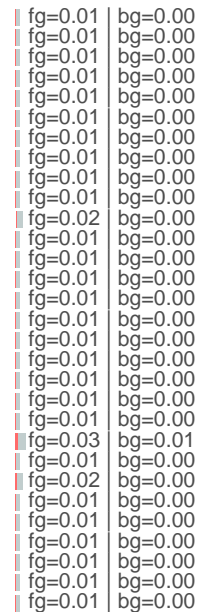


### **fraction genes in fg and bg**



fraction

n=98/83 input genes with annotations