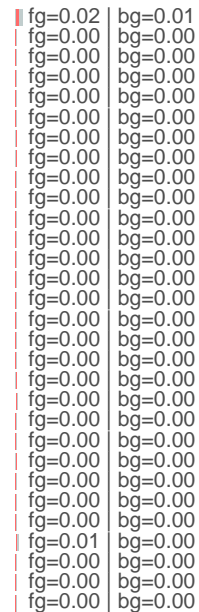


fraction genes in fg and bg



fraction

n=455/469 input genes with annotations