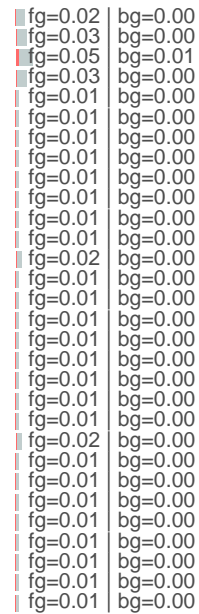


fraction genes in fg and bg



fraction

n=107/110 input genes with annotations