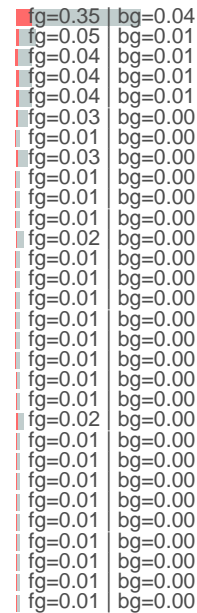


fraction genes in fg and bg



$-\log_{10}(p)$
n=40/104 input genes with annotations

fraction