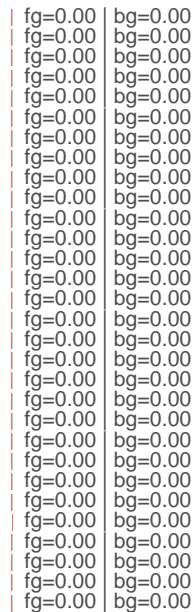


fraction genes in fg and bg



fraction

n=531/537 input genes with annotations