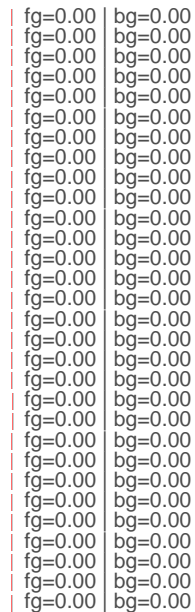


fraction genes in fg and bg



–log₁₀(p)
n=402/346 input genes with annotations