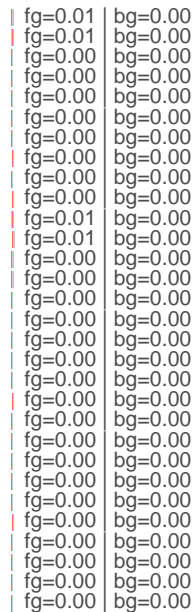


fraction genes in fg and bg



fraction

n=357/388 input genes with annotations