

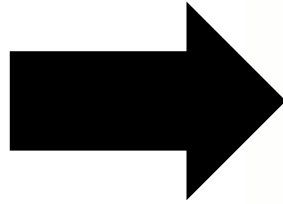
# Introduction to Packages



Introduction to R

 Package

=





CRAN

# GitHub



# Bioconductor

OPEN SOURCE SOFTWARE FOR BIOINFORMATICS



## CRAN Task Views

<a href="#"><u>Bayesian</u></a>	Bayesian Inference
<a href="#"><u>ChemPhys</u></a>	Chemometrics and Computational Physics
<a href="#"><u>ClinicalTrials</u></a>	Clinical Trial Design, Monitoring, and Analysis
<a href="#"><u>Cluster</u></a>	Cluster Analysis & Finite Mixture Models
<a href="#"><u>DifferentialEquations</u></a>	Differential Equations
<a href="#"><u>Distributions</u></a>	Probability Distributions
<a href="#"><u>Econometrics</u></a>	Econometrics
<a href="#"><u>Environmetrics</u></a>	Analysis of Ecological and Environmental Data
<a href="#"><u>ExperimentalDesign</u></a>	Design of Experiments (DoE) & Analysis of Experimental Data
<a href="#"><u>ExtremeValue</u></a>	Extreme Value Analysis
<a href="#"><u>Finance</u></a>	Empirical Finance
<a href="#"><u>FunctionalData</u></a>	Functional Data Analysis
<a href="#"><u>Genetics</u></a>	Statistical Genetics
<a href="#"><u>Graphics</u></a>	Graphic Displays & Dynamic Graphics & Graphic Devices & Visualization
<a href="#"><u>HighPerformanceComputing</u></a>	High-Performance and Parallel Computing with R
<a href="#"><u>MachineLearning</u></a>	Machine Learning & Statistical Learning
<a href="#"><u>MedicalImaging</u></a>	Medical Image Analysis
<a href="#"><u>MetaAnalysis</u></a>	Meta-Analysis
<a href="#"><u>Multivariate</u></a>	Multivariate Statistics
<a href="#"><u>NaturalLanguageProcessing</u></a>	Natural Language Processing
<a href="#"><u>NumericalMathematics</u></a>	Numerical Mathematics

## Task View

Bayesian

ChemPhys

ClinicalTrials

Cluster

DifferentialEquations

Distributions

Econometrics

Environmetrics

ExperimentalDesign

Finance

Genetics

Search all 14,381 CRAN, BioConductor and Github packages.

Search

Or explore packages in one of the [Task Views](#).

```
install.packages("ggplot2")
```

```
install.packages(c("ggplot2", "devtools", "lme4"))
```



The screenshot shows the RStudio interface. The top menu bar includes File, Edit, Code, View, Plots, Session, Build, Debug, Profile, Tools, and Help. The Tools menu is open, showing options: Install Packages..., Check for Package Updates..., Version Control, Shell..., Terminal, Addins, Keyboard Shortcuts Help, Modify Keyboard Shortcuts..., Project Options..., and Global Options... The console window is active, displaying the R version and copyright information, followed by the license text and the contributors list.

File Edit Code View Plots Session Build Debug Profile **Tools** Help

+ ▾ 📁 ▾ 💾 💾 🖨️ ➡️ Go to file/function 🗪 ▾ Addins ▾

**Console** **Terminal** x

/cloud/project/ ➡️

R version 3.4.4 (2018-03-15) -- "Someone to Lean On"  
Copyright (C) 2018 The R Foundation for Statistical Computing  
Platform: x86\_64-pc-linux-gnu (64-bit)

R is free software and comes with ABSOLUTELY NO WARRANTY.  
You are welcome to redistribute it under certain conditions.  
Type 'license()' or 'licence()' for distribution details.

R is a collaborative project with many contributors.  
Type 'contributors()' for more information and  
'citation()' on how to cite R or R packages in publications.

Install Packages...  
Check for Package Updates...  
Version Control ▶  
Shell...  
Terminal ▶  
Addins ▶  
Keyboard Shortcuts Help ⌘⌥K  
Modify Keyboard Shortcuts...  
Project Options... ⌘⌥  
Global Options... ⌘

## Install Packages

Install from: [? Configuring Repositories](#)

Repository (CRAN) ▾

Packages (separate multiple with space or comma):

Install to Library:

/home/rstudio-user/R/x86\_64-pc-linux-gnu-library/3.4 [De] ▾

☒ Install dependencies

Install

Cancel



```
source("https://bioconductor.org/biocLite.R")
```

```
biocLite()
```

```
biocLite("GenomicFeatures")
```



