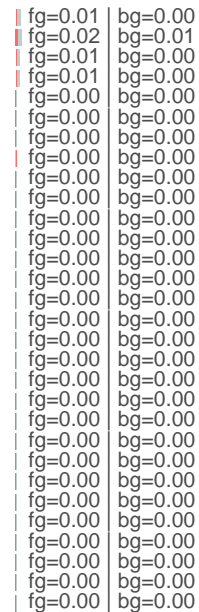


fraction genes in fg and bg



fraction

n=453/419 input genes with annotations