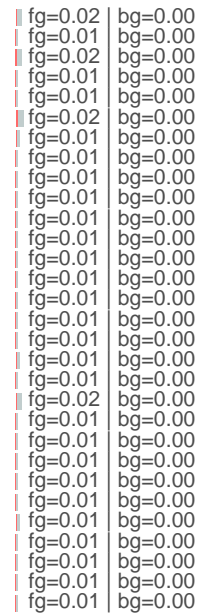


fraction genes in fg and bg



fraction

n=153/136 input genes with annotations