# Building R Packages An Introduction

David Diez Biostatistics Dept Harvard SPH

## Original version and source

Original author: David M Diez

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## Why build an R package?

#### Accessible

- Functions and objects contained in a package and installed on a machine can be easily loaded;
  - > library(myPackage)
- Many R users develop their own functions that they use regularly
- Even for a sole user, putting code into a package can be worthwhile

#### Reliable

- Documentation structure is familiar, and it is pretty easy to edit
- Testing can be built into the package itself

#### Clarity

- The process of organizing code and data into a package requires a project to become organized
- The result is less ambiguity about project goals and greater clarity about how the project will be completed



## Why every grad student should build a package

### Three important blocks of modern statistics

- Math: methodological development
- Science: applications to real world problems
- Computing: make statistical methods accessible

#### Fulfilling the computing block

- Traditional research focuses more on methods and applications
- Building an R package suggests competence in computing

#### **Employability**

- Not many grad students build an R package
- Display an ability to generalize code and make it user-friendly
- Potential employers can better understand what you've worked on

## Important software principles

#### Goal or mission

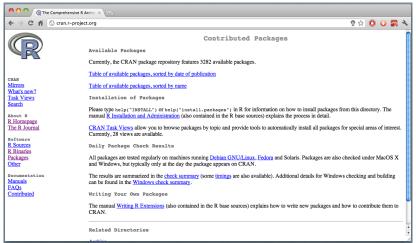
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### Coding principles

- Make high-quality software
- Implement clean coding practices so the code can be adequately reviewed and verified
- Provide helpful documentation

## Sharing data, functions, and an analysis online

Currently, CRAN features **3282** available packages (as of 9/15/2011, up from 2564 on 10/5/2010).





## What are all these packages?

#### Statistical and other methods

- Many R packages make accessible previously or newly developed statistical methods
- Graphical functions, complex numerical techniques, making it easier to work with big data sets, etc.

### Open research

- Publishing a paper for a new method does not make the technique open and accessible
- Ideally, researchers could try out a new method without prohibitive effort, i.e. not have to code it themselves

#### Data

- Sharing old, new, simulated, or research data sets
- Many of the best packages have both methods and data



## 3282 packages and counting

### Initially daunting

- If there are already so many packages, is there room for one more?
- Some might say the same about research: There are so many statistical methods, so can I really develop something both novel and helpful?
- The answer to each question better be yes for the sake of anyone wanting a PhD

## Keep an eye out

### If you are performing raw coding in R, one of the following is true:

- You are ignoring prebuilt functions in R or in an available package
- The method is too user-specific to have a general function
- This may be a place for a new R package

### Ultimate goal

Build a package to fulfill a need

#### Considerations

- The span of R users is wide: applied, software development, visualization, teaching, etc.
- Even if a method is already available, it doesn't mean it was written well or is accurate
- Some R user groups are ignored: find a niche

### Create a mission

### Ask important questions from the start

- What are you good at?
- What is needed?
- What part of that gap can you fill?

#### Identify a target audience

- Beginner or advanced?
- Researcher, student, or teacher?
- Do your target users have lots of data or none at all?

#### Avoid wasted time

- When a package isn't needed, identify this early on
- Sketch out what would be included in the ideal package that accomplishes your mission
- Identify at what stage the package (possibly preliminary) could and/or should be released



### Example: stockPortfolio

Offer a "starter" package for financial analysts who want to get into statistical modeling with R but have little background in statistical finance and/or R  $\,$ 

What is needed: a logical procedure to familiarize the process of collecting data, modeling, and obtaining results from models:

(1) Get the data

```
> tickers <- c('C','BAC', 'WFC', 'GS')
> financials <- getReturns(tickers, start='2004-01-01',
+ end='2008-12-31')</pre>
```

(2) Build the model

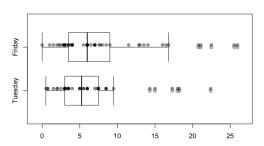
```
> sm <- stockModel(financials, model='CCM')</pre>
```

