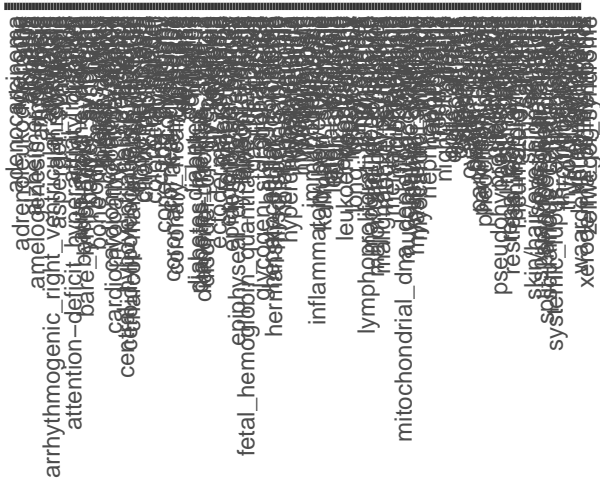
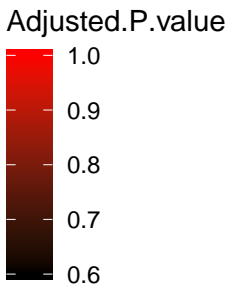


Number of genes overlap

30  
20  
10  
0



Module names

based on data from enrichR