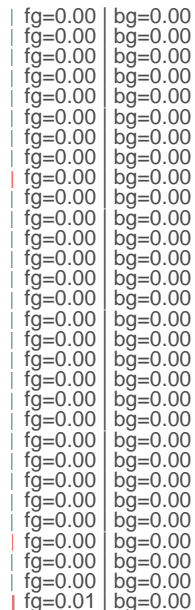


fraction genes in fg and bg



fraction

n=693/685 input genes with annotations