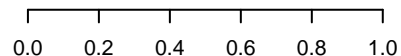
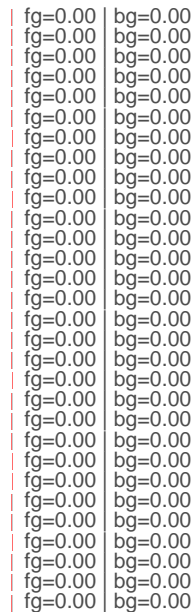


fraction genes in fg and bg



fraction

n=1042/1095 input genes with annotations