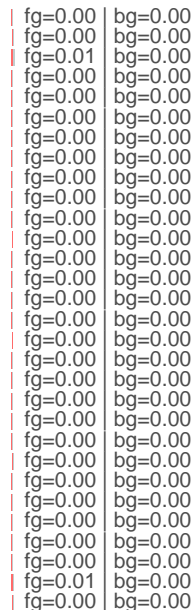


fraction genes in fg and bg



fraction

n=804/969 input genes with annotations