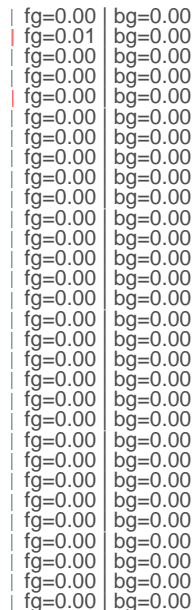


fraction genes in fg and bg



fraction

n=383/344 input genes with annotations