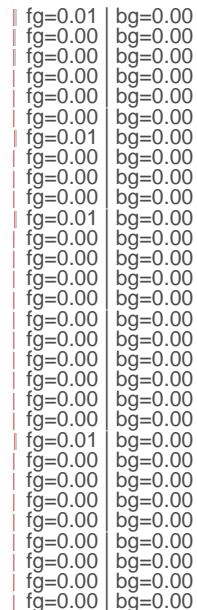


fraction genes in fg and bg



fraction

n=332/274 input genes with annotations