

# GANreproduce Quick Start Guide

Oct. 12, 2018

**Program** MATLAB (2016a for demonstration)

**Type** *Package (GANreproduce.m)*

**Title** *A package for reproducing the results reported in the manuscript entitled:*

*A trio of genes in germinal center B cells mediates the pathogenesis of  
blood cancers*

**Version** *1.4*

**Date** *2018/10/11*

**Authors** *Wei-Quan Fang and Ming-Jing Hwang*

**Maintainer** *Wei-Quan Fang <deleapoli@gmail.com>*

**Description** *In this work, we analyzed lymphoma-related gene expression and  
clinical data and identified a regulatory motif of germinal center  
B cell genes.*

**Repository** *GITHUB*

**Publication** *Submitted*

# Launch the Package

Three steps are needed to launch the *GANreproduce* Package.

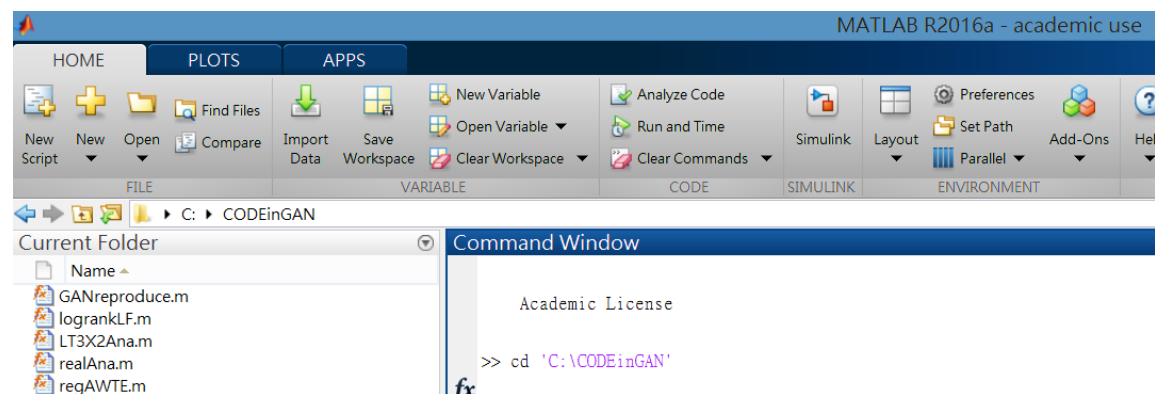
**Step1:** Download the two files, **DATAinGAN** and **CODEinGAN**, from the web-site <https://github.com/IBMSn121/GAN-paper1>.

GitHub, Inc. [US]   <a href="https://github.com/IBMSn121/GAN-paper1">https://github.com/IBMSn121/GAN-paper1</a>		
IBMSn121 Update README.md Latest commit 6f143b5 5 hours ago		
CODEinGAN	Create filenameCode.txt	6 hours ago
DATAinGAN	Create filenameData.txt	6 hours ago
LICENSE	Initial commit	3 months ago
README.md	Update README.md	5 hours ago

