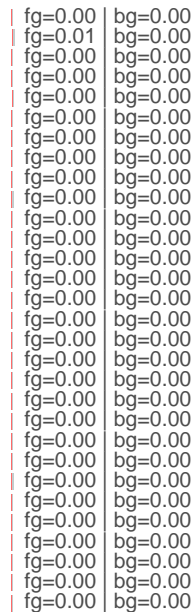


fraction genes in fg and bg



fraction

n=327/285 input genes with annotations