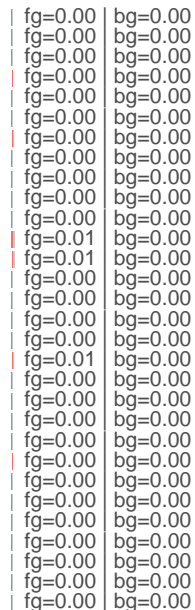


fraction genes in fg and bg



fraction

n=495/506 input genes with annotations