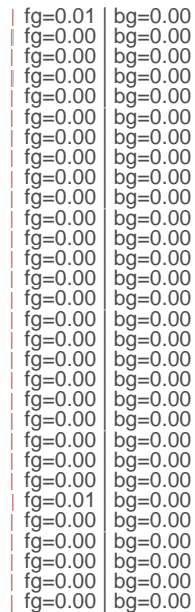


fraction genes in fg and bg



-log₁₀(p)
n=429/425 input genes with annotations