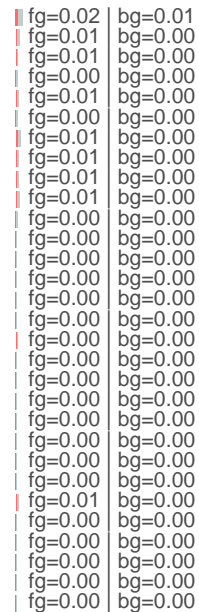


fraction genes in fg and bg



fraction

n=412/382 input genes with annotations