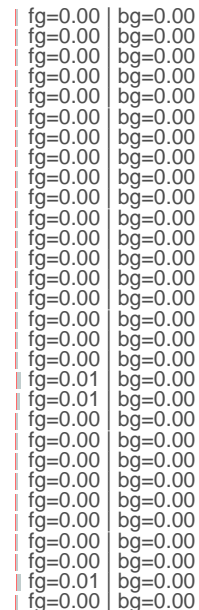


### **fraction genes in fg and bg**



fraction

n=182/153 input genes with annotations