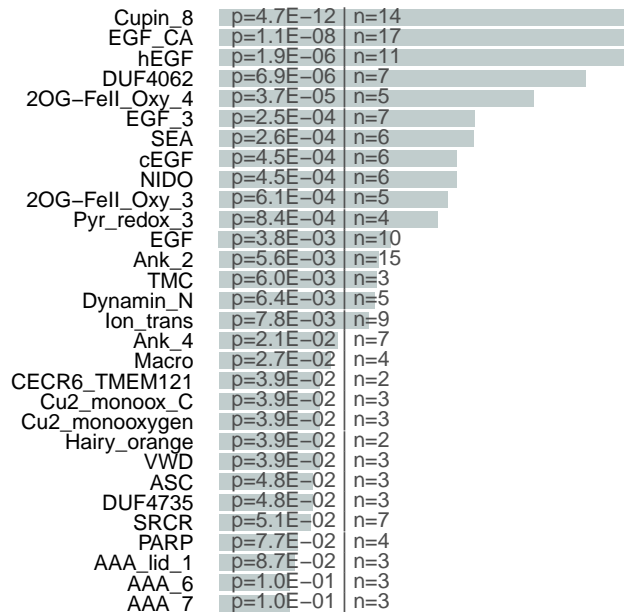


gc001a

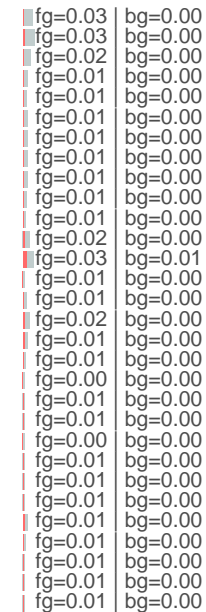


0 1 2 3 4 5

$-\log_{10}(p)$

n=250/342 input genes with annotations

fraction genes in fg and bg



0.0 0.2 0.4 0.6 0.8 1.0

fraction