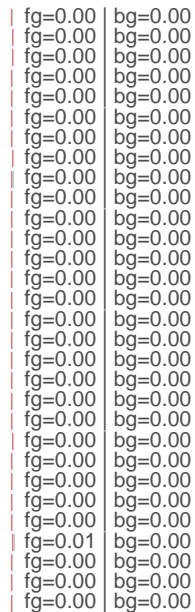


fraction genes in fg and bg



fraction

n=406/389 input genes with annotations