# GitHub

```
install.packages("devtools")
```

```
library(devtools)
```

```
install_github("author/package")
```

# Step 1: Install

```
install.packages("ggplot2")
```

```
Console Terminal x
/cloud/project/

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'citation()' on how to cite R or R packages in publications.

Type 'demo()' for some demos, 'help()' for on-line help, or
'help.start()' for an HTML browser interface to help.
Type 'q()' to quit R.

> install.packages("ggplot2")
Installing package into '/home/rstudio-user/R/x86_64-pc-linux-gnu-library/3.4'
(as 'lib' is unspecified)
also installing the dependencies 'glue', 'magrittr', 'stringi', 'colorspace', 'assertthat', 'utf8', 'Rcpp', 'stringr', 'RColorBrewer', 'dichromat', 'munsell', 'labeling', 'R6', 'viridisLite', 'cli', 'crayon', 'pillar', 'rlang', 'digest', 'gtable', 'plyr', 'reshape2', 'scales', 'tibble', 'lazyeval'

trying URL 'http://package-proxy/src/contrib/glue_1.2.0.tar.gz'
Content type 'application/x-tar' length 62719 bytes (61 KB)
=====
downloaded 61 KB
```

# Step 2: Load

```
library()
```

```
library(ggplot2)
```



FilesPlotsPackagesHelpViewer

InstallUpdatePackrat

NameDescriptionVersion

User Library

☐

assertthat

Easy Pre and Post Assertions

0.2.0

☐

cli

Helpers for Developing Command Line Interfaces

1.0.0

☐

colorspace

Color Space Manipulation

1.3-2

☐

crayon

Colored Terminal Output

1.3.4

☐

dichromat

Color Schemes for Dichromats

2.0-0

☐

digest

Create Compact Hash Digests of R Objects

0.6.15

☐

ggplot2

Create Elegant Data Visualisations Using the Grammar of Graphics

2.2.1

☐

glue

Interpreted String Literals

1.2.0

☐

gtable

Arrange 'Grobs' in Tables

0.2.0

☐

labeling

Axis Labeling

0.3

☐

lazyeval

Lazy (Non-Standard) Evaluation

0.2.1

☐

magrittr

A Forward-Pipe Operator for R

1.5

☐

munsell

Utilities for Using Munsell Colours

0.4.3

☐

pillar

Coloured Formatting for Columns

1.2.2

☐

plyr

Tools for Splitting, Applying and Combining Data

1.8.4

☐

R6

Classes with Reference Semantics

2.2.2

☐

RColorBrewer

ColorBrewer Palettes

1.1-2

☐

Rcpp

Seamless R and C++ Integration

0.12.17

☐

reshape2

Flexibly Reshape Data: A Reboot of the Reshape Package

1.4.3

☐

rlang

Functions for Base Types and Core R and 'Tidyverse' Features

0.2.0



# What packages are installed?

```
installed.packages() or library()
```

## Updating packages

```
old.packages()
```

```
update.packages()
```

```
install.packages("packagename")
```

Files

Plots

Packages

Help

Viewer

Install

Update

Packrat

Name

Description

Version

User Library

☐

assertthat

Easy Pre and Post Assertions

0.2.0

ⓧ

☐

cli

Helpers for Developing Command Line Interfaces

1.0.0

ⓧ

☐

colorspace

Color Space Manipulation

1.3-2

ⓧ

☐

crayon

Colored Terminal Output

1.3.4

ⓧ

☐

Color Schemes for Dichromats

2.0-0

ⓧ

☐

Create Compact Hash Digests of R Objects

0.6.15

ⓧ

☐

Create Elegant Data Visualisations Using the Grammar of Graphics

2.2.1

ⓧ

☐

Interpreted String Literals

1.2.0

ⓧ

☐

Arrange 'Grobs' in Tables

0.2.0

ⓧ

☐

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Axis Labeling

0.3

ⓧ

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Lazy (Non-Standard) Evaluation

