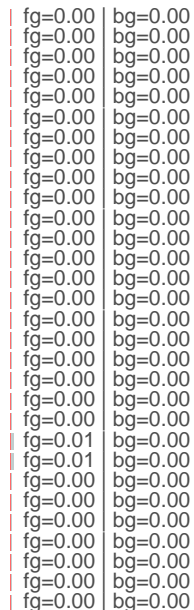


**fraction genes in fg and bg**



fraction

n=253/212 input genes with annotations