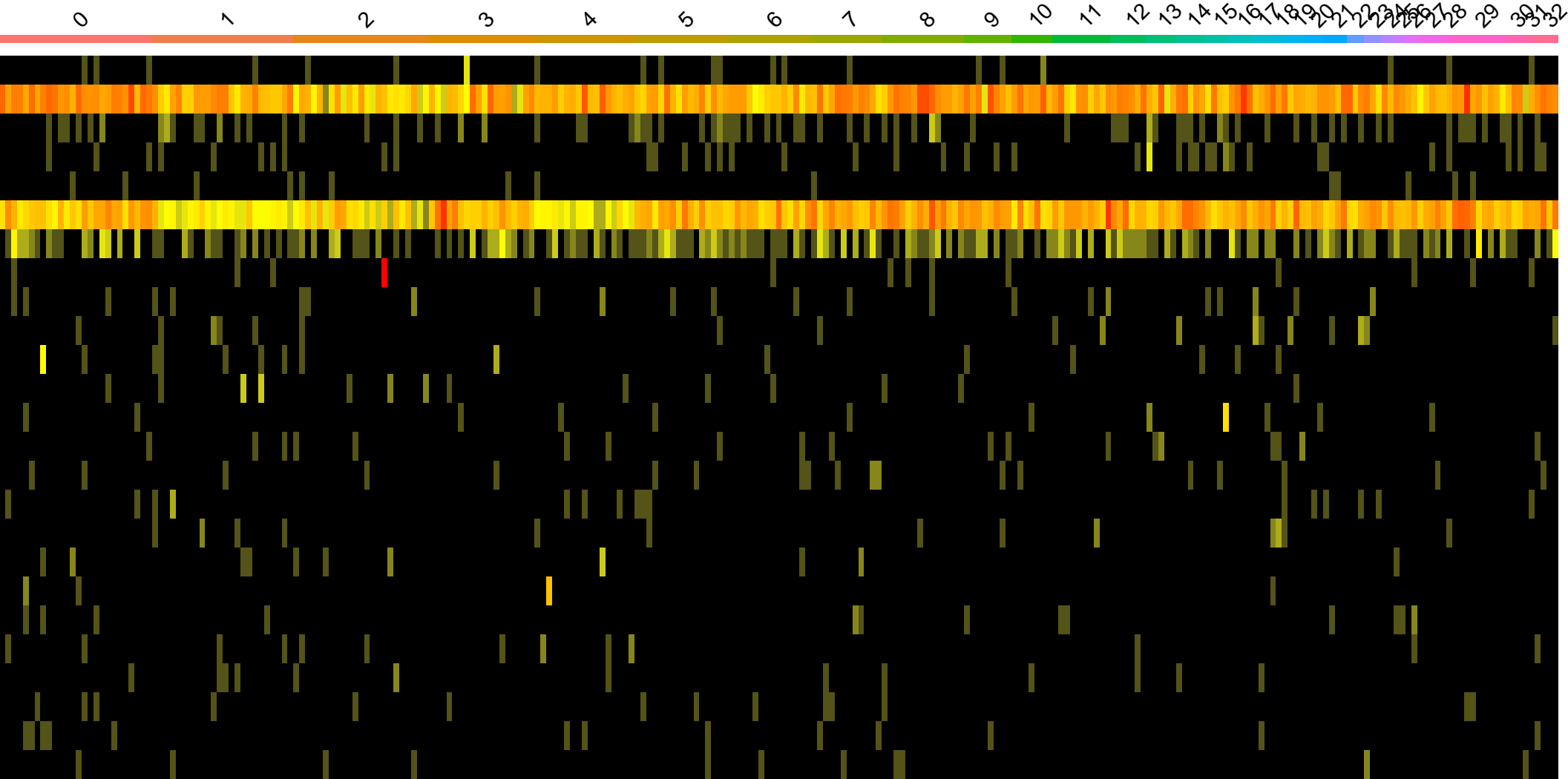


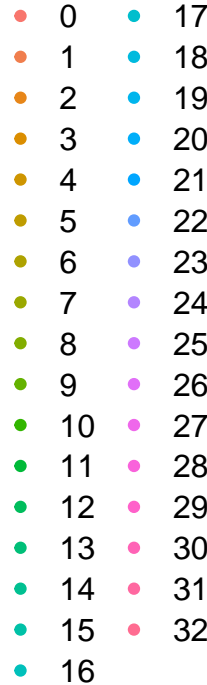
Cluster 1, Batch Correction: harmony_rmclusters_res4
265 out of 5175 total neurons, reclustered (USFC)
Gene List: Total Average Cluster Expression, Top 20



