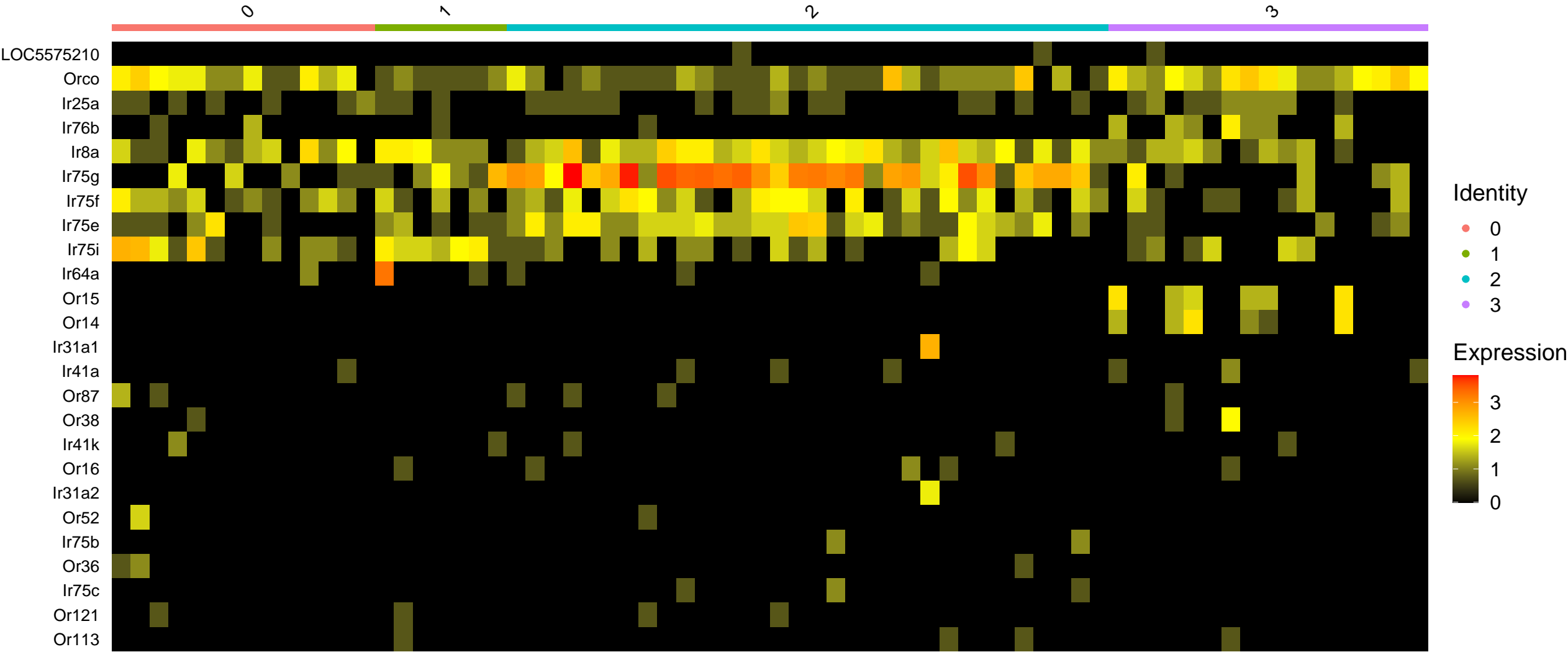


Cluster 29, Batch Correction: harmony\_rmclusters\_res4  
70 out of 5175 total neurons, reclustered (UFC)  
Gene List: Total Average Cluster Expression, Top 20



Cluster 29, Batch Correction: harmony\_rmclusters\_res4  
70 out of 5175 total neurons, reclustered (UFC)  
Gene List: Sub Cluster Average Expression, Top 5

