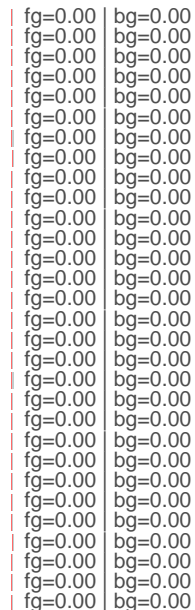


**fraction genes in fg and bg**



fraction

n=427/358 input genes with annotations