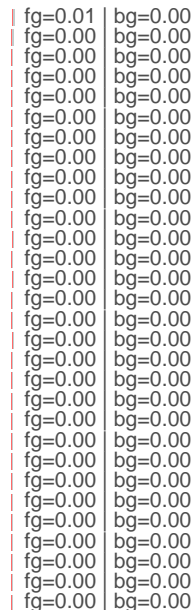


fraction genes in fg and bg



fraction

n=338/323 input genes with annotations