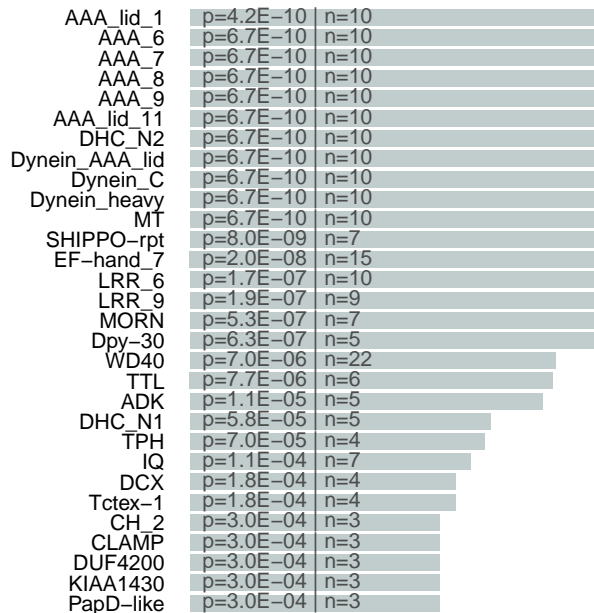


## gc030

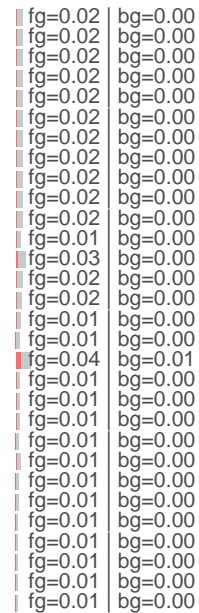
## fraction genes in fg and bg



0 1 2 3 4 5

$-\log_{10}(p)$

n=235/468 input genes with annotations



0.0 0.2 0.4 0.6 0.8 1.0

fraction