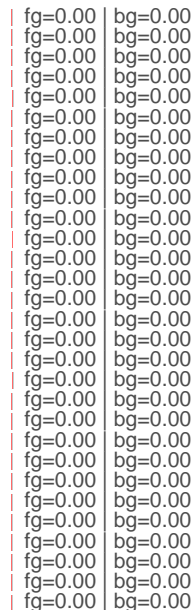


fraction genes in fg and bg



fraction

n=706/716 input genes with annotations