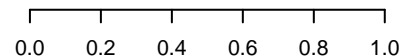
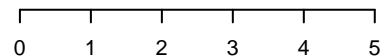
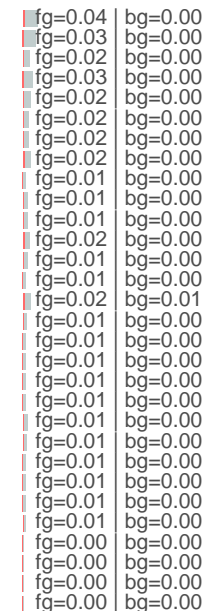
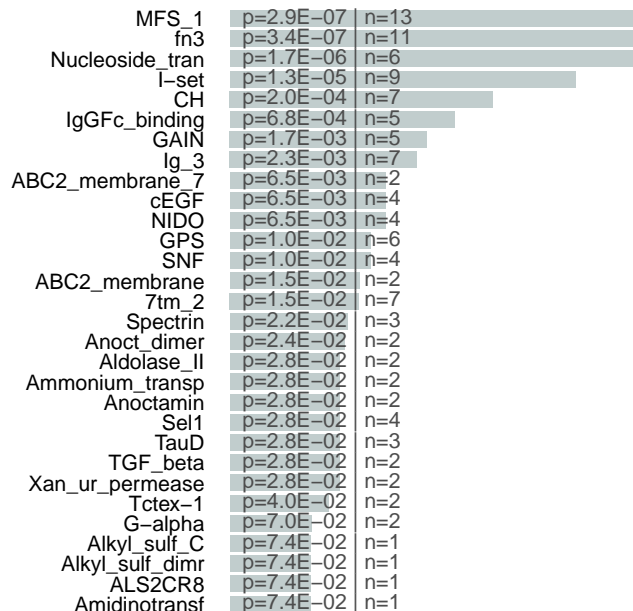


## gc025

## fraction genes in fg and bg



$-\log_{10}(p)$   
n=170/267 input genes with annotations

fraction