# 107-2 系統生物學概論軟體/套件安裝教學

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# 軟體/套件列表

- R
- RStudio
- R套件: Bioconductor(GEOquery, oligo, limma, DESeq2) pheatmap, ggfortify

- JAVA
- GSEA
- Cytoscape

#### Google: R → The R Project for Statistical Computing



## The R Project for Statistical Computing

**Getting Started** 

Download

R is a free software environment for statistical computinent of graphics. It compiles and runs on a wide variety of UNIX platforms, Windows and MacOS. To download R please choose your preferred CRAN

CRAN R Project

If you have questions about R like how to downlead and install the software, or what the license terms are, please read our answers to frequently asked questions before you send an email.

About R

找到"Taiwan"後 任選一個進入

#### Taiwan

https://ftp.yzu.edu.tw/CRAN/ http://ftp.yzu.edu.tw/CRAN/ http://cran.csie.ntu.edu.tw/

Department of Computer Science and Engineering, Yuan Ze University Department of Computer Science and Engineering, Yuan Ze University National Taiwan University, Taipei

The Comprehensive R Archive Network

Download and Install

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

- Download R for (Mac) OS X
- · Download R for Windows

#### 依據作業系統進行選擇

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

R for Windows

drectories

Binaries for base distribution. This is what you want to install R for the first time.

Binaries of contributed CRAN packages (for R >= 2.13.x; managed by Uwe Ligges). There is also information on third party software available for CRAN Windows services and corresponding environment and make variables.

Binaries of contributed CRAN packages for outdated versions of R (for R < 2.13.x; managed by Uwe Ligges).

Tools to build R and R packages. This is what you want to build your own packages on Windows, or to build R itself.

submit binaries to CRAN. Package developers might want to contact Uwe Ligges directly in case of questions / suggestions related to Windows binaries.

You may also want to read the R FAQ and R for Windows FAQ

Note: CRAN does some checks on these binaries for viruses, but cannot give guarantees. Use the normal precautions with downloaded executables.

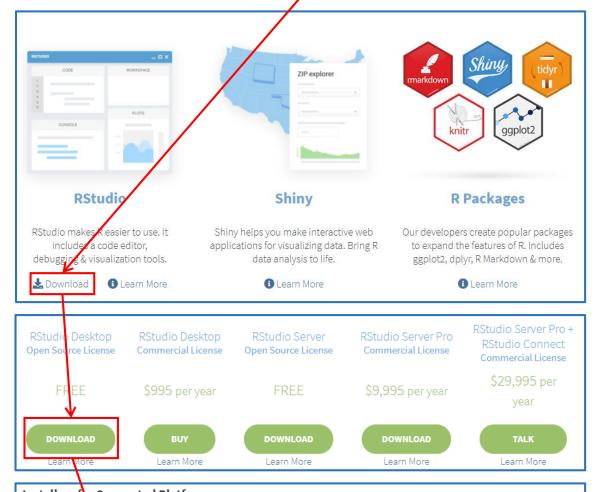
R-3.5.3 for Windows (32/64 bit)

Download R 3.5.3 for Windows (79 megabytes, 32/64 bit)

Installation and other instructions New features in this version

#### Google: RStudio → RStudio: Open source and ...

## RStudio



依據作業系統進行選擇

	Installers for Supported Platforms			
	Installers	Size	Date	MD5
	RStudio 1.1.463 - Windows Vista/7/8/10	85.8 MB	2018-10-29	58b3d796d8cf96fb8580c62f46ab64d4
•	RStudio 1.1.463 - Mac OS X 10.6+ (64-bit)	74.5 MB	2018-10-29	a79032ba4d7daaa86a8da01948278d94
	RStudio 1.1.463 - Ubuntu 12.04-15.10/Debian 8 (32-bit)	89.3 MB	2018-10-29	8a6755fa9fae2bafce289df3358aaf63
	RStudio 1.1.463 - Ubuntu 12.04-15.10/Debian 8 (64-bit)	97.4 MB	2018-10-29	bc50d6bd34926c1cc3ae4a209d67d649
	RStudio 1.1.463 - Ubuntu 16.04+/Debian 9+ (64-bit)	65 MB	2018-10-29	cfd659db18619cc78d1592fefaa7c753
	RStudio 1.1.463 - Fedora 19+/RedHat 7+/openSUSE 13.1+ (32-bit)	88.1 MB	2018-10-29	742f0bad60dfeaa3281576e14ad6699e
	RStudio 1.1.463 - Fedora 19+/RedHat 7+/openSUSE 13.1+ (64-bit)	90.6 MB	2018-10-29	c7303067a0ca99deea7e427b856952d1

# 安裝R套件

```
• 開啟RStudio或R
• 於左側Console內依序輸入以下指令:
# 安裝Bioconductor
source("http://www.bioconductor.org/biocLite.R")
biocLite()
# 安裝Bioconductor套件
source("http://www.bioconductor.org/biocLite.R")
biocLite(c("GEOquery", "oligo", "limma", "DESeq2"))
# 安裝非Bioconductor之R套件
install.packages(c("pheatmap", "ggfortify"))
```

# 呼叫/測試R套件

- 開啟Rstudio或R
- 於左側Console內依序輸入以下指令:

library(GEOquery)

library(oligo)

library(limma)

library(DESeq2)

library(pheatmap)

●請注意執行安裝、呼叫命令時所出現的提示訊息,若出現"Error:" 則代表安裝、呼叫失敗。通常是因為其套件需要其他尚未安裝的套件 (packages)來協助工作,請詳細閱讀提示訊息並將套件補齊後再執行 呼叫,直到不再出現Error為止。(安裝指令請參考上一頁)

