



무료 전자 책

배우기

R Language

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1: R

R Docs

R

- R .
- .
- 0 1 .
- R arrays() R " " dim .
- R (,) . , , , .
- .
- .

Examples

R

R RStudio . RStudio R .

Windows :

Visual Studio (2015 3) , IntelliSense R Tools . R .

Windows

1. CRAN Windows R R .
2. .
3. .
4. .

OSX / macOS

1

(0. XQuartz)

1. CRAN R .
- 2.

3..

R MacGUI . GUI / Applications / Folder R.app . Doc . () R.app R . R
/Library/Frameworks/R.Framework/Versions/ . RStudio R GUI R .

2

1. <https://brew.sh/> homebrew (macOS) .
2. brew install R

Mac R-SIG-Mac .

,

apt-get R . CRAN . "" CRAN .

```
sudo apt-get install r-base
```

CRAN CRAN . CRAN . install.packages() install.packages() . Linux .

```
sudo apt-get install r-base-dev
```

Red Hat Fedora

```
sudo dnf install R
```

Archlinux

R Extra .

```
sudo pacman -S r
```

Archlinux R [ArchWiki R](#) .

!

```
"Hello World!"
```

,
help() ? ? R . help.search() ?? .

```
#For help on the help function of R  
help()  
  
#For help on the paste function  
help(paste)      #OR  
help("paste")    #OR  
?paste          #OR  
?"paste"
```

<https://www.r-project.org/help.html> .

R

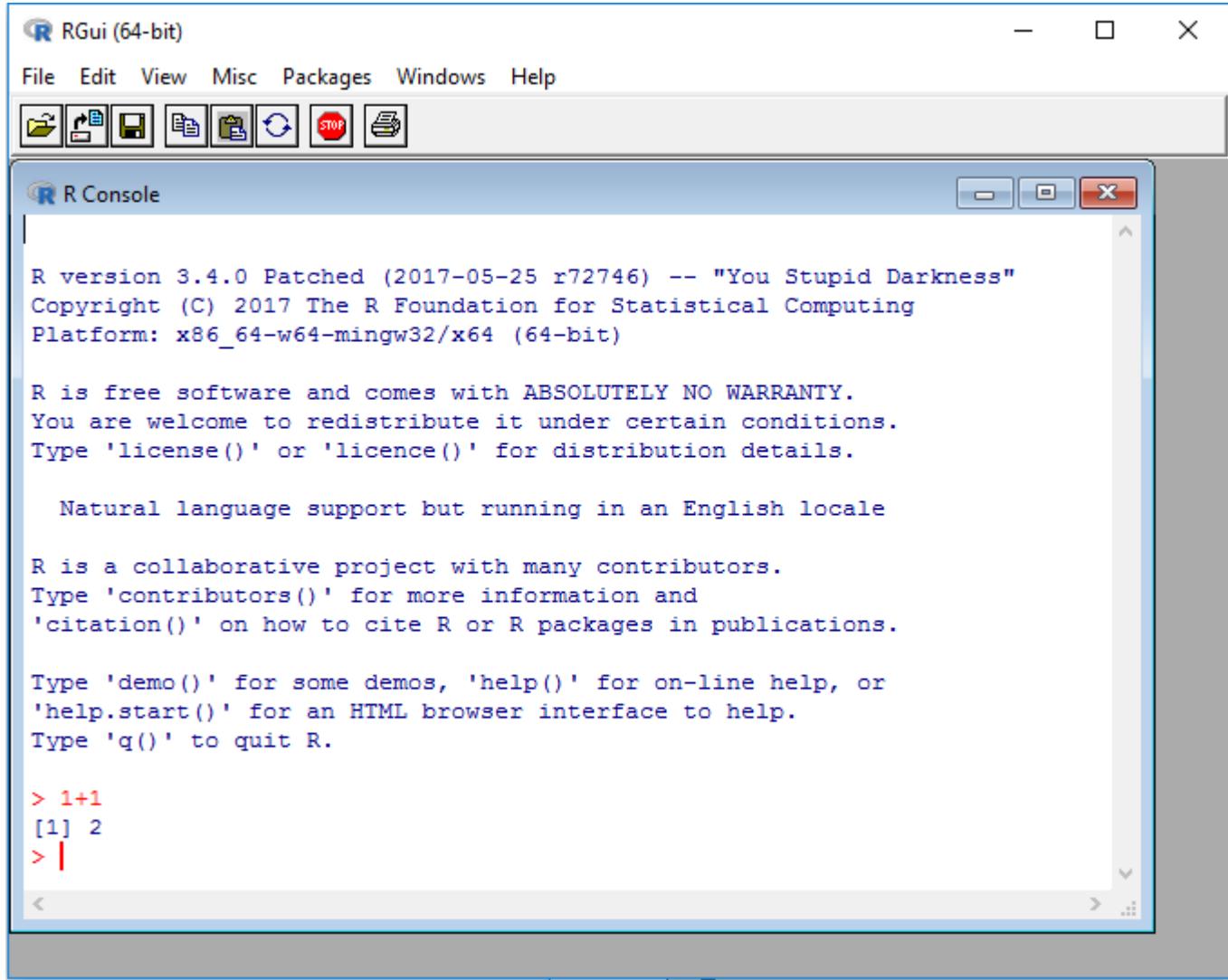
R . R .

