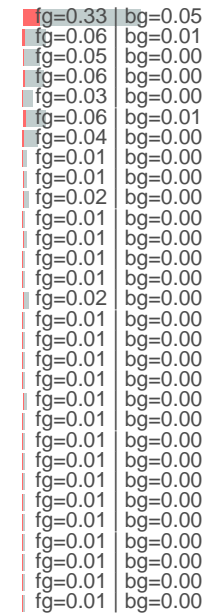


fraction genes in fg and bg



-log₁₀(p)
n=65/261 input genes with annotations

fraction