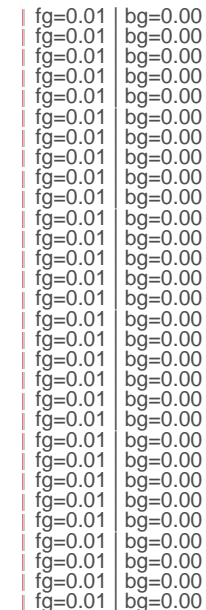


**fraction genes in fg and bg**



fraction

n=154/143 input genes with annotations