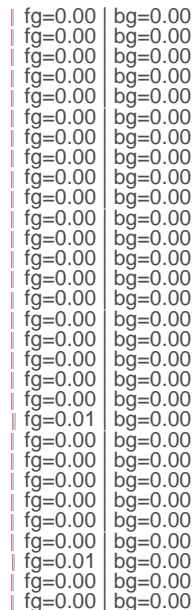


### **fraction genes in fg and bg**



fraction

n=227/196 input genes with annotations