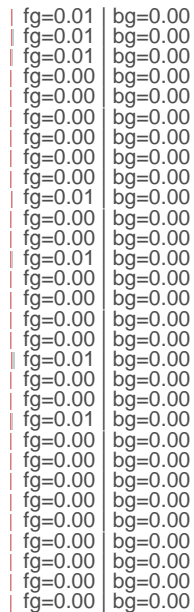


fraction genes in fg and bg



fraction

n=298/290 input genes with annotations