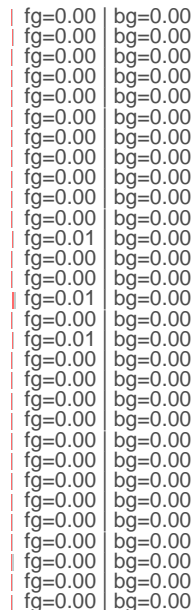


fraction genes in fg and bg



fraction

n=389/363 input genes with annotations