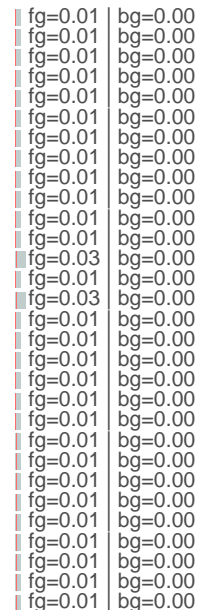


fraction genes in fg and bg



fraction

n=63/54 input genes with annotations