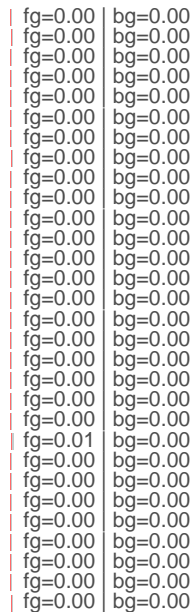


### **fraction genes in fg and bg**



fraction

n=547/526 input genes with annotations