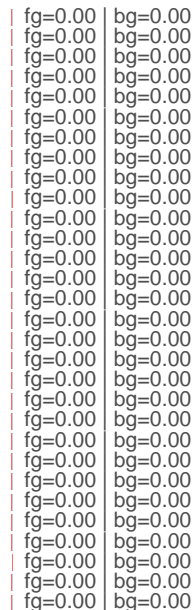


fraction genes in fg and bg



fraction

n=699/679 input genes with annotations