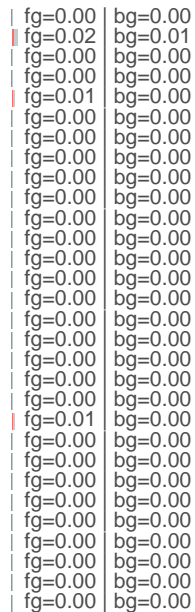


### **fraction genes in fg and bg**



fraction

n=243/187 input genes with annotations