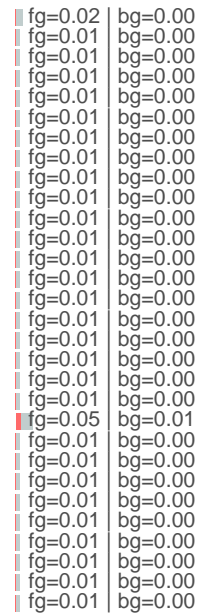


fraction genes in fg and bg



fraction

n=95/88 input genes with annotations