Fundamentals of Creating R Package

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System preparation

- Install the latest version of R and RStudio:
 - R (3.6.2): https://cran.r-project.org/
 - RStudio (1.2.5033): https://rstudio.com/products/rstudio/download/#download
- Install the following packages:
 - devtools
 - roxygen2
 - testthat
 - knitr

```
install.packages("devtools")
install.packages("roxygen2")
install.packages("testthat")
install.packages("knitr")
```

• Get updated version of devtools from GitHub:

```
devtools::install_github("r-lib/devtools")
```

• Check if the system is ready for package building:

```
devtools::has_devel()
```

If following message is shown, it suggests that the system is ready for package development. The C compiler installation part can be skipped.

Your system is ready to build packages!

Depending on the system, a C compiler may be required. If required:

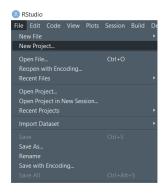
- For windows, install Rtools https://cran.r-project.org/bin/windows/Rtools/
- For Mac, get Xcode (available for free in the App Store)

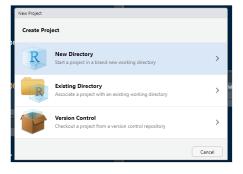
Create library

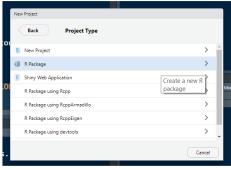
To create a new package, open RStudio

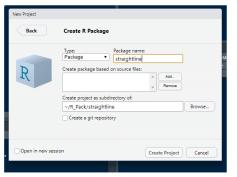
File >> New Project >> New Directory >> R Package

Give the name of the package straightline (or anything you like...)







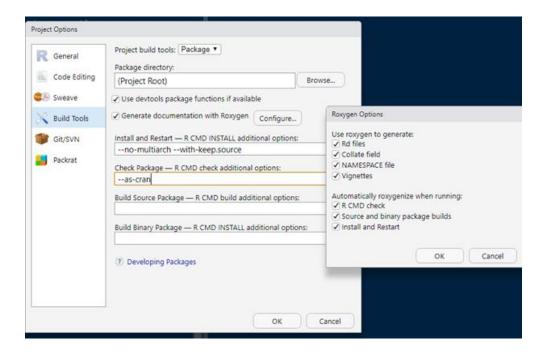


Package naming guidelines

- Unique name
- Matcehs with the type of problem the library deals with
 - Easy to google
 - Example neuralnet, NLP
- Case consistency
 - Same case preferred
 - Mixing not uncommon (randomForest)
- It is not uncommon to see an extra r in the name
 - Example: stringr, knitr
 - straightliner? Instead of straightline
- Formal requirements for an R package:
 - Contains only alphabetic letter, number and period
 - Cannot end with a period
 - Can start with only letter

Package configuration

To let roxygen to some heavylifting, go to Buid > Configure Build Tools and configure as follows:



Build and check package

RStudio includes a hello function with documentation in a default package when a package is created. To build the package, Build > Install and Restart (Cntl + Shift + B). This will install the package in the system and will be included in R session.

After building, check the package, Build > Check Package (Cnt1 + Shift + E). This command will check the package using best practices.

If everything left unaltered, the check should return with a warning for non-standard license.

```
-- R CMD check results
Duration: 22.8s

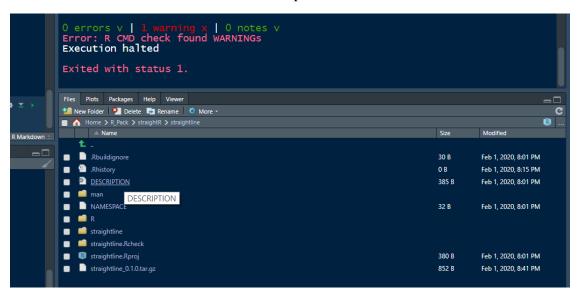
> checking DESCRIPTION meta-information ... WARNING
Non-standard license specification:
    What license is it under?
    Standardizable: FALSE

0 errors v | 1 warning x | 0 notes v
Error: R CMD check found WARNINGs
Execution halted

Exited with status 1.
```

Package metadata

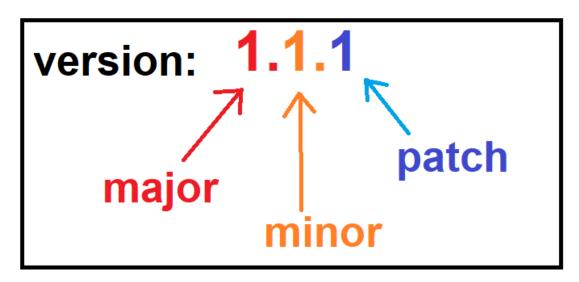
- Every package must have a description
- There will be an auto-generated DESCRIPTION file
- Using the file explorer-plot window of RStudio, go to the package folder and double click the DESCRIPTION file. This will open the file in RStudio editor window.



