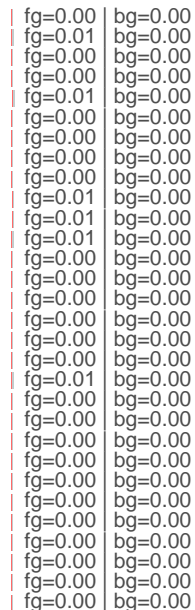


**fraction genes in fg and bg**



fraction

n=305/243 input genes with annotations