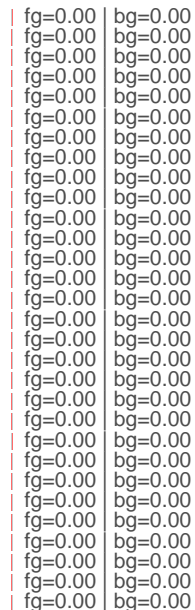


fraction genes in fg and bg



–log₁₀(p)
n=484/436 input genes with annotations