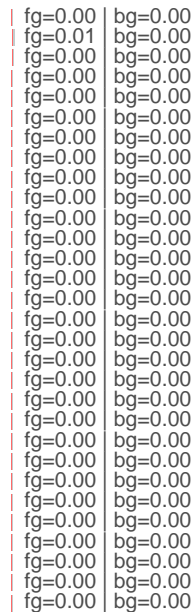


fraction genes in fg and bg



fraction

n=390/347 input genes with annotations