R

R Windows RGui R . Linux R .

```
user:~$ R  
  
R version 3.3.2 (2016-10-31) -- "Sincere Pumpkin Patch"  
Copyright (C) 2016 The R Foundation for Statistical Computing  
Platform: x86_64-pc-linux-gnu (64-bit)  
  
R ist freie Software und kommt OHNE JEGLICHE GARANTIE.  
Sie sind eingeladen, es unter bestimmten Bedingungen weiter zu verbreiten.  
Tippen Sie 'license()' or 'licence()' für Details dazu.  
  
R ist ein Gemeinschaftsprojekt mit vielen Beitragenden.  
Tippen Sie 'contributors()' für mehr Information und 'citation()',  
um zu erfahren, wie R oder R packages in Publikationen zitiert werden können.  
  
Tippen Sie 'demo()' für einige Demos, 'help()' für on-line Hilfe, oder  
'help.start()' für eine HTML Browserschnittstelle zur Hilfe.  
Tippen Sie 'q()', um R zu verlassen.  
  
> 1+1  
[1] 2  
> █
```

RGui Windows RGui.



> . R. R.

1+1

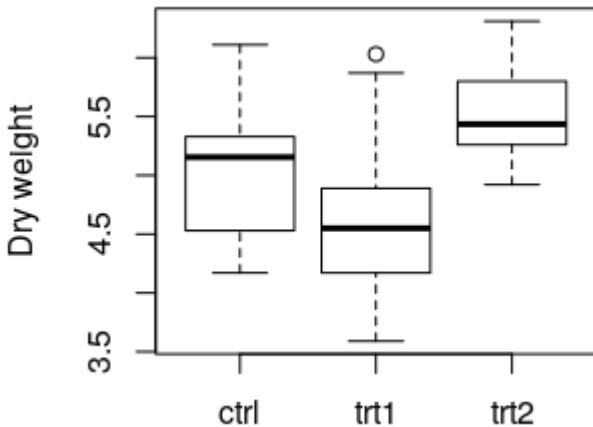
2. [1] R. (2).

R. PlantGrowth. PlantGrowth R

R. ## R.

```
data(PlantGrowth)
str(PlantGrowth)
## 'data.frame':   30 obs. of  2 variables:
## $ weight: num  4.17 5.58 5.18 6.11 4.5 4.61 5.17 4.53 5.33 5.14 ...
## $ group : Factor w/ 3 levels "ctrl","trtl",...: 1 1 1 1 1 1 1 1 1 1 ...
anova(lm(weight ~ group, data = PlantGrowth))
## Analysis of Variance Table
##
## Response: weight
##             Df  Sum Sq Mean Sq F value    Pr(>F)
## group        2  3.7663  1.8832  4.8461 0.01591 *
## Residuals  27 10.4921  0.3886
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
boxplot(weight ~ group, data = PlantGrowth, ylab = "Dry weight")
```



```
data(PlantGrowth)      PlantGrowth . PlantGrowth      () . PlantGrowth PlantGrowth . .
```

