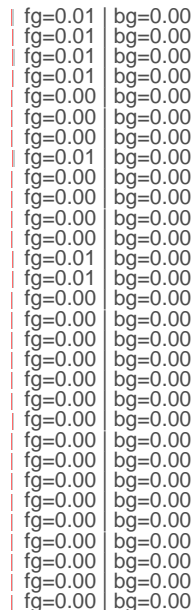


fraction genes in fg and bg



–log₁₀(p)
n=421/415 input genes with annotations