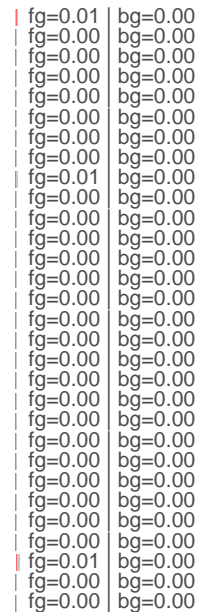


fraction genes in fg and bg



$-\log_{10}(p)$
n=280/248 input genes with annotations