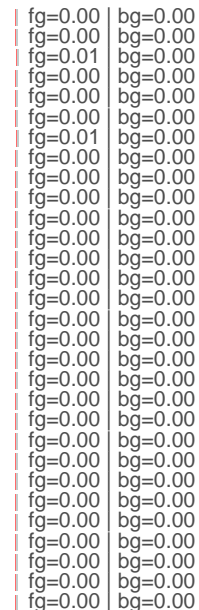


fraction genes in fg and bg



–log₁₀(p)
n=239/191 input genes with annotations