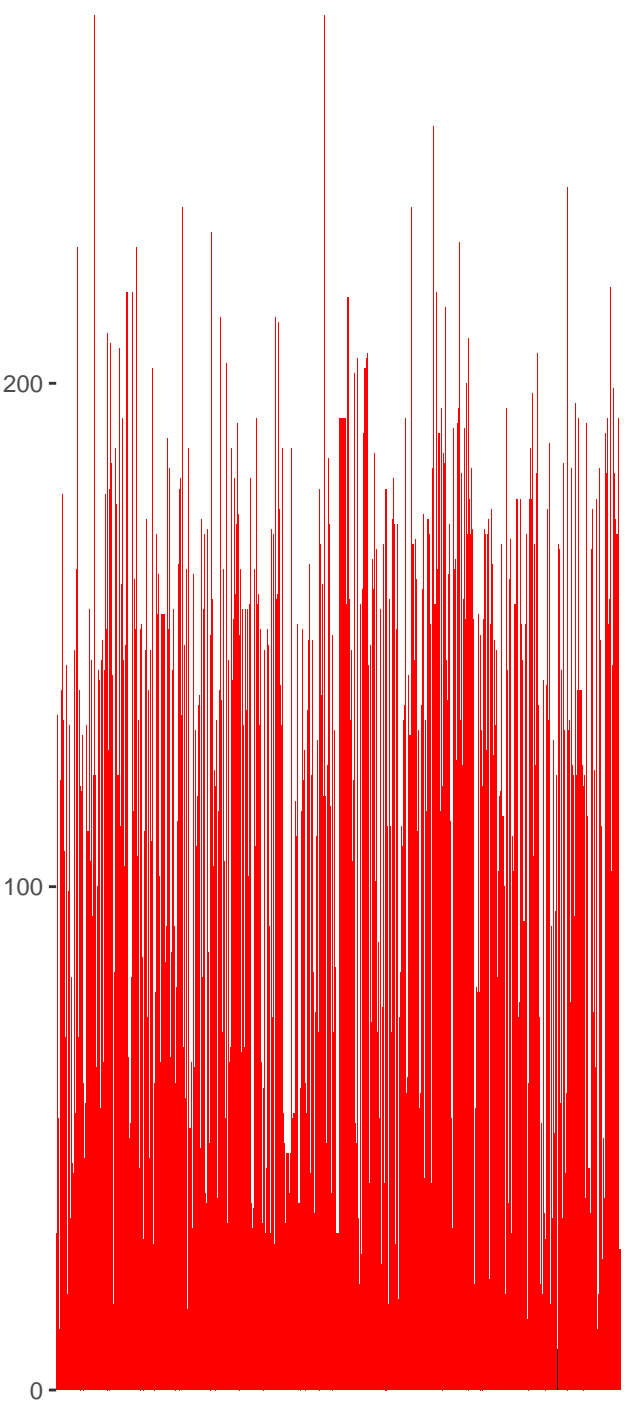
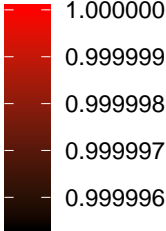


Number of genes overlap



Adjusted.P.value



Module names

based on data from enrichR