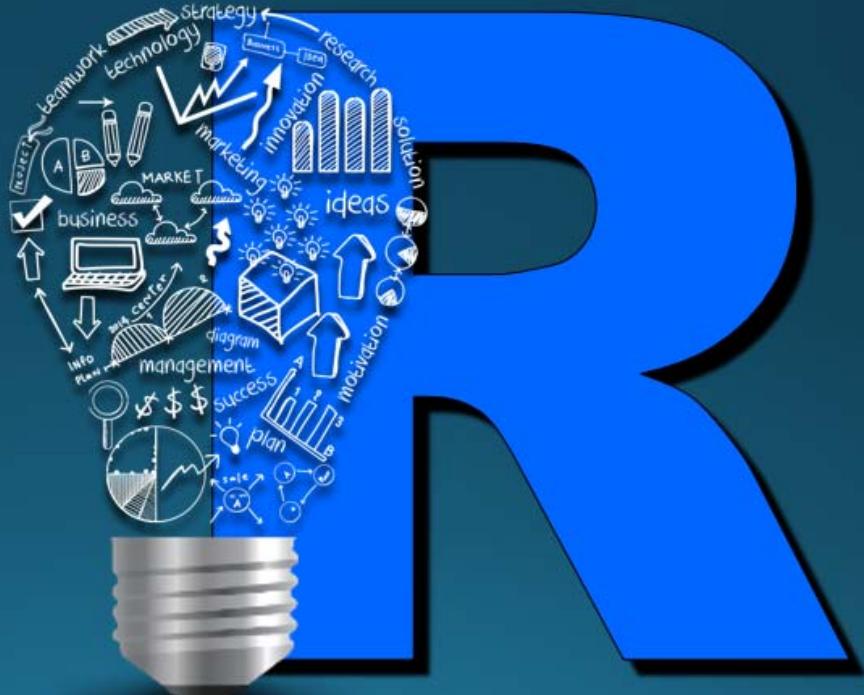


# R / RStudio 軟體環境介紹

吳漢銘  
國立政治大學 統計學系



<https://hmwu.idv.tw>



# 本章大綱&學習目標

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- R軟體環境簡介
  - R軟體安裝，R程式執行及套件(Package)
  - R相關檔案其副檔名的意義
  - 求助說明 `help()`、內訂Demo程式
- R程式IDE編輯器: RStudio
  - 建立專案、HELP、常用快速鍵
- 更新R軟體及R套件
- 常用查詢指令: `sessionInfo`, `.libPaths`, `R.Version`,  
`RStudio.Version`, `packageVersion`
- R的學習資源
- 安裝RStudio常見問題

# The R Project for Statistical Computing

<http://www.r-project.org>



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Developer Pages

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R Foundation

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Help With R

## The R Project for Statistical Computing

### Getting Started

R is a free software environment for statistical computing and graphics. It can be used on a variety of UNIX platforms, Windows and MacOS. To [download R](#), please choose a mirror.

If you have questions about R like how to download and install the software, or how to use it, please read our [answers to frequently asked questions](#) before you send an email to the [R help mailing list](#).

### News

- [R version 4.5.1 \(Great Square Root\)](#) has been released on 2025-06-13.
- [R version 4.5.0 \(How About a Twenty-Six\)](#) has been released on 2025-06-01.
- [R version 4.4.3 \(Trophy Case\)](#) (wrap-up of 4.4.x) was released on 2025-05-22.
- The [useR! 2025](#) conference will take place at Duke University, in Durham, NC, USA, on 2025-07-17 to 2025-07-20.
- We are deeply sorry to announce that our friend and colleague Friedrich (Fritz) Leisch has passed away. Please read our [tribute to Fritz here](#).
- You can support the R Foundation with a renewable subscription as a [supporter](#).

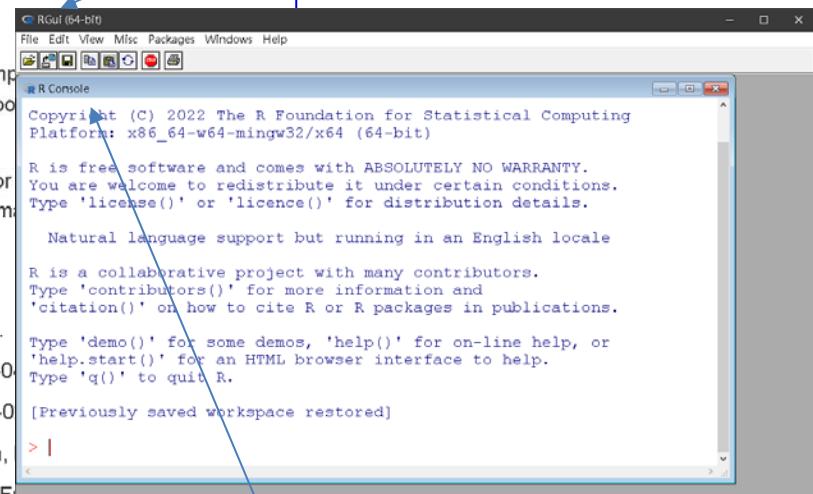
### News via Mastodon



R\_Foundation

R version 4.5.1 "Great Square Root" (source version) has been released. (You can find it in [cran.r-project.org/src/base/R-4/](https://cran.r-project.org/src/base/R-4/), or wait for CRAN to be updated.)  
Jun 13, 2025

Rgui



R Console



# R是什麼？

R is a language and environment for **statistical computing and graphics**.

- R 是一種程式語言與統計軟體環境，主要用於統計分析、資料探勘、機器學習與繪圖。
- 它最早(1985)由紐西蘭奧克蘭大學的 Ross Ihaka 和 Robert Gentleman 開發（名字 R 就來自他們的名字首字母）。
- R 有龐大的套件生態系 (CRAN)，研究人員、數據科學家和統計學家可以快速使用現成的套件做迴歸、分類、時間序列、文字探勘等分析。
- R 在學術界和產業界都非常普及，尤其是在需要大量統計建模和資料視覺化的領域。
- 簡單來說，R 就像是一個「專門為統計和資料分析打造的程式語言」，特別適合學術研究與數據處理。

## R 語言應用地圖

- 統計分析：迴歸、ANOVA、時間序列、貝葉斯模型
- 資料視覺化：ggplot2、互動式圖表、儀表板
- 機器學習：分類、分群、神經網路、隨機森林
- 商業與經濟分析：行銷數據、財務風險、顧客分析
- 生物資訊與醫學研究：基因資料、臨床試驗數據、生存分析
- 文字探勘與自然語言處理：文字雲、情感分析、語料庫研究
- 大數據與資料處理：SparkR、資料清理與轉換、大規模運算
- 學術研究與教育：教學、教材開發、學術論文分析

## 優點：

- 免費統計分析軟體' 完整的說明文件與討論區' 高品質的學術繪圖' 程式易根據需求修改

## 缺點：

- 不親切的使用者介面(Rgui)' 需詳知套件、函式名稱與程式編寫邏輯' 大量計算效能低。(已有改進方法)' its lack of security over the Web



# 我為什麼要使用R做為資料分析工具？

5/55

<http://www.r-project.org>

<https://posit.co/>

寫程式是資料分析的必要技能

<https://medium.com/datainpoint/9ee15b58cc>

Python or R, what should you learn first?

<https://read01.com/0ePnyD.html#.Wu66C3--kZY>

Why I use R for Data Science – An Ode to R

<https://www.r-bloggers.com/why-i-use-r-for-data-science-an-ode-to-r-2/>

選擇R開發數據分析平台的 4 個不錯的理由

<https://read01.com/660M4g.html>

做數據分析必須學R語言的4個理由

<https://read01.com/yyREB2.html>

Hadley Wickham : 一個改變了R的人

<https://read01.com/Mmy64J.html>

Hadley Wickham: "R is ... tailored to  
the problems of data science"



COPSS Presidents' Award: Hadley Wickham

SEPTEMBER 2, 2019

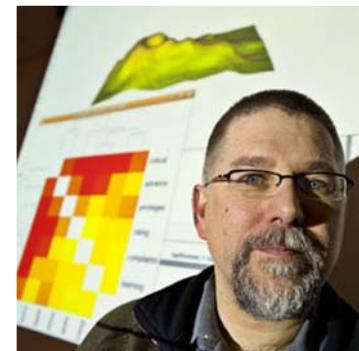
**Hadley Wickham wins the prestigious 2019 COPSS Presidents' Award**

Winners of the COPSS Presidents' Award [ edit ]

- 1981: Peter J. Bickel, University of California, Berkeley
- 1982: Stephen Fienberg, Carnegie Mellon University
- 1983: Tze Leung Lai, Stanford University
- 1984: David V. Hinkley, University of California, Santa Barbara
- 1985: James O. Berger, Duke University
- 1986: Ross L. Prentice, Fred Hutchinson Cancer Research Center
- 1987: C.F. Jeff Wu, Georgia Institute of Technology
- 1988: Raymond J. Carroll, Texas A&M University
- 1989: Peter Hall, Australian National University
- 1990: Peter McCullagh, University of Chicago
- 1991: Bernard Silverman, University of Oxford
- 1992: Nancy Reid, University of Toronto
- 1993: Wing Hung Wong, Stanford University
- 1994: David L. Donoho, Stanford University
- 1995: Iain M. Johnstone, Stanford University
- 1996: Robert J. Tibshirani, Stanford University
- 1997: Kathryn Roeder, Carnegie Mellon University
- 1998: Pascal Massart, Université de Paris-Sud
- 1999: Larry A. Wasserman, Carnegie Mellon University
- 2000: Jianqing Fan, Princeton University
- 2001: Xiao-Li Meng, Harvard University
- 2002: Jun Liu, Harvard University
- 2003: Andrew Gelman, Columbia University
- 2004: Michael A. Newton, University of Wisconsin
- 2005: Mark J. van der Laan, University of California, Berkeley
- 2006: Xihong Lin, Harvard University
- 2007: Jeff Rosenthal, University of Toronto
- 2008: T. Tony Cai, University of Pennsylvania
- 2009: Rafael Irizarry, Harvard University
- 2010: David Dunson, Duke University
- 2011: Nilanjana Chatterjee, Johns Hopkins University
- 2012: Samuel Kou, Harvard University
- 2013: Marc A. Suchard, UCLA
- 2014: Martin J. Wainwright, University of California, Berkeley
- 2015: John D. Storey, Princeton University
- 2016: Nicolai Meinshausen, ETH Zürich
- 2017: Tyler J. VanderWeele, Harvard University
- 2018: Richard J. Samworth, University of Cambridge
- 2019: Hadley Wickham, RStudio, Inc.
- 2020: Rina Foygel Barber, University of Chicago
- 2021: Jeffrey T. Leek, Johns Hopkins University
- 2022: Daniela Witten, University of Washington

Robert C. Gentleman

Department of Biostatistical Sciences,  
Dana Farber Cancer Institute



Ross Ihaka,

University of Auckland, 2017

# TIOBE 全球程式語言排名



<http://www.tiobe.com/tiobe-index/>

(共243種程式語言)

Aug 2025	Aug 2024	Change	Programming Language
1	1		Python
2	2		C++
3	3		C
4	4		Java
5	5		C#
6	6		JavaScript
7	8	▲	Visual Basic
8	9	▲	Go
9	25	▲	Perl
10	12	▲	Delphi/Object Pascal
11	10	▼	Fortran
12	7	▼	SQL
13	30	▲	Ada
14	19	▲	R
15	13	▼	PHP
16	11	▼	MATLAB
17	20	▲	Scratch
18	14	▼	Rust
19	18	▼	Kotlin
20	17	▼	Assembly language



# Vibe Coding

<https://www.bnnext.com.tw/article/82704/how-to-vibe-coding-2025>

**Vibe Coding是什麼？**

隨著生成式AI進入爆發生長時期，2025年初自美國矽谷颳起一股軟體開發新風潮，透過AI工具協助，軟體工程師可以在「不寫程式碼」的前提下開發產品，而這種近乎佛系、不用刻苦寫code的開發方式，被取了一個新穎的名字：**Vibe Coding**。

在Vibe Coding風潮中，其中Anthropic旗下的Claude系列模型被認為是擁有最強大寫程式能力的AI模型。近期外媒也訪問Anthropic高層，揭露其RLAIF學習技術、加強使用工具及記憶能力是勝出關鍵。

而除了Claude系列，好用的Vibe Coding工具有哪些？



<https://www.youtube.com/watch?v=8me0juJCpWM&t=40s>



# 軟體外觀 (Windows)

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RGui (64-bit)

File History Resize Windows

R Console

Type 'q()' to quit R.

[Previously saved workspace restored]

```
> names(iris)
[1] "Sepal.Length" "Sepal.Width"  "Petal.Length"
[4] "Petal.Width"   "Species"
> View(iris)
> attach(iris)
> plot(Sepal.Length, Sepal.Width, col = Species)
> x11()
> hist(Sepal.Length)
>
```

Data:iris

	Sepal.Length	Sepal.Width	Petal.Length	Petal.Width
1	5.1	3.5	1.4	0.2
2	4.9	3.0	1.4	0.2
3	4.7	3.2	1.3	0.2
4	4.6	3.1	1.5	0.2
5	5.0	3.6	1.4	0.2
6	5.4	3.9	1.7	0.4
7	4.6	3.4	1.4	0.3
8	5.0	3.4	1.5	0.2
9	4.4	2.9	1.4	0.2
10	4.9	3.1	1.5	0.1
11	5.4	3.7	1.5	0.2
12	4.8	3.4	1.6	0.2
13	4.8	3.0	1.4	0.1

R Graphics: Device 2 (inactive)

R Graphics: Device 3 (ACTIVE)

Histogram of Sepal.Length

# CRAN: The Comprehensive R Archive Network



New Zealand

<https://cran.stat.auckland.ac.nz/>

Norway

<https://cran.uib.no/>

Philippines

<https://cran.stat.upd.edu.ph/>

Poland

<https://cran.mi2.ai/>

Portugal

<https://cran.radicaldevelop.com/>

Russia

<https://mirror.truenetwork.ru/CRAN/>

South Africa

<https://cran.mirror.ac.za/>

Spain

<https://ftp.cixug.es/CRAN/>

<https://cran.rediris.es/>

Sweden

<https://ftp.acc.umu.se/mirror/CRAN/>

Switzerland

<https://stat.ethz.ch/CRAN/>

<https://mirror.metanet.ch/cran/>

Taiwan

<https://cran.csie.ntu.edu.tw/>

University of Auckland

University of Bergen

University of the Philippines and PREGINET

MI2.ai, Warsaw University of Technology

The Comprehensive R Archive Network

**Download and Install R**

Precompiled binary distributions of the base system and contributed packages, **Windows and Mac** users most likely want one of these versions of R:

- [Download R for Linux \(Debian, Fedora/Redhat, Ubuntu\)](#)
- [Download R for macOS](#)
- [Download R for Windows](#)

R is part of many Linux distributions, you should check with your Linux package management system in addition to the link above.

**Source Code for all Platforms**

Windows and Mac users most likely want to download the precompiled binaries listed in the upper box, not the source code. The sources have to be compiled before you can use them. If you do not know what this means, you probably do not want to do it!

- The latest release (2025-06-13, Great Square Root) [R-4.5.1.tar.gz](#), read [what's new](#) in the latest version.
- The CRAN directory [src/base-prerelease](#) contains R alpha, beta, and rc releases as daily snapshots in time periods before a planned release.
- Between releases, the same directory [src/base-prerelease](#) contains snapshots of current patched and development versions. Please read about [new features and bug fixes](#) before filing corresponding feature requests or bug reports.
- Alternatively, daily snapshots are [available here](#).
- Source code of older versions of R is [available here](#).
- Contributed extension [packages](#).

`repos="https://cran.csie.ntu.edu.tw/"`

<https://hmwu.idv.tw>



# R軟體下載

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The Comprehensive R Archive Network

Download and Install R

Precompiled binary distributions of the base system and contributed packages, Windows and Mac users most likely want one of these versions of R:

- [Download R for Linux \(Debian, Fedora/Redhat, Ubuntu\)](#)
- [Download R for macOS](#)
- [Download R for Windows](#)

R is part of many Linux distributions, you should check with your Linux package management system in addition to the link above.

