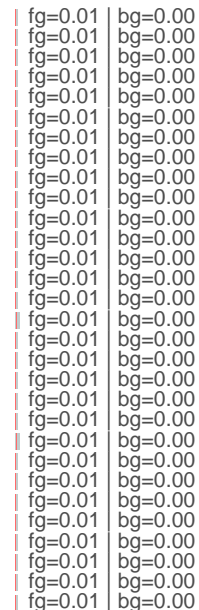


fraction genes in fg and bg



fraction

n=148/100 input genes with annotations