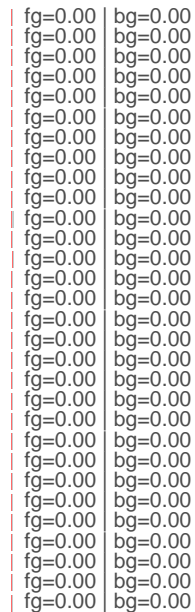


**fraction genes in fg and bg**



fraction

n=362/310 input genes with annotations