

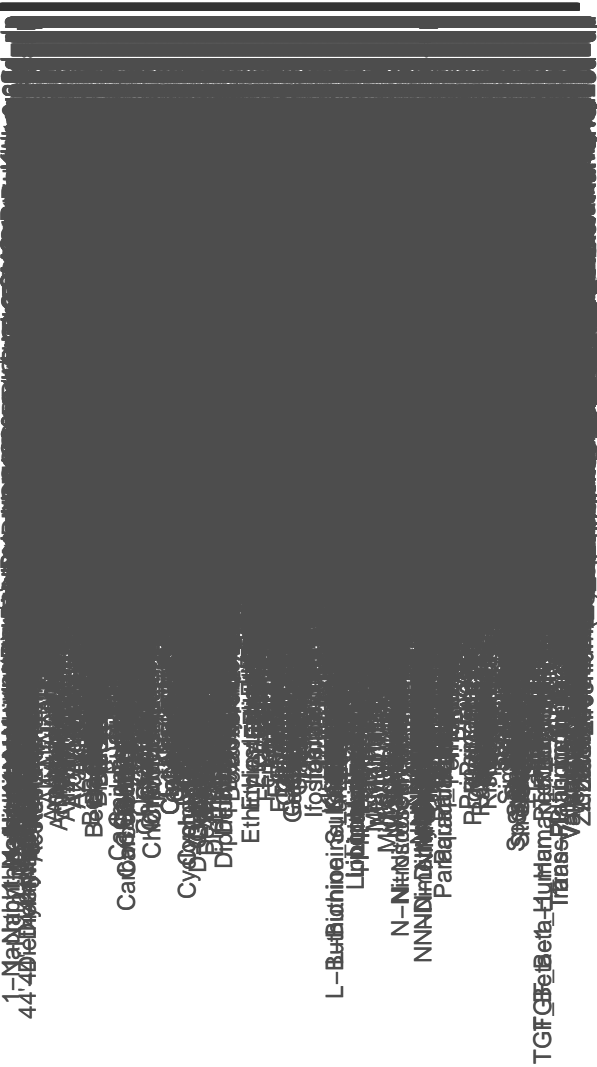
Number of genes overlap

30 -
20
10
0

Adjusted.P.value



1



Module names

based on data from enrichR