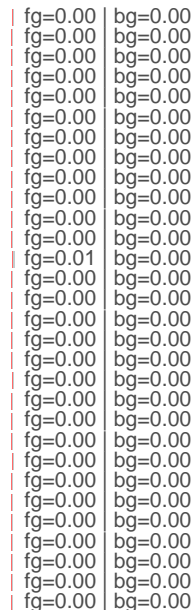


### **fraction genes in fg and bg**



fraction

n=302/309 input genes with annotations