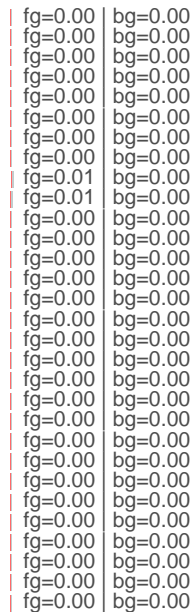


fraction genes in fg and bg



$-\log_{10}(p)$
n=292/258 input genes with annotations