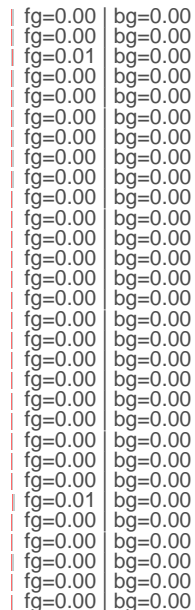


fraction genes in fg and bg



fraction

n=409/359 input genes with annotations