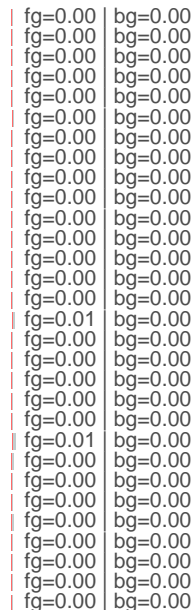


fraction genes in fg and bg



fraction

n=404/344 input genes with annotations