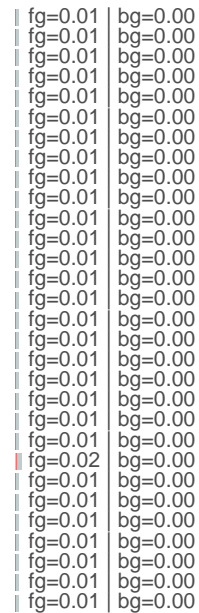


### **fraction genes in fg and bg**



fraction

n=113/88 input genes with annotations