Input: .txt file with header, the first column is the gene ID, the 2-6 columns are the gene expression in each sample

g1：Group A



g2：Group B



Parameter:

1) Group color: default

2) X-axis title: None

3) Y-axis title: gene expression

4) Graph title: None

5) Scattered points: don’t display

Output:

1. Result file compression package



Contains two OS tool source files, PDF and PNG format result graphs, excel statistical result table



1. Result graph



1. Result table