- [\(CSV, TSV\)](#)
- [\(Excel, SAS, SPSS, Stata\) I/O](#)

```
str(PlantGrowth)      . PlantGrowth A data.frame R . data.frame 30 . , . $ . weight ( num , ) . , group . R factor . .
```

```
anova(lm( ... )) . weight ~ group "weight" group ". R . data = ... .
```

```
(Column Pr(>F) ), p = 0.01591 ) . Tukey 's Test post-hoc .
```

```
boxplot(...) . . weight ~ group :" group " ylab = ... y :
```

```
q() Ctrl - D R .
```

R

```
. R . R R .
```

```
plants.R .
```

```
data(PlantGrowth)

anova(lm(weight ~ group, data = PlantGrowth))

png("plant_boxplot.png", width = 400, height = 300)
boxplot(weight ~ group, data = PlantGrowth, ylab = "Dry weight")
dev.off()
```

```
( R !).
```

```
R --no-save <plant.R >plant_result.txt
```

plant_result.txt R . .

```
png dev.off() . . . png("FILENAME", width = ..., height = ...) , ( png("FILENAME", width =  
..., height = ...) PNG . dev.off() . dev.off() . .
```

R : <https://riptutorial.com/ko/r/topic/360/r-->

2: * ()

```
*apply  for .for  *apply  .
```

1..

2..

3..

```
for , *apply  .
```

```
for  *apply  for  .
```

```
*apply  *apply
```

```
*apply  *apply  .
```

apply	matrix , data.frame array	()
sapply	list	()
lapply	list	list
vapply	'	
mapply	, lists	list

''' .

Examples

```
apply  ( )  .
```

```
iris . iris  3 150  .
```

```
> head(iris)
```

	Sepal.Length	Sepal.Width	Petal.Length	Petal.Width	Species
1	5.1	3.5	1.4	0.2	setosa
2	4.9	3.0	1.4	0.2	setosa
3	4.7	3.2	1.3	0.2	setosa
4	4.6	3.1	1.5	0.2	setosa
5	5.0	3.6	1.4	0.2	setosa
6	5.4	3.9	1.7	0.4	setosa

```
. for , R apply  ( ).
```

```
> apply(iris[1:4], 2, mean)

Sepal.Length Sepal.Width Petal.Length Petal.Width
      5.843333     3.057333     3.758000     1.199333
```

- mean 4 iris .
- 2 (rxC) . 1 .

```
# standard deviation
apply(iris[1:4], 2, sd)
# variance
apply(iris[1:4], 2, var)
```

:R colMeans colMeans rowMeans .

. 0.5 . mean .

```
> our.mean.function <- function(x) { mean(x[x > 0.5]) }
> apply(iris[1:4], 2, our.mean.function)

Sepal.Length Sepal.Width Petal.Length Petal.Width
      5.843333     3.057333     3.758000     1.665347
```

(Petal.Width .)

?

```
apply(iris[1:4], 2, function(x) { mean(x[x > 0.5]) })
```

, apply .

:apply . *apply ().

- paste0()

```
files <- paste0("file_", 1:100, ".rds")
```

- list.files() () .

```
files <- list.files("./", pattern = "\\.rds$", full.names = TRUE)
```

X .

lapply .

```

readRDS .rds

my_file_list <- lapply(files, readRDS)

for
  . . .
    library() . . .

lapply(c("jsonlite", "stringr", "igraph"), library, character.only=TRUE)

```

`data.frames` (`lapply`, `mapply`)

4

```

library(broom)

#* Create the bootstrap data sets
BootData <- lapply(1:4,
                    function(i) mtcars[sample(1:nrow(mtcars),
                                              size = nrow(mtcars),
                                              replace = TRUE), ])

#* Fit the models
Models <- lapply(BootData,
                  function(BD) lm(mpg ~ qsec + wt + factor(am),
                                 data = BD))

#* Tidy the output into a data.frame
Tidied <- lapply(Models,
                 tidy)

#* Give each element in the Tidied list a name
Tidied <- setNames(Tidied, paste0("Boot", seq_along(Tidied)))

