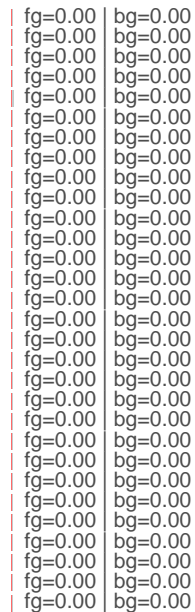


### **fraction genes in fg and bg**



fraction

n=484/431 input genes with annotations