



NCAR



Introduction to R

Siparcs

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National Center for Atmospheric Research



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[About R](#)

[Contributors](#)

[What's New?](#)

[Mailing Lists](#)

[Bug Tracking](#)

[Conferences](#)

[Search](#)

R Foundation

[Foundation](#)

[Board](#)

[Members](#)

[Donors](#)

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Documentation

[Manuals](#)

[FAQs](#)

The R Project for Statistical Computing

Getting Started

R is a free software environment for statistical computing and graphics. It compiles and runs on a wide variety of UNIX platforms, Windows and MacOS. To [download R](#), please choose your preferred [CRAN mirror](#).

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

News

- **R version 3.2.0** (Full of Ingredients) has been released on 2015-04-16.
- **R version 3.1.3** (Smooth Sidewalk) has been released on 2015-03-09.
- **The R Journal Volume 6/2** is available.
- **useR! 2015**, will take place at the University of Aalborg, Denmark, June 30 - July 3, 2015.
- **useR! 2014**, took place at the University of California, Los Angeles, USA June 30 - July 3, 2014.

<http://www.R-project.org>

Some advanced topics relevant to Atmospheric Sciences



- Reading and writing NetCDF files

<http://www.image.ucar.edu/Software/Netcdf>

- A climate related precipitation example

<http://www.image.ucar.edu/~nychka/FronrangePrecip/>

- Extreme Value Analysis Applied to Weather and Climate

[http://www.ral.ucar.edu/staff/ericg/extRemes/
extRemes2.pdf](http://www.ral.ucar.edu/staff/ericg/extRemes/extRemes2.pdf)

- Forecast verification and CI's

[http://opensky.library.ucar.edu/collections/TECH-
NOTE-000-000-000-846](http://opensky.library.ucar.edu/collections/TECH-NOTE-000-000-000-846)

How R operates

- A function is implemented from the R prompt using the function name and () even if no arguments are passed.
 - For example, to see the current date and time:
`Sys.time()`
- A (visible) function's source code is listed with just the name at the R prompt.
 - For example, to see the source code for the `Sys.time` function:
`Sys.time`

How R operates

- All work is performed in a “dot” file called `.RData`, referred to as the workspace.
- `.RData` exists in the working directory.
- `getwd()` will show the path for this directory (see also `setwd`).
- Use `save.image()` to save your work.
- `q()` to quit an R session (will prompt you to save the directory or not). Use `q(“yes”)` to avoid the prompt and save, or `q(“no”)` to avoid the prompt and not save.

Packages

- Thousands of user-supplied packages freely available on CRAN, and accessible within an R session.
- Installing a package:
 - `install.packages("package name")`
- Updating already installed packages:
 - `update.packages()`
- Loading an installed package:
 - `library("package name")`

Some Atmospheric Science Oriented Packages



- Spatial Statistics
 - fields, LatticeKrig
 - spatstat
 - sp
 - inla
- Maps
 - maps
- Extreme Value Analysis
 - extRemes
 - SpatialExtremes
 - See also: <http://www.ral.ucar.edu/staff/ericg/softextreme.php>
- Forecast Verification
 - verification
 - SpatialVx
- Handling NetCDF files:
 - ncdf4, ncdf, netCDF
- Skew-T plots
 - RadioSonde

Citing R

- To find out how to cite R itself in a paper, from the R prompt, use:

`citation()`

- To find out how to cite a user-supplied package, e.g., the package called fields, use:

`citation("fields")`

Getting help

- Various search engines and documentation found on the R project web site.
- Within an R session, you can use the help function, or the ? short cut to learn about most functions, packages and data sets.
Examples:
 - ?fields
 - ?Sys.time
- To create a help file template for your own function, package or data set, use the prompt function.

Object oriented

- Both S3 and S4 methods can be used.
- plot, summary, and print are three very standard *generic* functions that can be used for most R objects.
- `methods(plot)`
- `methods(class = "lm")`

Logical Operators

- and: &, &&
 - use "?"&" to get help
- or: |, ||
- equal: ==
- >=, >, <, <=
- !=
- is.na
- is.finite
- is.numeric
- is.logical
- is.element
- many more

Other Operators

- Arithmetic operators:
 - `+`, `-`, `*`, `/`
 - `^`, `**` (power operators, `x^2` or `x**2` give x^2)
- Matrix operations:
 - multiplication: `%*%`
 - Use `?"%*%"` to get help.
 - transpose: `?t`
- Assignment operators
 - `<-`
 - e.g., `tiid <- Sys.time()`
 - `->`
 - e.g., `Sys.time() -> tiid`
 - `?assign`
 - `=`

R objects and classes

- Atomic objects (all components of same mode: numeric, complex, logical, character or raw)
 - vectors
- Recursive objects
 - lists
 - functions
 - expressions
- Classes
 - data frames
 - matrices
 - arrays
 - other ...