and put them in C:\

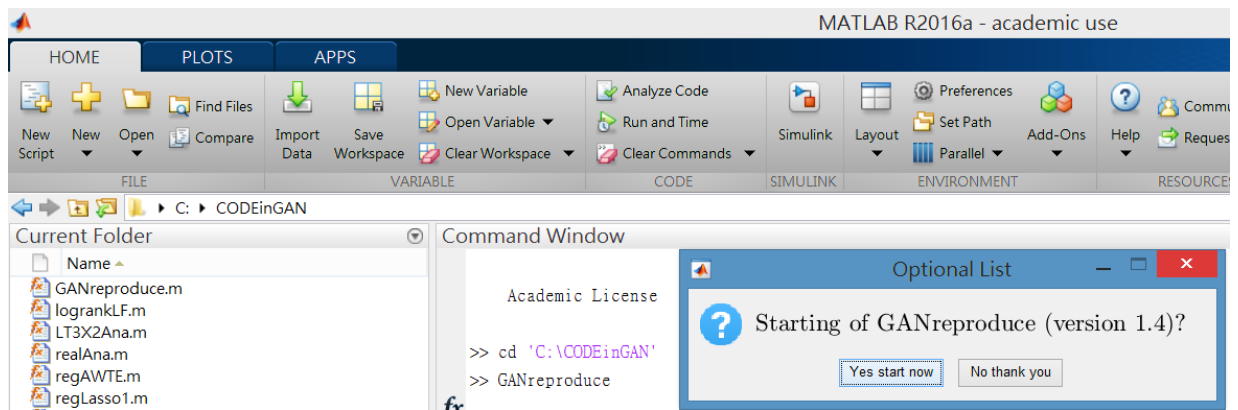
CODEinGAN	2018/10/1'
DATAinGAN	2018/10/9
Program Files	2017/5/18
Program Files (x86)	2018/5/17

**Step2:** Open MATLAB program window and type **cd 'C:\CODEinGAN'** in it, then press **ENTER**.



(*Please check* that GANreproduce Package **GANreproduce.m** can be seen in the MATLAB panel of **Current Folder** after this step)

**Step3:** Type **GANreproduce** and press **ENTER** to launch the GANreproduce Package and start *dialog boxes* for reproducing the results.

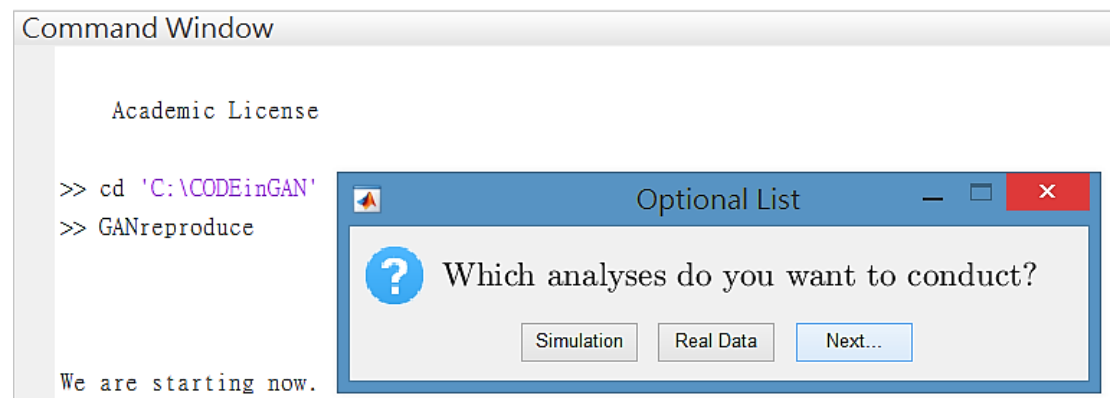


## Examples

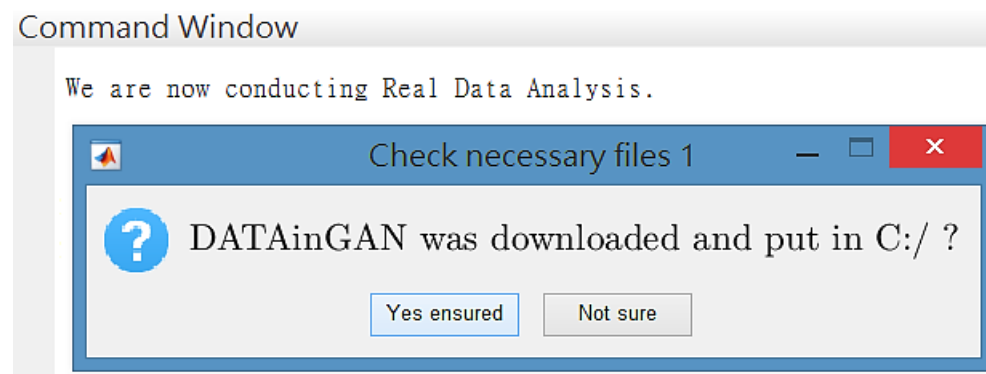
Three examples, concerning **Real Data Analysis**, **Linear Trend Analysis** and **Clinical Controversy Analysis**, were provided for reproducing the results via GANreproduce Package with selections of *user friendly dialog boxes*.

