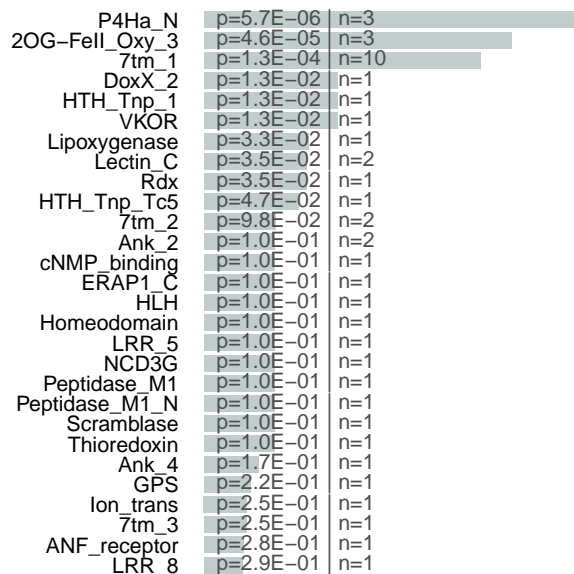


gc010

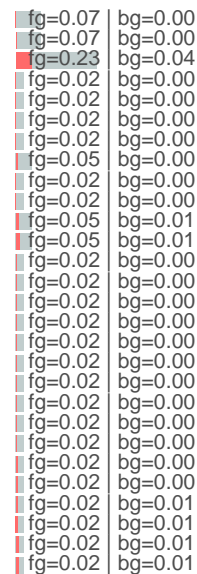
fraction genes in fg and bg



0 1 2 3 4 5

-log10(p)

n=28/50 input genes with annotations



0.0 0.2 0.4 0.6 0.8 1.0

fraction