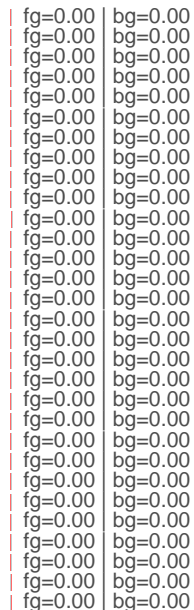


### **fraction genes in fg and bg**



fraction

n=757/771 input genes with annotations