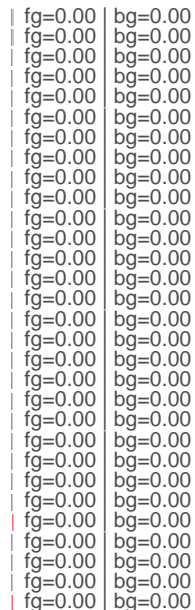


fraction genes in fg and bg



fraction

n=477/438 input genes with annotations