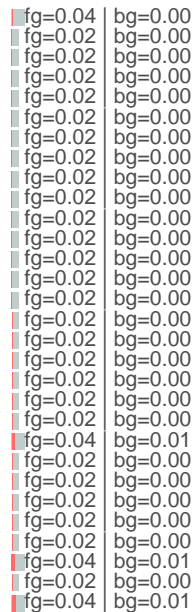


### **fraction genes in fg and bg**



fraction

n=47/45 input genes with annotations