0.2.1

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ⓧ

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2.2.2

ⓧ

☐

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ColorBrewer Palettes

1.1-2

ⓧ

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Rcpp

Seamless R and C++ Integration

0.12.17

ⓧ

☐

reshape2

Flexibly Reshape Data: A Reboot of the Reshape Package

1.4.3

ⓧ

☐

rlang

Functions for Base Types and Core R and

0.2.0

ⓧ

Check for Updates

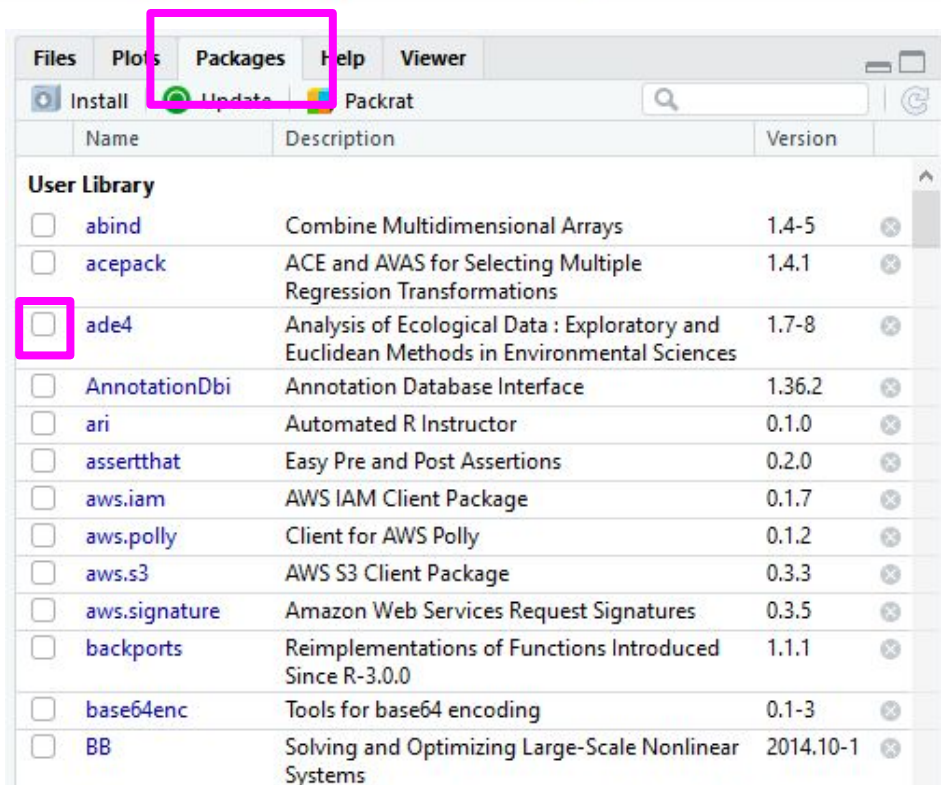
i

All packages are up to date.

OK

```
detach()
```

```
detach("package:ggplot2", unload=TRUE)
```



```
remove.packages()
```

```
remove.packages("ggplot2")
```


Files


Plots


Packages


Help


Viewer








 Install

 Update

 Packrat





Name	Description	Version	
User Library			
<input type="checkbox"/> assertthat	Easy Pre and Post Assertions	0.2.0	
<input type="checkbox"/> cli	Helpers for Developing Command Line Interfaces	1.0.0	
<input type="checkbox"/> colorspace	Color Space Manipulation	1.3-2	
<input type="checkbox"/> crayon	Colored Terminal Output	1.3.4	
<input type="checkbox"/> dichromat	Color Schemes for Dichromats	2.0-0	
<input type="checkbox"/> digest	Create Compact Hash Digests of R Objects	0.6.15	
<input checked="" type="checkbox"/> ggplot2	Create Elegant Data Visualisations Using the Grammar of Graphics	2.2.1	



```
Console Terminal x
/cloud/project/ ↵

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Platform: x86_64-pc-linux-gnu (64-bit)

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'citation()' on how to cite R or R packages in publications.

Type 'demo()' for some demos, 'help()' for on-line help, or
'help.start()' for an HTML browser interface to help.
Type 'q()' to quit R.

> |
```

version

sessionInfo()





```
> sessionInfo()
```

```
R version 3.4.4 (2018-03-15)
```

```
Platform: x86_64-pc-linux-gnu (64-bit)
```

```
Running under: Ubuntu 16.04.3 LTS
```

```
Matrix products: default
```

```
BLAS: /usr/lib/atlas-base/atlas/libblas.so.3.0
```

```
LAPACK: /usr/lib/atlas-base/atlas/liblapack.so.3.0
```

```
locale:
```

```
[1] LC_CTYPE=C.UTF-8
```

```
LC_NUMERIC=C
```

```
LC_TIME=C.UTF-8
```

```
[4] LC_COLLATE=C.UTF-8
```

```
LC_MONETARY=C.UTF-8
```

```
LC_MESSAGES=C.UTF-8
```

```
[7] LC_PAPER=C.UTF-8
```

```
LC_NAME=C
```

```
LC_ADDRESS=C
```

```
[10] LC_TELEPHONE=C
```

```
LC_MEASUREMENT=C.UTF-8
```

```
LC_IDENTIFICATION=C
```

```
attached base packages:
```

```
[1] stats
```

```
graphics
```

```
grDevices
```

```
utils
```

```
datasets
```

```
methods
```

```
base
```

```
other attached packages:
```

```
[1] ggplot2_2.2.1
```

```
loaded via a namespace (and not attached):
```

```
[1] colorspace_1.3-2 scales_0.5.0
```

```
compiler_3.4.4
```

```
lazyeval_0.2.1
```

```
[5] plyr_1.8.4
```

```
tools_3.4.4
```

```
pillar_1.2.2
```

```
gtable_0.2.0
```

```
[9] tibble_1.4.2
```

```
Rcpp_0.12.17
```

```
grid_3.4.4
```

