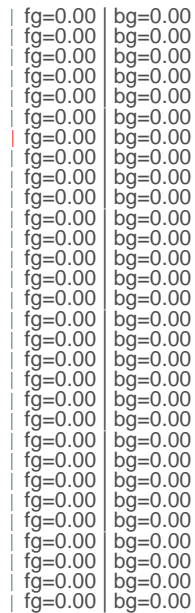


**fraction genes in fg and bg**



fraction

n=434/395 input genes with annotations