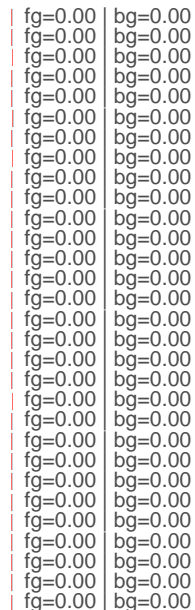


### **fraction genes in fg and bg**



fraction

n=562/534 input genes with annotations