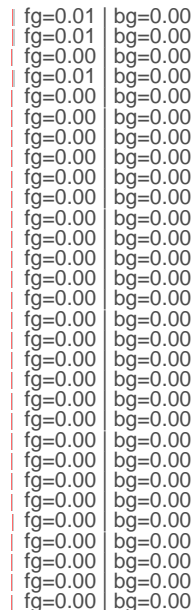


fraction genes in fg and bg



fraction

n=388/356 input genes with annotations