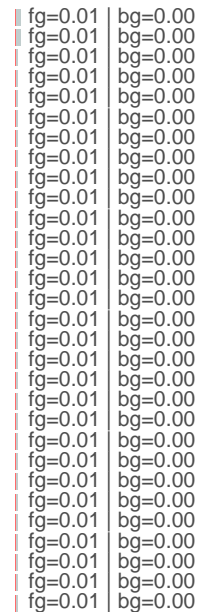


### **fraction genes in fg and bg**



fraction

n=144/118 input genes with annotations