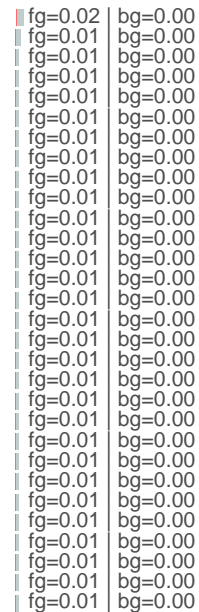


### **fraction genes in fg and bg**



fraction

n=124/113 input genes with annotations