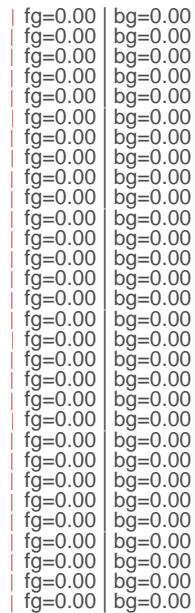


### **fraction genes in fg and bg**



fraction

n=951/996 input genes with annotations