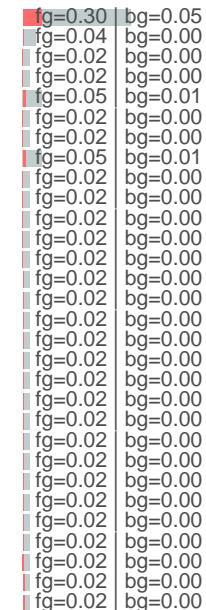


### **fraction genes in fg and bg**



fraction

n=35/64 input genes with annotations