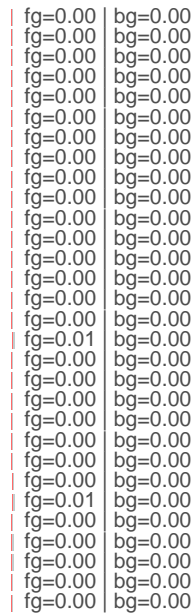


fraction genes in fg and bg



fraction

n=412/373 input genes with annotations