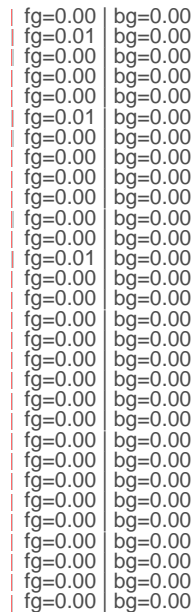


fraction genes in fg and bg



fraction

n=405/365 input genes with annotations