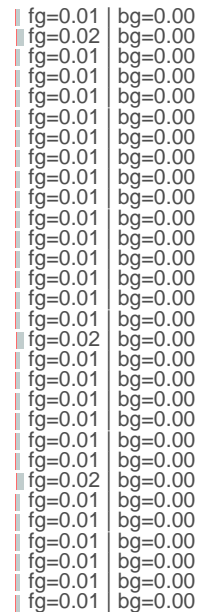


fraction genes in fg and bg



fraction

n=95/84 input genes with annotations