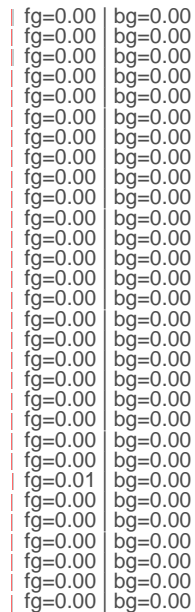


fraction genes in fg and bg



–log₁₀(p)
n=434/429 input genes with annotations