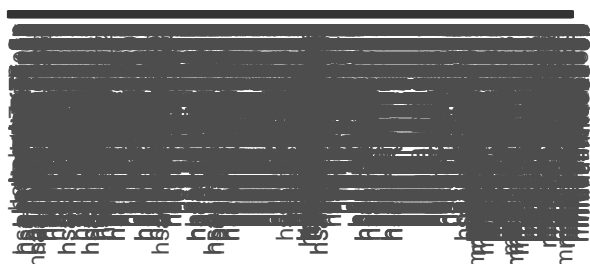
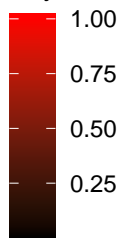


Number of genes overlap

400 -
300 -
200 -
100 -
0

Adjusted.P.value



Module names

based on data from enrichR