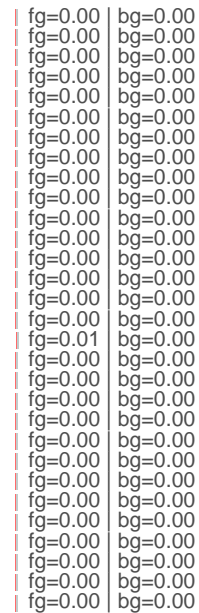


### **fraction genes in fg and bg**



fraction

n=219/183 input genes with annotations