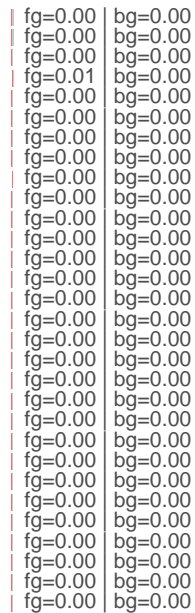


**fraction genes in fg and bg**



fraction

n=541/570 input genes with annotations