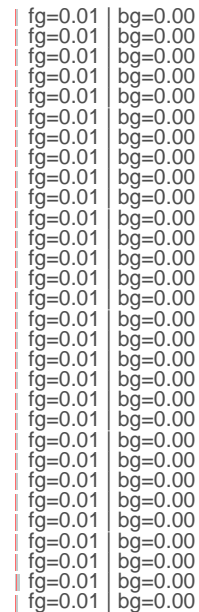


fraction genes in fg and bg



–log₁₀(p)
n=152/144 input genes with annotations