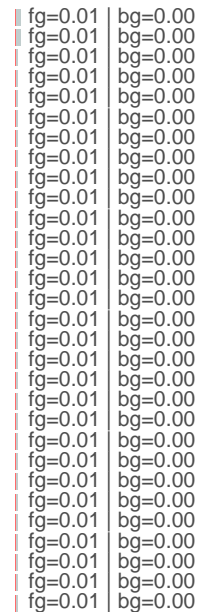


**fraction genes in fg and bg**



$-\log_{10}(p)$   
n=144/118 input genes with annotations