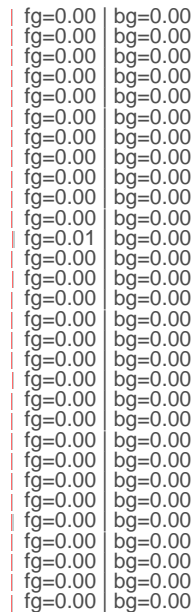


**fraction genes in fg and bg**



fraction

n=374/346 input genes with annotations