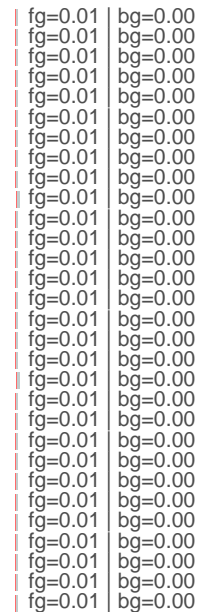


**fraction genes in fg and bg**



fraction

n=168/156 input genes with annotations