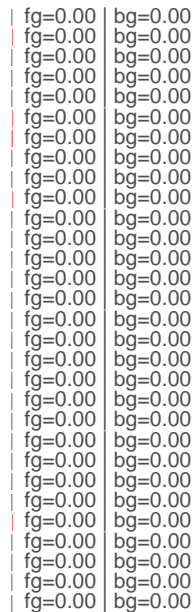


fraction genes in fg and bg



–log₁₀(p)
n=852/911 input genes with annotations