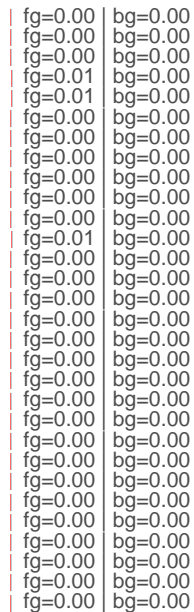


fraction genes in fg and bg



fraction

n=334/272 input genes with annotations