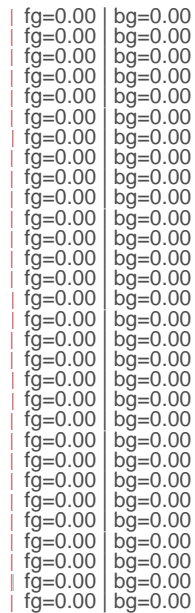


fraction genes in fg and bg



fraction

n=355/304 input genes with annotations