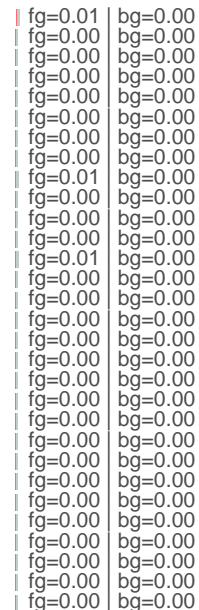


fraction genes in fg and bg



fraction

n=252/198 input genes with annotations