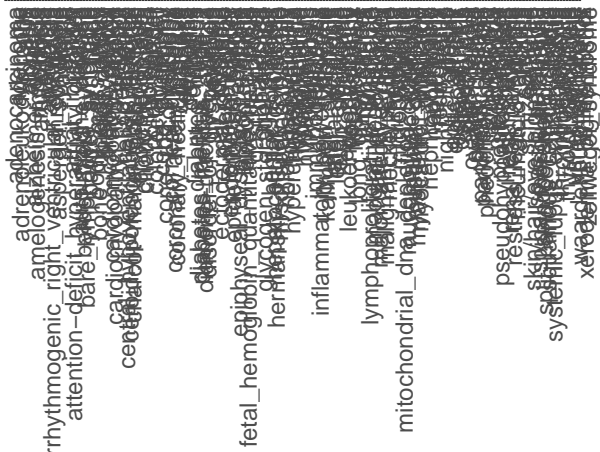
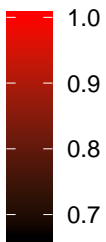


Number of genes overlap

10  
5  
0

Adjusted.P.value



Module names  
based on data from enrichR