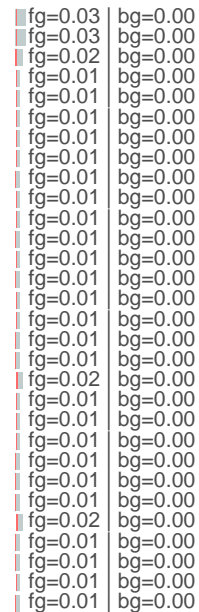


fraction genes in fg and bg



fraction

n=82/76 input genes with annotations