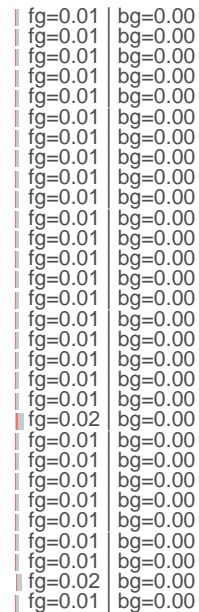


### **fraction genes in fg and bg**



fraction

n=115/95 input genes with annotations