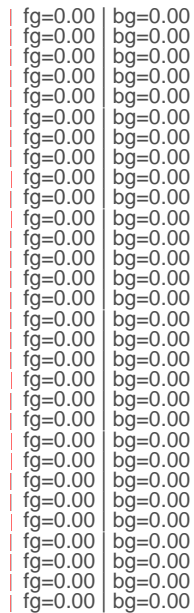


**fraction genes in fg and bg**



fraction

n=1010/1073 input genes with annotations