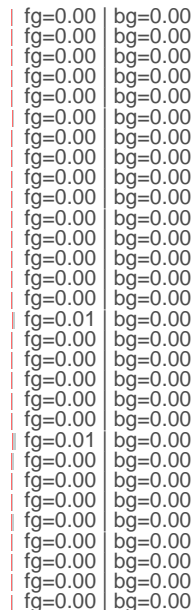


### **fraction genes in fg and bg**



fraction

n=404/344 input genes with annotations