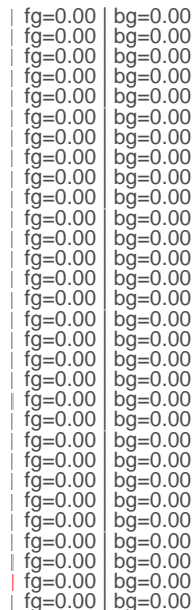


fraction genes in fg and bg



fraction

n=419/384 input genes with annotations