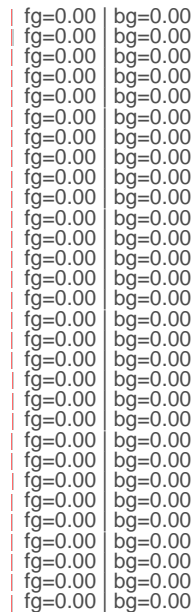


fraction genes in fg and bg



fraction

n=337/309 input genes with annotations