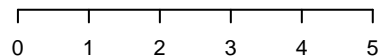
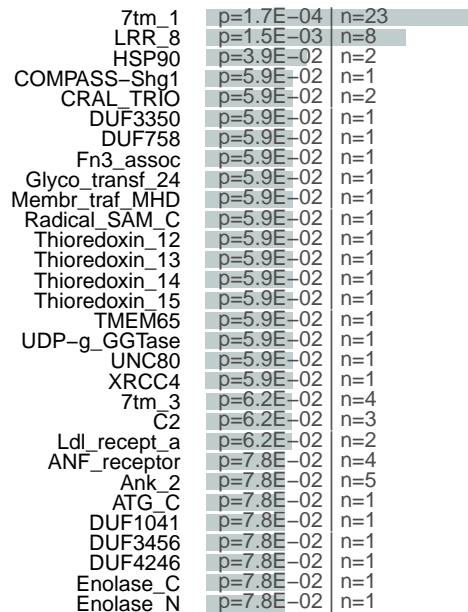
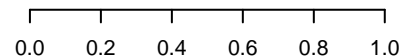
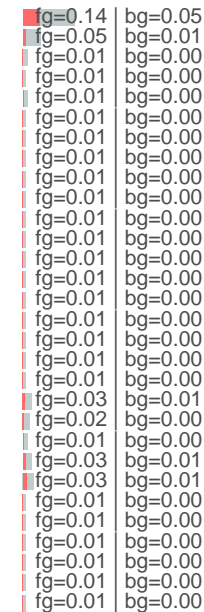


gc006

fraction genes in fg and bg



$-\log_{10}(p)$
n=106/209 input genes with annotations



fraction