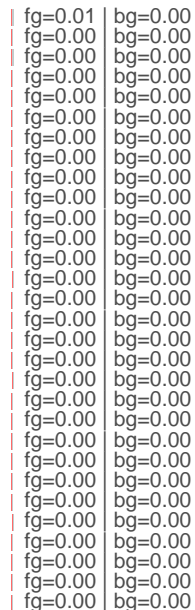


fraction genes in fg and bg



fraction

n=435/368 input genes with annotations