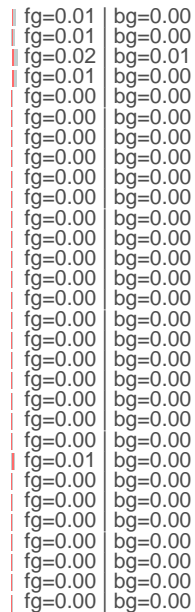


fraction genes in fg and bg



-log₁₀(p)
n=615/633 input genes with annotations