### Example 1 (Real Data Analysis)

Click **Yes start now** after the launch of the Package, then click **Real Data**.

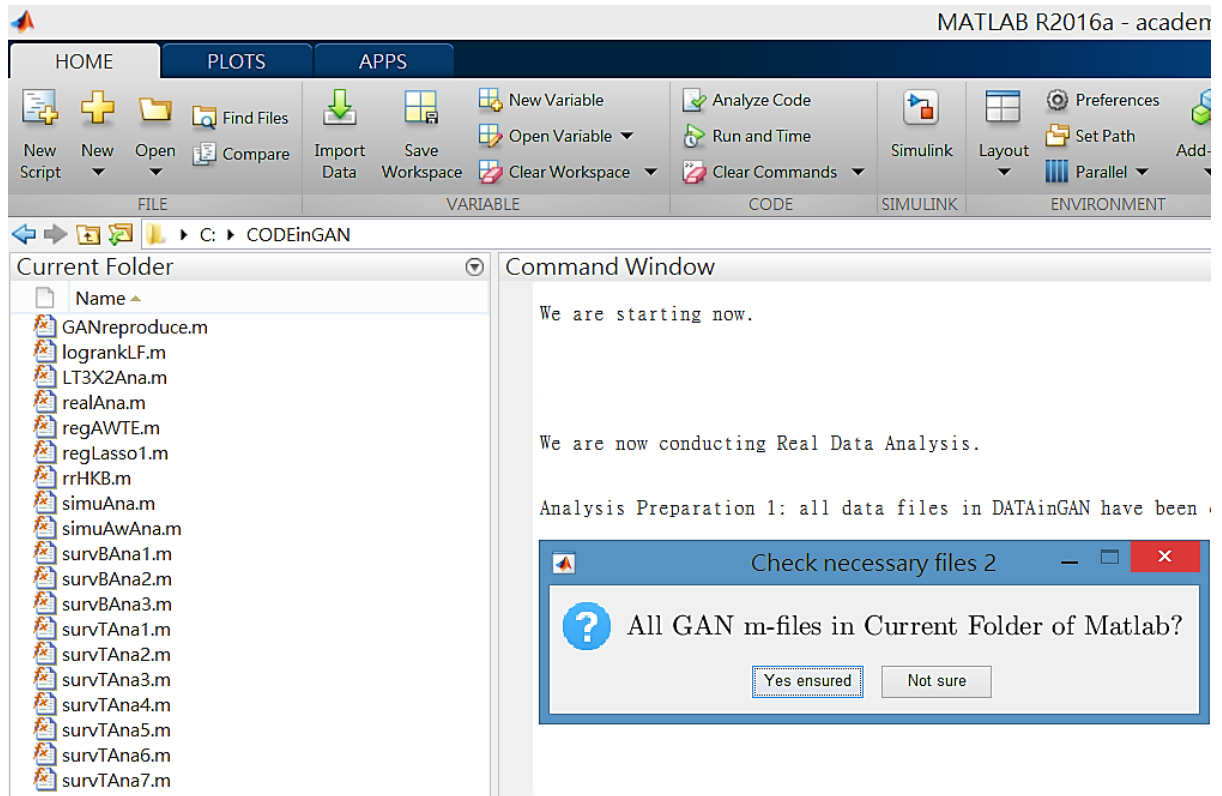


Ensure all the data in DATAinGAN were set up right and click **Yes ensured**.



