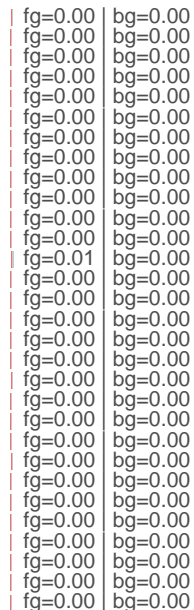


fraction genes in fg and bg



fraction

n=302/309 input genes with annotations