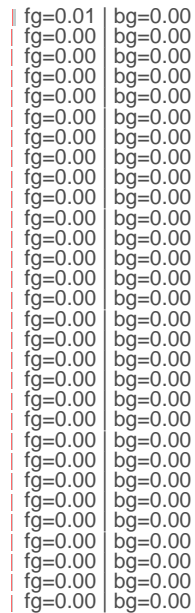


**fraction genes in fg and bg**



fraction

n=258/252 input genes with annotations