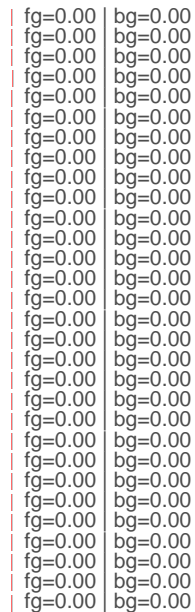


fraction genes in fg and bg



-log₁₀(p)
n=693/674 input genes with annotations