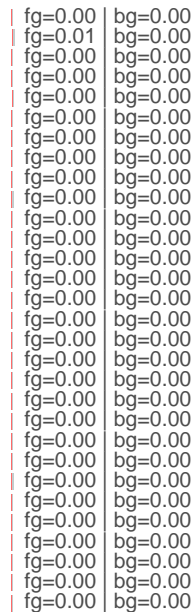


### **fraction genes in fg and bg**



fraction

n=327/285 input genes with annotations