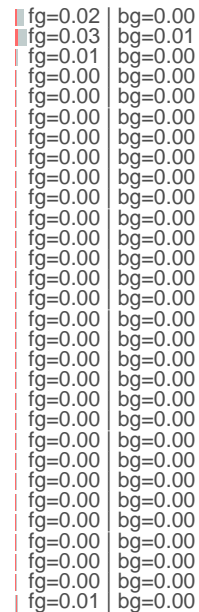


fraction genes in fg and bg



–log₁₀(p)
n=276/269 input genes with annotations