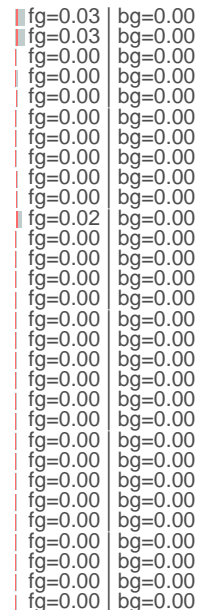


fraction genes in fg and bg



fraction

n=324/351 input genes with annotations