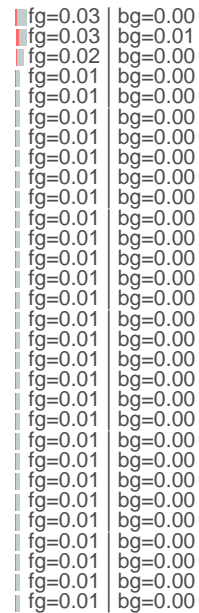


fraction genes in fg and bg



fraction

n=82/70 input genes with annotations