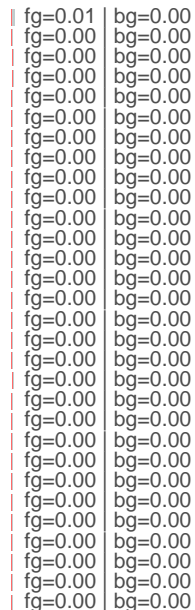


fraction genes in fg and bg



fraction

n=391/343 input genes with annotations