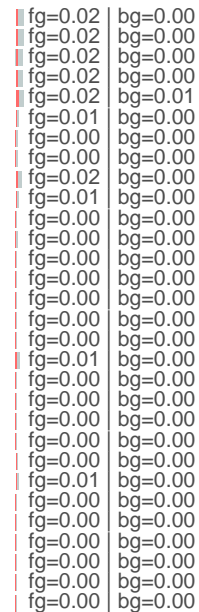


fraction genes in fg and bg



–log₁₀(p)
n=283/354 input genes with annotations