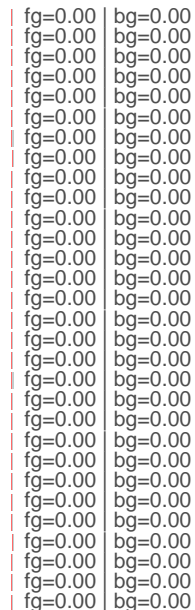


fraction genes in fg and bg



fraction

n=427/358 input genes with annotations