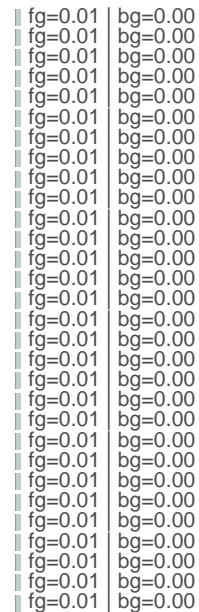


fraction genes in fg and bg



fraction

n=73/61 input genes with annotations