(3) Obtain the optimal portfolio

```
> opSM <- optimalPort(sm)</pre>
```

## Example: openintro

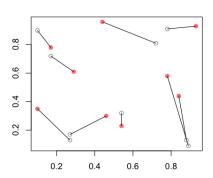
Provide data and simple graphical functions for reproducing results and figures in the **OpenIntro Statistics** textbook



## Example: ppMeasures

Provide basic functions for implementing new methods and reproducing major results from dissertation work

```
> data(pattEx2)
> x <- pattEx2[pattEx2[,1] == 1,c(2,3)]
> y <- pattEx2[pattEx2[,1] == 2,c(2,3)]
> (hold2 <- stDist(x, y, 2))
[1] 5.54
> summary(hold2)
Algorithm: IMA
Max branch: 4
9 points were matched
Distance: 5.54
> plot(hold2)
```



## Find what already exists

#### Examine CRAN

- cran.r-project.org
- Look for similar topics
- Identify the audience of other packages
- Check if overlapping packages are adequate

### Other repositories to check/consider

- R Forge: rforge.net
- Bioconductor: bioconductor.org
- This list is not exhaustive!

## Writing and generalizing code

### Balance simplicity with complexity

- Want to offer diverse set of options
- It should be in a form that it will not be too complex
- Target audience is a guide to the right balance
- Offer many arguments and choose appropriate defaults

#### Example: stockPortfolio

- The models implemented are generally basic
- Intended for folks breaking into stock modeling, i.e may not be familiar with R
- Result: 3-step procedure for implementing any of the models from only a few functions
- Advanced options are made available for users who are interested in learning more

## General R coding advice

#### Performance

- Initialize an entire object rather than grow it slowly
- Compute unchanging values only once

#### Functionality

- Choose variable and function names carefully
- Create default values and use ... in functions when it's helpful
- Outputting a list? Give each list item a name

#### **Aesthetics**

- Align assignment characters
- Use tabs and white space for alignment or when it is meaningful
- If including comments, do so in a style that is not obstructive
- Avoid all caps
- No more than one assignment per line of code

## Evaluating and re-evaluating

#### Build a foundation of diverse examples

• Look for ways to improve speed, accuracy, and usability

### Sufficiently general

- Have a colleague/friend look at the function
- Does it work well for the original problem?
- Is it easy to apply to similar problems?
- Can it be further generalized, or would that be too confusing?

#### Example

- Røme glm wasn't built in a day
- Developers could have made one function for each scenario
- Instead they simplified everything: different scenarios are addressed by modifying arguments, and these arguments have good defaults

## Picking data sets

### Which examples highlight the package?

- If the package is function-centric, choose examples that highlight the performance and graphics
- If your method might be known to be a poor choice in some instances, it would be helpful to point this out to researchers, possibly with an example
- For data-centric packages, use basic functions to show off the data
- Be clear if data are not real or were collected in a haphazard fashion
- Real data are strongly preferred

#### Common knowledge worth repeating

 Don't release data unless you have permissions to or the data are public

### Why use classes?

### Classes make it easy to apply general R functions

- We can change the class of an object in R to be 'ourClass';
   class(myObject) <- 'ourClass')</li>
- Next we build special methods, e.g.
  - o print.ourClass
  - summary.ourClass
  - plot.ourClass
- When we apply plot to a function of class 'ourClass', R actually applies the function plot.ourClass

#### Classes are useful for communication and experimentation

 Allowing the user to connect new functions with old functions is helpful

Downside: classes can mask what is actually contained in an object

### How to create classes

### An object's class can be assigned:

```
stockModel <- function (stockReturns, # other args omitted
){
  # some code omitted
          <- list()
                          # tM = The Model to be returned
    t.M
    class(tM) <- "stockModel" # assign the class !</pre>
    tM$model <- model[1]
  # lots of amazing R code
   return(tM)
```

