

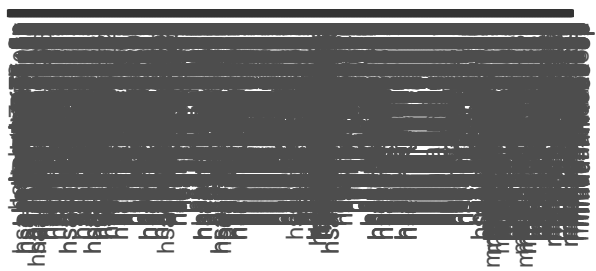
Number of genes overlap

150 -  
100 -  
50 -  
0

Adjusted.P.value



1



Module names

based on data from enrichR