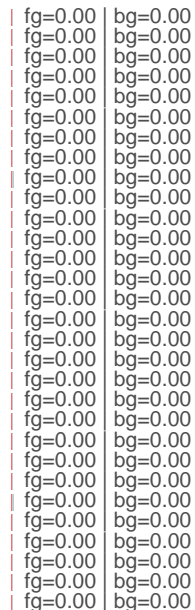


fraction genes in fg and bg



fraction

n=373/321 input genes with annotations