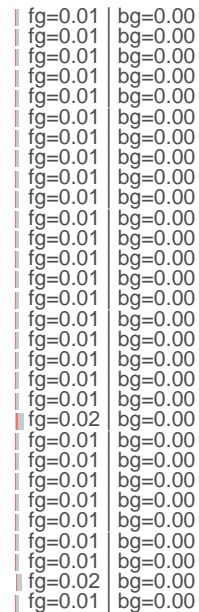


fraction genes in fg and bg



-log₁₀(p)
n=115/95 input genes with annotations