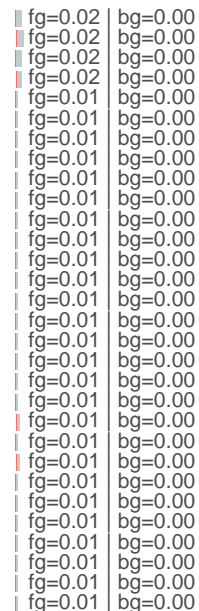
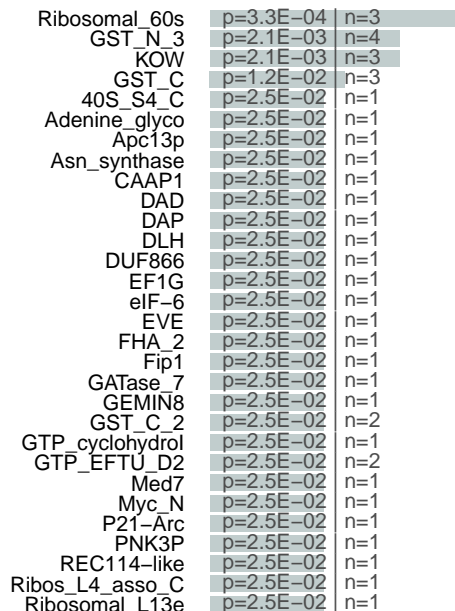


gc007a

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=170/159 input genes with annotations