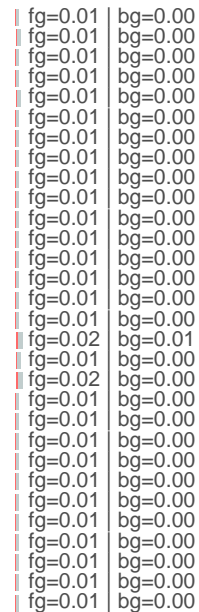


fraction genes in fg and bg



fraction

n=138/110 input genes with annotations