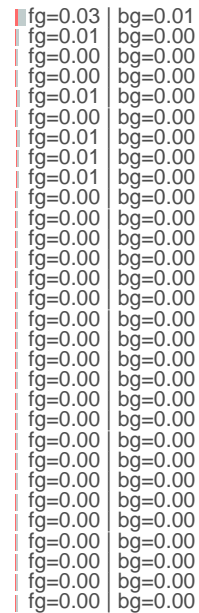


fraction genes in fg and bg



fraction

n=234/208 input genes with annotations