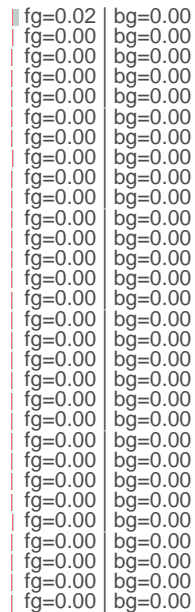


fraction genes in fg and bg



fraction

n=318/334 input genes with annotations