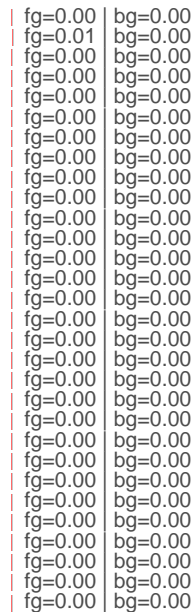


fraction genes in fg and bg



fraction

n=695/665 input genes with annotations