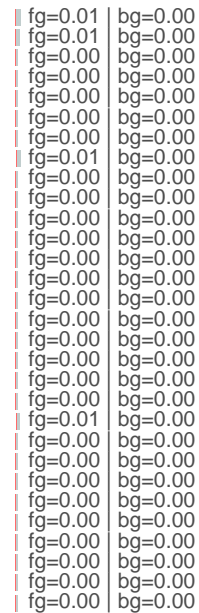


### **fraction genes in fg and bg**



fraction

n=181/167 input genes with annotations