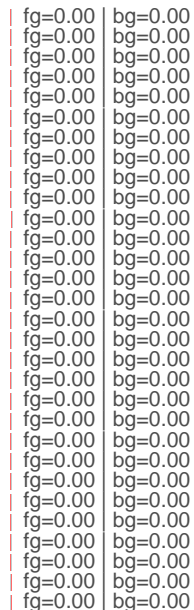


**fraction genes in fg and bg**



$-\log_{10}(p)$   
n=757/771 input genes with annotations