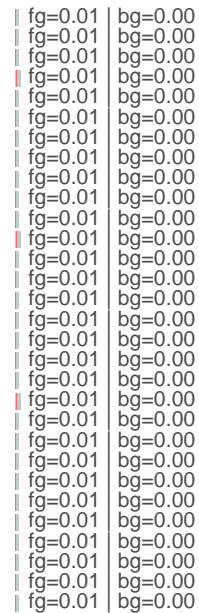


### **fraction genes in fg and bg**



fraction

n=121/89 input genes with annotations