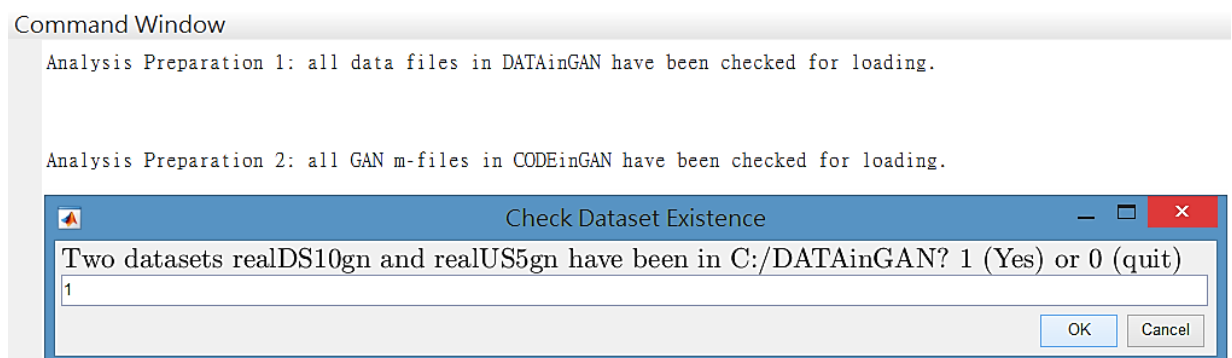


Ensure all the codes in CODEinGAN were set up right and click **Yes ensured**.



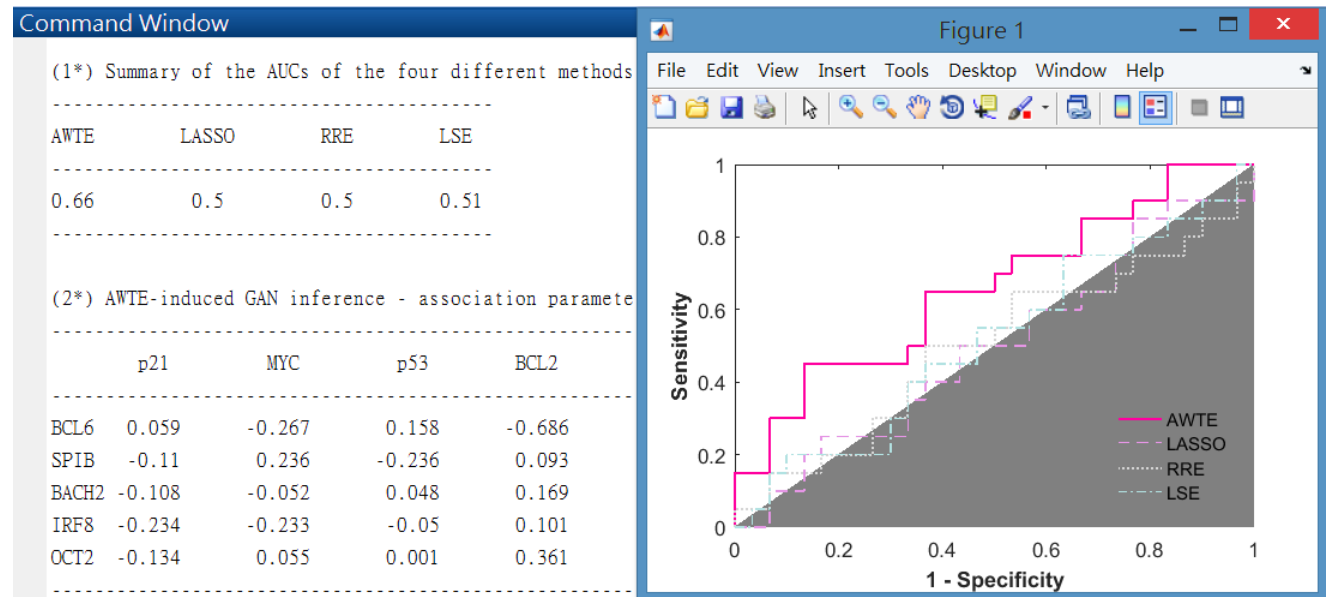
Check the two datasets realDS10gn and realUS5gn used in real data analysis

were set up right, then input **1** and click **Ok** to confirm the setting.





