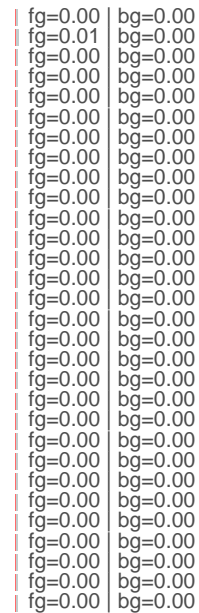


### **fraction genes in fg and bg**



fraction

n=230/177 input genes with annotations