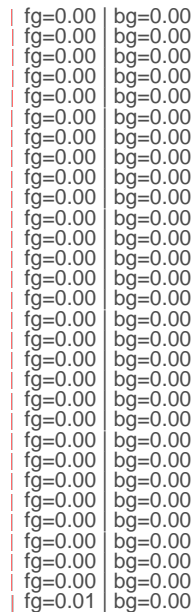


**fraction genes in fg and bg**



fraction

n=502/482 input genes with annotations