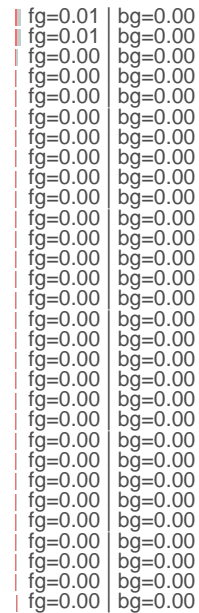


fraction genes in fg and bg



fraction

n=574/591 input genes with annotations