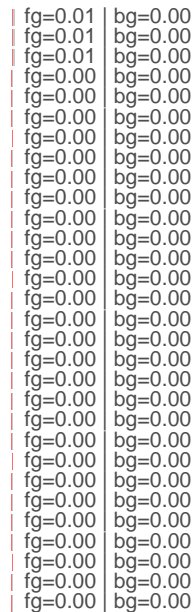


fraction genes in fg and bg



fraction

n=520/519 input genes with annotations