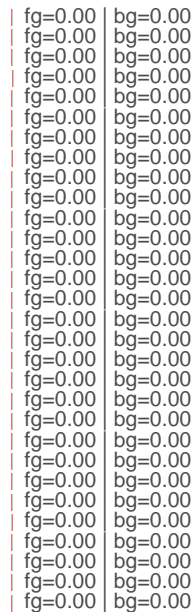


fraction genes in fg and bg



fraction

n=736/722 input genes with annotations