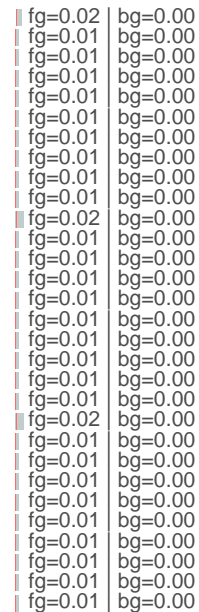


fraction genes in fg and bg



fraction

n=118/104 input genes with annotations