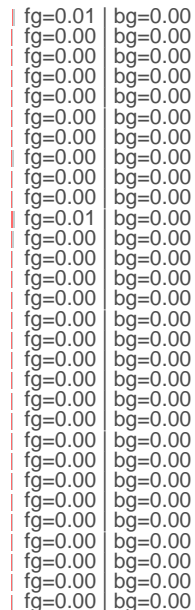


### **fraction genes in fg and bg**



fraction

n=457/452 input genes with annotations