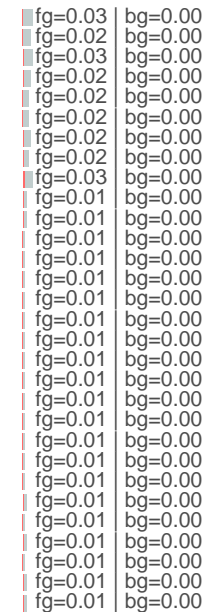
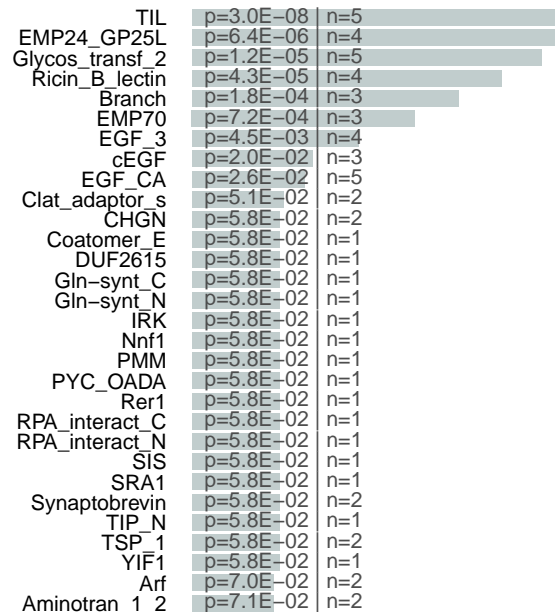


gc008

fraction genes in fg and bg



0 1 2 3 4 5

0.0 0.2 0.4 0.6 0.8 1.0

-log10(p)

fraction

n=137/170 input genes with annotations