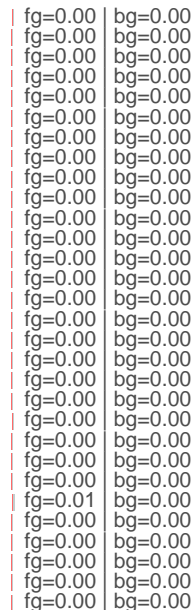


fraction genes in fg and bg



fraction

n=378/330 input genes with annotations