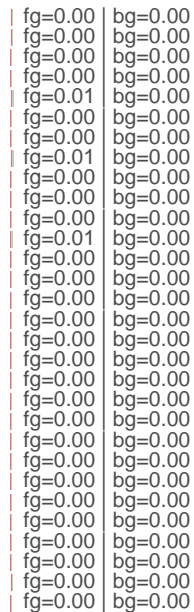


fraction genes in fg and bg



-log₁₀(p)
n=425/426 input genes with annotations