## Building methods

```
print.stockModel <- function (x, ...){</pre>
    cat("Model:", x$model, "\n")
    cat(x$n, "observations, each one", x$period, "apart\n")
  # some code omitted
    colnames(hold) <- theNames
    temp
                   <- format(hold, digits = 2, scientific = FALSE)
    print(temp)
plot.stockModel <- function (x, xlab = "Risk", # some args omitted</pre>
){
    # code for plotting a stockModel object, x
    # ...
    # want to return some object?
    invisible(objectToReturn)
}
```

### Considerations

#### Pros of classes

- Users can apply familiar R functions to new objects
- Allows output to be formatted for user digestion
- Saves the user time in finding or visualizing important information

#### Cons of classes

- Using methods for classes especially for print takes the user one step away from the true R object
- Some users are unsure how to explore all the attributes of new objects

#### General tip: see what's in a list object via subsetting or str:

- > objName[1:5] # prints first five list items
- > str(objName) # prints summary information

### Overview

#### Step 1: Create the package files

- Package all data and objects from an R session;package.skeleton('packageName')
- See ?package.skeleton for additional options

#### Step 2: Edit the package files

- The DESCRIPTION and help files (man > .Rd) need to be filled in
- Changes to functions should be done directly to the package files
- C or other non-R source code is placed in its own src folder

#### Step 3: Build, check, and install the package

- Run a few Unix commands to build, check, and install the package
- Usually errors arise when checking the package, so return to step 2 as needed

## Step 1: The package files

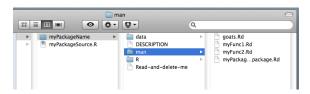
### Create the package files:

```
> myFunc1 <- function(x){ }
> myFunc2 <- function(xy, o5){ }</pre>
>
> goats <- data.frame(beards = rexp(50),
                      tails = rnorm(50, 10)
+
>
> package.skeleton("myPackageName")
Creating directories ...
Creating DESCRIPTION ...
Creating Read-and-delete-me ...
Saving functions and data ...
Making help files ...
Done.
Further steps are described in './myPackageName/Read-and-delete-me'.
```

## Step 1: The package files

#### Folders within the newly created myPackageName folder

- data Contains .rda files of each data object
- R Contains .R files for each function
- man Help files for each function, data set, and the function



- src Create this folder for any C or FORTRAN source code
- tests Create this folder for any test code
- o Other folders with special meanings: demo, exec, inst, po

## Step 2: Edit the package files

#### Edit the DESCRIPTION file

- Update all information
- Choose your license (e.g. GPL-3)



## Step 2: Edit the package files

### Edit each help file in the man folder

- Use \code{ } to write in Courier
- Link to other help files via \link{ }: \code{\link{myFunc2}}
- May create new help files via prompt function in R

```
mvFunc2.Rd
\name{myFunc2}
\alias{myFunc2}
%- Also NEED an '\alias' for EACH other topic documented here.
%% ~~function to do ... ~~
\description{
XX --- A concise (1-5 lines) description of what the function does. ---
\usaae{
myFunc2(xy, o5)
%- maybe also 'usage' for other objects documented here.
\arauments{
  \item{xy}{
     ~Describe \code{xv} here~
  \item{o5}{
     ---Describe \code{o5} here---
XX --- If necessary, more details than the description above ---
\value{
XX ~Describe the value returned
XX If it is a LIST, use
%% \item{comp1 }{Description of 'comp1'}
%% \item{comp2 }{Description of 'comp2'}
xx ...
```

## Step 2: If documentation is not important

#### DESCRIPTION file

• Choose your license (e.g. GPL-2)

#### In the man folder

- Make sure all help files have some title that is not commented out
- In the package help file (man > myPackageName-package.Rd), leave the examples section empty or put in only working R code

### Caution: If you don't build adequate help files...

- Will the package be clear when you return to it in a year?
- Is saving time now worth the chance of spending more later?

