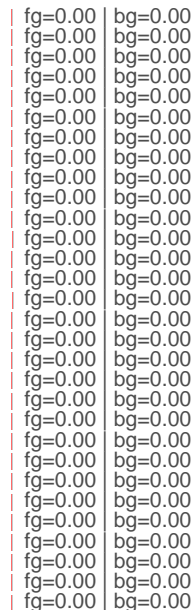


fraction genes in fg and bg



fraction

n=361/332 input genes with annotations