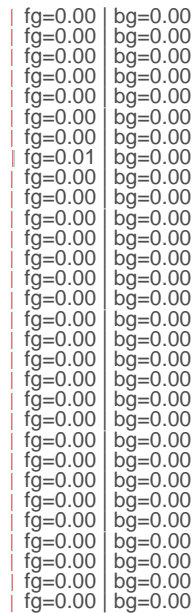


### **fraction genes in fg and bg**



fraction

n=513/466 input genes with annotations