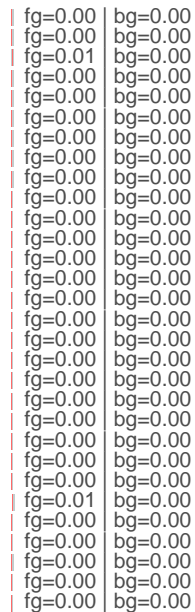


### **fraction genes in fg and bg**



fraction

n=409/359 input genes with annotations