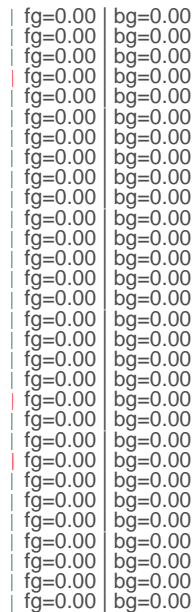


### **fraction genes in fg and bg**



fraction

n=1193/1446 input genes with annotations