Source Code for all Platforms

Windows and Mac users most likely want to download the precompiled binaries listed in the upper box, not the source code. The sources have to be compiled before you can use them. If you do not know what this means, you probably do not want to do it!

**R-4.5.1 for Windows**

[Download R-4.5.1 for Windows](#) (86 megabytes, 64 bit)

[README on the Windows binary distribution](#)

[New features in this version](#)

<https://cran.csie.ntu.edu.tw/bin/windows/base/R-4.2.2-win.exe>

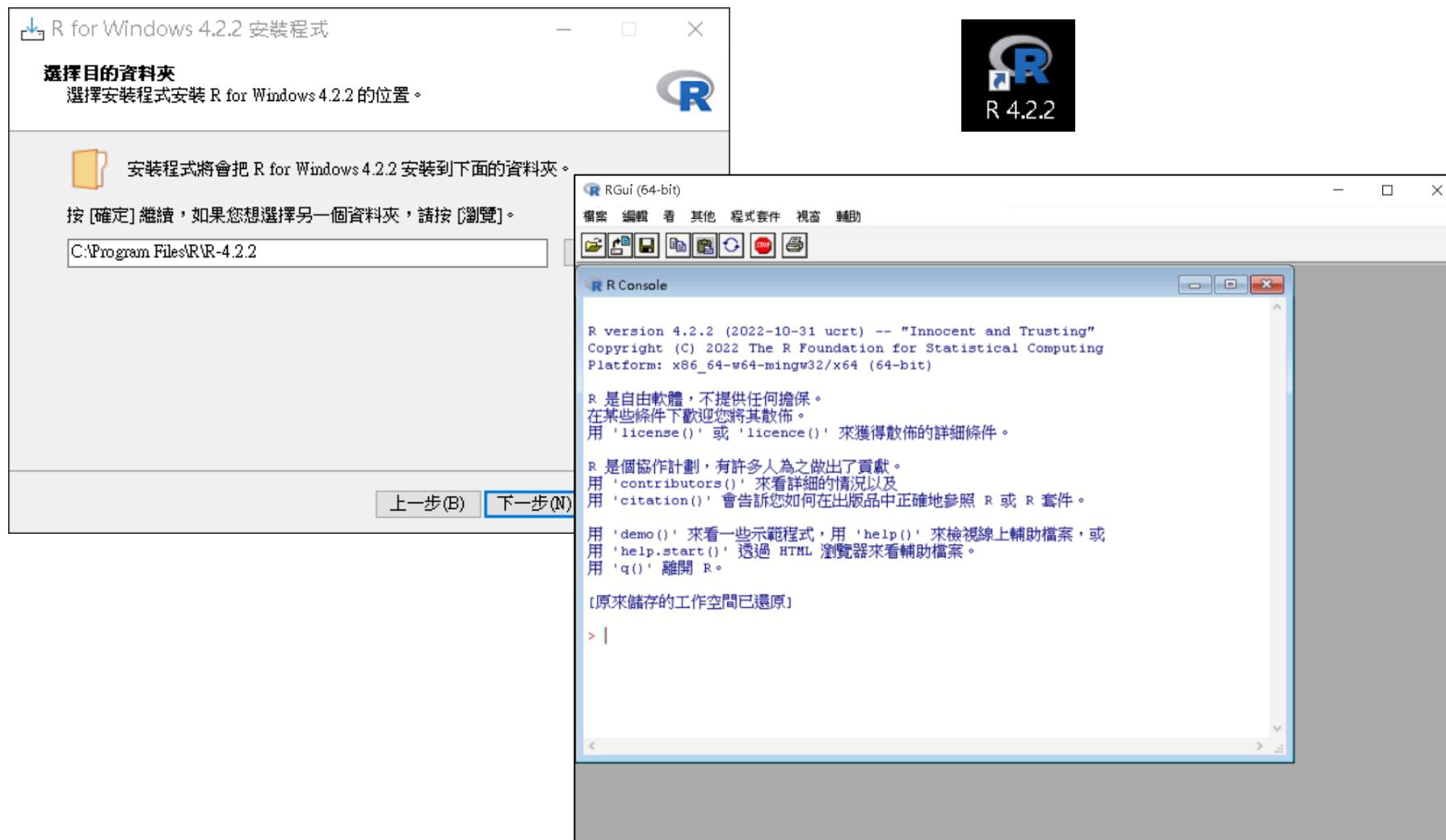
<https://hmwu.idv.tw>



# 軟體安裝；啟動R

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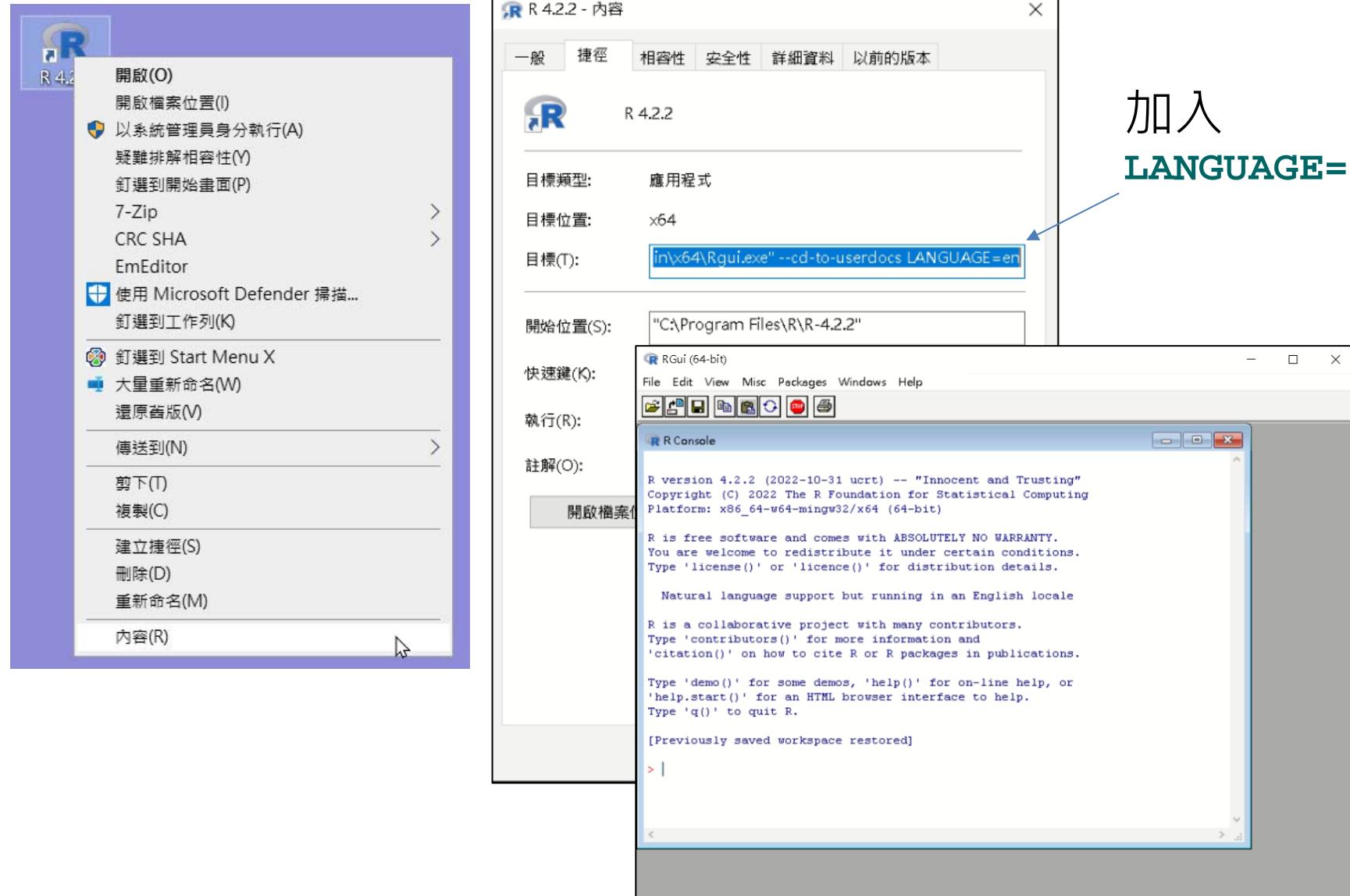
注意: R軟體安裝目錄，使用者目錄、R套件安裝目錄的路徑





# 想把中文介面改成英文介面

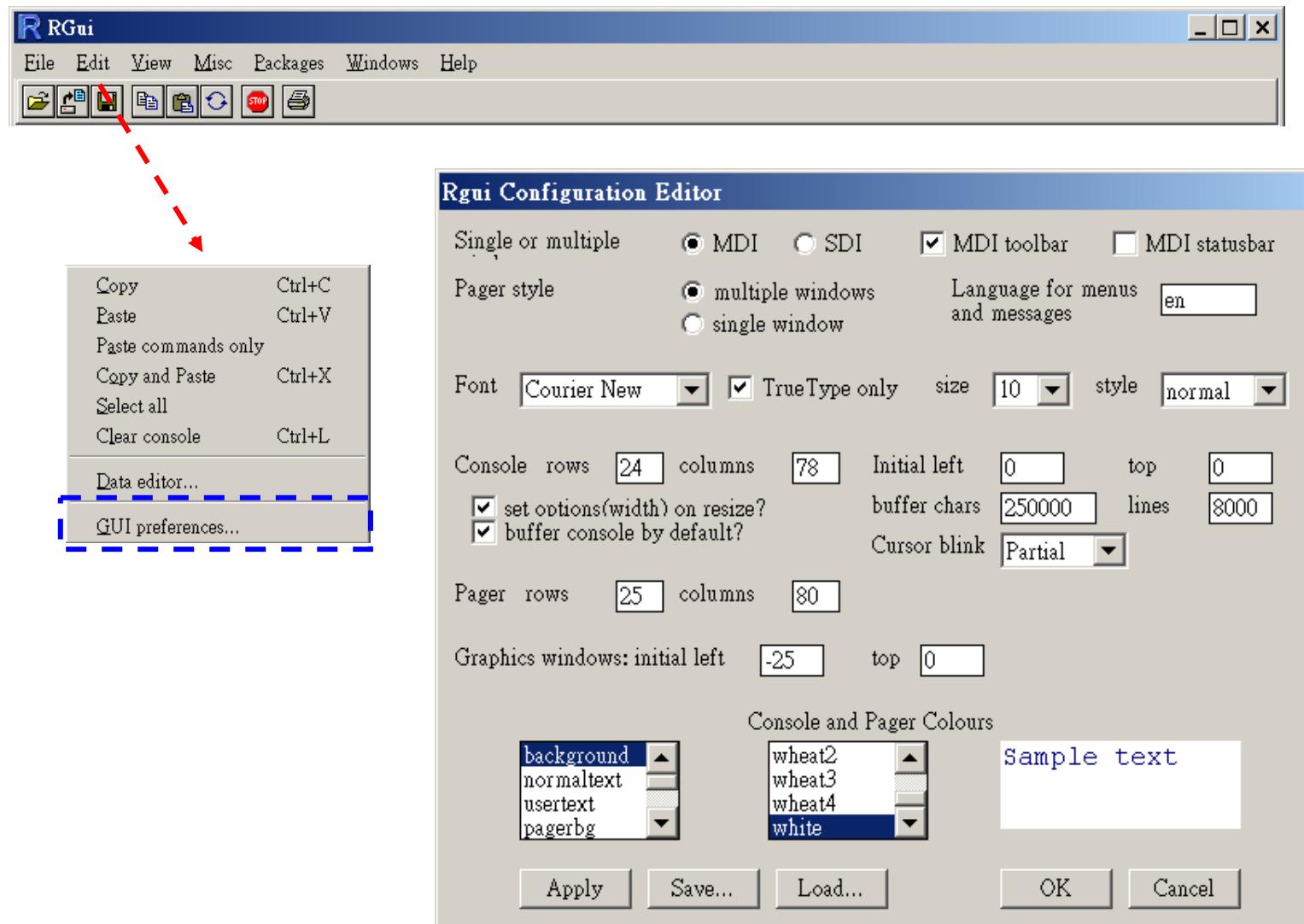
12/55



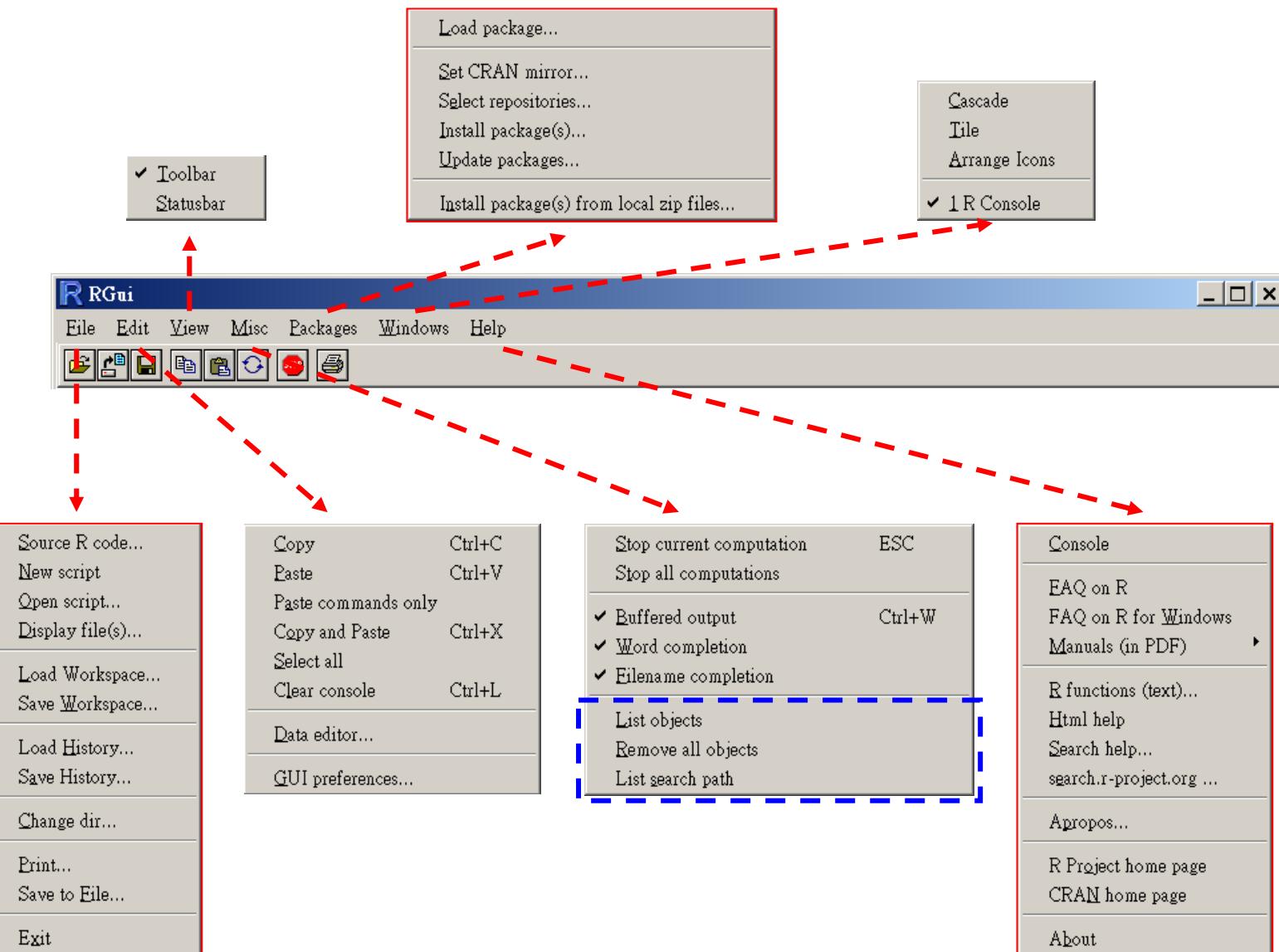


# 喜好設定

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# R 選單



# 寫程式及執行

The screenshot shows the RGui (64-bit) interface with three callouts illustrating different ways to run R code:

- (1) 執行程式**: A red arrow points from the "Run line or selection Ctrl+R" option in the context menu of the R Editor to the R console window.
- (2) 執行程式**: A red arrow points from the "Save Ctrl+S" option in the File menu to the R Editor window.
- (3) 利用source 執行程式**: A red arrow points from the "> source("myRcode.R")" command in the R console to the "Save as" option in the File menu.

