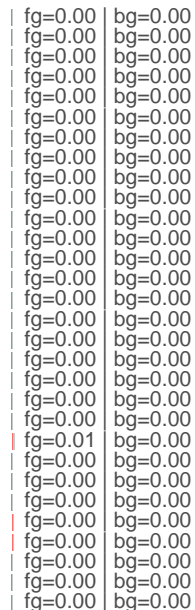


**fraction genes in fg and bg**



fraction

n=461/421 input genes with annotations