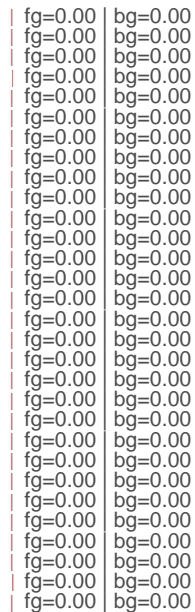


fraction genes in fg and bg



fraction

n=693/683 input genes with annotations