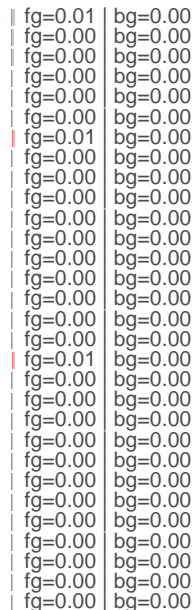


fraction genes in fg and bg



–log₁₀(p)
n=327/267 input genes with annotations