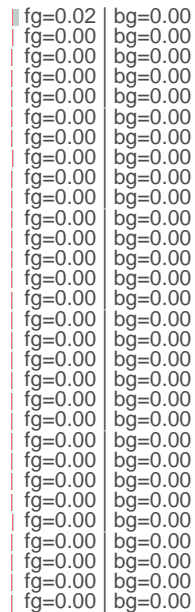


fraction genes in fg and bg



–log₁₀(p)
n=318/334 input genes with annotations