## Step 3: Build, check, and install the package

#### There are a few ways to do this. One way:

- Drag/drop package file to the Desktop
- Open Unix (e.g. Terminal in Mac OS X), navigate to the desktop, and type
  - R CMD build myPackageName
  - R CMD check myPackageName
  - R CMD install myPackageName

```
Terminal — bash — 74x13

Last Login: Tue Oct 12 15:30:18 on tty:000
David-Diezs-MacBook-Pro:~ ddiezS cd Desktop
David-Diezs-MacBook-Pro:Desktop ddiezS R CMD build myPackageName

* checking for file leyPackageName/DESCRIPTION' ... OK

* preparing PECRIPTION OF THE CONTROL OF T
```

## Step 3: Build, check, and install the package

### Remarks (for check)

- Warnings and errors are very common in the check stage
- Sometimes the package will install even if check returns an error
- Package only for personal use? Consider skipping the check stage
- CRAN will not accept a package that has warnings or errors from check

### Other useful UNIX commands

- R CMD REMOVE packName
  - Remove a package
- R CMD BUILD --binary packName
  - Creates a binary archive of a package
- R CMD Rd2pdf packName
  - Make a PDF manual for a package

## Recap on building the package

#### Step 1: Create the package files

Packaging all data and objects in an R session is easy;
 package.skeleton('packageName')

#### Step 2: Edit the package files

- Fill in DESCRIPTION and man files
- May edit functions, but make corresponding changes in help files

#### Step 3: Build, check, and install the package

- If a package is being submitted to CRAN, it must pass check
- Warning: installing a package will overwrite any previous version of the package

### Potential trouble

# Packages A and B have different functions but these functions share the same name, fcnName

- One of your functions relies on fcnName from package A
- If user loads your package (which also loads package A), that user might also load package B
- If your package doesn't have a namespace but relies on fcnName, the function from package B might be called instead of the function from package A

### Namespaces help prevent such errors

### Namespaces

Namespaces manage how the user can interact with a package, and it also facilitates high-level communication among packages

- A NAMESPACE file is optional, but if added it goes in the main directory of the package
- Contains instructions for what is imported from other packages
- Describes what files should be easily accessed by other packages

Most packages can get by without a namespace, but occasionally trouble can arise

Tip: use a namespace when publishing a package to CRAN whenever your package relies on another package

Tip: build your namespace after you have stopped adding or removing functions from the package to be released

## Submitting to CRAN

#### Verbatim from CRAN:

To "submit" to CRAN, simply upload to ftp://cran.r-project.org/incoming and send email to cran@r-project.org. Please do not attach submissions to emails, because this will clutter up the mailboxes of half a dozen people.

Note that we generally do not accept submissions of precompiled binaries due to security reasons. All binary distribution listed above are compiled by selected maintainers, who are in charge for all binaries of their platform, respectively.

## Submitting to CRAN

### Before submitting

- Install the package on your computer and ensure the help files and examples look proper and run as expected
- Verify one last time that R CMD check comes with no warnings or errors

### Uploading files

Use an FTP client to upload files

#### Keep in mind

 CRAN personel post packages for free, so be especially considerate of their time

### Remarks

#### Packages can lead to papers

- Initially a package may provide support for an applied and methodological paper in the name of open research
- A robust package can have its own paper

#### Two journals to consider, both with free access

- Journal of Statistical Software www.jstatsoft.org
- R Journal journal.r-project.org

#### Find the source of packages on their CRAN pages



## Helpful references

### Software for Data Analysis

John Chambers

Springer, 2008

### Creating R Packages: A Tutorial

Friedrich Leisch

Department of Statistics Ludwig-Maximilians-Universität München

R Development Core Team

http://cran.r-project.org/doc/contrib/Leisch-CreatingPackages.pdf Friedrich.Leisch@R-project.org