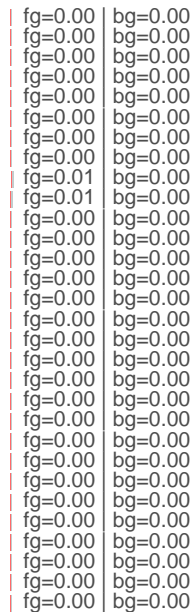


### **fraction genes in fg and bg**



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