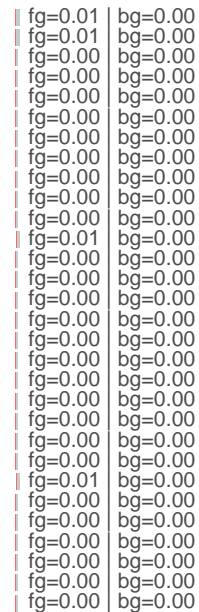


fraction genes in fg and bg



–log₁₀(p)
n=181/171 input genes with annotations