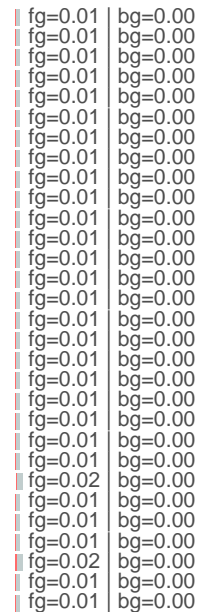


fraction genes in fg and bg



-log₁₀(p)
n=89/85 input genes with annotations