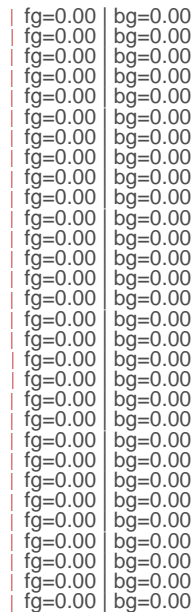


**fraction genes in fg and bg**



fraction

n=432/450 input genes with annotations