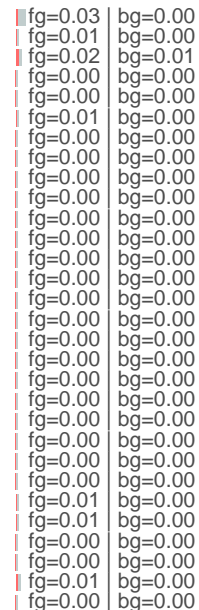


### **fraction genes in fg and bg**



fraction

n=183/187 input genes with annotations