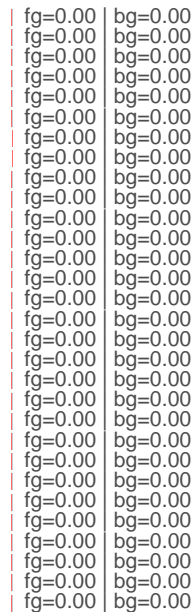


fraction genes in fg and bg



fraction

n=1024/983 input genes with annotations