```

data.frame

```

#* Insert the element name into the summary with `lapply`
#* Requires passing the names attribute to `lapply` and referencing `Tidied` within
#* the applied function.
Described_lapply <-
  lapply(names(Tidied),
        function(nm) cbind(nm, Tidied[[nm]]))

Combined_lapply <- do.call("rbind", Described_lapply)

#* Insert the element name into the summary with `mapply`
#* Allows us to pass the names and the elements as separate arguments.
Described_mapply <-
  mapply(
    function(nm, dframe) cbind(nm, dframe),
    names(Tidied),
    Tidied,
    SIMPLIFY = FALSE)

Combined_mapply <- do.call("rbind", Described_mapply)

```

```
magrittr ,      (      ).
```

```
library(magrittr)
library(broom)
Combined <- lapply(1:4,
  function(i) mtcars[sample(1:nrow(mtcars),
    size = nrow(mtcars),
    replace = TRUE), ])
lapply(function(BD) lm(mpg ~ qsec + wt + factor(am), data = BD)) %>%
lapply(tidy) %>%
setNames(paste0("Boot", seq_along(.))) %>%
mapply(function(nm, dframe) cbind(nm, dframe),
  nm = names.,
  dframe = .,
  SIMPLIFY = FALSE) %>%
do.call("rbind", .)
```

: lapply(), sapply() mapply()

R , . . .

- lapply() = .
- sapply() = lapply() .
 - vapply() = sapply() .
- mapply() = mapply() lapply() . . sapply() . .
 - Map() mapply() SIMPLIFY = FALSE mapply() .

lapply()

lapply() . .

- lapply(variable, FUN)
- lapply(seq_along(variable), FUN)

```
# Two ways of finding the mean of x
set.seed(1)
df <- data.frame(x = rnorm(25), y = rnorm(25))
lapply(df, mean)
lapply(seq_along(df), function(x) mean(df[[x]]))
```

sapply()

sapply() . .

```
# Two examples to show the different outputs of sapply()
sapply(letters, print) ## produces a vector
x <- list(a = 1:10, beta = exp(-3:3), logic = c(TRUE,FALSE,FALSE,TRUE))
sapply(x, quantile) ## produces a matrix
```

mapply ()

mapply() lapply() (m).

```
mapply(sum, 1:5, 10:6, 3) # 3 will be "recycled" by mapply
```

. Hadley Wickham Functionals .

```
randomise <- function(f) f(runif(1e3))

lapply2 <- function(x, f, ...) {
  out <- vector("list", length(x))
  for (i in seq_along(x)) {
    out[[i]] <- f(x[[i]]), ...
  }
  out
}
```

, randomise f Uniform . set.seed .

```
set.seed(123)
randomise(mean)
#[1] 0.4972778

set.seed(123)
mean(runif(1e3))
#[1] 0.4972778

set.seed(123)
randomise(max)
#[1] 0.9994045

set.seed(123)
max(runif(1e3))
#[1] 0.9994045
```

functionals (x) (f) base:::lapply mean na.rm f .

```
lapply(list(c(1, 3, 5), c(2, NA, 6)), mean)
# [[1]]
# [1] 3
#
# [[2]]
# [1] NA

lapply2(list(c(1, 3, 5), c(2, NA, 6)), mean)
# [[1]]
# [1] 3
#
# [[2]]
# [1] NA

lapply(list(c(1, 3, 5), c(2, NA, 6)), mean, na.rm = TRUE)
```

```
# [[1]]
# [1] 3
#
# [[2]]
# [1] 4

lapply2(list(c(1, 3, 5), c(2, NA, 6)), mean, na.rm = TRUE)
# [[1]]
# [1] 3
#
# [[2]]
# [1] 4
```

* () : [https://riptutorial.com/ko/r/topic/3567/-----](https://riptutorial.com/ko/r/topic/3567/)

3: .

R

Examples

.Rprofile -

```
.Rprofile .Rprofile R R .R Rprofile.site R .Rprofile.site .Rprofile R .  
: RStudio RStudio .Rprofile .
```

.Rprofile .