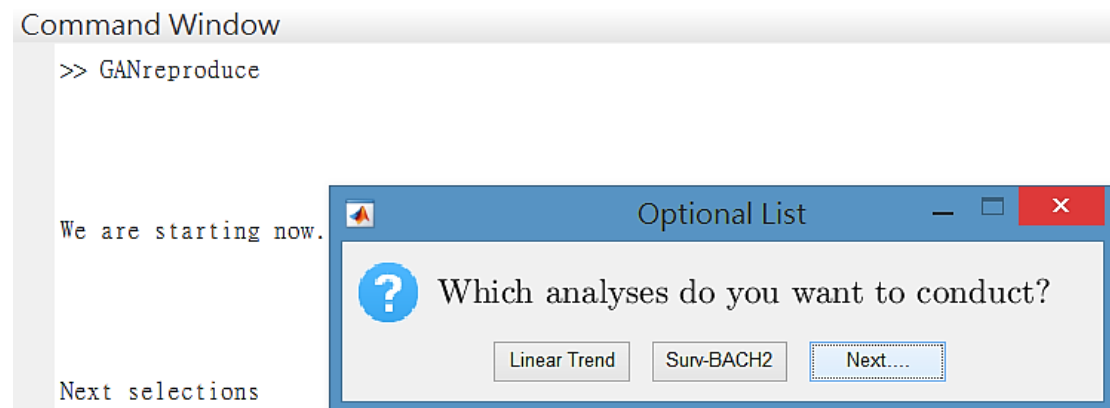
Results for Real Data Analysis were reproduced on the **Figure** and **MATLAB Command Window**.



### Example 2 (Linear Trend Analysis)

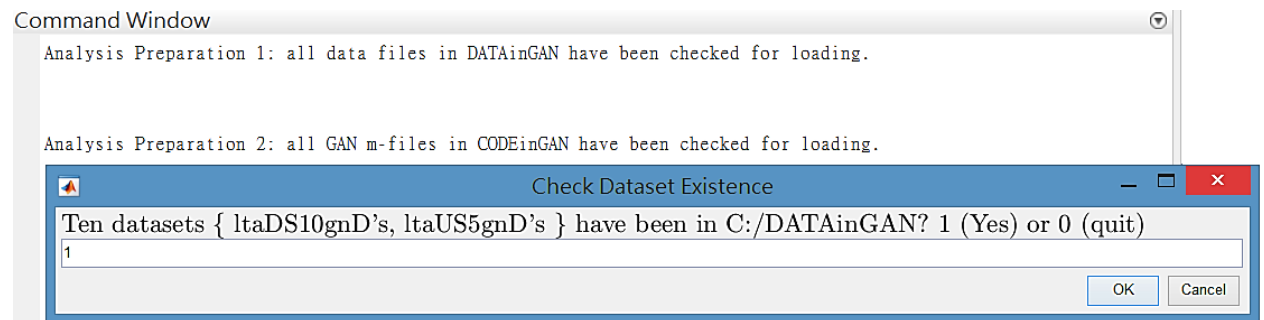
Click **Yes start now** after the launch of the Package, then click **Next...** and

