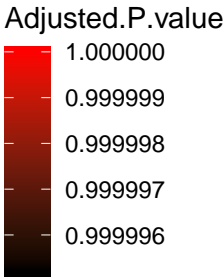


Number of genes overlap

150 -
100 -
50 -
0 -



Module names

based on data from enrichR