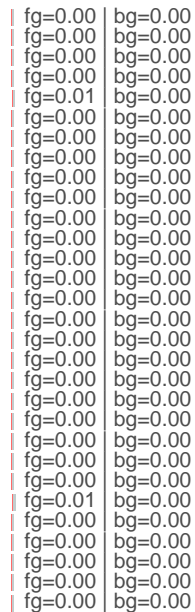


### **fraction genes in fg and bg**



fraction

n=217/182 input genes with annotations