Edit Title, Version, Author, Maintainer, Description and License in the DESCRIPTION file. An example:

```
Package: straightline
Type: Package
Title: An Example R Package for Straight Line in 2d Coordinate System
Version: 1.1.1
Author: Riaz Khan
Maintainer: Riaz Khan <rk@sdsu.net>
Description: This package will do basic calculations of a straight
    line in 2D coordinate system. This package will do basic
    calculations of a straight line in 2D coordinate system. This
    package will do basic calculations of a straight line in 2D
    coordinate system.
License: GPL-3
Encoding: UTF-8
LazyData: true
RoxygenNote: 7.0.2
```

Version



Multiple authors

If there are multiple authors/contributors, then Authors@R field can be used:

Following roles can be added in the Authors@R filed:

- cre: maintainer
- aut: author having significant contributions
- ctb: contributors with smaller contributions
- cph: copyright holder, in case where the copyright holder is other than the author (e.g employer of the author)

License

There are three that can be used in an R package:

 MIT: Simple permissive. Anyone can do anything as long as the original copyright and license notice included. See https://tldrlegal.com/license/mit-license • GPL-2, GPL-3: If the code is distributed in a bundle, then the whole bundle must be in GPL license. In case of code modification, the source code must be available. GPL-3 is little stricter than GPL-2. See https://tldrlegal.com/license/gnu-general-public-license-v2, https://tldrlegal.com/license/gnu-general-public-license-v3-(gpl-3)

Dependencies

Frequently, a new package uses the functionality of other packages. For this package dependency, Imports field must be used. If a package is used, but not required (likely for examples), then those should be declared as Suggests. For example:

Package: straightline

Type: Package

• • •

Imports: ggplot2
Suggests: MASS

Package documentation

So far, we have only one function in our package, called hello. Let us see the documentation of this function.

?hello

A description, usage and example is given in the function Help section. This documentation is created by the default Rd file. Go to ~/man/hello.Rd to see the actual codes that generated this documentation.



Note:

• Documentation and R files live in separate folders

- ~/man and ~/R folders respectively
- R and Rd documents need to be written separately
- Roxygen- codes and documentation simultaneously
- Roxygen auto generates the Rd files
 - When we configured at the beginning, we told roxygen to handle the Rd files along with other things

Go to the manual directory and delete the hello.Rd file. Then build the package (ctrl+Shift+B) and run the following again.

?hello

Now there is no documentation for the hello function. Now we will try to re-create the documentation.

• Go to the hello.R, put the cursor on hello where the function is defined. Then click Code > Insert Roxygen Skeleton. This will create some roxygen style field for function documentation. These fields can be used for function documentation.

Note: The title can be documented using the @title tag.

Now, build the package again. Examine the Rd file that is generated by roxygen. Now, documentation will be available for the hello function.

Note: It is recommended that we check the package periodically (ctrl+Shift+E). If checking gives warning that the NAMESPACE will not be overwritten, then delete the

NAMESPACE and build+check again. Roxygen generates the NAMESPACE and the Rd files each time we build package.

Writing own function

Functions are main drivers of specific tasks that the package is intended for. Now we will make our own function and document it.

We will write a function (let us call it straighten) that

Takes three inputs a, b, and c, which are the parameters of a straight line of form ax + by + c = 0 in 2D coordinate system. The function will throw error message if a and b both are zero. We want the function to return the followings:

- the value of a
- the value of b
- the value of c
- the slope of the straight line
- x intercept of the straight line
- y intercept of the straight line
- the angle between x axis and the straight line in degrees

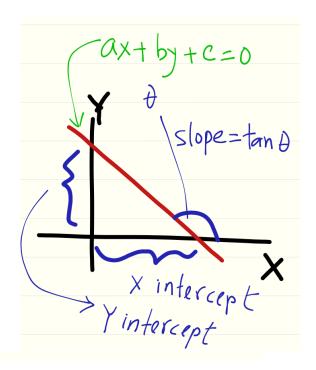
Additionally, we want the return object of the function of its own class, stline

straight line 101

In 2D coordinate system, if a straight line has the

form of ax + by + c = 0, then:

- slope = -a/b
- x intercept = -c/a
- y intercept = -c/b
- angle with x axis = $tan^{-1}(slope)$
- line parallel to x axis, if a = 0
- line parallel to y axis, if b = 0
- passes through the origin, if c = 0



