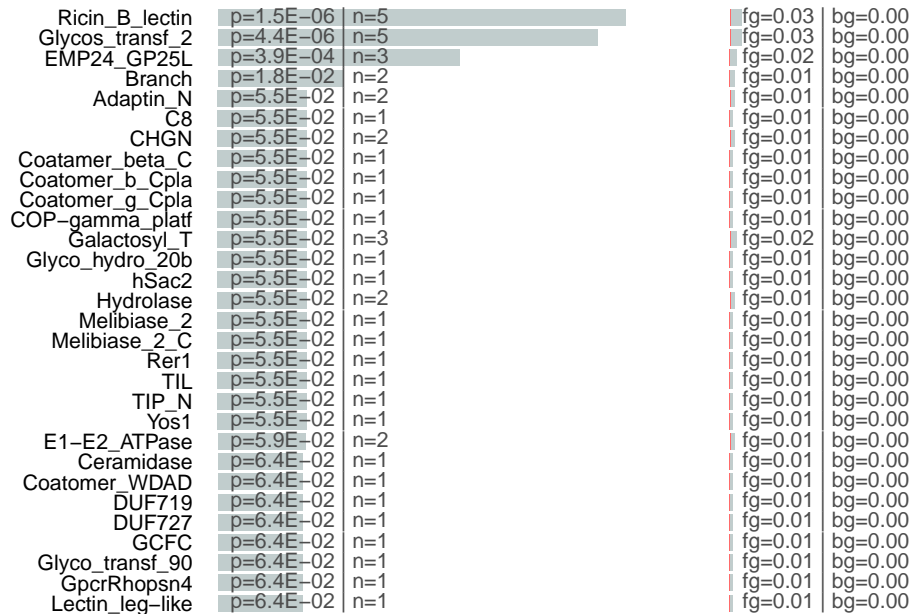


gc008

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=114/156 input genes with annotations