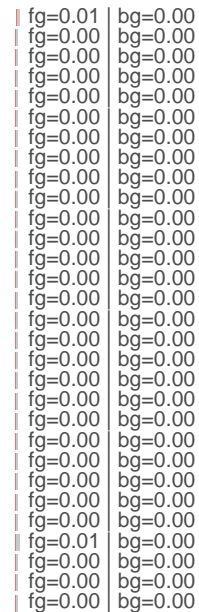


**fraction genes in fg and bg**



fraction

n=185/155 input genes with annotations