**注意工作目錄及print**

**R Gui (64-bit)**

File Edit View Misc Packages Windows Help

New script

Open script...

Display file(s)...

Load Workspace...

Save Workspace...

Load History...

Save History...

Change dir...

Print...

Save to File...

Exit

Type 'demo()' for some demos, 'help()' for help, 'help.start()' for an HTML browser i

Type 'q()' to quit R.

```

> x <- 1+4
> x
[1] 5
> (y <- 1:100)
[1] 1 2 3 4 5 6 7
[19] 19 20 21 22 23 24 25 2
[37] 37 38 39 40 41 42 43 4
[55] 55 56 57 58 59 60 61 6
[73] 73 74 75 76 77 78 79 8
[91] 91 92 93 94 95 96 97 9
> sum(y)
[1] 5050
> sample(1:42, 6)
[1] 4 11 39 27 35 10
>

```

**R Untitled - R Editor**

```

##### My testing R code #####
# Han-Ming Wu
# 2024/07/16
#####
cat("Use iris data to test")
data(iris)
View(iris)
head(iris, 3)
summary(iris)
attach(iris)
plot(Sepal.Length, Sepal.Width, col=Species)

```

Run line or selection Ctrl+R

Undo Ctrl+Z

Cut Ctrl+X

Copy Ctrl+C

Paste Ctrl+V

Delete

Select all Ctrl+A

> source("myRcode.R")

**R Gui (64-bit)**

File Edit Packages Windows Help

New script Ctrl+N

Open script... Ctrl+O

Save Ctrl+S

Save as

Print...

Close Ctrl+P



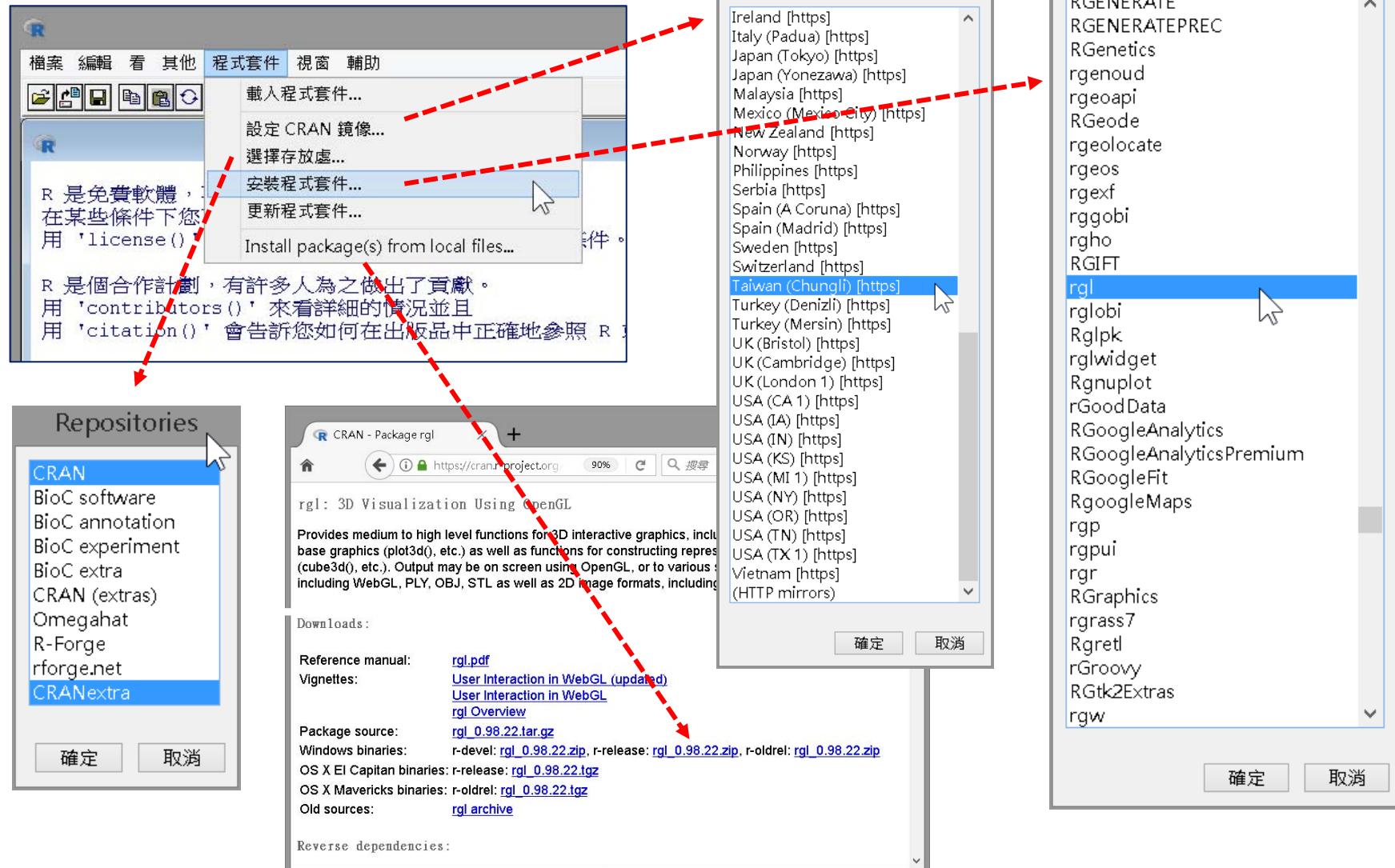
# R相關檔案副檔名

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- **.R** : 標準 R 程式碼檔案
- **.Rhistory** : 指令操作紀錄檔
- **.Rprofile** : 啟動時自動執行的設定檔
- **.RData** : 整個工作環境 ( workspace )
- **.rda / .Rda** : 多個 R 物件
- **.rds** : 單一 R 物件 ( 資料交換常用 )
- **.tar.gz** : Linux/Unix R 套件壓縮檔
- **.zip** : Windows R 套件壓縮檔
- **.dll / .so** : 編譯後的函式庫
- **.Rmd** : R Markdown 文件
- **.Rnw** : Sweave ( R + LaTeX 文件 )
- **.Renvironment** : R 環境變數設定檔
- **.Rproj** : RStudio 專案檔
- **.Rd** : R 套件文件檔 ( help )

# 安裝R套件 (RGui選單)

- Packages => Install package(s) => CRAN mirror => installr, rgl, scatterplot3d (按Ctrl可多重選取) => OK





# 安裝R套件 (指令方式)

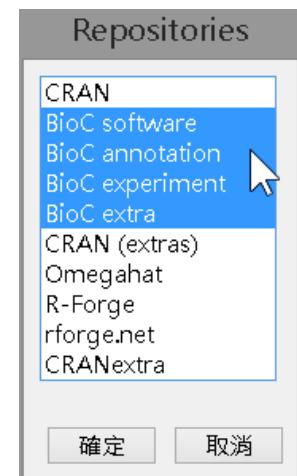
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```
> setRepositories()  
> chooseCRANmirror()  
> install.packages("rgl") # 安裝套件  
> library(rgl) # 載入套件  
> # same as > library("rgl")  
> detach("package:rgl", unload=TRUE) # 卸載套件  
> remove.packages("rgl") # 移除套件  
> .libPaths() # 列出library安裝目錄
```

```
> # 指定repositories  
> install.packages("rgl", repos = "http://cran.csie.ntu.edu.tw")
```

```
> options(repos="http://cran.csie.ntu.edu.tw")  
> pks <- c("rgl", "ggplot2", "maps", "e1071")  
> install.packages(pks)  
> lapply(pks, library, character.only = TRUE)
```

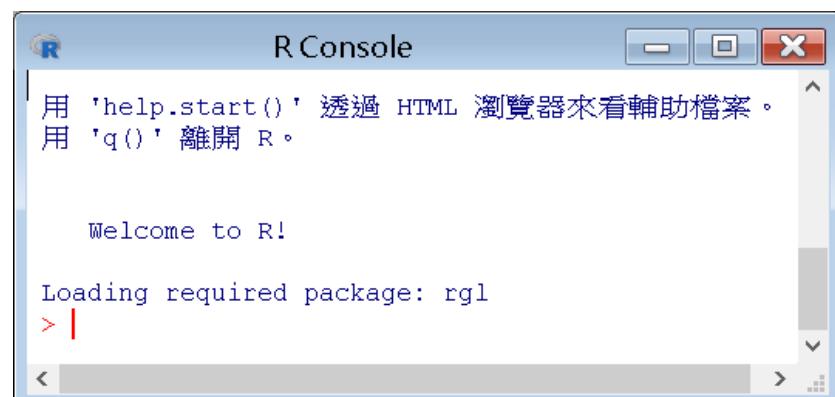
```
> # 安裝 BioC 的套件  
> # setRepositories(graphics = F, ind = 2)  
> # options("repos")  
> source("https://bioconductor.org/biocLite.R")  
> biocLite("EBIImage")
```



## Automatically load packages in the Startup of R:

- call `.First()` in the `Rprofile` file.
- 編輯檔案(Windows) C:\Program Files\R\R-3.3.1\library\base\Rprofile

```
.First <- function(){  
  cat("\n  Welcome to R!\n\n")  
  require(rgl)  
}
```





# 安裝R套件 (RStudio選單)

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The screenshot shows the RStudio interface. The 'Tools' menu is open, and the 'Install Packages...' option is highlighted. A modal dialog titled 'Install Packages' is displayed, containing fields for 'Install from:' (set to 'Repository (CRAN, CRANextra)'), 'Packages' (containing 'rgl scatterplot3d ggplot2'), 'Install to Library:' (set to 'C:/Users/userpc/Documents/R/win-library/3.4 [Default]'), and a checked 'Install dependencies' option. At the bottom are 'Install' and 'Cancel' buttons.

```
Console Terminal x
E:\06-NTPU_Course_Now\106-1-高維度資料分析\小考\106-1-HDDA-exam2/
downloaded 302 KB

trying URL 'https://cran.rstudio.com/bin/windows/contrib/3.4/ggplot2_2.2.1.zip'
Content type 'application/zip' length 2784537 bytes (2.7 MB)
downloaded 2.7 MB

package 'rgl' successfully unpacked and MD5 sums checked
package 'scatterplot3d' successfully unpacked and MD5 sums checked
package 'ggplot2' successfully unpacked and MD5 sums checked

The downloaded binary packages are in
  C:\Users\userpc\AppData\Local\Temp\Rtmpwz9041\downloaded_packages
> |
```

```
Rcode - RStudio
File Edit Code View Plots Session Build Debug Profile Tools
+ + - Addins Go to file/function Addins
rcode.R x test.R x
Source on Save
! Packages c1Valid and rgl required but are not installed. Install Don't Show Again
1 library(rgl)
2 library(c1Valid)
3
```



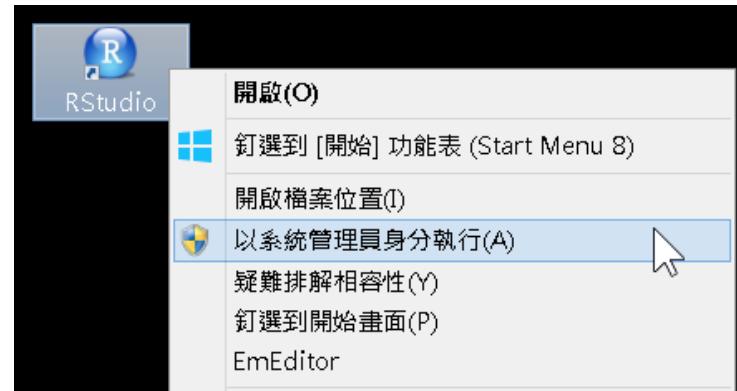
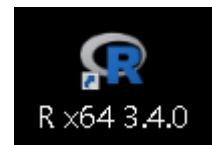
# 安裝R套件常見問題

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```
Console Terminal x
~/Documents/my first R/ ↵
x/el-capitan/contrib/3.4/rgl_0.98.22.tgz'
Warning in install.packages :
  URL 'https://cran.rstudio.com/bin/macosx/e
l-capitan/contrib/3.4/rgl_0.98.22.tgz': stat
us was 'Couldn't connect to server'
Error in download.file(url, destfile, method
, mode = "wb", ...):
  無法開啟 URL 'https://cran.rstudio.com/bin/
macosx/el-capitan/contrib/3.4/rgl_0.98.22.tg
z'
Warning in install.packages :
  download of package 'rgl' failed
> install.packages("rgl")
嘗試 URL 'https://cran.rstudio.com/bin/macos
x/el-capitan/contrib/3.4/rgl_0.98.22.tgz'
Content type 'application/x-gzip' length 383
0751 bytes (3.7 MB)
=====
=====

downloaded 3.7 MB

The downloaded binary packages are in
/var/folders/zb/lqb9zm_j0bdgmfj24glp38dh0000
gn/T//Rtmp006ZKM/downloaded_packages
> |
```



Installing R on OS X

<https://www.r-bloggers.com/installing-r-on-os-x/>

Why mac OS failed to install rgl: rgl package needs X11 to show its 3D plots.

- (1) Download/install XQuartz <https://www.xquartz.org>
- (2) Check the library /usr/X11/lib/libGLU.1.dylib is loaded
- (3) install.packages("rgl")



# library 和 require 的差別

21/55

- 狀況(1): 載入已安裝的套件，且 using the command outside of a function definition，則：
  - "require" or "library" 相同。
- 狀況(2): 載入未安裝的套件，則：
  - **library(foo)**: 程式停止 with the message "**Error in library(foo): there is no package called 'foo'.**"
  - **require(foo)**: get a warning, but not an error. Your program will continue to run, only to crash later when you try to use a function from the library "**foo**"
- **require(package)** returns (invisibly) TRUE if the package is available

```
> # install "package" if it doesn't exist, and then load it.  
> if (!require(package)) install.packages("package")  
> library(package)
```



# CRAN Task Views

<https://cran.r-project.org/web/views/>

<a href="#">ActuarialScience</a>	Actuarial Science
<a href="#">Agriculture</a>	Agricultural Science
<a href="#">Bayesian</a>	Bayesian Inference
<a href="#">CausalInference</a>	Causal Inference
<a href="#">ChemPhys</a>	Chemometrics and Computational Physics
<a href="#">ClinicalTrials</a>	Clinical Trial Design, Monitoring, and Analysis
<a href="#">Cluster</a>	Cluster Analysis & Finite Mixture Models
<a href="#">CompositionalData</a>	Compositional Data Analysis
<a href="#">Databases</a>	Databases with R
<a href="#">DifferentialEquations</a>	Differential Equations
<a href="#">Distributions</a>	Probability Distributions
<a href="#">DynamicVisualizations</a>	Dynamic Visualizations and Interactive Graphics
<a href="#">Econometrics</a>	Econometrics
<a href="#">Environmetrics</a>	Analysis of Ecological and Environmental Data
<a href="#">Epidemiology</a>	Epidemiology
<a href="#">ExperimentalDesign</a>	Design of Experiments (DoE) & Analysis of Experimental Data
<a href="#">ExtremeValue</a>	Extreme Value Analysis
<a href="#">Finance</a>	Empirical Finance
<a href="#">FunctionalData</a>	Functional Data Analysis
<a href="#">GraphicalModels</a>	Graphical Models
<a href="#">HighPerformanceComputing</a>	High-Performance and Parallel Computing with R
<a href="#">Hydrology</a>	Hydrological Data and Modeling
<a href="#">MachineLearning</a>	Machine Learning & Statistical Learning
<a href="#">MedicalImaging</a>	Medical Image Analysis
<a href="#">MetaAnalysis</a>	Meta-Analysis
<a href="#">MissingData</a>	Missing Data
<a href="#">MixedModels</a>	Mixed, Multilevel, and Hierarchical Models in R
<a href="#">ModelDeployment</a>	Model Deployment with R
<a href="#">NaturalLanguageProcessing</a>	Natural Language Processing
<a href="#">NetworkAnalysis</a>	Network Analysis
<a href="#">NumericalMathematics</a>	Numerical Mathematics
<a href="#">OfficialStatistics</a>	Official Statistics & Survey Statistics
<a href="#">Omics</a>	Genomics, Proteomics, Metabolomics, Transcriptomics, and Other Omics
<a href="#">Optimization</a>	Optimization and Mathematical Programming
<a href="#">Paleontology</a>	Paleontology
<a href="#">Pharmacokinetics</a>	Analysis of Pharmacokinetic Data
<a href="#">Phylogenetics</a>	Phylogenetics
<a href="#">Psychometrics</a>	Psychometric Models and Methods
<a href="#">ReproducibleResearch</a>	Reproducible Research
<a href="#">Robust</a>	Robust Statistical Methods
<a href="#">Spatial</a>	Analysis of Spatial Data
<a href="#">SpatioTemporal</a>	Handling and Analyzing Spatio-Temporal Data
<a href="#">SportsAnalytics</a>	Sports Analytics
<a href="#">Survival</a>	Survival Analysis
<a href="#">TeachingStatistics</a>	Teaching Statistics
<a href="#">TimeSeries</a>	Time Series Analysis
<a href="#">Tracking</a>	Processing and Analysis of Tracking Data
<a href="#">WebTechnologies</a>	Web Technologies and Services

```
> install.packages("ctv")
> library(ctv)
>
> available.views()
```

## CRAN Task Views

---

Name:

**Bayesian**

Topic: Bayesian Inference

Maintainer: Jong Hee Park

Repository: <https://cran.fhcrc.org>

...

```
> ## install MachineLearning view
> install.views("MachineLearning")
> ## only with core packages
> install.views("MachineLearning",
coreOnly = TRUE)
> ## update MachineLearning view
> update.views("MachineLearning")
```

What is the difference between **require()** and **library()**? Not much difference in everyday work. **require** is used inside functions, as it outputs a warning and continues if the package is not found, whereas **library** will throw an error.



