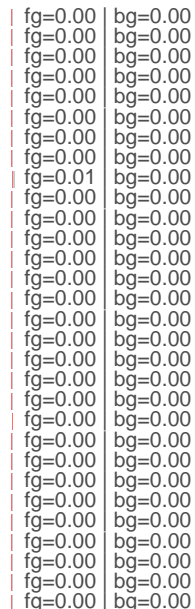


fraction genes in fg and bg



fraction

n=574/530 input genes with annotations