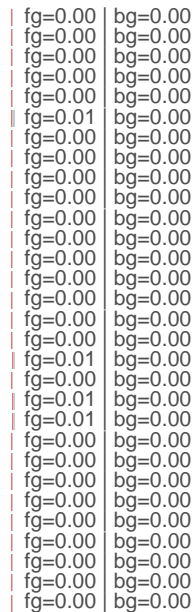


### **fraction genes in fg and bg**



fraction

n=253/205 input genes with annotations