**Linear Trend**.

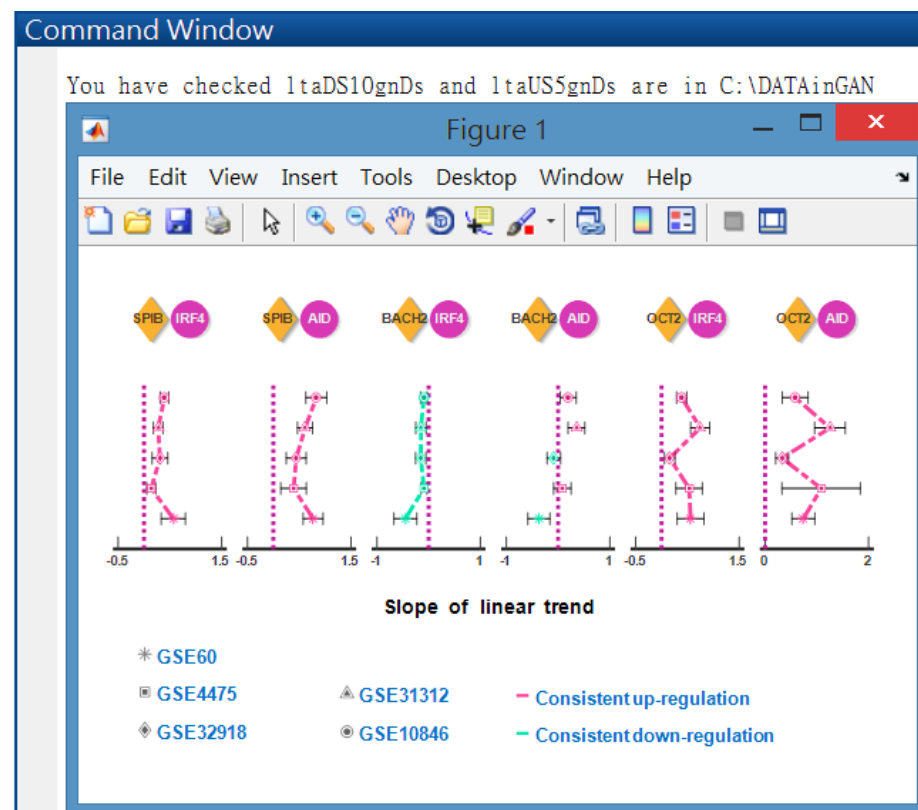




Click **Yes ensured** twice and check the necessary files, datasets and codes, as in Example 1, then input **1** and click **Ok** to confirm the setting.

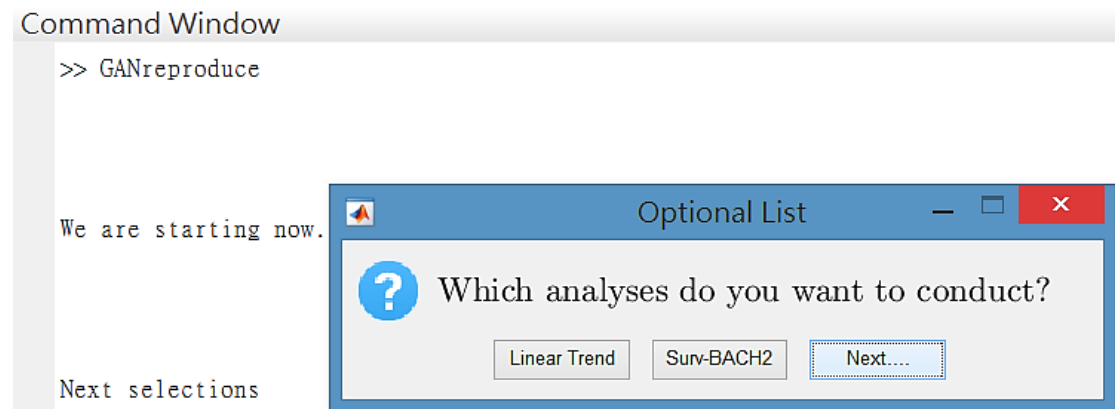


Results for Linear Trend Analysis were reproduced on the **Figure**.

