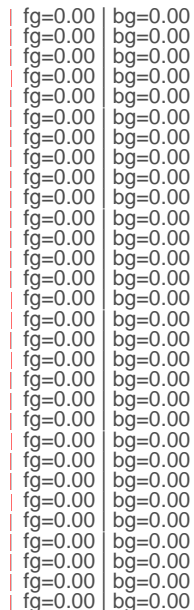


### **fraction genes in fg and bg**



fraction

n=851/879 input genes with annotations