## R套件列表

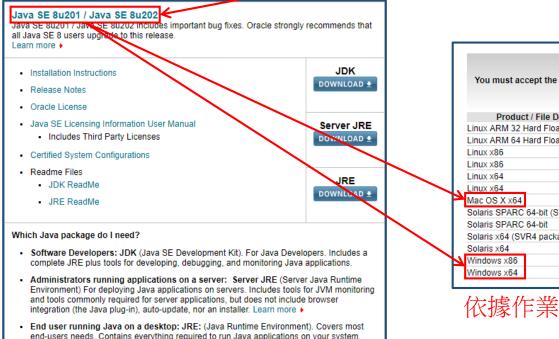
- Bioconductor套件列表:
  https://www.bioconductor.org/packages/release/bioc/
- R套件(非Bioconductor套件)列表: https://cran.r-project.org/web/packages/available\_packages\_by\_name.html

## JAVA

• 下載GSEA、Cytoscape前,請先確定電腦 是否已安裝JAVA 8(建議安裝JAVA 8 SE版)

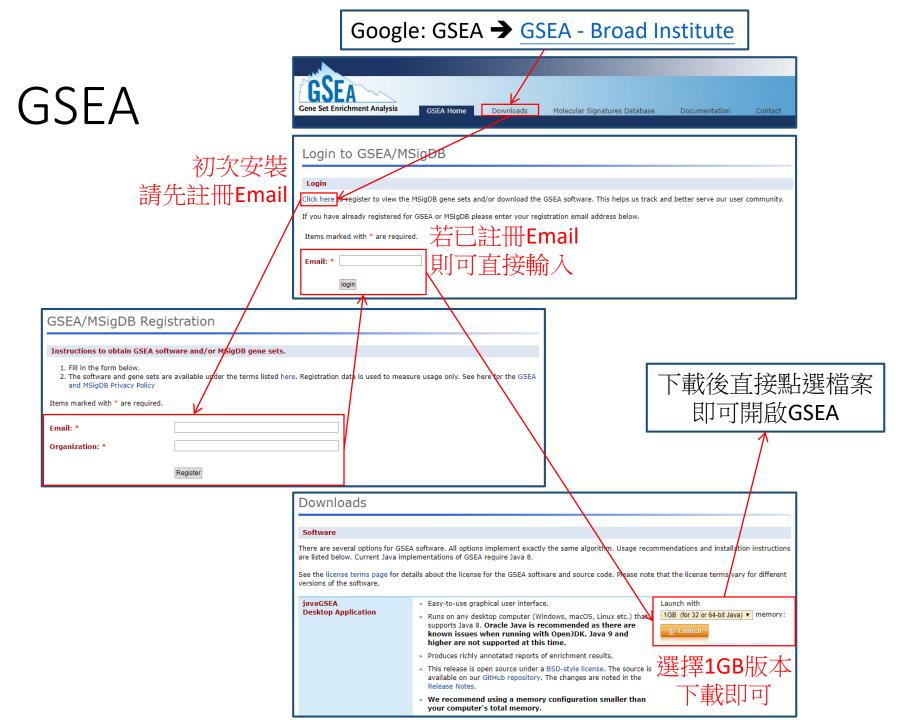
Google: Java SE → Java SE - Downloads | Oracle Technology Network | Oracle

#### 請勿安裝8u以外的版本

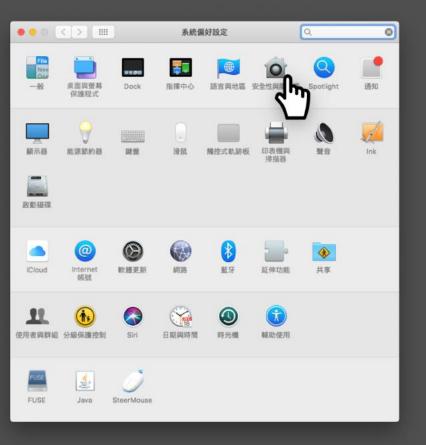


Java SE Development Kit 8u201 You must accept the Oracle Binary Code License Agreement for Java SE to download this software. Accept License Agreement
 Decline License Agreement File Size Product / File Description Download Linux ARM 32 Hard Float ABI 72.98 MB Jidk-8u201-linux-arm32-vfp-hflt.tar.gz Linux ARM 64 Hard Float ABI 69.92 MB →jdk-8u201-linux-arm64-vfp-hflt.tar.gz 170.98 MB ₱jdk-8u201-linux-i586.rpm 245.92 MB ₱jdk-8u201-macosx-x64.dmg Solaris SPARC 64-bit (SVR4 package) 125.33 MB ₹jdk-8u201-solaris-sparcv9.tar.Z 88.31 MB ₱jdk-8u201-solaris-sparcv9.tar.gz Solaris x64 (SVR4 package) 133.99 MB ₱jdk-8u201-solaris-x64.tar.Z 92.16 MB ₱jdk-8u201-solaris-x64.tar.gz 197.66 MB ₹jdk-8u201-windows-i586.exe 207.46 MB €jdk-8u201-windows-x64.exe

依據作業系統進行選擇



### macOS可能會遇到如右圖的問題 解決方法如下







## Cytoscape

Google: Cytoscape → Cytoscape: An Open Source ...



## Cytoscape - EnrichementMap

Manage Sites...

Install

Close

**EnrichmentMap** 

Visualizes enrichments of pathways as an

enrichment map, a network representing overlaps among enriched pathways.

3.2.0

View on App Store

