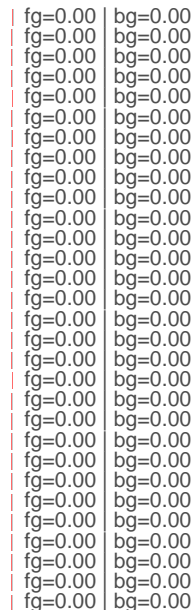


fraction genes in fg and bg



fraction

n=942/987 input genes with annotations