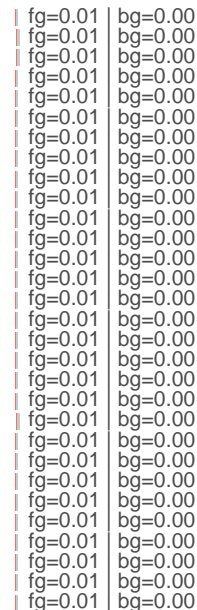


**fraction genes in fg and bg**



fraction

n=171/121 input genes with annotations