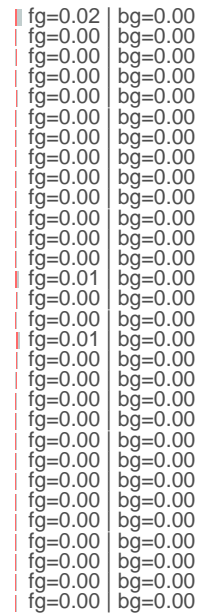


fraction genes in fg and bg



–log₁₀(p)
n=439/468 input genes with annotations