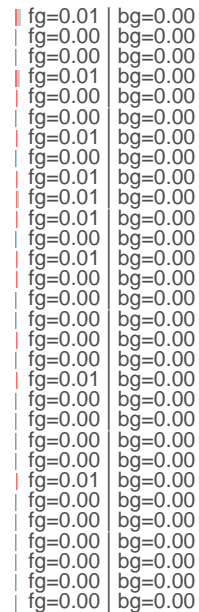


fraction genes in fg and bg



fraction

n=593/648 input genes with annotations