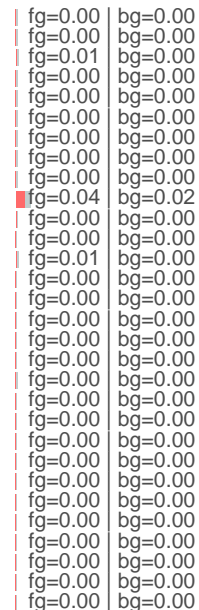


fraction genes in fg and bg



fraction

n=378/359 input genes with annotations