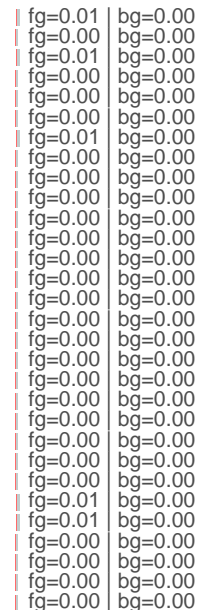


fraction genes in fg and bg



fraction

n=165/157 input genes with annotations