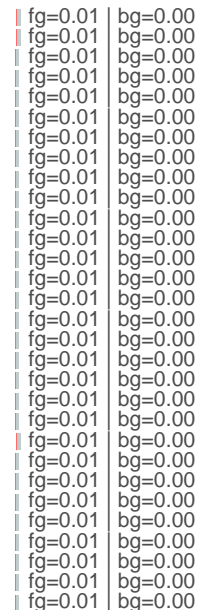


fraction genes in fg and bg



fraction

n=134/120 input genes with annotations