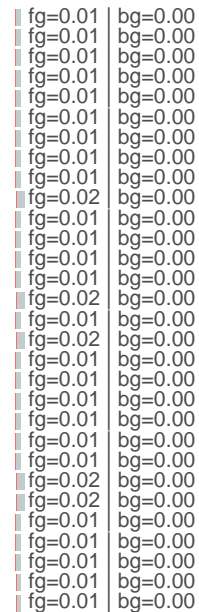


fraction genes in fg and bg



fraction

n=71/67 input genes with annotations