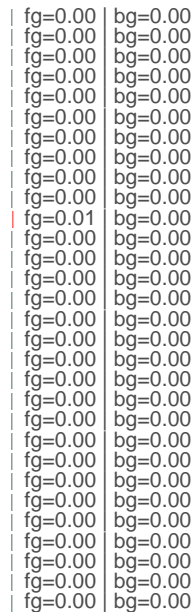


fraction genes in fg and bg



fraction

n=556/537 input genes with annotations