R

```
# set R_home  
Sys.setenv(R_USER="c:/R_home") # just an example directory  
# but don't confuse this with the $R_HOME environment variable.
```

```
options(papersize="a4")  
options(editor="notepad")  
options(pager="internal")
```

```
options(help_type="html")
```

```
.Library.site <- file.path(chartr("\\", "/", R.home()), "site-library")
```

```
local({r <-getOption("repos")  
      r["CRAN"] <- "http://my.local.cran"  
      options(repos=r)})
```

R .

```
# library location
```

```
.libPaths("c:/R_home/Rpackages/win")
```

R . .Last.value .

```
makeActiveBinding(".", function(){.Last.value}, .GlobalEnv)
```

.Rprofile R R .

```
help(Startup) . Profile . , Rprofile, , Profile.site . R ${RHOME}/etc .
~/.Renvironment Renvironment Renvironment.site .
```

.profile

```
# Load library setwidth on start - to set the width automatically.
.First <- function() {
  library(setwidth)
  # If 256 color terminal - use library colorout.
  if (Sys.getenv("TERM") %in% c("xterm-256color", "screen-256color")) {
    library("colorout")
  }
}
```

```
# Select default CRAN mirror for package installation.
options(repos=c(CRAN="https://cran.gis-lab.info/"))
```

```
# Print maximum 1000 elements.
options(max.print=1000)
```

```
# No scientific notation.
options(scipen=10)
```

```
# No graphics in menus.
options(menu.graphics=FALSE)
```

```
# Auto-completion for package names.
utils::rc.settings(ipck=TRUE)
```

```
# Invisible environment to mask defined functions
.env = new.env()
```

```
# Quit R without asking to save.
.env$q <- function (save="no", ...) {
  quit(save=save, ...)
}
```

```
# Attach the environment to enable functions.
attach(.env, warn.conflicts=FALSE)
```

. : <https://riptutorial.com/ko/r/topic/4166/>--

4: ANOVA

Examples

aov ()

(aov)

(). aov() Wilkinson-Rogers y~f lm() . y () f aov() ANOVA

aov() Type I () Sum of Squares . (Sum of Squares) () . . ().

Type I Sum of Squares Type II Type III Sum of Squares . Type II Sum of Squares
Type II Sum of Squares .

III . Type III Sum of Squares .

II III Anova() .

mtcars .

```
mtCarsAnovaModel <- aov(wt ~ factor(cyl), data=mtcars)
```

ANOVA :

```
summary(mtCarsAnovaModel)
```

lm()

```
coefficients(mtCarsAnovaModel)
```

Anova ()

/ Type II Type III Sum of Squares . car Anova() . Type II Sum of Squares . Type III .

Anova() lm() .

mtcars II III .

```
> Anova(lm(wt ~ factor(cyl)*factor(am), data=mtcars), type = 2)
Anova Table (Type II tests)
```

```
Response: wt
          Sum Sq Df F value    Pr(>F)
factor(cyl)   7.2278  2 11.5266 0.0002606 ***
factor(am)    3.2845  1 10.4758 0.0032895 **
```

```

factor(cyl):factor(am) 0.0668  2  0.1065  0.8993714
Residuals                 8.1517 26
---
Signif. codes:  0 '****' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

> Anova(lm(wt ~ factor(cyl)*factor(am), data=mtcars), type = 3)
Anova Table (Type III tests)

Response: wt
            Sum Sq Df F value    Pr(>F)
(Intercept) 25.8427  1 82.4254 1.524e-09 ***
factor(cyl)   4.0124  2  6.3988  0.005498 **
factor(am)    1.7389  1  5.5463  0.026346 *
factor(cyl):factor(am) 0.0668  2  0.1065  0.899371
Residuals      8.1517 26
---
Signif. codes:  0 '****' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

```

ANOVA : <https://riptutorial.com/ko/r/topic/3610/anova>

5: data.table

Data.table R . data.table .

