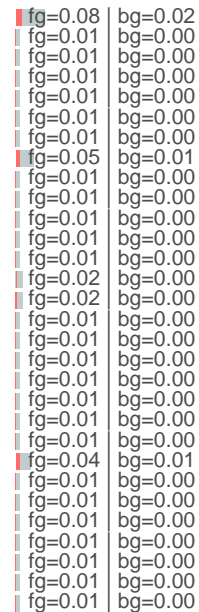


fraction genes in fg and bg



-log₁₀(p)
n=77/71 input genes with annotations