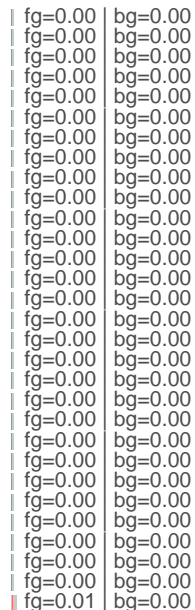


**fraction genes in fg and bg**



fraction

n=172/158 input genes with annotations