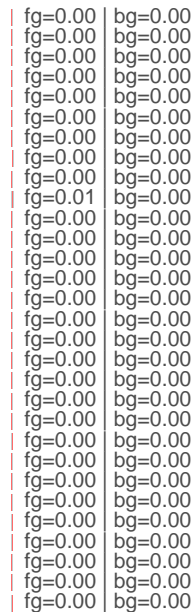


### **fraction genes in fg and bg**



fraction

n=511/505 input genes with annotations