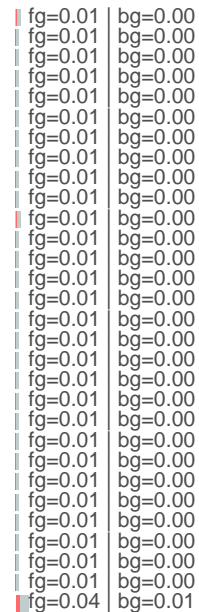


fraction genes in fg and bg



–log₁₀(p)
n=119/91 input genes with annotations