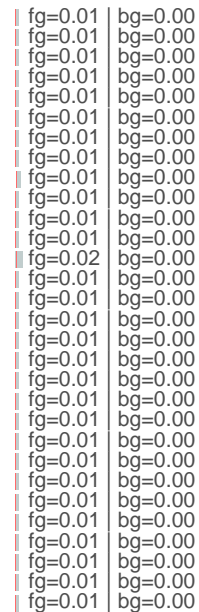


fraction genes in fg and bg



–log₁₀(p)
n=132/92 input genes with annotations

fraction