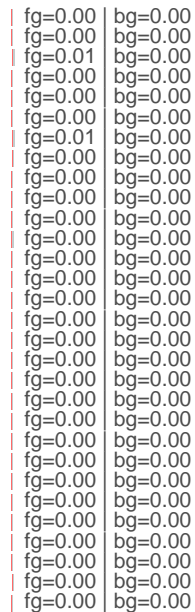


fraction genes in fg and bg



fraction

n=440/470 input genes with annotations