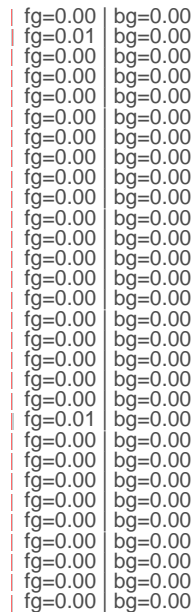


**fraction genes in fg and bg**



fraction

n=295/251 input genes with annotations