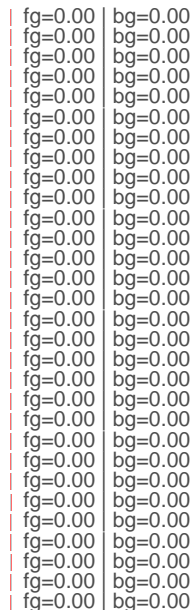


fraction genes in fg and bg



fraction

n=525/497 input genes with annotations