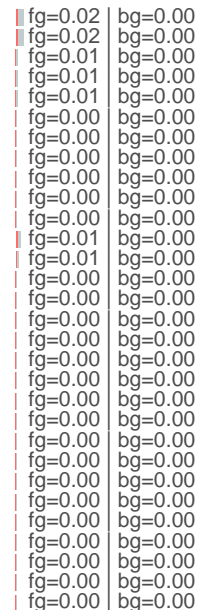


fraction genes in fg and bg



fraction

n=271/286 input genes with annotations