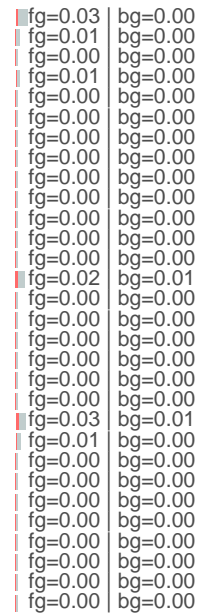


fraction genes in fg and bg



fraction

n=161/170 input genes with annotations