# Popular Packages

- Top 25 R Packages (You Need To Learn In 2024) | R-bloggers

The screenshot shows the R-bloggers homepage with a navigation bar including 'HOME', 'ABOUT', 'RSS', 'ADD YOUR BLOG!', 'LEARN R', 'R JOBS', and 'CONTACT US'. The main content area features a dark blue header with the title 'Top 25 R Packages (You Need To Learn In 2024)' in white. Below the title, a sub-header reads 'Posted on August 25, 2024 by Business Science in R bloggers | 0 Comments'.

<https://www.r-bloggers.com/2024/08/top-25-r-packages-you-need-to-learn-in-2024/>

- Top 15 R Libraries for Data Science in 2023

<https://www.knowledgehut.com/blog/data-science/top-r-libraries-for-data-science>

dplyr, tidyr, readr, stringr, lubridate, jsonlite, Shiny, tseries, Prophet, RColorBrewer, githubinstall, ggmap, sqldf, caret

- Quick list of useful R packages

<https://support.posit.co/hc/en-us/articles/201057987-Quick-list-of-useful-R-packages>

Categories: To Flood data, To manipulate data, To visualize data, To model data, To report results, For Spatial data, For Time Series and Financial data, To write high performance R code, To work with the web, To write your own R packages

# <https://github.com/qinwf/awesome-R>



qinwf / awesome-R Public

Code Issues 4 Pull requests 15 Actions Projects Security ...

master Go to file Code

qinwf Claude PR Assistant workflow (#257) 47e0c62 · last month

.github/workflows Claude PR Assistant ... last month

misc move files 7 years ago

README.md Update README.md: ... 9 months ago

README

## Awesome R

awesome

A curated list of awesome R packages and tools. Inspired by [awesome-machine-learning](#).

for [Top 50](#) CRAN downloaded packages or repos with 400+ ⭐

- [Awesome R](#)
  - [2023](#)
  - [2020](#)

About

A curated list of awesome R packages, frameworks and software.

data-science list awesome r  
rstats data-analysis awesome-list

Readme Activity 6.3k stars 415 watching 1.5k forks Report repository

Releases No releases published

Packages No packages published

Contributors 91

## Data Formats

Packages for reading and writing data of different formats.

- [arrow](#) - An interface to the Arrow C++ library.
- [feather](#) - Fast, interoperable binary data frame storage for Python, R, and more powered by Apache Arrow.
- [fst](#) - Lightning Fast Serialization of Data Frames for R.
- [haven](#) - Improved methods to import SPSS, Stata and SAS files in R.
- [jsonlite](#) - A robust and quick way to parse JSON files in R.
- [gs](#) - Quick serialization of R objects.
- [readxl](#) - Read excel files (.xls and .xlsx) into R.
- [readr](#) - A fast and friendly way to read tabular data into R.
- [rio](#) - A Swiss-Army Knife for Data I/O.
- [readODS](#) - Read OpenDocument Spreadsheets into R as data.frames.
- [RcppTOML](#) - Rcpp Bindings to C++ parser for TOML files.
- [vroom](#) - Fast reading of delimited files.
- [writexl](#) - Portable, light-weight data frame to xlsx exporter for R.
- [yaml](#) - R package for converting objects to and from YAML.

## Graphic Displays

Packages for showing data.

- [ggplot2](#) - An implementation of the Grammar of Graphics.
- [gfortify](#) - A unified interface to ggplot2 popular statistical packages using one line of code.
- [ggrepel](#) - Repel overlapping text labels away from each other.
- [ggaft](#) - Extra Coordinate Systems, Geoms and Statistical Transformations for ggplot2.
- [ggstatsplot](#) - ggplot2 Based Plots with Statistical Details



# R Related Project

## Related Projects

<https://www.r-project.org/other-projects.html>

- **Bioconductor: Bioinformatics with R**

<http://www.bioconductor.org>

- Rgeo: Spatial Statistics with R

<https://geodacenter.asu.edu/projects/rsp>

- Robust Statistics with R

<http://www.statistik.tuwien.ac.at/rsr/>

- Rmetrics: Financial Market Analysis with R

<http://www.rmetrics.org>

- Omegahat: Distributed Statistical Computing

<http://www.omegahat.org>

- Graphical User Interfaces for R

[http://www.sciviews.org/\\_rgui/](http://www.sciviews.org/_rgui/)

- ESS: Emacs speaks Statistics

<http://ess.R-project.org/>

- R for Mediawiki

[http://mars.wiwi.hu-berlin.de/mediawiki/sk/index.php/R\\_Extension\\_for\\_MediaWiki](http://mars.wiwi.hu-berlin.de/mediawiki/sk/index.php/R_Extension_for_MediaWiki)

- PowerShell R Interop Cmdlet

<http://powershellinterop.codeplex.com>

- TANGO/ALGENCAN

<http://www.ime.usp.br/~ebirgin/tango/>

The screenshot shows the Bioconductor website at <http://www.bioconductor.org>. The header features the Bioconductor logo and navigation links for Home, Install, Help, Developers, and About. A search bar is also present. The main content area includes a message about BioC 2016, sections for About Bioconductor, News, and various user and developer guides.

**About Bioconductor**

Bioconductor provides tools for the analysis and comprehension of high-throughput genomic data. Bioconductor uses the R statistical programming language, and is open source and open development. It has two releases each year, [1211 software packages](#), and an active user community. Bioconductor is also available as an [AMI](#) (Amazon Machine Image) and a series of [Docker](#) images.

**News**

- Bioconductor [3.3](#) is available.
- Bioconductor [F1000 Research Channel](#) launched.
- Orchestrating high-throughput genomic analysis with [Bioconductor](#) ([abstract](#)) and other [recent literature](#).
- Read our latest [newsletter](#) and [course material](#).
- Use the [support site](#) to get help installing, learning and using Bioconductor.

**Install »**

Get started with Bioconductor

- [Install Bioconductor](#)
- [Explore packages](#)
- [Get support](#)
- [Latest newsletter](#)
- [Follow us on twitter](#)
- [Install R](#)

**Learn »**

Master Bioconductor tools

- [Courses](#)
- [Support site](#)
- [Package vignettes](#)
- [Literature citations](#)
- [Common work flows](#)
- [FAQ](#)
- [Community resources](#)
- [Videos](#)

**Use »**

Create bioinformatic solutions with Bioconductor

- [Software](#), [Annotation](#), and [Experiment](#) packages
- [Amazon Machine Image](#)
- [Latest release announcement](#)
- [Support site](#)

**Develop »**

Contribute to Bioconductor

- [Developer resources](#)
- [Use Bio 'devl'](#)
- ['Devel' Software, Annotation and Experiment packages](#)
- [Package guidelines](#)
- [New package submission](#)
- [Build reports](#)

<http://www.bioconductor.org>



# 說明手冊 (Help Manuals)

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Windows Help

Console

FAQ on R  
FAQ on R for Win  
Manuals (in PDF)

R functions (text),  
**Html help**  
Search help...  
search.r-project.or

Apropos...

R Project home pa  
CRAN home page

About

The R Language

Statistical Da

An Introduction to R  
Writing R Extensions  
R Data Import/Export

Packages

Miscell

About R  
License  
NEWS

Frequen

Material speci

CHANGES up to R 2.15.0

R: Package Index

mvtnorm Multivariate Nor  
nlme Linear and Nonli  
nnet Feed-forward N  
Models

parallel Support for Para  
pcaPP Robust PCA by  
plyr Tools for splitting  
proto Prototype object  
R2HTML HTML exportatio  
Rcmdr R Commander  
RColorBrewer ColorBrewer pa  
Rcpp Seamless R and C  
reshape2 Flexibly reshape  
rgl 3D visualization  
robustbase Basic Robust St  
RODBC ODBC Database  
rpanel Simple interactiv  
rpart Recursive Partition  
rrcov Scalable Robust  
RSDA RSDA - R to Sy  
SQLite SQLite interface  
SQLite.extfun Math and String  
rstudio Tools and Utilitie  
scales Scale functions

R: 3D visualization device system ...

3D visualization device system (OpenGL)

Documentation for package ‘rgl’ version 0.93.1098

- [DESCRIPTION file](#).
- [Code demos](#). Use [demo\(\)](#) to run them.
- [Package NEWS](#).

Help Pages

A B C D E G I L M N O P Q R S T V W misc

rgl-package 3D visualization device system

-- A --

abclines3d Lines intersecting the bounding box  
addNormals Add normal vectors to objects so they render more smoothly.  
addNormals.mesh3d Add normal vectors to objects so they render more smoothly.  
addNormals.shapelist3d Add normal vectors to objects so they render more smoothly.  
asEuclidean Work with homogeneous coordinates  
asHomogeneous Work with homogeneous coordinates  
aspect3d Set the aspect ratios of the current plot  
axes3d Draw boxes, axes and other text outside the data  
axis3d Draw boxes, axes and other text outside the data

# 求助範例

```
> help.search("t test")
> ?t.test
```

```
> help(solve)
> ?solve
> help("[[")
> help.start()
> ?help
```

**NOTE:**

```
> ?'if'
> ?Control
> ?'while'
> ?'%'
```

R: Search Results

The search string was "t test"

Vignettes:

- [Rcpp::Rcpp-unitTests](#) Rcpp-unitTests PDF source R code
- [survival::tests](#) Cox models and ``type 3'' Tests PDF source R code

Code demonstrations:

- [fda::create.test](#) run some tests on basis functions (Run demo in console)
- [gsubfn::gsubfn-unitTests](#) Run gsubfn unit test suite. (Run demo in console)
- [sqldf::sqldf-unitTests](#) Run unit tests using svUnit package. (Run demo in console)
- [tcltk::tkttest](#) t-test example of GUI interface to a function call. (Run demo in console)

Help pages:

- [ade4::RV.rtest](#) Monte-Carlo Test on the sum of eigenvalues of a co-inertia analysis (in R).
- [ade4::gearymoran](#) Moran's I and Geary's randomization tests for spatial and phylogenetic autocorrelation

R: Search Results

stats::fisher.test FISHER'S EXACT TEST FOR COUNT DATA

stats::fligner.test Fligner-Killeen Test of Homogeneity of Variances

stats::friedman.test Friedman Rank Sum Test

stats::kruskal.test Kruskal-Wallis Rank Sum Test

stats::ks.test Kolmogorov-Smirnov Tests

stats::mantelhaen.test Cochran-Mantel-Haenszel Chi-Squared Test for Count Data

stats::mauchly.test Mauchly's Test of Sphericity

stats::mcnemar.test McNemar's Chi-squared Test for Count Data

stats::mood.test Mood Two-Sample Test of Scale

stats::pairwise.t.test Pairwise t tests

stats::pairwise.wilcox.test Pairwise Wilcoxon Rank Sum Tests

stats::poisson.test Exact Poisson tests

stats::power.anova.test Power Calculations for Balanced One-Way Analysis of Variance Tests

stats::power.prop.test Power Calculations for Two-Sample Test for Proportions

stats::power.t.test Power calculations for one and two sample t tests

stats::PP.test Phillips-Perron Test for Unit Roots

stats::quade.test Quade Test

stats::shapiro.test Shapiro-Wilk Normality Test

stats::t.test Student's t-Test

stats::var.test F Test to Compare Two Variances

stats::wilcox.test Wilcoxon Rank Sum and Signed Rank Tests

survival::plot.cox.zph Graphical Test of Proportional Hazards

survival::survobrien O'Brien's Test for Association of a Single Variable with Survival

tools::RdTextFilter Select text in an Rd file.

utils::file\_test Shell-style Tests on Files

?package.name

?package.name::function.name

apropos("norm") #search all function names for the "norm" key



# 求助範例 (cont.)

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t.test {stats}

Student's t-Test

Description

Performs one and two sample t-tests on vectors of data.

Usage 語法

t.test(x, ...)

## Default S3 method:

```
t.test(x, y = NULL,  
       alternative = c("two.sided", "less", "greater"),  
       mu = 0, paired = FALSE, var.equal = FALSE,  
       conf.level = 0.95, ...)
```

## S3 method for class 'formula'

```
t.test(formula, data, subset, na.action, ...)
```

Arguments 參數

x a (non-empty) numeric vector of data values.

y an optional (non-empty) numeric vector of data values.

alternative a character string specifying the alternative hypothesis, must be one "less". You can specify just the initial letter.

mu a number indicating the true value of the mean (or difference in mean).

paired a logical indicating whether you want a paired t-test.

var.equal a logical variable indicating whether to treat the two variances as being used to estimate the variance otherwise the Welch (or Satterthwaite) used.

Details

## 說明

The formula interface is only applicable for the 2-sample tests.

alternative = "greater" is the alternative that x has a larger mean than y.

If paired is TRUE then both x and y must be specified and they must be length 1 or length equal. If var.equal is TRUE then the pooled estimate of the variance is used; if FALSE then the variance is estimated separately for both groups and weighted averages of the two sample variances are used for the degrees of freedom.

回傳值

Value

A list with class "htest" containing the following components:

statistic the value of the t-statistic.

parameter the degrees of freedom for the t-statistic.

p.value the p-value for the test.

conf.int a confidence interval for the mean appropriate to the specified alternative hypothesis.

estimate the estimated mean or difference in means depending on whether it was a one-sample or two-sample test.

null.value the specified hypothesized value of the mean or mean difference depending on whether it was a one-sample or two-sample test.

alternative a character string describing the alternative hypothesis.

method a character string indicating what type of t-test was performed.

data.name a character string giving the name(s) of the data.

