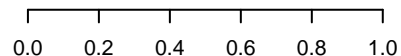
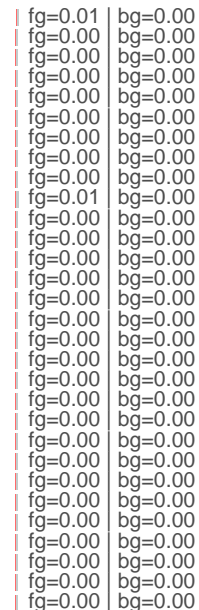


fraction genes in fg and bg



fraction

n=234/199 input genes with annotations