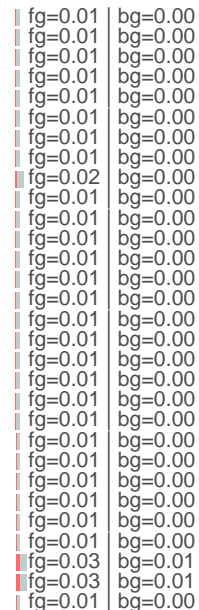


fraction genes in fg and bg



fraction

n=78/72 input genes with annotations