See Also

[prop.test](#)

## 範例

Examples

```
require(graphics)
```

```
t.test(1:10, y = c(7:20)) # P = .00001855
```

```
t.test(1:10, y = c(7:20, 200)) # P = .1245 -- NOT significant anymore
```

```
## Classical example: Student's sleep data
```

```
plot(extra ~ group, data = sleep)
```

```
## Traditional interface
```

```
with(sleep, t.test(extra[group == 1], extra[group == 2]))
```

```
## Formula interface
```

```
t.test(extra ~ group, data = sleep)
```

# Search R



<http://www.rdocumentation.org/>

Rdocumentation powered by datacamp



Learn R Programming

Search from 30,811 R packages on CRAN and Bioconductor

For example, try 'ggplot2' or 'geom\_point'

Search

Explore learning paths with DataCamp

R Fundamentals

Big Data with R

Machine Learning with R

Course

Introduction to R  
Data Visualization with R  
Tutorial

R Basics Cheat Sheet  
Linear Regression in R  
Histograms in R

Course

Practicing Interview Questions in R  
Data Manipulation with R  
Tutorial

Importing Data into R  
Principal Component Analysis in R  
Contingency Tables in R

Course

Machine Learning in the Tidyverse  
Supervised Machine Learning in R  
Tutorial

Decision Trees in R  
Hierarchical Clustering in R

RSeek:  
<https://rseek.org/>



**pkgsearch:** Search and Query CRAN R Packages

<https://cran.r-project.org/web/packages/pkgsearch/index.html>

# `ps()` is an alias to `pkg_search()`

```
> install.packages("pkgsearch")
> library(pkgsearch)
> ps("survival")
> ps("networks")
> ps("visualization")
```



## &gt; demo( )

```
R demos

Demos in package 'base':

error.catching      More examples on catching and handling errors
is.things            Explore some properties of R objects and
                     is.FOO() functions. Not for newbies!
recursion            Using recursion for adaptive integration
scoping              An illustration of lexical scoping.

Demos in package 'graphics':

Hershey               Tables of the characters in the Hershey
                      vector fonts
Japanese              Tables of the Japanese characters in the
                      Hershey vector fonts
graphics              A show of some of R's graphics capabilities
image                 The image-like graphics builtins of R
persp                 Extended persp() examples
plotmath              Examples of the use of mathematics annotation

Demos in package 'grDevices':

colors                A show of R's predefined colors()
hclColors             Exploration of hcl() space

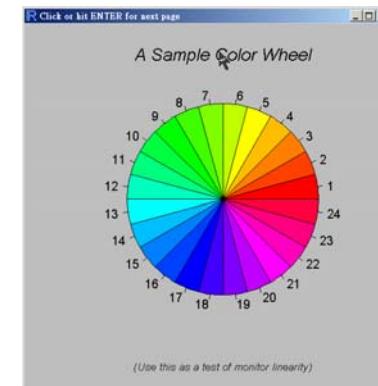
Demos in package 'stats':

glm.vr                Some glm() examples from V&R with several
                      predictors
lm.glm                 Some linear and generalized linear modelling
                      examples from 'An Introduction to Statistical
                      Modelling' by Annette Dobson
nlm                   Nonlinear least-squares using nlm()
smooth                'Visualize' steps in Tukey's smoothers

Use 'demo(package = .packages(all.available = TRUE))'
to list the demos in all *available* packages.
```

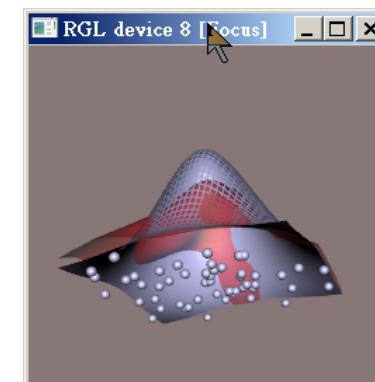
```
demo(package = .packages(all.available = TRUE))
```

```
> demo(graphics)
```



```
> library(rgl)
```

```
> demo(rgl)
```





# 演示程式碼 (Demo Code)

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The screenshot shows the EmEditor application open with the file `graphics.R`. The code in the editor window is as follows:

```
1 # Copyright (C) 1997-2009 The R Core Team
2 #
3 require(datasets)
4 require(grDevices); require(graphics)
5 #
6 ## Here is some code which illustrates some of the differences between
7 ## R and S graphics capabilities. Note that colors are generally specified
8 ## by a character string name (taken from the X11 rgb.txt file) and that
9 ## textures are given similarly. The parameter "bg" sets the background
10 ## parameter for the plot and there is also an "fg" parameter which sets
11 ## the foreground color.
12 #
13 #
14 x <- stats::rnorm(50)
15 opar <- par(bg = "white")
16 plot(x, ann = FALSE, type = "n")
17 abline(h = 0, col = gray(.90))
18 lines(x, col = "green4", lty = "dotted")
19 points(x, bg = "limegreen", pch = 21)
20 title(main = "Simple Use of Color In a Plot",
21       xlab = "Just a Whisper of a Label",
22       col.main = "blue", col.lab = gray(.8),
23       cex.main = 1.2, cex.lab = 1.0, font.main = 4, font.lab = 3)
24 #
25 #
26 ## A little color wheel. This code just plots equally spaced hues
27 ## a pie chart. If you have a cheap SVGA monitor (like me) you will
28 ## probably find that numerically equispaced does not mean visually
29 ## equispaced. On my display at home, these colors tend to cluster at
```

The status bar at the bottom left indicates `4.92 KB (5,039 字節), 146 行。`. The status bar at the bottom right indicates `R 行 8, 欄 56 繁體中文 (Big5)`.

To the right of the editor, a file browser window titled `demo` is open, showing the directory structure:

- Top level: `graphics.R`, `Hershey.R`, `image.R`, `Japanese.R`, `persp.R`, `plotmath.R`
- Second level: `Files ▾ R ▾ R-3.1.0 ▾ library ▾ rgl ▾ demo`
- Third level: `abundance.r`, `bivar.r`, `envmap.r`, `flag.R`, `hist3d.r`, `lollipop3d.R`, `lsystem.r`, `mouseCallbacks.R`, `regression.r`, `rgl.r`, `shapes3d.R`, `stereo.R`, `subdivision.r`

# R程式IDE編輯器: RStudio



Download RStudio Desktop  
<https://posit.co/>

The screenshot shows the Posit website's main navigation bar with links for PRODUCTS, SOLUTIONS, LEARN & SUPPORT, EXPLORE MORE, PRICING, and a search icon. A prominent blue button labeled 'DOWNLOAD RSTUDIO' is visible. Below the navigation, the text 'Build better with Posit Workbench' is displayed in large blue font. To the right of the text are three overlapping images: a code editor window showing R code, a 'New Session' dialog box, and a scatter plot.

Code in R. Code in Python. Develop in the environments you prefer: Jupyter, VSCode, and the RStudio IDE. Create the data products your stakeholders need: applications, reports, dashboards, and more.

IDE: integrated development environment

The screenshot shows the RStudio Desktop landing page. It features a large 'DOWNLOAD' button at the top. Below it, the text 'RStudio Desktop' is prominently displayed in a large serif font. A descriptive paragraph explains that RStudio is used by millions of people weekly to help them be more productive with R and Python. The page also includes the Posit navigation bar at the top.

Used by millions of people weekly, the RStudio integrated development environment (IDE) is a set of tools built to help you be more productive with R and Python.

# 客制化環境



R code - RStudio

File Edit Code View Plots Session Build Debug Profile Tools Help

rcode.R x Source on Save Go to file/function Run Source

```

49 h <- hist(ScoreData$V1, breaks=seq(0, 100, 10), right=F, plot = F)
50 a <- max(h$counts)
51
52 no.take <- nrow(ScoreData)
53 no.give.up <- length(which(is.na(ScoreData$V1)))
54 no.100 <- length(which(ScoreData$V1 == 100))
55 no.pass <- length(which(ScoreData$pass))
56 no.fail <- no.take - no.pass - no.give.up
57 pass.rate <- no.pass/(no.take-no.give.up)
58 Average <- round(mean(ScoreData$V1, na.rm = T), 2)
59 Median <- round(median(ScoreData$V1, na.rm = T), 2)
60 SD <- round(sd(ScoreData$V1, na.rm = T), 2)
61
62 ggplot(ScoreData, aes(x = V1, fill = pass)) +
63   geom_histogram(color="black", breaks = seq(0, 100, 10), closed = "left") +
64   labs(title = paste0("NCCU ", year, " ", subject, " 學期總成績"), x = "分數", y = "人數",
65       caption = "http://www.hmwu.idv.tw") +
66   scale_x_continuous(breaks = seq(0, 100, 10)) +
67   scale_y_continuous(breaks = 1:a) +
68   theme(legend.position = "none") +
69   annotate(geom="text", hjust = 0, x = 30, y = a, label = paste0("修課人數: ", no.take))
70   annotate(geom="text", hjust = 0, x = 30, y = a-1, label = paste0("棄修人數: ", no.give.up))
71   annotate(geom="text", hjust = 0, x = 30, y = a-2, label = paste0("100分人數: ", no.100))
72   annotate(geom="text", hjust = 0, x = 30, y = a-3, label = paste0("及格人數: ", no.pass))
73   annotate(geom="text", hjust = 0, x = 30, y = a-4, label = paste0("不及格人數: ", no.fail))
74   annotate(geom="text", hjust = 0, x = 30, y = a-5, label = paste0("及格率: ", round(pass.r)))
75   annotate(geom="text", hjust = 0, x = 30, y = a-6, label = paste0("分數平均數: ", Average))
76   annotate(geom="text", hjust = 0, x = 30, y = a-7, label = paste0("分數中位數: ", Median))
77

```

74:67 (Top Level) : R Script

Environment History Files Connections Tutorial

New Folder New Blank File Delete Rename More

D: \OS-NCCU\_Course\_Now\ R Code

Name	Size	Modified
..	3.6 KB	Jan 23, 2023, 7:38 PM
.RData	29.8 KB	Jan 23, 2023, 7:38 PM
.Rhistory	356 B	Jan 27, 2022, 1:28 AM
1101-Reg.txt	1.1 kB	Jan 27, 2022, 1:28 AM
1101-Stat.txt	617 B	Jun 29, 2022, 3:39 PM
1102-Reg.txt	391 B	Jun 26, 2022, 11:40 PM
1102-Stat.txt	58 kB	Jan 23, 2023, 1:09 AM
1111-Reg-histogram2.png	273 B	Jan 22, 2023, 4:22 PM
1111-Reg.txt	61 kB	Jan 23, 2023, 1:04 AM
1111-Stat-histogram.png	372 B	Feb 22, 2023, 10:57 PM
1111-Stat.txt	226.5 kB	Jan 27, 2022, 1:23 AM
128823-000359031.doc	71.9 kB	Jan 27, 2022, 1:24 AM
128823-300807001.doc	3.5 kB	Feb 22, 2023, 10:57 PM
png	217 B	Feb 22, 2023, 10:51 PM
rcode.R		
Rcode.Rproj		

Console Background Jobs

R 4.2.2 D:\OS-NCCU.Course\_Now\RCode\

```

+ annotate(geom="text", hjust = 0, x = 0, y = a, label = paste0("小考(2次): ", quiz.p)) +
+ annotate(geom="text", hjust = 0, x = 0, y = a-1, label = paste0("期中考: ", midterm.p)) +
+ annotate(geom="text", hjust = 0, x = 0, y = a-2, label = paste0("期末考: ", final.p)) +
+ annotate(geom="text", hjust = 0, x = 0, y = a-3, label = paste0("作業(無): ", HW.p)) +
+ annotate(geom="text", hjust = 0, x = 0, y = a-4, label = paste0("點名(無): ", attend.p)) +
+ annotate(geom="text", hjust = 0, x = 0, y = a-5, label = paste0("實習課(TA): ", TA.p)) +
+ annotate(geom="text", hjust = 0, x = 0, y = a-6, label = paste0("程式加分考(自由參加): ", bonus.p)) +
+ annotate(geom="text", hjust = 0, x = 0, y = a-7, label = paste0("閱讀心得(自由參加): ", other.p)) +
+ annotate(geom="text", hjust = 0, x = 0, y = a-8, label = paste0("總分: ", total.p))
Warning messages:
1: package 'ragg' is not available; using default graphics backend instead
2: Removed 9 rows containing non-finite values (stat_bin()).
> |

```

Plots Packages Help Viewer Presentation

NCCU 111-1 統計學(一) 學期總成績

22-	小考(2次): 30%	修課人數: 85
21-	期中考: 30%	棄修人數: 9
20-	期末考: 40%	100分人數: 0
19-	作業(無): 0%	及格人數: 61
18-	點名(無): 0%	不及格人數: 15
17-	實習課(TA): 0%	及格率: 80.26%
16-	程式加分考(自由參加): 20%	分數平均數: 67.49
15-	閱讀心得(自由參加): 10%	分數中位數: 73.5
14-	總分: 130%	分數標準差: 22.39

人數 分數

# Tools => Global Options...

The image shows two overlapping dialog boxes from the RStudio Global Options interface.

**Left Dialog: R Sessions**

- General Tab:**
  - R version: [Default] [64-bit] C:\Program Files\R\R-4.2.2 (Change...)
  - Default working directory (when not in a project): ~ (Browse...)
  - Restore most recently opened project at startup
  - Restore previously open source documents at startup
  - Workspace**
  - Restore .RData into workspace at startup
  - Save workspace to .RData on exit: Ask
  - History**
  - Always save history (even when not saving .RData)
  - Remove duplicate entries in history
  - Other**
  - Wrap around when navigating to previous/next tab
  - Automatically notify me of updates to RStudio
  - Send automated crash reports to RStudio
- Buttons:** OK, Cancel, Apply

**Right Dialog: Options**

Choose the layout of the panels in RStudio by selecting from the controls in each panel. Add up to three additional Source Columns to the left side of the layout. When a column is removed, all saved files within the column are closed and any unsaved files are moved to the main Source Pane.

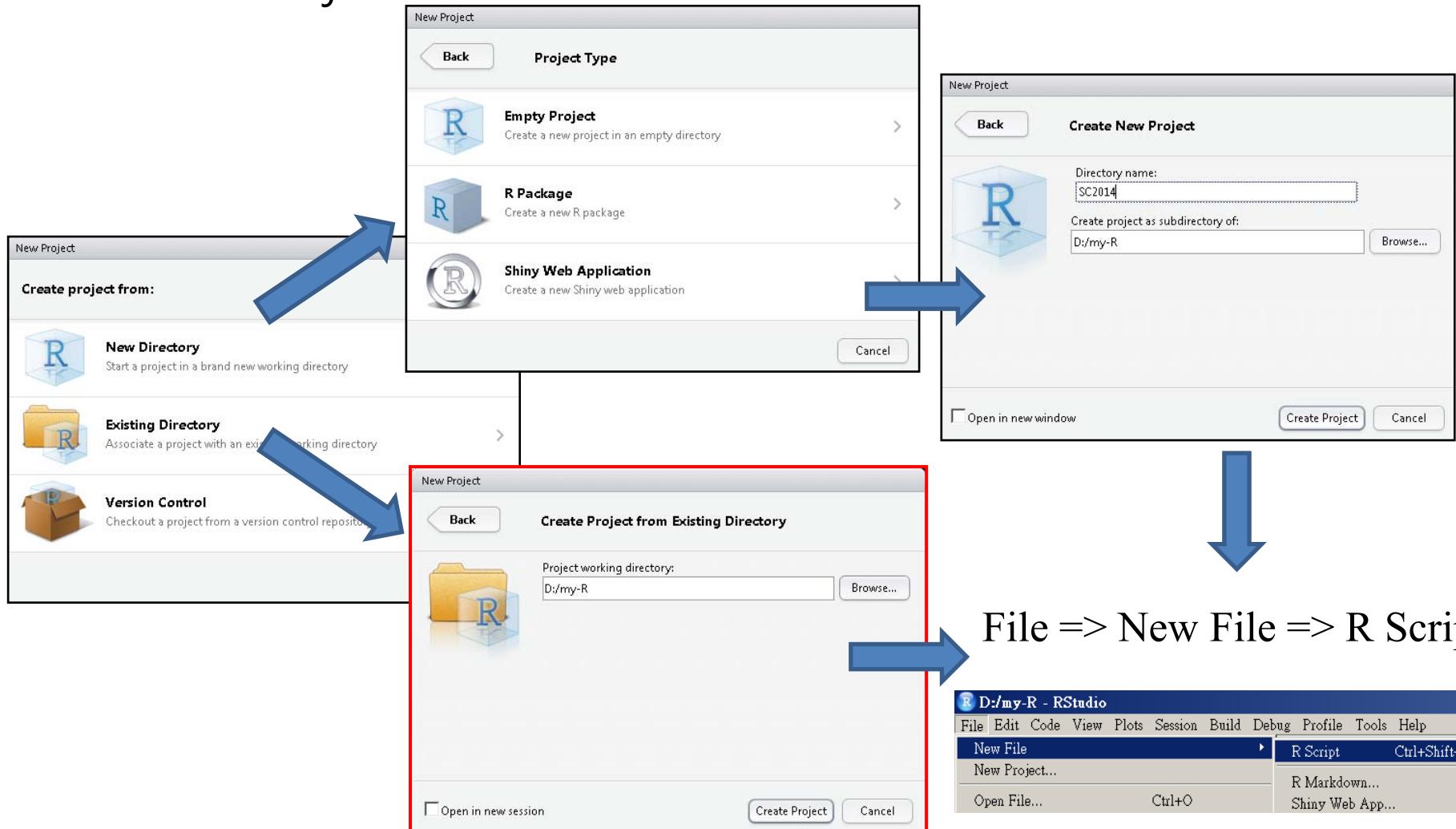
- Add Column** (+) | Remove Column (-)
- Source Panel:** Source
- Console Panel:** Console
- Panel Layout Options:**
  - Environment, History, Files, Colors:** Environment, History, Files (checked), Plots, Connections, Packages, Help, Build, VCS, Tutorial
  - Plots, Packages, Help, Viewer, etc.:** Environment, History, Files, Plots, Connections, Packages, Help, Build, VCS, Tutorial
- Buttons:** OK, Cancel, Apply



