

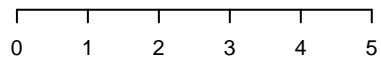
CniBil_loss

fraction genes in fg and bg

Histone

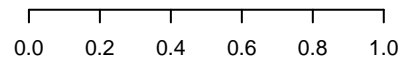
p=2.3E-03 | n=1

fg=1.00 | bg=0.00



$-\log_{10}(p)$

n=1/1 input genes with annotations



fraction