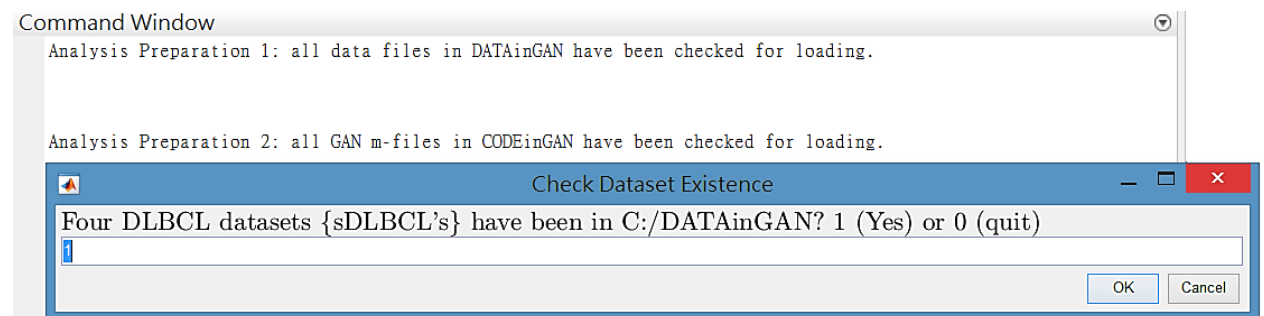


### Example 3 (Clinical Controversy Analysis)

Click **Yes start now** after the launch of the Package, then click **Next...** and **Surv-BACH2**.



Click **Yes ensured** twice and check the necessary files, datasets and codes, as in Example 1, then input **1** and click **Ok** to confirm the setting.

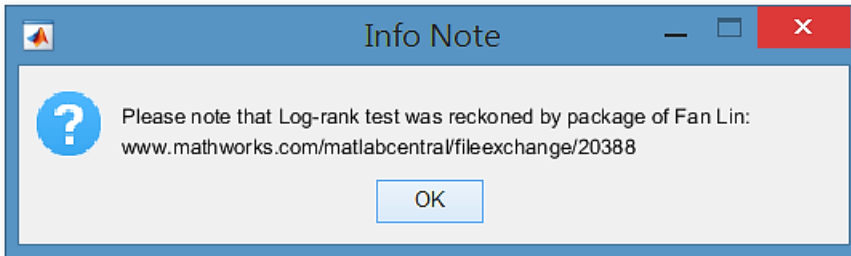


Receive the information on the package used for log-rank statistical testing and click **Ok**.



## Command Window

You have checked 4 DLBCL datasets are in C:\DATAinGAN

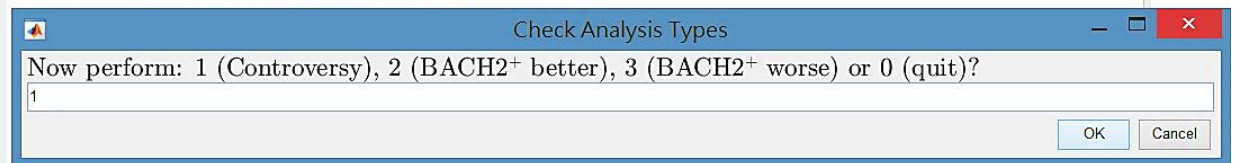


Select analytical type (e.g., Controversy), input **1** and click **Ok**.

## Command Window

You have checked 4 DLBCL datasets are in C:\DATAinGAN

Note that Log-rank test was reckoned by package of Fan Lin



Results for Clinical Controversy Analysis were reproduced as shown below.

## Command Window

You have checked 4 DLBCL datasets are in C:\D

Note that Log-rank test was reckoned by packa

Now we perform Clinical Controversy for BACH2

(1.1\*) Controversy of BACH2's role - GSE4475

Summary of the Number of Censored and Uncenso

GROUP	Total	Failed	Censored
X	124	59	65
Y	31	11	20
Total	155	70	85

Chi-square:1.0630

P:0.3025