Examples

```
x <- numeric(0)
mode( x )
class( x )
length( x )
x
x[ 1 ]
any( is.na( x ) )
x <- c( x, 3 )
length( x )
x[ 1 ]
x <- c( x, rnorm( 99, mean = 3, sd = 25 ) )
x
length( x )
```

Examples

```
plot( x )
```

```
summary( x )
```

```
x <- c("Mary", x )
```

```
class( x )
```

```
mode( x )
```

```
length( x )
```

```
x[ x == "Mary" ] <- NA
```

```
mode( x )
```

```
x <- as.numeric( x )
```

```
x <- x[ !is.na( x ) ]
```

Examples

```
x[ 2:4 ]  
x[ x < 0 ] <- 0
```

```
plot( x, type = "l", col = "darkblue" )
```

```
x <- rexp( length( x ) )
```

```
y <- rgamma( 100, shape = 1 )  
hist( y )
```

```
dat <- data.frame( x = x, y = y, z = rnorm( 100 ) )
```


Examples

```
names( dat )  
colnames( dat )  
rownames( dat )
```

```
class( dat )  
mode( dat )
```

```
plot( y ~ x + z, data = dat )
```

```
par( mfrow = c(1, 2) )  
plot( y ~ x + z, data = dat )
```

```
plot( dat )
```

Examples

```
dim( dat )  
dim( t( dat ) )
```

```
summary( dat )
```

```
dat[1:3, 2] <- 0
```

```
dat$w <- dat$x * dat$y
```

```
plot( dat )
```

```
par( mfrow = c(1,1) )  
boxplot( dat )
```

Examples



```
dat[[ "x" ]]
```

```
dat$x
```

```
dat[, 1]
```

```
dat[, "x" ]
```

```
dat[, -(2:dim( dat)[ 2 ] ) ]
```

```
dat[1:10, ]
```

```
dat[ 1:10, "x"]
```

Examples

```
boxplot( dat, notch = TRUE, col = "darkred" )
```

```
dat$ind <- rep( 1:10, each = 10 )
```

```
X <- as.matrix( dat )
```

```
class( X )
```

```
mode( X )
```

```
dim( X )
```

```
names( X )
```

```
colnames( X )
```

```
X[ , "ind" ]
```

```
plot( X )
```

Examples

```
colnames( X ) <- NULL
```

```
plot( X )
```

```
diag( X )
```

```
Y <- cbind( 1:100, 0, rnorm( 100 ), NA, rexp( 100 ) )
```

```
X <- t( X ) %*% Y
```

```
dim( X )
```

```
X
```

```
summary( X )
```

Examples

```
Y <- X[ -c(2,3), -1]
```

```
sum( is.na( X ) )
```

```
colSums( X )
```

```
rowSums( X )
```

```
sum( X, na.rm = TRUE )
```

```
mean( X )
```

```
mean( X, na.rm = TRUE )
```

```
dat$lab <- rep( c("Moanne", "Tiiwusdei", "Woansdei",  
  "Tongersdei", "Freed", "Sneon", "Snein", "Sko",  
  "Sinne", "Hjerst" ), each = 10 )
```

Examples

?par

```
plot( y ~ x, data = dat, xlab = "Example data",  
      ylab = "Response", pch = 22, cex = 2, bg = "lightblue",  
      col = "darkblue", cex.lab = 2,  
      main = "Example", cex.main = 2 )
```

```
plot( x ~ ind, data = dat, xaxt = "n", pch = 23,  
      bg = "lightblue", col = "darkblue", cex = 1.5 )
```

```
axis( 1, at = unique( dat$ind ), labels = unique( dat$lab ) )
```

Examples

```
plot( x ~ ind, data = dat, xaxt = "n", pch = 23,  
      bg = "lightblue", col = "darkblue", cex = 1.5,  
      xlab = "" )
```

```
axis( 1, at = unique( dat$ind ), labels = unique( dat$lab ),  
      las = 2 )
```


More plotting features

See also:

- ?lines, ?points (for adding to an existing plot)
 - use “lty” argument for different line types
 - use “lwd” argument for different line widths
- ?legend (add a legend to a plot)
- ?title, ?mtext
- Use “xlim” and “ylim” to set the plotting region.
- Use “mar”, “oma”, etc. in call to “par” to change inner and outer margin areas.
- To write a plot to a file, see ?pdf, ?png, ?jpeg
- ?dev.new, ?dev.off

Avoiding loops in R

- Use of logical vectors
- colSums, colMeans, rowSums, rowMeans
- apply, lapply, sapply

Avoiding loops in R

```
laundry <- list()
```

```
laundry[[ 1 ]] <- rbind( 1:3, c(4, -1, 0) )  
laundry$clean <- rexp( 100 )
```

```
laundry
```

```
lapply( laundry, summary )
```

```
summary( laundry )
```

Avoiding loops in R

```
laundry <- list()
```

```
laundry[[ 1 ]] <- rbind( 1:3, c(4, -1, 0) )  
laundry$clean <- rexp( 100 )
```

```
laundry
```

```
lapply( laundry, summary )
```

```
summary( laundry )
```

