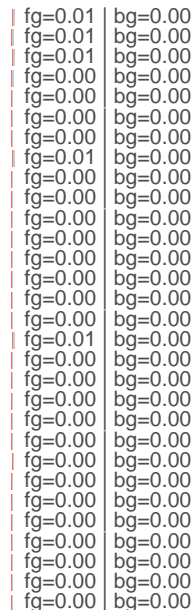


fraction genes in fg and bg



fraction

n=433/404 input genes with annotations