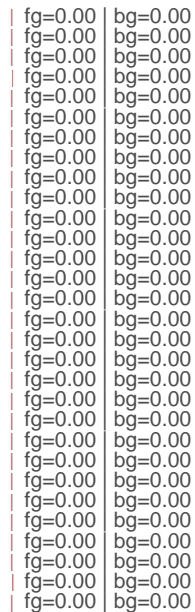


### **fraction genes in fg and bg**



fraction

n=693/683 input genes with annotations