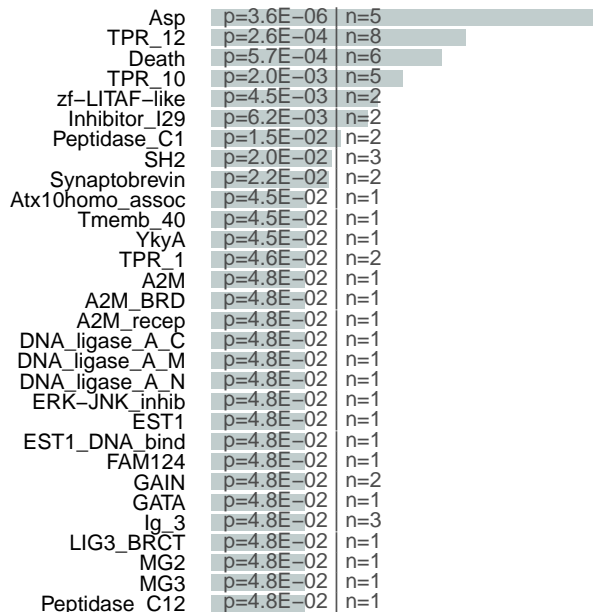


gc005

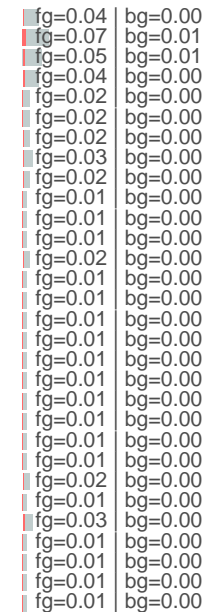
fraction genes in fg and bg



0 1 2 3 4 5

-log₁₀(p)

n=75/141 input genes with annotations



0.0 0.2 0.4 0.6 0.8 1.0

fraction