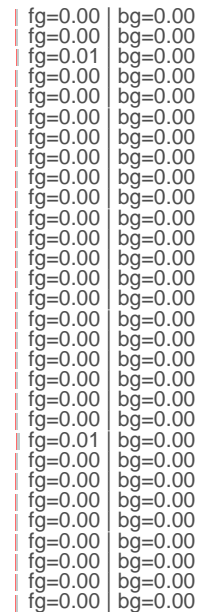


fraction genes in fg and bg



fraction

n=237/226 input genes with annotations