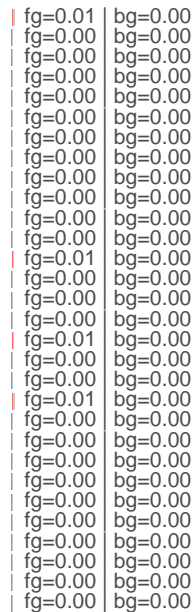


fraction genes in fg and bg



fraction

n=316/278 input genes with annotations