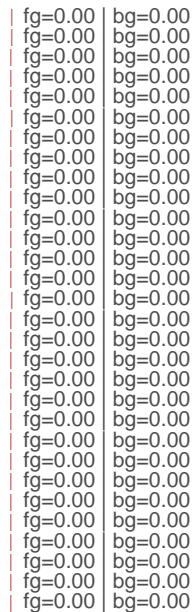


fraction genes in fg and bg



fraction

n=372/333 input genes with annotations