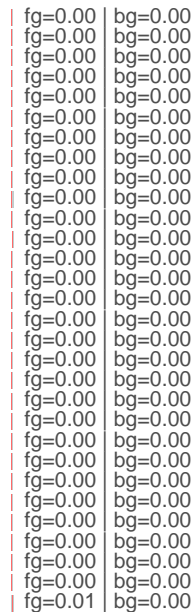


fraction genes in fg and bg



fraction

n=395/335 input genes with annotations