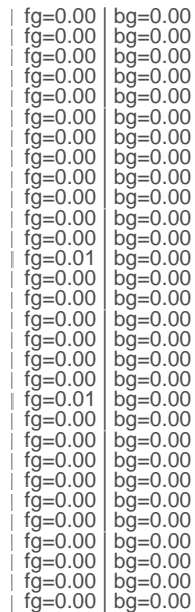


fraction genes in fg and bg



fraction

n=294/256 input genes with annotations