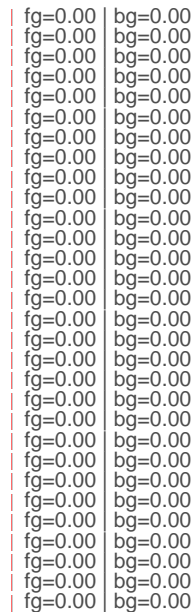


fraction genes in fg and bg



fraction

n=571/539 input genes with annotations