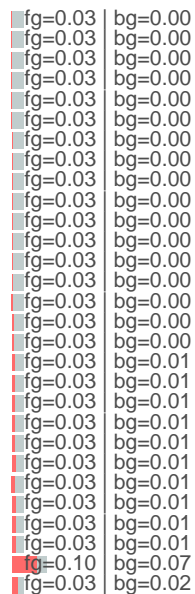


### **fraction genes in fg and bg**



fraction

n=29/19 input genes with annotations