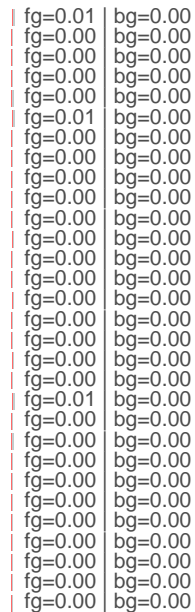


fraction genes in fg and bg



fraction

n=351/309 input genes with annotations