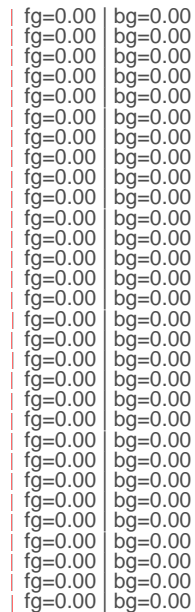


**fraction genes in fg and bg**



fraction

n=504/484 input genes with annotations