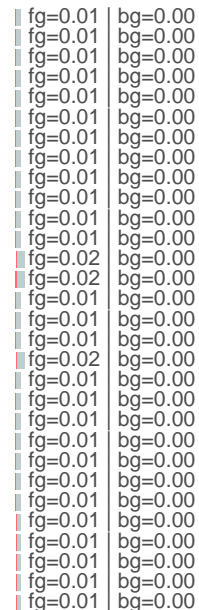


fraction genes in fg and bg



fraction

n=69/55 input genes with annotations