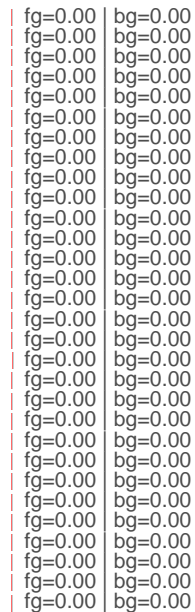


### **fraction genes in fg and bg**



fraction

n=581/538 input genes with annotations