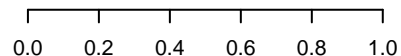
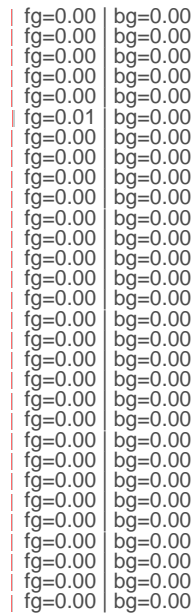


fraction genes in fg and bg



fraction

n=611/541 input genes with annotations