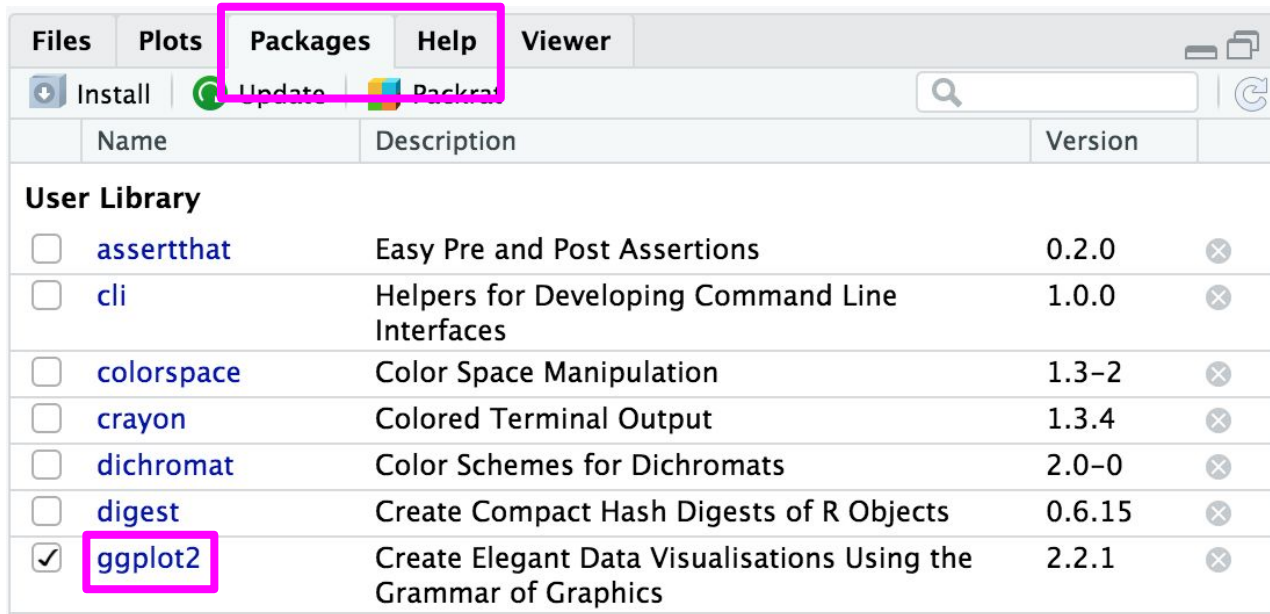
```
rlang_0.2.0
```












```
[13] munsell_0.4.3
```



```
help()
```

```
help(package = "ggplot2")
```



Files	Plots	Packages	Help	Viewer	
 Install	 Update	 Packrat			 <input type="text"/>
	Name	Description	Version		
<b>User Library</b>					
<input type="checkbox"/>	<a href="#">assertthat</a>	Easy Pre and Post Assertions	0.2.0		
<input type="checkbox"/>	<a href="#">cli</a>	Helpers for Developing Command Line Interfaces	1.0.0		
<input type="checkbox"/>	<a href="#">colorspace</a>	Color Space Manipulation	1.3-2		
<input type="checkbox"/>	<a href="#">crayon</a>	Colored Terminal Output	1.3.4		
<input type="checkbox"/>	<a href="#">dichromat</a>	Color Schemes for Dichromats	2.0-0		
<input type="checkbox"/>	<a href="#">digest</a>	Create Compact Hash Digests of R Objects	0.6.15		
<input checked="" type="checkbox"/>	<a href="#">ggplot2</a>	Create Elegant Data Visualisations Using the Grammar of Graphics	2.2.1		

```
browseVignettes()
```

```
browseVignettes("ggplot2")
```

**Vignettes found by "browseVignettes("ggplot2")"**

**Vignettes in package ggplot2**

- Aesthetic specifications - [HTML](#) [source](#) [R code](#)
- Extending ggplot2 - [HTML](#) [source](#) [R code](#)



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