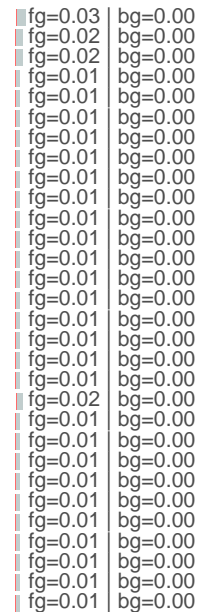


fraction genes in fg and bg



fraction

n=100/91 input genes with annotations