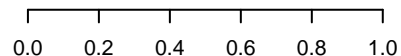
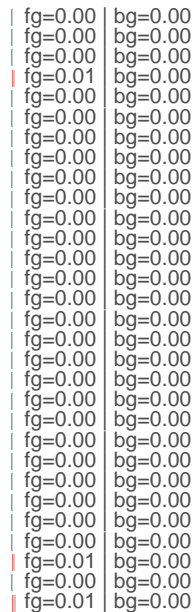


fraction genes in fg and bg



fraction

n=268/228 input genes with annotations