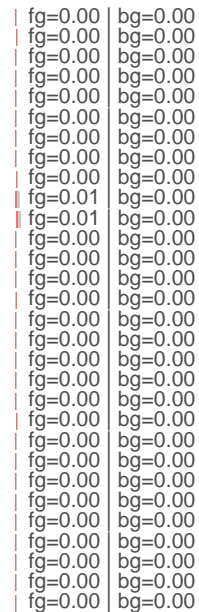


fraction genes in fg and bg



fraction

n=448/514 input genes with annotations