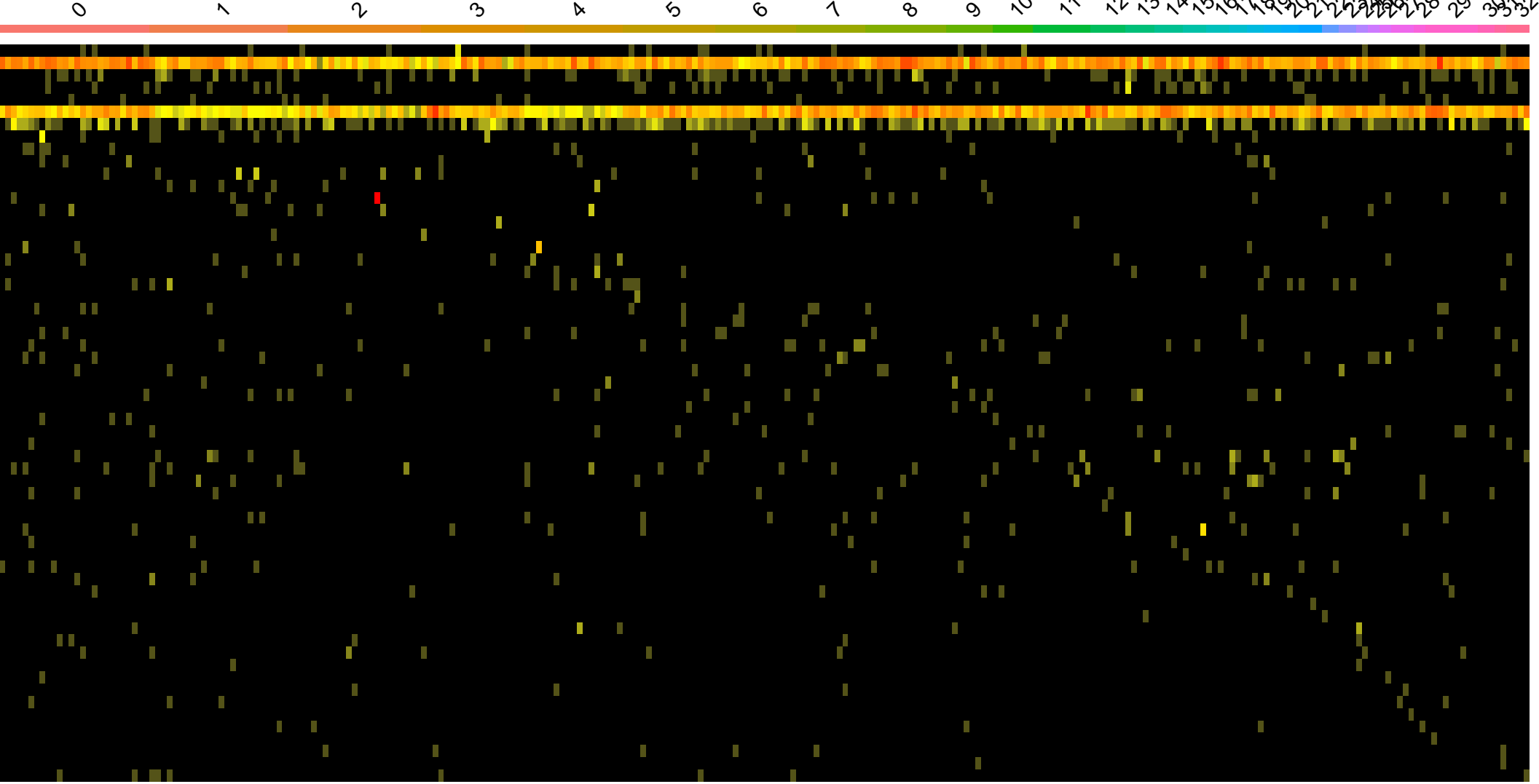
Expression



Identity



Cluster 1, Batch Correction: harmony_rmclusters_res4
265 out of 5175 total neurons, reclustered (USFC)
Gene List: Sub Cluster Average Expression, Top 5



Expression



Identity

