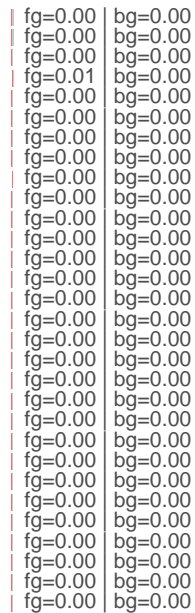


fraction genes in fg and bg



fraction

n=541/570 input genes with annotations