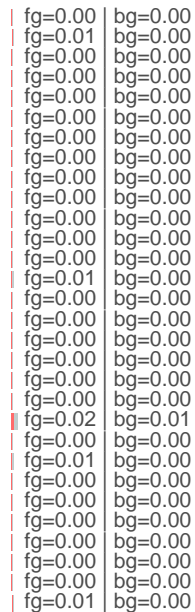


fraction genes in fg and bg



fraction

n=301/263 input genes with annotations