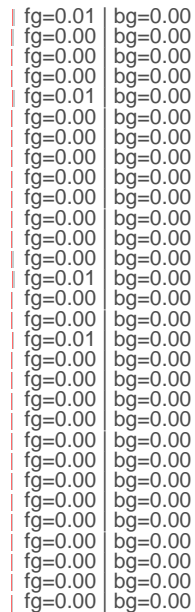


fraction genes in fg and bg



fraction

n=441/401 input genes with annotations