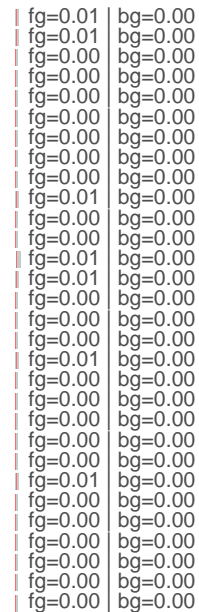


**fraction genes in fg and bg**



fraction

n=190/161 input genes with annotations