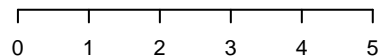
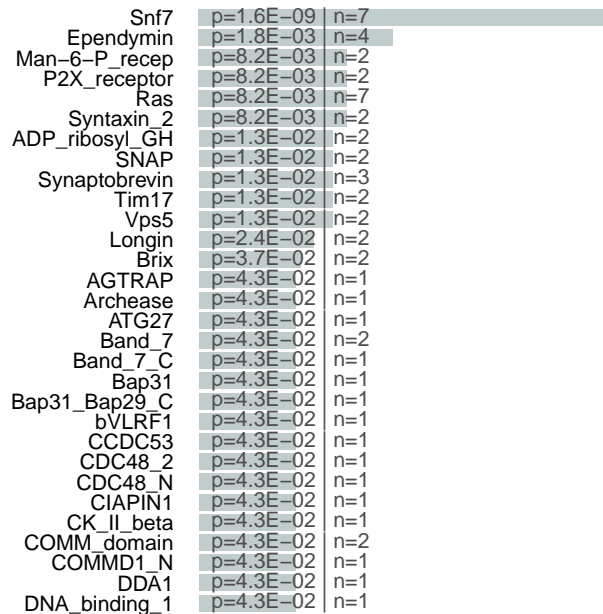
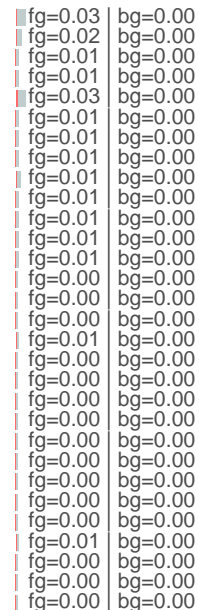


gc032

fraction genes in fg and bg



-log₁₀(p)
n=196/218 input genes with annotations



fraction