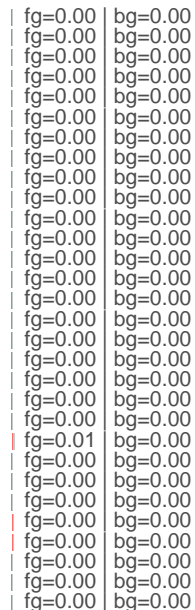


fraction genes in fg and bg



fraction

n=461/421 input genes with annotations