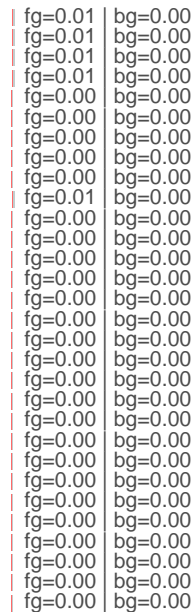


**fraction genes in fg and bg**



fraction

n=308/236 input genes with annotations