Function straighten

- Function in ~/R folder
- file: straighten.R

```
straighten <- function(a, b, c){</pre>
  if(a==0 & b==0){
    # error message
    stop("a and b both cannot be zero")
  }
  # calculations
  slope <- - a / b
  x_intercept <- - c / a
  y_intercept <- - c / b
  angle_x <- atan(slope) * (180 / pi)</pre>
  # outputs in a list
  value <- as.list(c(a = a, b = b, c = c,
             slope = slope,
             x_intercept = x_intercept,
             y_intercept = y_intercept,
             angle_x = angle_x)
  # output class
  class(value) <- "stline"</pre>
  # return output
  return(value)
}
```

Add roxygen comments to document the function.

```
#' @title Straight Line Measures
#' @param a Parameter \eqn{a}
#' @param b Parameter \eqn{b}
#' @param c Parameter \eqn{c}
#' and angle with x axis of a straight line
#' and angle with x axis of a straight line
#' and angle with x axis of a straight line
#' and angle with x axis of a straight line
#' and angle with x axis of a straight line
#' and angle with x axis are given by \eqn{ax+by+c=0}
#' then slope, x intercept, y intercept and the angle made
#' by the line with x axis are given by \eqn{-a/b}, \eqn{-c/a},

#' \eqn{-c/b} and \eqn{arctan(slope)}
#'
#' ard measures described in details.
#'
#' author Jon Smith
#'
#' author Jon Smith
#'
#' author Jon Smith
#'
#' aseealso \code{\link{hello}}
#'
#' aseealso \code{\link{hello}}
#'
#' asource \url{https://github.com/riazakhan94/straightline}
#' asource \url{https://github.com/riazakhan94/st
```



Function intersection_stline:

```
intersect_stline <- function(x, y){</pre>
  if(class(x) != "stline"){
    stop("x must be of class 'stline'")
  if(class(y) != "stline"){
    stop("y must be of class 'stline'")
  A <- matrix(c(x\$a, y\$a, x\$b, y\$b), nrow = 2)
  determinant_A <- det(A)</pre>
  if(determinant_A == 0){
    stop("The straightlines are either parallel or identical")
  }
  b \leftarrow matrix(c(-x$c, -y$c), nrow = 2)
  solution <- solve(A, b)</pre>
  solution <- c(x_intersect = solution[1,1],</pre>
                 y_intersect = solution[2,1])
  return(solution)
}
```

Create documentation for the intersect_stline function.



Coding style: best practices

- Lower case recommended for function and object, with efficient use of underscore (_)
- There should be spaces before and after all infix operators (e.g: +, -, *, /, =, ==, <-, ->)
 - Space after comma, not before
- Use of <- for assignment (not =)
- Spelling out TRUE and FALSE, as opposed to just using T and F
- Opening curly brace followed by a new line
- Closing curly brace gets its own line
 - Exception: use of else
- Limited number of characters per line
- Use of unique name to avoid conflicts
- Proper indentations for easy understanding

For indentation, select the code block, then Code > Reindent Lines

S3 method: print for class stline

It is recommended that print method can be applied to an object of user defined class. Here we will define the print function applicable for stline object.

```
#' @title Print \code{"stline"} Object
#' @param object An object of class \code{"stline"}
#'
#' @param ... NULL; supplied for S3 method consistency
#'
#' @examples
#' line <- straighten(1, 0, 5)</pre>
#' print(line)
#' line2 <- straighten(7, -5, 0)
#' print(line2)
#'
#' @method print stline
#'
#' @export
print.stline <- function(x, ... = NULL){</pre>
  char_a <- paste0("(", as.character(round(x$a, 3)), ")", "*x")
char_b <- paste0("(", as.character(round(x$b, 3)), ")", "*y")
char_c <- paste0("(", as.character(round(x$c, 3)), ")")</pre>
  Equation <- paste(char_a, "+", char_b, "+", char_c)</pre>
  message("Straight Line Equation : ", Equation)
                                                 , x$slope)
  message("Slope
                                                 , x$x_intercept)
  message("x intercept
  message("y intercept
                                              : ", x$y_intercept)
  message("Angle with x axis (degrees) : ", x$angle_x)
  if(x$a == 0){
    message("Line parallel to x axis")
  if(x$b == 0){
    message("Line parallel to y axis")
  if(x$c == 0){
    message("Line passes through origin")
  }
```

External data

- External data lives in the ~/data folder, with extension .RData
- Data can be saved in .RData format using save
- Example

```
set.seed(10)
a <- runif(100, 0, 100)
set.seed(20)
b <- runif(100, 0, 100)
set.seed(30)
c <- runif(100, 0, 100)
example_data <- data.frame(cbind(a = a, b = b, c = c))
save(example_data, file = "example_data.RData")</pre>
```

