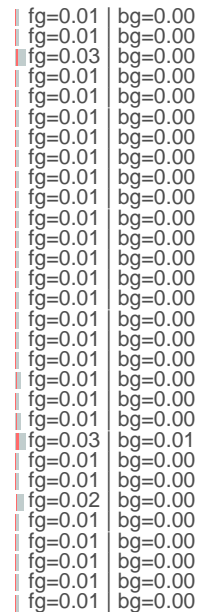


fraction genes in fg and bg



fraction

n=119/122 input genes with annotations