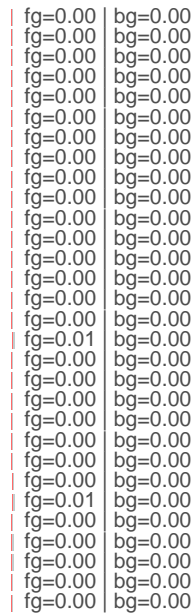


### **fraction genes in fg and bg**



fraction

n=412/373 input genes with annotations