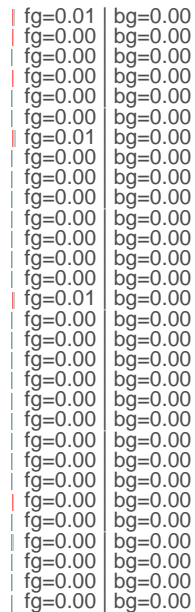


fraction genes in fg and bg



fraction

n=323/331 input genes with annotations