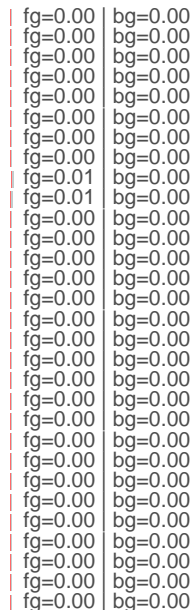


fraction genes in fg and bg



fraction

n=292/258 input genes with annotations