# 新增專案及R程式檔

35/55

- File => New Project... => New Directory => Empty Project => Create Project



# 執行程式

The screenshot shows the RStudio interface with the following components:

- Script Editor (myRcode-chap01.R):**

```

1 #####
2 # My testing R code
3 # Han-Ming Wu
4 # 2024/07/16
5 #####
6
7
8 cat ("Use iris data to test")
9 data(iris)
10 View(iris)
11 head(iris, 3)
12 summary(iris)
13 attach(iris)
14 plot(Sepal.Length, Sepal.Width, col=Species)
15

```
- Console:**

```

Median :5.000 Median :3.000 Median :4.350
Mean   :5.843 Mean   :3.057 Mean   :3.758
3rd Qu.:6.400 3rd Qu.:3.300 3rd Qu.:5.100
Max.   :7.900 Max.   :4.400 Max.   :6.900
Petal.Width Species
Min.   :0.100 setosa  :50
1st Qu.:0.300 versicolor:50
Median :1.300 virginica:50
Mean   :1.199
3rd Qu.:1.800
Max.   :2.500
> attach(iris)
> plot(Sepal.Length, Sepal.Width, col=Species)
>

```
- Environment Browser:**
  - Disk: D: > my-R > SC2014
  - File: myRcode-chap01.R (384 B, Jul 9, 2014, 12:05 PM)
  - File: SC2014.Rproj (218 B, Jul 9, 2014, 12:03 PM)
- Plots:**

A scatter plot showing Sepal.Length on the x-axis (ranging from 4.5 to 8.0) and Sepal.Width on the y-axis (ranging from 2.0 to 4.0). The data points are colored by species: setosa (light blue), versicolor (orange), and virginica (green).

以鍵盤選取程式碼:

配合Home, End, Shift, 上下左右鍵  
執行: Ctrl + Enter 或 Ctrl + R

# 安排RStudio 專案目錄結構

The screenshot shows the RStudio interface with the following components:

- Title Bar:** X:/08-MyProjects/07-exploreSDA/MyPackage/exploreSDA - RStudio
- File Menu:** File Edit Code View Plots Session Build Debug Tools Help
- Toolbar:** Source on Save Go to file/function Run Source
- Code Editor:** Displays R code for the ESDA-Demo.R script.
- Console:** Shows R command history and output.
- Project Explorer:** Shows the project structure under X:/08-MyProjects/07-exploreSDA/MyPackage/exploreSDA. It includes folders like src-x64, src-i386, src, raw-data, R, man, inst, extdata, demo, data, and files like README.txt, exploreSDA.dll, exploreSDA.Rproj, .Rhistory, .Rbuildignore, .RData, NAMESPACE, and DESCRIPTION.
- Project Menu:** A context menu for the project named "exploreSDA" is open, with the "exploreSDA" option highlighted.

制作一R套件的目錄結構

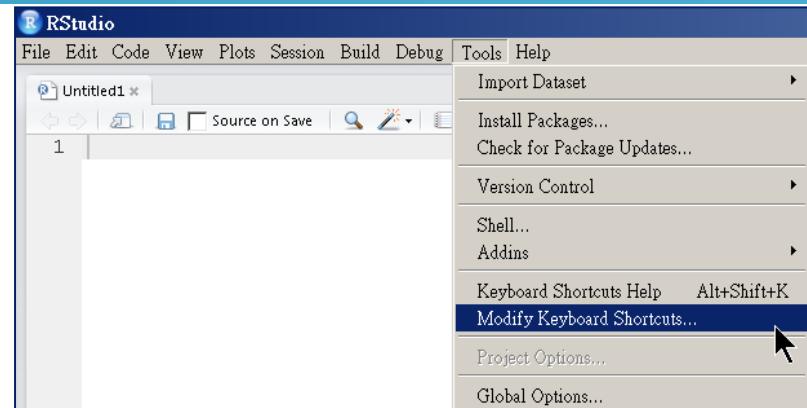


# RStudio 常用快速鍵 (Windows)

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## 控制台(Console)

- 游標移動到控制台 (Move cursor to Console): **Ctrl+2**
- 清除控制台 (Clear console): **Ctrl+L**
- 移動游標到行首 (Move cursor to beginning of line): **Home**
- 移動游標到行尾 (Move cursor to end of line): **End**
- 調出歷史指令 (Navigate command history): **上箭頭/下箭頭**
- 彈出歷史指令 (Popup command history): **Ctrl+上箭頭**
- 中斷正在執行的指令 (Interrupt currently executing command): **Esc**



RStudio 0.99.893

<C:\Program Files\RStudio\www\docs\keyboard.htm>

## 原始碼編輯區(Source)

- 游標移動到原始檔編輯區 (Move cursor to Source Editor): **Ctrl+1**
- 存檔 (Save active document): **Ctrl+S**
- 關閉當前檔案 (Close active document): **Ctrl+W**
- 執行當前行或者選擇的行 (Run current line/selection): **Ctrl+Enter**
- 執行當前檔 (Run current document): **Ctrl+Alt+R**
- 從檔案開頭執行到該行 (Run from document beginning to current line): **Ctrl+Alt+B**
- 行縮排 (Reindent lines) : **Ctrl+I**
- 註解(或消除)當前行或者所選程式碼 (Comment/uncomment current line/selection): **Ctrl+Shift+C**
- 跳轉到配對的括弧 (Jump to Matching Brace/Parentheses): **Ctrl+P**
- 刪除行 (Delete Line): **Ctrl+D**
- 選擇 (Select): **Shift+箭頭**

RStudio , 重新載入R:  
選單: Session => Restart R  
快速鍵: ctrl + shift + F10  
指令: > **.rs.restartR()**

## 一般

- 尋找和替換 (Find and Replace): **Ctrl+F**; 尋找上一個 (Find Previous): **Shift+F3**; 尋找下一個(Find Next): **F3**
- 上一步 (Undo ): **Ctrl+Z**; 剪下 (Cut): **Ctrl+X**; 複制 (Copy): **Ctrl+C**; 貼上 (Paste): **Ctrl+V**; 全選 (Select All): **Ctrl+A**

# RStudio Help

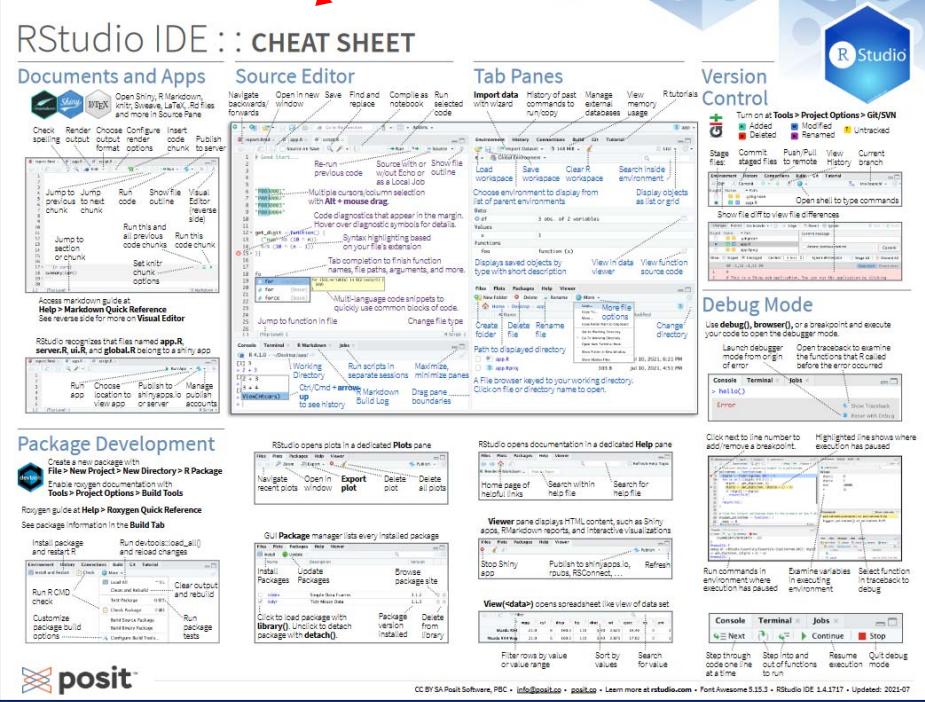


Tools Help

- R Help
- Search R Help Ctrl+Alt+F1
- About RStudio
- Check for Updates
- Accessibility
- RStudio Docs
- RStudio Community Forum
- Cheat Sheets
- Keyboard Shortcuts Help Alt+Shift+K
- Markdown Quick Reference
- Roxygen Quick Reference
- Diagnostics

RStudio IDE Cheat Sheet

Data Transformation with dplyr  
Data Visualization with ggplot2  
List manipulation with purrr  
Package Development with devtools  
Web Applications with shiny  
Interfacing Spark with sparklyr  
R Markdown Cheat Sheet  
R Markdown Reference Guide  
Browse Cheat Sheets...



## Posit Documentation





# Posit Cheatsheets

<https://posit.co/resources/cheatsheets/>

# Posit Cheatsheets

Data transformation with dplyr :: CHEAT SHEET

dplyr functions work with pipes and expect tidy data. In tidy data:

- Each variable is in its own column
- Each observation, or case, is in its own row
- x %>% f(y) becomes f(x, y)

**Manipulate Cases**

**EXTRACT CASES**

Row functions return a subset of rows as a new table.

- filter(data, ..., preserve = FALSE) Extract rows that meet logical criteria.  
filter(mtcars, mpg > 20)
- distinct(data, ..., keep\_all = FALSE) Remove rows with duplicate values.  
distinct(mtcars, gear)
- slice(data, ..., preserve = FALSE) Select rows by position.  
slice(mtcars, 1:15)
- slice\_sample(data, ..., n, prop, weight\_by = NULL, replace = FALSE) Randomly select rows to sum a fraction of rows.  
slice\_sample(mtcars, n = 5, replace = TRUE)
- slice\_min(data, order\_by, ..., n, prop, with\_ties = TRUE) and slice\_max() Select rows with the lowest and highest values.  
slice\_min(mtcars, mpg = 25)
- slice\_head(data, ..., n, prop) and slice\_tail() Select the first or last rows.  
slice\_head(mtcars, n = 5)

**Logical and boolean operators to use with filter()**

- `=` `<` `=>` `is.na()` `%in%` `|` `xor()`
- `!=` `>` `=>` `is.na()` `!` `&`

See ?base::Logic and ?Comparison for help.

**ARRANGE CASES**

Order rows by values of a column or columns (low to high), use with desc() to order from high to low.

- arrange(data, ..., by\_group = FALSE)
- arrange(data, ..., by\_group = TRUE)
- arrange(mtcars, mpg)
- arrange(mtcars, desc(mpg))

**ADD CASES**

Add one or more rows to a table.

- add\_row(data, ..., before = NULL, after = NULL)
- add\_col(data, ..., before = 1, after = 1)

**Manipulate Variables**

Column functions return a set of columns as a new vector or table.

- pull(data, var = -1, name = NULL, ...) Extract column values as a vector, by name or index.  
pull(mtcars, wt)
- select(data, ...) Extract columns as a table.  
select(mtcars, mpg:wt)
- relocate(data, ..., before = NULL, after = NULL) Move columns to new position.  
relocate(mtcars, mpg, cyl, after = last\_col())

Use these helpers with select() and across()  
e.g. select(mtcars, mpg:cyl)

- contains(match)
- num\_range(prefix, range)
- ends\_with(match)
- all\_of(...)
- any\_of(..., vars)
- starts\_with(match)
- matches(match)
- everything()

**MANIPULATE MULTIPLE VARIABLES AT ONCE**

- across(cols, funs, ..., names = NULL) Summarise or mutate multiple columns in the same way.  
summarise(mtcars, across(everything(), mean))
- c\_across(cols) Compute across columns in row-wise data.  
transmute(iris, sepal.Length = c\_across(1:2))

**MAKE NEW VARIABLES**

Apply **vectorized functions** to columns. Vectorized functions take vectors as input and return vectors of the same length as output (see back).

- vectorized function
- mutate(data, ..., keep = "all", before = NULL, after = NULL) Compute new column(s). Also, add\_column(), add\_count(), and add\_tally().  
mutate(mtcars, gpm = 1 / mpg)
- transmute(data, ...) Compute new column(s), drop others.  
transmute(mtcars, gpm = 1 / mpg)
- rename(data, ...) Rename columns. Use rename\_with() to rename with a function.  
rename(mtcars, distance = dist)

CC BY SA Posit Software, PBC • [info@posit.co](mailto:info@posit.co) • [posit.co](#) • Learn more at [dplyr.tidyverse.org](https://dplyr.tidyverse.org) • dplyr 1.0.7 • Updated: 2021-07

**posit**

- Sparklyr Cheat Sheet
- R Markdown Cheat Sheet
- RStudio IDE Cheat Sheet
- Shiny Cheat Sheet
- Data Visualization Cheat Sheet
- Package Development Cheat Sheet
- Data Wrangling Cheat Sheet
- R Markdown Reference Guide
- Contributed Cheatsheets
- Base R
- Advanced R
- Regular Expressions
- How big is your graph? (base R graphics)



