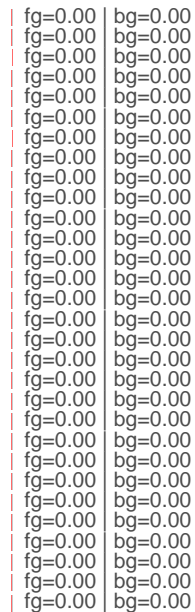


fraction genes in fg and bg



fraction

n=823/857 input genes with annotations