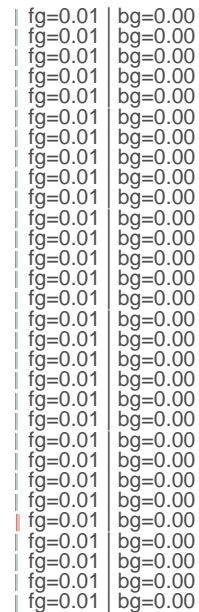


fraction genes in fg and bg



fraction

n=170/156 input genes with annotations