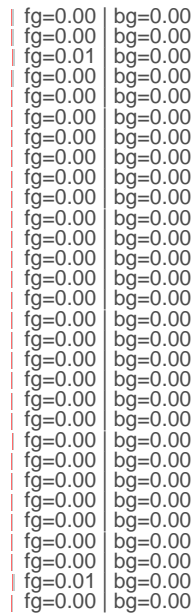


fraction genes in fg and bg



fraction

n=321/303 input genes with annotations