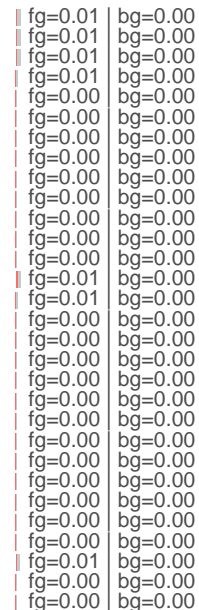


**fraction genes in fg and bg**



fraction

n=260/232 input genes with annotations