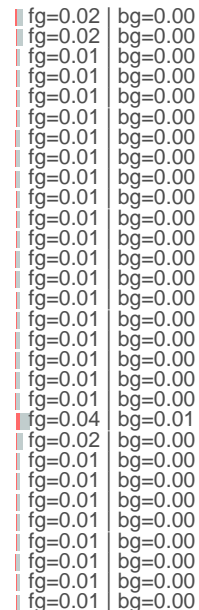


**fraction genes in fg and bg**



fraction

n=89/83 input genes with annotations