PHIONION

CRAN & Bioconductor

Major repositories for **R packages** that extend R functionality

CRAN: Comprehensive R Archive Network

- CRAN is a network of mirrored servers around the world that administer and distribute R itself, R documentation and R packages (basically add on functionality!)
- There are currently ~9,000 packages on CRAN in the areas of finance, bioinformatics, machine learning, high performance computing, multivariate statistics, natural language processing, etc. etc.

https://cran.r-project.org/

Side-note: R packages come in all shapes and sizes











R packages can be of variable quality and often there are multiple packages with overlapping functionality.

Refer to relevant publications, package citations, update/maintenance history, documentation quality and your own tests!

The journal has sufficient experience with CRAN and Bioconductor resources to endorse their use by authors. We do not yet provide any endorsement for the suitability or usefulness of other solutions.

From: "Credit for Code". Nature Genetics (2014), 46:1

https://cran.r-project.org



Contributed

Contributed Packages

13. C

Available Packages

Currently, the CRAN package repository features 8952 available packages.

@ cran.r-project.org

Table of available packages, sorted by date of publication

Table of available packages, sorted by name

Installation of Packages

Please type help("INSTALL") or help("install.packages") in R for information on how to install packages from this repository. The manual R Installation and Administration (also contained in the R base sources) explains the process in detail.

<u>CRAN Task Views</u> allow you to browse packages by topic and provide tools to automatically install all packages for special areas of interest. Currently, 33 views are available.

Package Check Results

All packages are tested regularly on machines running Debian GNU/Linux, Fedora, OS X, Solaris and Windows.

The results are summarized in the check summary (some timings are also available). Additional details for Windows checking and building can be found in the Windows check summary.

Writing Your Own Packages

The manual <u>Writing R Extensions</u> (also contained in the R base sources) explains how to write new packages and how to contribute them to CRAN.

Repository Policies

The manual CRAN Repository Policy [PDF] describes the policies in place for the CRAN package repository.

Installing a package

RStudio > Tools > Install Packages

- > install.packages("bio3d")
 - > library("bio3d")

Pick a package to explore and install

Rmarkdown

Reports, websites, documenting etc.: Promoting reproducibility.

ggplot2

 Popular graphics package: We have already explored this.

bio3d

 Widely used and highly cited structural bioinformatics package.

Bioconductor

R packages and utilities for working with high-throughput genomic data

http://bioconductor.org



More pragmatic:

Bioconductor is a software repository of R packages with some rules and guiding principles.

Version 3.3 had 1211 software packages.

Bioconductor has emphasized

Reproducible Research since its start, and has been an early adapter and driver of tools to do this. "Bioconductor: open software development for computational biology and bioinformatics" Gentleman et al Genome Biology 2004, 5:R80

"Orchestrating high-throughput genomic analysis with Bioconductor"

Huber et al

Nature Methods 2015, 12:115-121