# R Packages Developed by RStudio

<https://posit.co/products/open-source/rpackages/>

R PACKAGES

Inspired by R and our community

Analyze and explore



Connect and Integrate



Communicate and interact



Model and predict





# 更新R軟體及R套件

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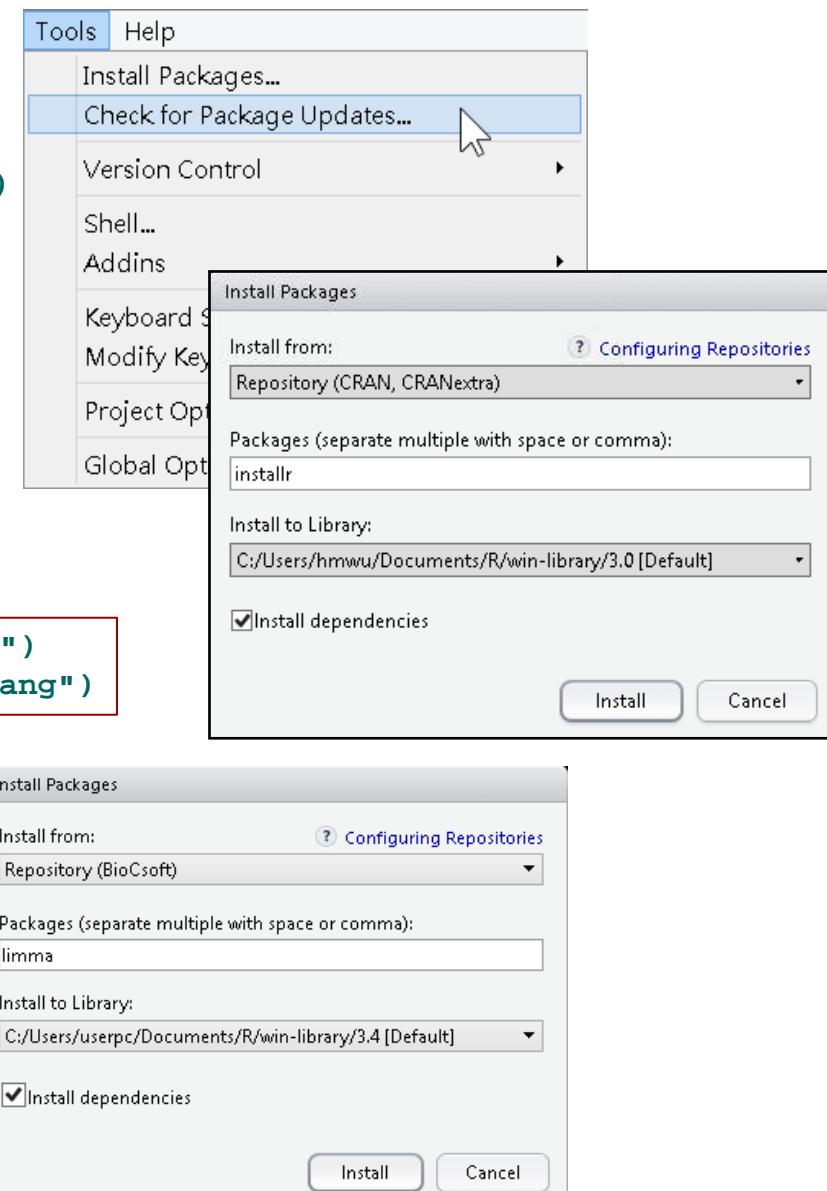
- 安裝R套件
- 利用installr套件更新R軟體及R套件

```
> update.packages(checkBuilt=TRUE, ask=FALSE)
  ■ RGui => Packages => Update packages
  ■ RStudio => Tools => Check for Package Updates...
```

```
> setRepositories()
--- Please select repositories for use in this
session ---
1: + CRAN
2: BioC software
3: BioC annotation
4: BioC experiment
5: BioC extra
6: CRAN (extras)
7: Omegahat
8: R-Forge
9: rforge.net

Enter one or more numbers separated by
spaces, or an empty line to cancel
1: 2

  > update.packages("rlang")
  > uninstall.packages("rlang")
```

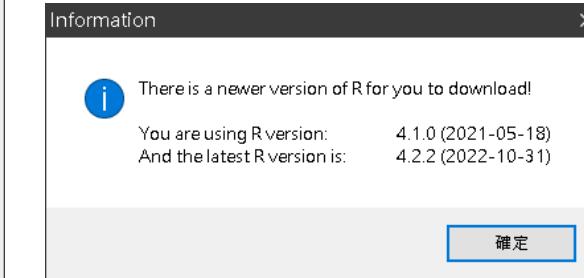


Note: 先暫時不移除舊版的R及套件!

# 利用installr套件更新R軟體



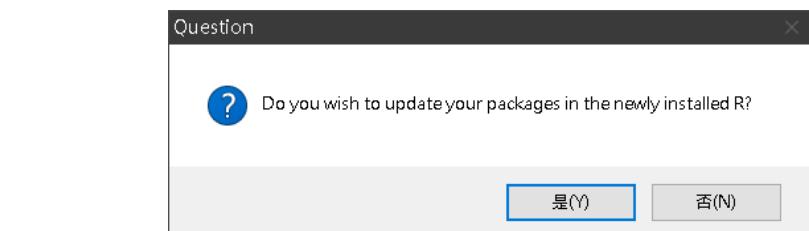
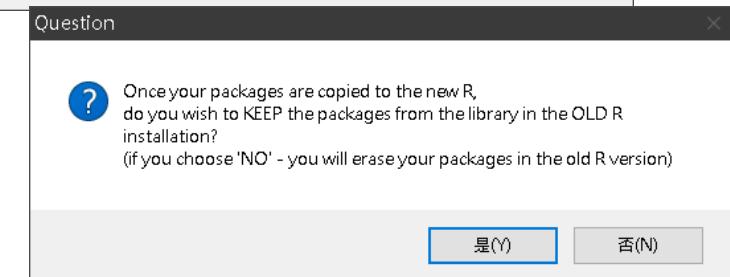
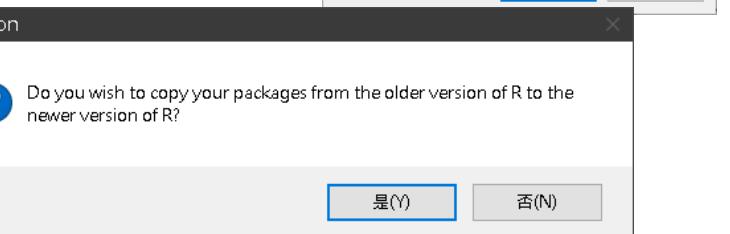
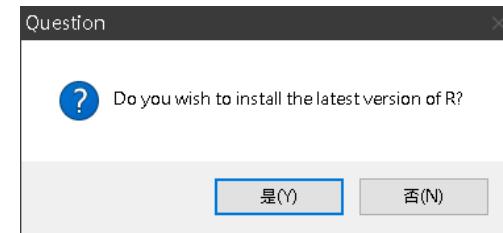
```
> library(installr)
> installr()
```



```
> sessionInfo()
```

```
> version

platform      x86_64-w64-mingw32
arch          x86_64
os            mingw32
crt           ucrt
system        x86_64, mingw32
status
major         4
minor         2.2
year          2022
month         10
day           31
svn rev       83211
language      R
version.string R version 4.2.2 (2022-10-31 ucrt)
nickname      Innocent and Trusting
```





# 學習資源

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<http://www.rdatamining.com/>
- 中華R軟體學會?  
中華R軟體研發暨應用協會?

R and Data Mining

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datacamp Learn R by doing Start For Free

This website presents documents, examples, tutorials and resources on R and data mining.

Documents on Data Mining with R



# R-bloggers • R Cheatsheet

**R-bloggers** R news and tut

HOME ABOUT RSS ADD YOUR BLOG! LEARN R R JOBS CONTACT US

## Forecasting Disney Stock Prices as the Latest Earnings Beat Estimates

February 21, 2023 | Selcuk Disci

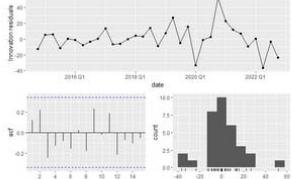
Walt Disney (NYSE: DIS) recently announced significant labor cuts to ease shareholders' pressure on reducing costs due to rising streaming investment. These cuts and some structural changes in the company have provided some boost to stock prices. We will examine these price changes based on earnings per share (EPS) and ...

[Read more...]

## Calculating Log Likelihood Ratios (LLR module)

February 21, 2023 | ["Peter M.B. Cahusac"]

tl;dr Ever wanted to try doing an evidential analysis? You may have found it difficult to find a statistical platform to do it. Now there is the jamovi module jeva which can provide log likelihood ratios for a range of common statistical tests.

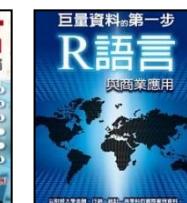
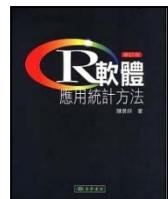


<https://www.r-bloggers.com>

## R Cheatsheet:

- <https://pyoflife.com/wp-content/uploads/2022/01/R-Cheatsheet-.pdf>
- [https://www.rforecology.com/uploads/The\\_essential\\_R\\_Cheatsheet\\_v1\\_0.pdf](https://www.rforecology.com/uploads/The_essential_R_Cheatsheet_v1_0.pdf)
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- <https://www.rstudio.com/resources/cheatsheets/>
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- <https://www.amelia.mn/Syntax-cheatsheet.pdf>
- <https://cheatography.com/dipakk/cheat-sheets/r-basic/pdf/>
- <https://cheatography.com/noella/cheat-sheets/introduction-to-package-r-sheet/pdf/>
- <https://cheatography.com/bwaldo/cheat-sheets/r-subsetting/pdf/>

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您輸入的關鍵字: R軟體

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搜尋結果共 1108 筆, 頁數 1 / 47

排序依 準確度 呈現:

應用機器學習: R軟體實務演練	R軟體: 應用統計方法(二版)(附光碟1片)	R軟體統計應用分析實務	R軟體在決策樹的實務應用	R軟體統計進階分析實務	心理測驗理論與應用：含RT與R軟體分析



# R/RStudio on the Phone

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## R Programming Compiler

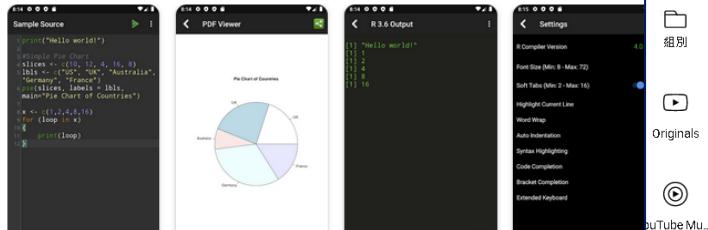
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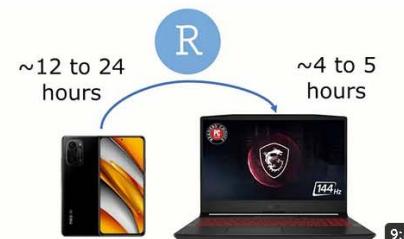
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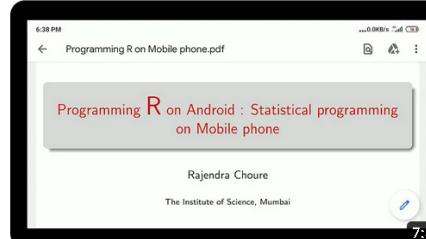
YouTube Premium TW

### How to install RStudio on Android

How to install RStudio on Android 視看次數：1630次 • 8 個月前 Franz Chandi Timestamps: 0:00 - Intro 0:31 - Reasons to install RStudio on Android 2:09 - How to install RStudio on Android 11 個章節



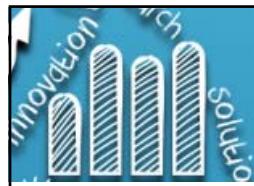
Programming R on mobile phone: Statistical programming on mobile 視看次數：1萬次 • 2 年前 Rajendra Choure programming #rprogrammingonandroid #rcompiler #rprogrammingonmobilephon This video discusses use of mobile for ...



### How to run R Programming in Mobile 📱/Free/R Programming

How to run R Programming in Mobile 📱/Free/R Programming 視看次數：5415次 • 1 年前 Om Ojha Hello guys Subscribe kar agar kuch information mili ho to Share Like #R Programming Mobile # Coding in mobile #coding ...





# Rcmdr: R Commander

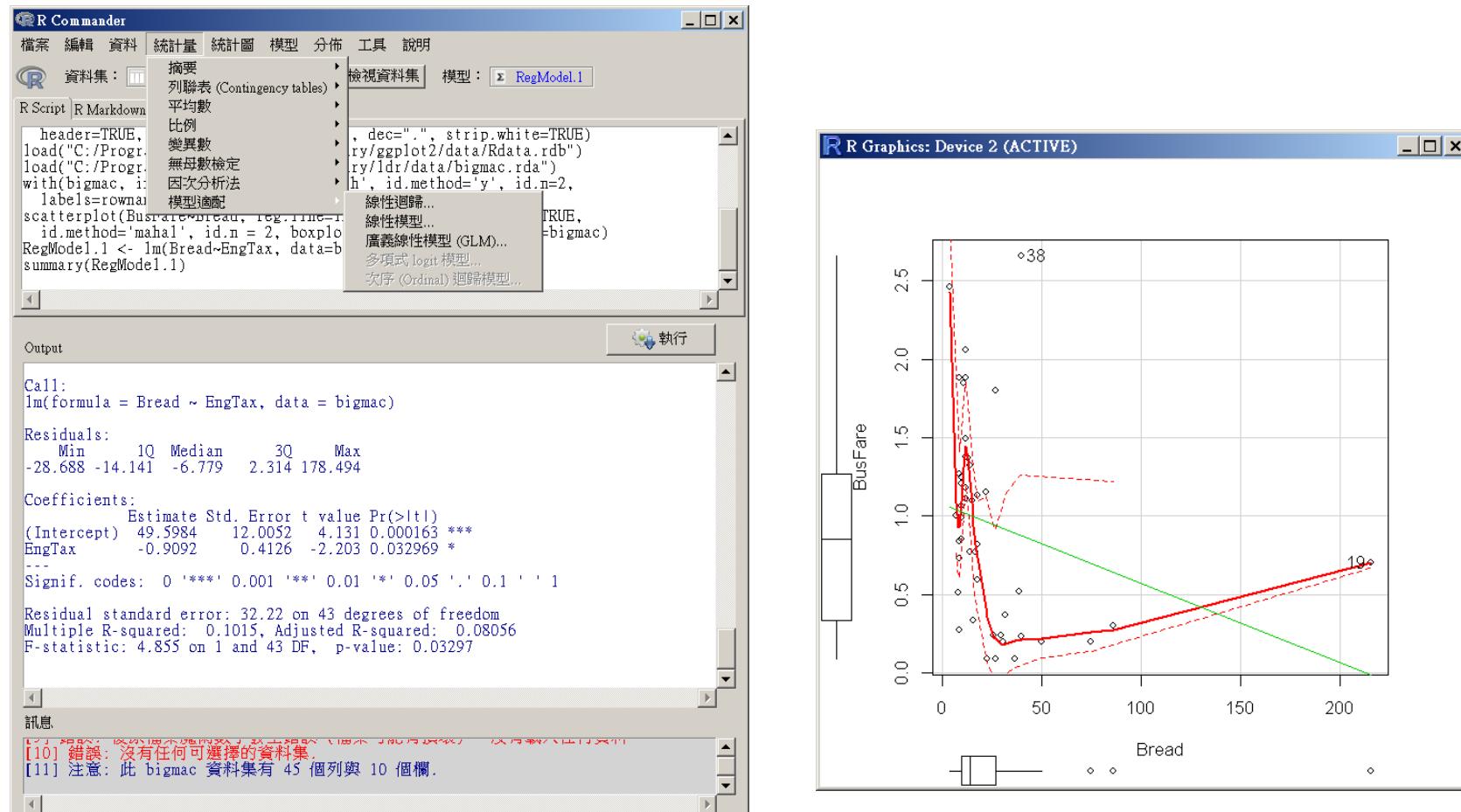
## The R Commander: A Basic-Statistics GUI for R

Current Version: 2.7-x

John Fox

<https://cran.r-project.org/web/packages/Rcmdr/index.html>

<https://socialsciences.mcmaster.ca/jfox/Misc/Rcmdr/>





# R AnalyticFlow

EN JP

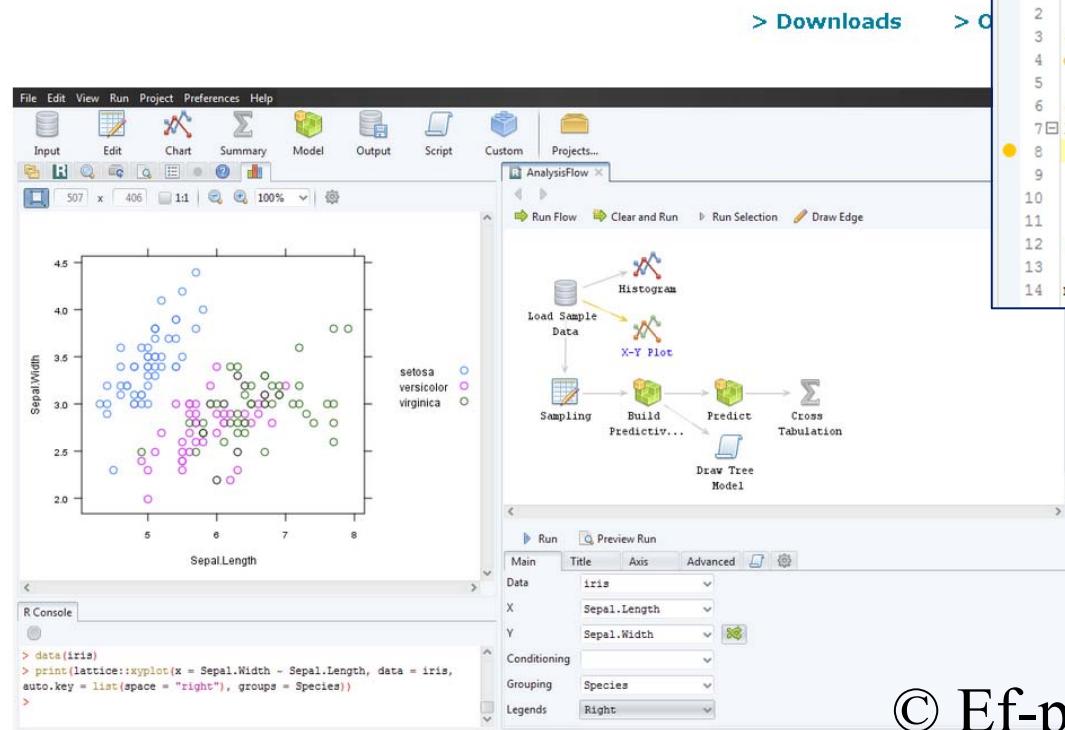
[Overview](#)[Downloads](#)[Documents](#)[Help & Support](#)

Designed for data analysis. Great for everyone.