- DT[i, j, by]
DT [where, select | update | do, by]
- DT[...] [...]
#
- ##### Shortcuts, special functions and special symbols inside DT[...]
- .()
- list () #
- J()
i, list () .
- :=
in j,
- .
in i,
in j,
- .
in j, (i)
- .SD
in j,
.SDcols #
- .GRP
in j,
- .
in j,
- V1, V2, ...
j
- ##### Joins inside DT[...]
- DT1 [DT2, on, j]
#
- .
join DT2 .
- by = .EACHI
.
- DT1 [! DT2, on, j]
-
- DT1 [DT2, on, roll, j]
, on =
- ##### Reshaping, stacking and splitting
- (DT, id.vars, measure.vars)

measure.vars = patterns (...) .
- dcast (DT,)
#

- rbind (DT1, DT2, ...)
stack data.tables
 - rbindlist (DT_list, idcol)
.
 - (DT, by)
data.table .
 - ##### Some other functions specialized for data.tables
 - #
 - #
 - #
 - fintersect, fsetdiff, funion, fsetequal, , , anyDuplicated
set-theory
 - N
#
 - rowidv (DT, cols)
cols ID (1 ~ N)
 - rleidv (DT, cols)
cols ID (1 .GRP)
 - shift (DT, n, type = c ("lag", "lead"))
.
 - setorder, setcolorder, setnames, setkey, setindex, setattr
#
-

data.table .

```
# install from CRAN
install.packages("data.table")

# or install development version
install.packages("data.table", type = "source", repos =
"http://Rdatatable.github.io/data.table")

# and to revert from devel to CRAN, the current version must first be removed
remove.packages("data.table")
install.packages("data.table")
```

. StackOverflow .

data.table . , library(data.table) , data.table::fread fread . help("fread") ?fread .
?data.table::fread .

Examples

data.table

data.table base R **data.frame** . class() vector "data.table" "data.frame" **data.table** . **data.table**

```
data.table .
```

```
library(data.table)
```

```
DT <- data.table(  
  x = letters[1:5],  
  y = 1:5,  
  z = (1:5) > 3  
)  
#   x y     z  
# 1: a 1 FALSE  
# 2: b 2 FALSE  
# 3: c 3 FALSE  
# 4: d 4  TRUE  
# 5: e 5  TRUE
```

```
data.frame , data.table
```

```
sapply(DT, class)  
#           x          y          z  
# "character" "integer" "logical"
```

```
dt <- fread("my_file.csv")
```

```
read.csv  fread .
```

data.frame

```
data.table data.frame list ( ) data.table .
```

```
# example data.frame  
DF <- data.frame(x = letters[1:5], y = 1:5, z = (1:5) > 3)  
# modification  
setDT(DF)
```

```
DF <- . data.frame .
```

```
sapply(DF, class)  
#           x          y          z  
# "factor" "integer" "logical"
```

data.table .

```
list , data.frame , data.table , setDT A data.table ( as.data.table ). . .
```

```
( :) R as.data.table A data.table .
```

```
mat <- matrix(0, ncol = 10, nrow = 10)  
DT <- as.data.table(mat)  
# or  
DT <- data.table(mat)
```

```
DT[where, select|update|do, by] data.table .
```

- "where" i .
- "select | update | do" j .

```
mtcars = data.table(mtcars, keep.rownames = TRUE)
```

```
j := .
```

```
mtcars[, mpg_sq := mpg^2]
```

```
NULL :
```

```
mtcars[, mpg_sq := NULL]
```

```
:= .
```

```
mtcars[, `:=` (mpg_sq = mpg^2, wt_sqrt = sqrt(wt)) ]  
# or  
mtcars[, c("mpg_sq", "wt_sqrt") := .(mpg^2, sqrt(wt)) ]
```

```
mtcars[, c("mpg_sq", "mpg2_hp") := .(temp1 <- mpg^2, temp1/hp)]
```

```
.() LHS := RHS .
```

```
vn = "mpg_sq"  
mtcars[, (vn) := mpg^2]
```

```
set
```

```
set(mtcars, j = "hp_over_wt", v = mtcars$hp/mtcars$wt)
```

```
i "" .
```

```
mtcars[1:3, newvar := "Hello"]
# or
set(mtcars, j = "newvar", i = 1:3, v = "Hello")
```

```
data.frame
```

```
. i "" , .
```

```
levels<- names<- . data.table .
```

```
setnames data.table data.frame setattr .
```

```
# Print a message to the console whenever the data.table is copied
tracemem(mtcars)
mtcars[, cyl2 := factor(cyl)]
```

```
# Neither of these statements copy the data.table
setnames(mtcars, old = "cyl2", new = "cyl_fac")
setattr(mtcars$cyl_fac, "levels", c("four", "six", "eight"))
```

```
# Each of these statements copies the data.table
names(mtcars)[names(mtcars) == "cyl_fac"] <- "cf"
levels(mtcars$cf) <- c("IV", "VI", "VIII")
```

```
# This function also changes the levels in the global environment
edit_levels <- function(x) setattr(x, "levels", c("low", "med", "high"))
edit_levels(mtcars$cyl_factor)
```

