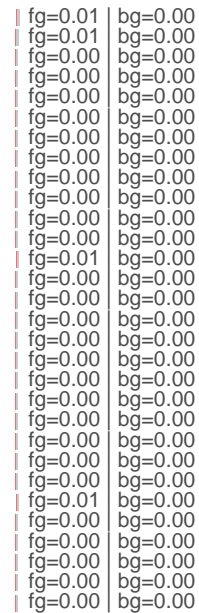


### **fraction genes in fg and bg**



fraction

n=237/199 input genes with annotations