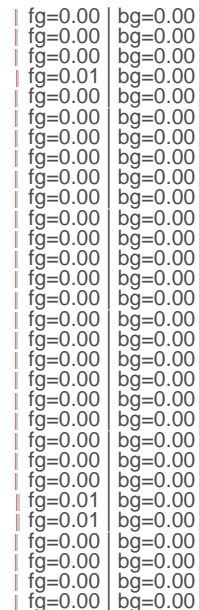


### **fraction genes in fg and bg**



fraction

n=237/204 input genes with annotations