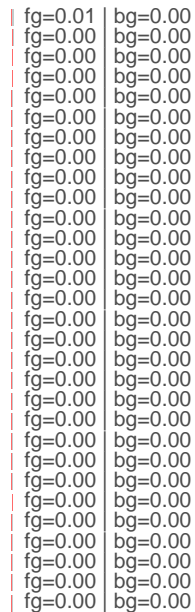


### **fraction genes in fg and bg**



fraction

n=508/527 input genes with annotations