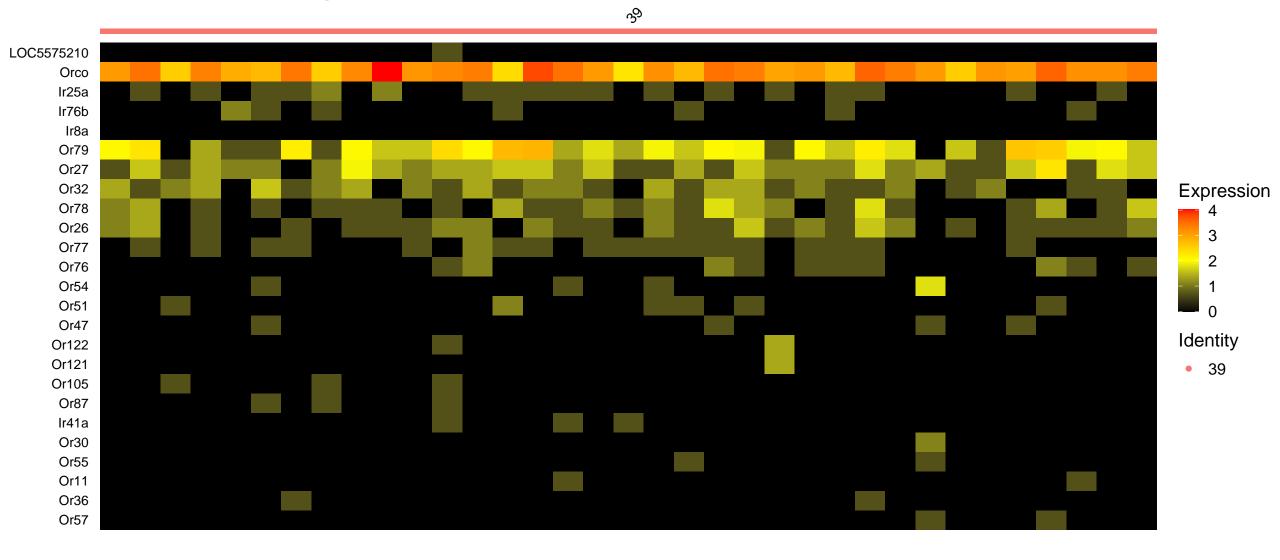
Cluster 39, Batch Correction: harmony_rmclusts_res4 35 out of 5175 total neurons, reclustered (UF) Gene List: Total Average Cluster Expression, Top 20



Cluster 39, Batch Correction: harmony_rmclusts_res4 35 out of 5175 total neurons, reclustered (UF) Gene List: Sub Cluster Average Expression, Top 5

