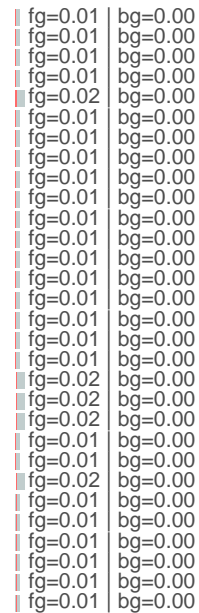


fraction genes in fg and bg



fraction

n=69/66 input genes with annotations