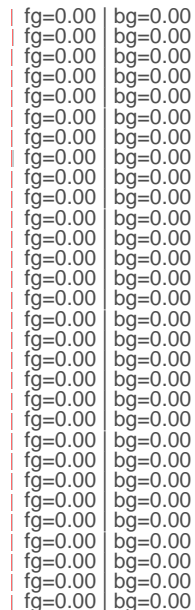


fraction genes in fg and bg



fraction

n=351/307 input genes with annotations