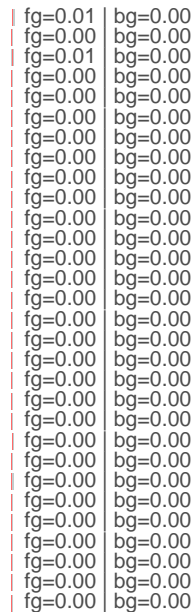


### **fraction genes in fg and bg**



fraction

n=402/368 input genes with annotations