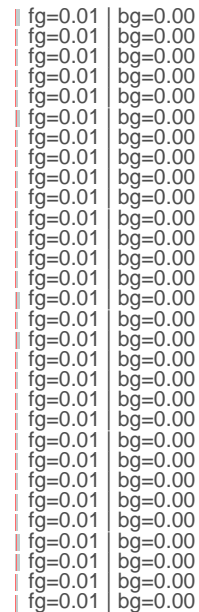


fraction genes in fg and bg



fraction

n=153/126 input genes with annotations