- Create a folder called data and keep a copy of the .RData file in it
- Create documentation of the data

```
#'
  @title \code{"example_data"}: An Example Data
#'
#' @description A data containing information about 900 borrowers. It is a
#' modified version of publicly available real data.
#'
#'
#'
#' @format A data frame with 100 rows and 3 variables:
#' \describe{
#'
     \item{a}{A series of parameter \eqn{a}}
#'
     \item{b}{A series of parameter \eqn{b}}
#'
     \item{c}{A series of parameter \eqn{c}}
#'}
#'
#' @examples
#' data("example data")
#' plot(example_data$a~example_data$b)
#' grid()
#'
#'
#' @source \url{https://github.com/riazakhan94/straightline}
"example data"
```

Note: Adding .RData may require to add dependency. If so, edits need to be made in the DESCRIPTION file (Depends: R (>= 3.5.0)).

Dependency

```
#' @title Dependency Example
#'
#' @param x A numeric data series
#' @description This is an exmple to show the use of dependencies
#'
#'
#' @examples
#' df <- MASS::Boston</pre>
#' age <- df$age</pre>
#' example_dependency(age)
#'
#'
#'
#' @export
example_dependency <- function(x){</pre>
  ggplot2::qplot(x, geom = "histogram", binwidth = 10)
}
```

Package vignette

- Inclusion of package vignette recommended.
- Long guide of the package
 - What is does
 - How to use
- Markdown file
- In the vignettes folder
- To create:
 - create a folder called vignette and create an Rmd file (for example, stline.Rmd)
 - add knitr and rmarkdown to the Suggests field in DESCRIPTION
 - add VignetteBuilder field in the DESCRIPTION

```
Suggests: MASS, knitr, rmarkdown
VignetteBuilder: knitr
```

Add the following at the top of the vignette, before starting the document:

See RStudio guide of R Markdown

Package testing

Automated tests are useful for robust code, provides all the tests in an organized way. To set up:

- Make a ~/tests/testthat folder
- Add testthat to the Suggests field of DESCRIPTION
- Make a ~/tests/testthat.R file

A test lives in the ~/tests/testthat directory. All files must start with test. Tests in ~/tests/testthat.R file are checked during the package check.

Test Exmaple

Create ~/tests/testthat/test_classtype.R file:

```
# file: test_classtype.R
context("Class of function straighten")
library(straightline)

myline <- straighten(pi, exp(1), log(7))

test_that("Class of the function return is stline", {
    expect_equal(class(myline), "stline")
})</pre>
```

```
# direct access
test_that("Class of the function return is stline", {
  expect_equal(class(straighten(1,1,0)), "stline")
  expect_equal(class(straighten(2,15,7)), "stline")
  expect_equal(class(straighten(0,1,5)), "stline")
})
```

• Create ~/tests/testthat/test_anlge.R file:

```
# file: test_anlge.R
context("Angle of a straight line")
library(straightline)

test_that("Angle of a straight line 45/-45", {
    expect_equal(straighten(1, -1, 0)$angle_x, 45)
    expect_equal(straighten(1, 1, 0)$angle_x, -45)
})

test_that("Angle of a straight line 60/-60", {
    expect_equal(straighten(1, 1/sqrt(3), 0)$angle_x, -60)
    expect_equal(straighten(1, -1/sqrt(3), 0)$angle_x, 60)
})

test_that("Angle of a straight line 30/-30", {
    expect_equal(straighten(1, sqrt(3), 0)$angle_x, -30)
    expect_equal(straighten(1, sqrt(3), 0)$angle_x, 30)
})
```

• Create ~/tests/testthat/test_error.R file:

```
# file: test_error.R

context("Error message if both a = b = 0")
library(straightline)

test_that("a, b both cannot be zero", {
   expect_error(straighten(0, 0, 3), "a and b both cannot be zero")
})
```

- More test functions:
 - expect length
 - expect message
 - expect match
 - expect null
 - See testthat documentation

- Add all the tests in ~/tests/testthat.R file:
 - the library needs to added once for all the tests

After writing all the tests in ~/tests/testthat directory, run all the tests by running devtools::test(), or use Ctrl+Shit+T.

Check the package (Ctrl+Shift+E)

Package release

If checking returns 0 errors $v \mid 0$ warnings $v \mid 0$ notes then we are ready for submission

README file

- README is usually aimed for the new users.
- README should contain some high level information about the package and how to use the package.
- README has extension .md.
- Possible to use R Markdown (.Rmd) to create the .md file.
- Example using . Rmd
 - Copy the contents from the README.Rmd file and run. It will create the README.md file. Include the README.md file in the package directory.

NEWS file

- Aimed for the old users.
- Brief description of changes made in the new version
- An example NEWS.md for a new package

Release

- Run devtools::release()
- Iterative process

Go through all the process.

Congratulations!:)

Reference: R packages by Hadley Wickham