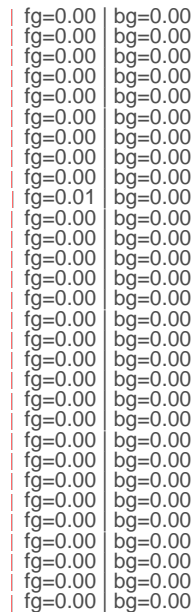


**fraction genes in fg and bg**



fraction

n=438/422 input genes with annotations