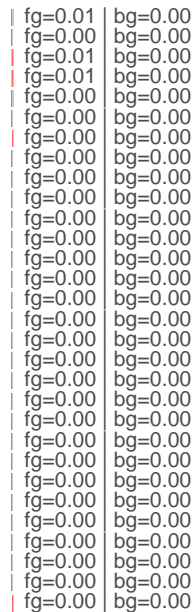


fraction genes in fg and bg



fraction

n=397/354 input genes with annotations