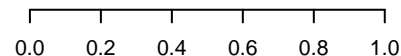
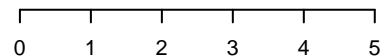
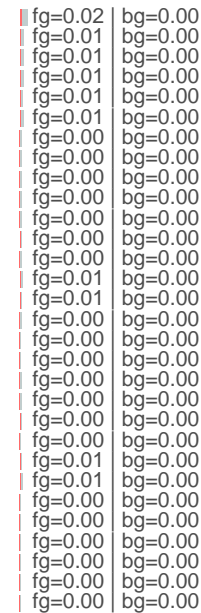
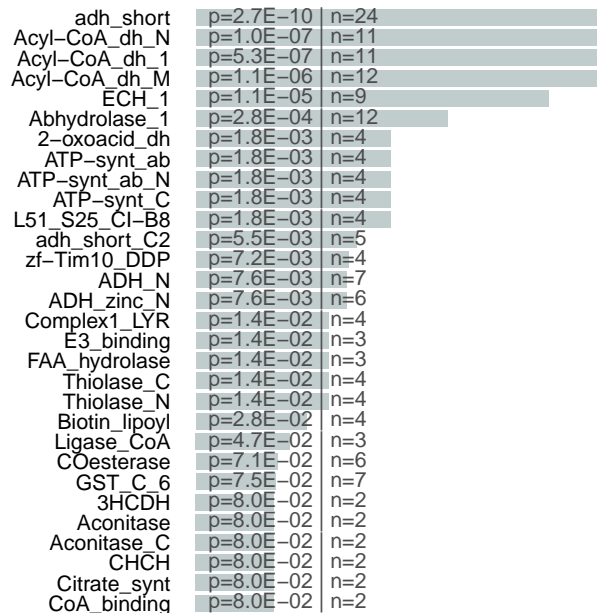


## gc002a

## fraction genes in fg and bg



$-\log_{10}(p)$   
n=735/947 input genes with annotations

fraction