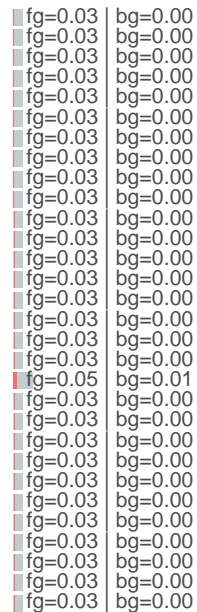


fraction genes in fg and bg



$-\log_{10}(p)$
n=38/30 input genes with annotations