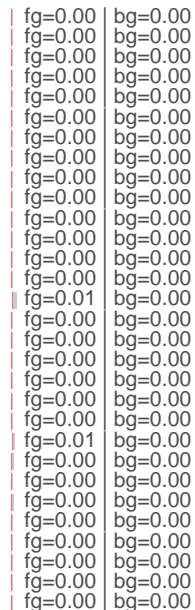


### **fraction genes in fg and bg**



fraction

n=404/364 input genes with annotations