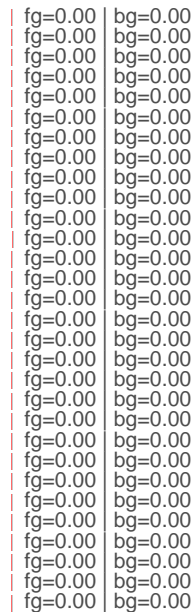


fraction genes in fg and bg



-log₁₀(p)
n=747/760 input genes with annotations