Avoiding loops in R

```
apply( X, 2, max )
```

```
apply( X, 2, max, na.rm = TRUE )
```

```
apply( X, 1, min )
```

```
apply( X, 1, min, na.rm = TRUE )
```

```
apply( X, 2, range )
```

```
apply( X, 2, range, finite = TRUE )
```

Avoiding loops in R

```
Z <- array( NA, dim = c(3, 4, 2) )
```

```
class( Z )
```

```
mode( Z )
```

```
dim( Z )
```

```
Z[, , 1 ] <- cbind( c(1, 0, -1), rep(1, 3), 1:3, 4 )
```

```
Z[, , 2 ] <- rbind( 1:4, c(-1, 0, 1, 0), 0 )
```

```
Z[, , 1, drop = FALSE ]
```

Avoiding loops in R

```
apply( Z, 1:2, sum )
```

```
apply( Z, 3, sum )
```

```
Z[1,2,1] <- NA
```

```
apply( Z, 1:2, sum )
```

```
apply( Z, 1:2, sum, na.rm = TRUE )
```

Writing Functions in R



```
summy <- function( x, y ) (x + y) / (x - y)
```

```
summy( 4, 2 )
```

```
summy( 1:10, c(-1, -2, 0, 7:1) )
```

```
summy(1, 1)
```


Writing Functions in R



```
summy <- function(x, y, ... ) {  
  
  cat("Hello, I am summy.\n")  
  
  mx <- mean( x, ... )  
  my <- mean( y, ... )  
  n <- x^2 + y^2  
  
  return( list( mx = mx, my = my, n = n ) )  
  
}
```

Writing Functions in R



```
summy(4, 2)
```

```
summy( X, 1 )
```

```
summy( X, 1, na.rm = TRUE )
```

Writing Functions in R



Writing a generic function

```
summy <- function(x, ... ) {  
  UseMethod("summy")  
}
```

```
summy( X )
```

Writing Functions in R



Writing a method function

```
summy.default <- function(x, ... ) {  
  res <- sum( x, ... )  
  class( res ) <- "summied"  
  invisible( res )  
}
```

Writing Functions in R



Writing a method function

```
methods( summy )
```

```
summy( X )
```

```
x2 <- summy( X, na.rm = TRUE )
```

```
x2
```

```
class( x2 )
```

```
mode( x2 )
```

Writing Functions in R



Writing a method function

```
summy.matrix <- function(x, ... ) {  
  
  res <- colSums( x, ... )  
  
  class( res ) <- "summied"  
  
  invisible( res )  
  
}
```

Writing Functions in R



Writing a method function

```
x3 <- summy( X, na.rm = TRUE )
```

```
x3
```

```
x2
```

```
sum( x3 )
```

Writing Functions in R



Writing a method function

```
print.summied <- function( x, ... ) {  
  
  cat("Hello, \n")  
  cat("and thank you for using the summy function.\n\n")  
  
  print(c(x))  
  
  invisible()  
  
}
```


Writing Functions in R



Writing a method function

x2

x3

Writing Functions in R



```
summy.matrix <- function(x, ... ) {  
  theCall <- match.call()  
  res <- colSums( x, ... )  
  class( res ) <- "summied"  
  attr( res, "call" ) <- theCall  
  invisible( res )  
}
```

Writing Functions in R



```
print.summied <- function( x, ... ) {  
  
  cat("Hello, \n")  
  cat("and thank you for using the summy function.\n\n")  
  
  a <- attributes( x )  
  if( !is.null( a$call ) ) print( a$call )  
  
  print(c(x))  
  
  invisible()  
  
}
```

Writing Functions in R



```
x4 <- summy( rnorm( 10 ) )
```

```
x5 <- summy( X )
```

```
x4
```

```
x5
```

S4 Methods



Everything I've showed so far is an S3 method. I know less about S4 methods, but see:

<http://cran.r-project.org/doc/contrib/Genolini-S4tutorialV0-5en.pdf>

Statistical Models



- Base/Stats packages
 - Linear Regression/ANOVA
 - ?lm
 - General linear models
 - ?glm
 - Time series (see also ?lm)
 - ?acf, ?pacf
 - arima, arima.sim
 - Nonlinear least squares
 - ?nls

Also note the method
function for making
predictions from these
models.
?predict

Other

- package: boot
 - ?boot, ?tsboot, ?boot.ci
- package: cluster, fastcluster
 - ?hclust, many more ...
- package: fields, LatticeKrig
 - ?Krig, ?Tps, ?mKrig, ?LKrig
 - ?image.plot, ?poly.image
- package: ggplot2

Reading Data into R



- ?scan
- ?read.table
- ?read.csv
- ?read.fwf
- ?source

Writing data from R to computer



- ?write
- ?write.table
- ?dump

Attaching a workspace

Because `.RData` can become too large if you're using large data sets, it can be more efficient to attach other R workspaces that contain the data. See `?attach`, `?detach` and `?search` to learn more.