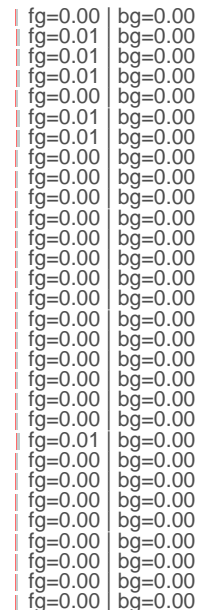


**fraction genes in fg and bg**



fraction

n=175/138 input genes with annotations