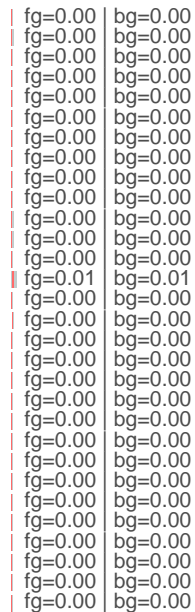


fraction genes in fg and bg



fraction

n=325/254 input genes with annotations