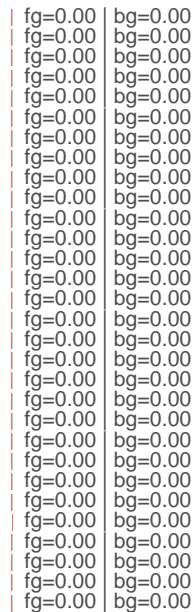


### **fraction genes in fg and bg**



fraction

n=531/537 input genes with annotations