R AnalyticFlow is a data analysis software that utilizes the R environment for statistical computing.

In addition to intuitive user interface, it also provides advanced features for R experts. These features enable you to share the processes of data analysis between users with differing levels of proficiency.

AnalyticFlow works on Windows, Mac OS X, and Linux and is free for any use.



## R Programming Support

The figure shows a code editor window titled '\*New Flow'. The code is an R script for building a predictive model using the rpart package. The code includes data loading, sampling, and model building steps. A yellow box highlights the line of code where the variable 'x' is assigned to the 'iris' dataset.

```
## Build and Validate Predictive Model
# Load Data
data(iris)

# Sampling
local({
  x <- iris
  smpl <- iris[iris3, , drop = FALSE]
  assign(x = smpl, value = x[smpl, , drop = FALSE], pos = parent.frame(n))
  assign(x = "test", value = x[-smpl, , drop = FALSE], pos = parent.frame(n))
})

model <- rpart::rpart(formula = Species ~ ., data = train)
```

© Ef-prime, Inc.



# 系統資訊/參數

50/55

```
> # R.Version: Version Information
> as.data.frame(R.Version())
  platform    arch     os      system status major minor year month day svn.rev
1 x86_64-w64-mingw32 x86_64 mingw32 x86_64, mingw32            3   3.1 2016   06  21  70800
  language      version.string      nickname
1          R R version 3.3.1 (2016-06-21) Bug in Your Hair
>
> # Sys.info: Extract System and User Information
> Sys.info()
  sysname   release   version   nodename   machine   login   user   effective_user
"Windows" "8.1 x64" "build 9600" "HMWU-HOME" "x86-64" "userpc" "userpc" "userpc"
>
> # .Platform: Platform Specific Variables
> as.data.frame(.Platform)
  OS.type file.sep dynlib.ext   GUI endian   pkgType path.sep r_arch
1 windows        /       .dll Rgui little win.binary ;      x64
>
> # .Machine {base}: Numerical Characteristics of the Machine
> as.data.frame(.Machine)
  double.eps double.neg.eps  double.xmin  double.xmax double.base double.digits
1 2.220446e-16 1.110223e-16 2.225074e-308 1.797693e+308         2           53
  double.rounding double.guard double.ulp.digits double.neg.ulp.digits double.exponent
1             5              0            -52            -53           11
  double.min.exp double.max.exp integer.max sizeof.long sizeof.longlong sizeof.longdouble
1        -1022          1024  2147483647            4              8           16
  sizeof.pointer
1              8
```

# ChatGPT for R, or in RStudio



## 8 ChatGPT packages for R

<https://www.infoworld.com/article/2338386/8-chatgpt-tools-for-r-programming.html>  
**air, TheOpenAIR, RTutor, CodeLingo, askgpt, gptstudio, gpttools, gptchatteR**

<https://cran.r-project.org/web/packages/chattr/index.html>

chattr: Interact with Large Language Models in 'RStudio'

Enables user interactivity with large-language models ('LLM') inside the 'RStudio' integrated development environment (IDE). The user can interact with the model using the 'shiny' app included in this package, or directly in the 'R' console. It comes with back-ends for 'OpenAI', 'GitHub' 'Copilot', and 'LlamaGPT'.

Version: 0.3.1

Published: 2023-10-03

To integrate a ChatGPT chat window directly into RStudio we can use a package called chattr.

Run ChatGPT in RStudio with Chattr

To integrate a ChatGPT chat window directly into RStudio we can use a package called chattr.

See video description for a link to these notes

Run ChatGPT in RStudio with chattr - Quick Setup

Susan B. 8750位訂閱者

觀看次數：2665次 6 個月前

How to Integrate a ChatGPT window directly into RStudio using the R package chattr. Includes details on auto-starting ChatGPT when RStudio loads.

...更多內容

<https://www.youtube.com/watch?v=AvRPsxnqzAM>



gptr: An R Interface with the ChatGPT API

Author: Wanjun Gu

gptr is an R package that provides a convenient interface with the OpenAI ChatGPT API. It allows you to interact with ChatGPT, a powerful language model, for various natural language processing tasks.

The gptr R package makes talking to ChatGPT in R super easy. It helps researchers and data folks by simplifying the complicated stuff, like asking questions and getting answers. With gptr, you can use ChatGPT in R without any hassle, making it simpler for everyone to do cool things with language!

### Installation

You can install gptr directly from CRAN:

```
install.packages("gptr")
```

<https://cran.r-project.org/web/packages/gptr/readme/README.html>

R

Analysis with R

作者 : Emirhan Duran

An expert in data analysis using the R programming language.

註冊以開始聊天

註冊或登入以開始聊天

<https://chatgpt.com/g/g-WUWF2eOOb-analysis-with-r>



# sessionInfo( )

```
> sessionInfo()
R version 4.2.2 (2022-10-31 ucrt)
Platform: x86_64-w64-mingw32/x64 (64-bit)
Running under: Windows 10 x64 (build 19044)
```

```
Matrix products: default
```

```
locale:
[1] LC_COLLATE=Chinese (Traditional)_Taiwan.utf8
[2] LC_CTYPE=Chinese (Traditional)_Taiwan.utf8
[3] LC_MONETARY=Chinese (Traditional)_Taiwan.utf8
[4] LC_NUMERIC=C
[5] LC_TIME=Chinese (Traditional)_Taiwan.utf8
```

```
attached base packages:
[1] stats      datasets  grid      lattice
```

```
loaded via Rcpp:
[1] compiler  tools     Rcpp     RcppEigen
```

```
> # 檢視所有套件安裝路徑:
```

```
> .libPaths()
[1] "C:/Users/hanmi/AppData/Local/R/win-library/4.5"
[2] "C:/Program Files/R/R-4.5.0/library"
```

```
>
```

```
> # 檢視特定套件的安裝目錄:
```

```
> system.file(package = "ggplot2")
[1] "C:/Users/hanmi/AppData/Local/R/win-library/4.5/ggplot2"
```

```
>
```

```
> # 檢視所有已安裝的套件清單與路徑:
```

```
> # installed.packages()
> installed.packages()[, c("Package", "LibPath")]
   Package           LibPath
1  abind      "abind"    "C:/Users/hanmi/AppData/Local/R/win-library/4.5"
2 admisc      "admisc"   "C:/Users/hanmi/AppData/Local/R/win-library/4.5"
3   arm        "arm"     "C:/Users/hanmi/AppData/Local/R/win-library/4.5"
4   ...
```

```
> RStudio.Version()
$citation
...
$version
[1] '2022.7.2.576'
```

# 檢視套件版本

```
> packageVersion("rlang")
[1] '0.4.5'
```



## The Most Frequently Used Data Set

### UCI Machine Learning Repository <https://archive.ics.uci.edu/ml/>

Welcome to the UC Irvine Machine Learning Repository!

We currently maintain 292 data sets as a service to the machine learning community. You may [view all data sets](#) through our searchable interface. Our [old web site](#) is still available, for those who prefer the old format. For a general overview of the Repository, please visit our [About page](#). For information about citing data sets in publications, please read our [citation policy](#). If you wish to donate a data set, please consult our [donation policy](#). For any other questions, feel free to [contact the Repository librarians](#). We have also set up a [mirror site](#) for the Repository.

Supported By: In Collaboration With:

Latest News:	Newest Data Sets:	Most Popular Data Sets (hits since 2007):
2013-04-04: Welcome to the new Repository admins Kevin Bache and Moshe Lichman	2014-06-18:  Gesture Phase Segmentation	583104:  Iris
2010-03-01: Note from donor regarding Netflix data	2014-06-12:  Parkinson Speech Dataset with Multiple Types of Sound Recordings	407709:  Adult
2009-10-16: Two new data sets have been added.	2014-06-01:  Tennis Major Tournament Match Statistics	349110:  Wine
2009-09-14: Several data sets have been added.		294819:  Breast Cancer
2008-07-23: Repository mirror has been set up		

The sepal length, sepal width, petal length, and petal width are measured in centimeters on 50 iris specimens from each of three species, *Iris setosa*, *I. versicolor*, and *I. virginica*. Fisher (1936)



資料編輯器					
	Sepal.Length	Sepal.Width	Petal.Length	Petal.Width	Species
1	5.1	3.5	1.4	0.2	setosa
2	4.9	3	1.4	0.2	setosa
3	4.7	3.2	1.3	0.2	setosa
4	4.6	3.1	1.5	0.2	setosa
5	5	3.6	1.4	0.2	setosa
6	5.4	3.9	1.7	0.4	setosa
7	4.6	3.4	1.4	0.3	setosa
8	5	3.4	1.5	0.2	setosa
9	4.4	2.9	1.4	0.2	setosa
10	4.9	3.1	1.5	0.1	setosa
11	5.4	3.7	1.5	0.2	setosa
12	4.8	3.4	1.6	0.2	setosa
13	4.8	3	1.4	0.1	setosa
14	4.3	3	1.1	0.1	setosa
15	5.8	4	1.2	0.2	setosa
16	5.7	4.4	1.5	0.4	setosa
17	5.4	3.9	1.3	0.4	setosa
18	5.1	3.5	1.4	0.3	setosa
19	5.7	3.8	1.7	0.3	setosa
20	5.1	3.8	1.5	0.3	setosa
21	5.4	3.4	1.7	0.2	setosa

2014 IASC Data Analysis Competition: <http://www.iasc-isi.org/node/227>

- World Bank - <http://data.worldbank.org>
- United Nations - <http://www.un.org/en/databases/#stats>
- World Health Organization - <http://www.who.int/research/en/>

- StatLib: <http://lib.stat.cmu.edu/>
- 政府資料開放平台  
<http://data.gov.tw/>
- 開放資料Open Data  
<http://www.opendata.tw/>



# RStudio 套件安裝問題 (1)

54/55

```
> install.packages("CVD")
WARNING: Rtools is required to build R packages but is not currently installed. Please download and install the appropriate version of Rtools before proceeding:

https://cran.rstudio.com/bin/windows/Rtools/
warning in install.packages :
  cannot create dir 'C:\Users\xáñçò\AppData\Local\Temp\Rtmpw9xEQw\downloaded_packages', reason 'No such file or directory'
Error in install.packages : unable to create temporary directory 'C:\Users\xáñçò\AppData\Local\Temp\Rtmpw9xEQw\downloaded_packages'
> |
```

## 中文帳戶名問題

### ■ My favorite RStudio tips and tricks

[http://datacornering.com/my-favorite-rstudio-tips-and-tricks/?fbclid=IwAR2ZK9Fu3Q5j\\_wUQVo-tFuX0lxP4Zp2TwcfWKXHyV2R-C6ElgG1Gn-t3uJg](http://datacornering.com/my-favorite-rstudio-tips-and-tricks/?fbclid=IwAR2ZK9Fu3Q5j_wUQVo-tFuX0lxP4Zp2TwcfWKXHyV2R-C6ElgG1Gn-t3uJg)

## 套件安裝問題，可能的解決方法：

- (1) 以「系統管理員身份執行」開啟RStudio。
- (2) 改變R套件安裝資料夾「.libPaths」  
<https://www.twblogs.net/a/5cc0edfc9eee397113dede>
- (3) 解決RStudio不支持Windows系統中文用戶名問題  
[https://blog.csdn.net/qq\\_16146103/article/details/105445198](https://blog.csdn.net/qq_16146103/article/details/105445198)
- (4) win10用戶名為中文導致RStudio繪圖錯誤的解決方法  
<https://www.cnblogs.com/yanjiamin/p/12064048.html>

### ■ (5) Windows中文使用者與RStudio的環境變數調校

<http://shorturl.at/AMNQ9>

- (6) 解決RStudio中的亂碼  
<https://itw01.com/5XYZENB.html>

- (7) 如何更改R的預設語系  
<https://psmethods.postach.io/post/ru-he-geng-gai-rde-yu-she-yu-xi>

## 中文路徑名問題

```
Error in utils::shortPathName(path[i]) : 無效的多位元組字串於 '<e6><88><91>?□~1/DOCUME~1/R/WIN-LI~1/4.0/RMARKD~1/rmd/h/DEFAUL~1.HTM'
Calls: <Anonymous> ... do.call -> <Anonymous> ->
  pandoc_path_arg -> <Anonymous>
停止執行
```

```
# 列出已安裝套件 ? installed.packages
mypks <- as.data.frame(installed.packages()[, 1:3])
mypks
```



# RStudio 套件安裝問題 (2)

55/55

```
> install.packages("gridExtra")
```

將程式套件安載入 'C:/Users/User/Documents/R/win-library/4.1'  
(因為 'lib' 沒有被指定)

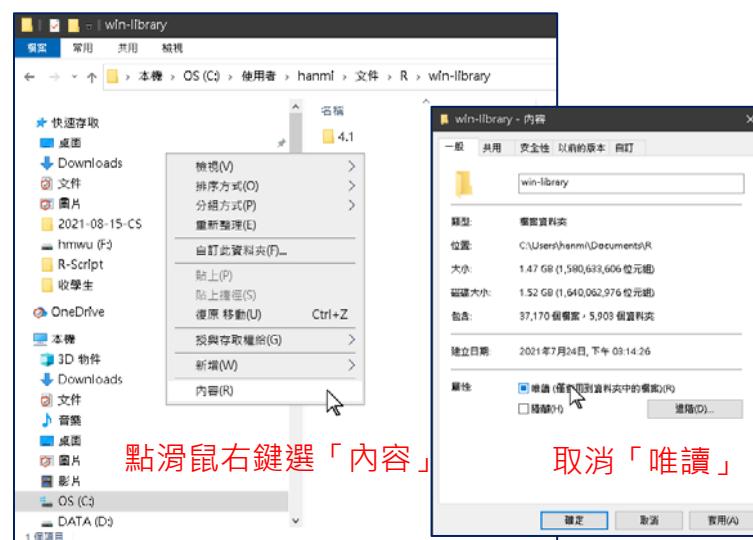
Warning in install.packages :

```
'lib = "C:/Users/User/Documents/R/win-library/4.1"' is not writable  
嘗試 URL 'https://cran.rstudio.com/bin/windows/contrib/4.1/gridExtra_2.3.zip'  
Content type 'application/zip' length 1109407 bytes (1.1 MB)  
downloaded 1.1 MB
```

Warning in install.packages :

無法建立目錄 'C:\Users\User\Documents\R\win-library\4.1\file2b905b04620'，原因是 'No such file or directory'

Error in install.packages : unable to create temporary directory  
'C:\Users\User\Documents\R\win-library\4.1\file2b905b04620'



點滑鼠右鍵選「內容」

取消「唯讀」，按確定。

