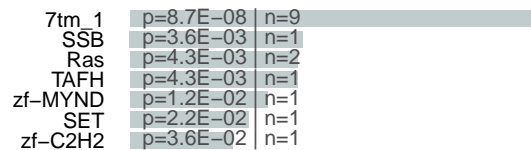


gc003a

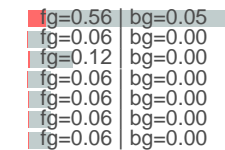
fraction genes in fg and bg



0 1 2 3 4 5

 $-\log_{10}(p)$ 

n=7/20 input genes with annotations



0.0 0.2 0.4 0.6 0.8 1.0

fraction