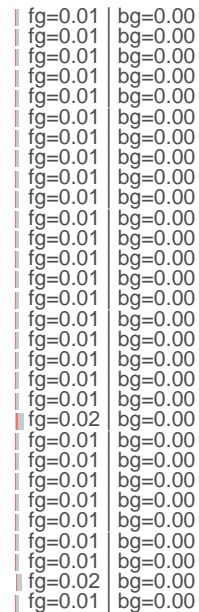


fraction genes in fg and bg



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