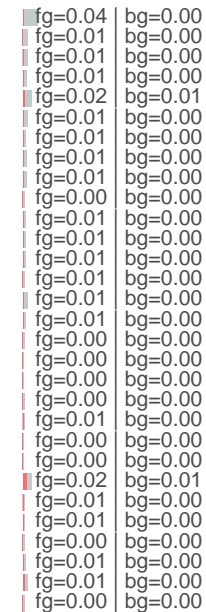
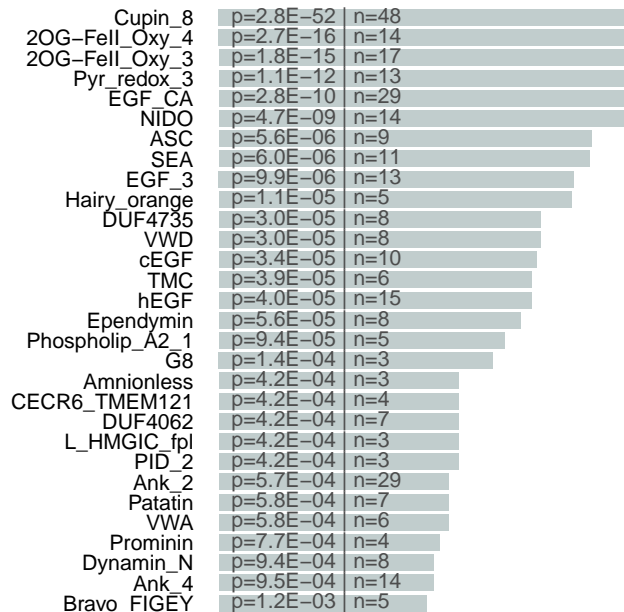


## gc001a

## fraction genes in fg and bg



0 1 2 3 4 5

0.0 0.2 0.4 0.6 0.8 1.0

$-\log_{10}(p)$

fraction

n=342/965 input genes with annotations