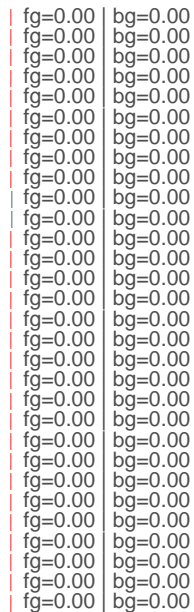


fraction genes in fg and bg



fraction

n=529/513 input genes with annotations