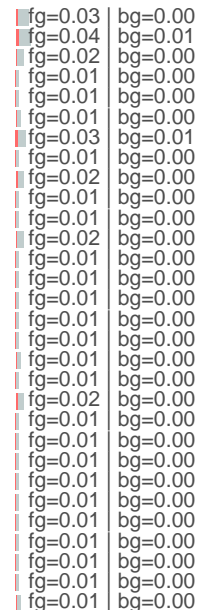


**fraction genes in fg and bg**



fraction

n=115/106 input genes with annotations