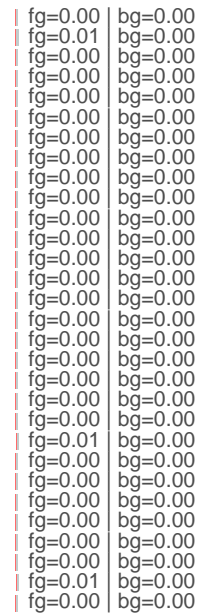


### **fraction genes in fg and bg**



fraction

n=206/156 input genes with annotations