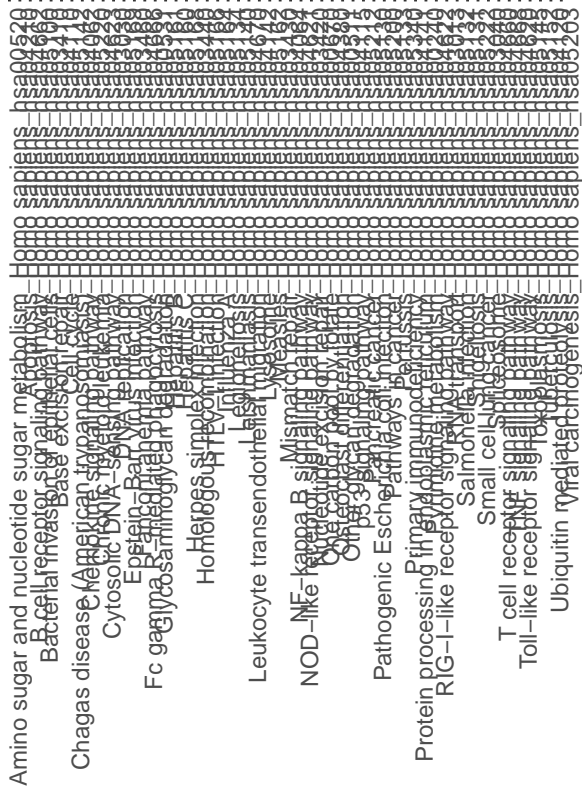
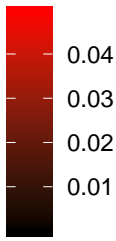


Number of genes overlap

100
75
50
25
0

Adjusted.P.value



Module names

based on enrichR database