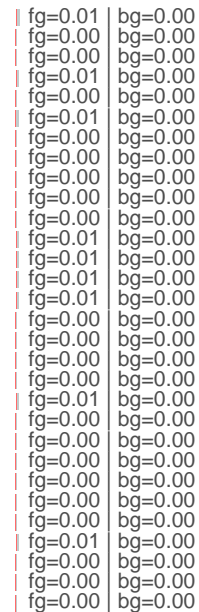


### **fraction genes in fg and bg**



fraction

n=253/202 input genes with annotations