data.table

.SD

```
.SD data.table by .
```

```
.SD lapply A data.table
```

```
mtcars .
```

```
mtcars = data.table(mtcars) # Let's not include rownames to keep things simpler
```

```
, cyl :
```

```

mtcars[ , lapply(.SD, mean), by = cyl]

#   cyl      mpg      disp       hp      drat      wt      qsec      vs      am      gear
carb
#1:  6 19.74286 183.3143 122.28571 3.585714 3.117143 17.97714 0.5714286 0.4285714 3.857143
3.428571
#2:  4 26.66364 105.1364  82.63636 4.070909 2.285727 19.13727 0.9090909 0.7272727 4.090909
1.545455
#3:  8 15.10000 353.1000 209.21429 3.229286 3.999214 16.77214 0.0000000 0.1428571 3.285714
3.500000

```

cyl vs , am , gear carb . mean . . . SDcols .

.SDcols

.SDcols data.table .SD .

() gear cyl gear cyl :

```

# All the continuous variables in the dataset
cols_chosen <- c("mpg", "disp", "hp", "drat", "wt", "qsec")

mtcars[order(gear, cyl), lapply(.SD, mean), by = .(gear, cyl), .SDcols = cols_chosen]

#   gear cyl      mpg      disp       hp      drat      wt      qsec
#1:  3     4 21.500 120.1000 97.00000 3.700000 2.465000 20.0100
#2:  3     6 19.750 241.5000 107.50000 2.920000 3.337500 19.8300
#3:  3     8 15.050 357.6167 194.16670 3.120833 4.104083 17.1425
#4:  4     4 26.925 102.6250  76.00000 4.110000 2.378125 19.6125
#5:  4     6 19.750 163.8000 116.50000 3.910000 3.093750 17.6700
#6:  5     4 28.200 107.7000 102.00000 4.100000 1.826500 16.8000
#7:  5     6 19.700 145.0000 175.00000 3.620000 2.770000 15.5000
#8:  5     8 15.400 326.0000 299.50000 3.880000 3.370000 14.5500

```

mean . . by .

```

mtcars[ , lapply(.SD, mean), .SDcols = cols_chosen]

#      mpg      disp       hp      drat      wt      qsec
#1: 20.09062 230.7219 146.6875 3.596563 3.21725 17.84875

```

:

- cols_chosen . . SDcols .
- SDcols SDcols . mtcars[, lapply(.SD, mean), .SDcols = c(1,3:7)]

.N . N .

```

iris[, .(count=.N), by=Species]

```

```
#      Species count
#1:    setosa     50
#2: versicolor   50
#3: virginica    50
```

data.frame data.table

data.table R data.frame , matrix (2D) array . A[rows, cols] .

(A) matrix data.frame data.table :

```
ma <- matrix(rnorm(12), nrow=4, dimnames=list(letters[1:4], c('X', 'Y', 'Z')))
df <- as.data.frame(ma)
dt <- as.data.table(ma)

ma[2:3] #---> returns the 2nd and 3rd items, as if 'ma' were a vector (because it is!)
df[2:3] #---> returns the 2nd and 3rd columns
dt[2:3] #---> returns the 2nd and 3rd rows!
```

```
ma[2:3, ] # \
df[2:3, ] # }---> returns the 2nd and 3rd rows
dt[2:3, ] # /
```

```
ma[, 2:3] # \
df[, 2:3] # \
dt[, 2:3] # }---> returns the 2nd and 3rd columns
ma[, c("Y", "Z")] # /
df[, c("Y", "Z")] # /
dt[, c("Y", "Z")] # /
```

```
mycols <- 2:3
ma[, mycols] # \
df[, mycols] # }---> returns the 2nd and 3rd columns
dt[, mycols, with = FALSE] # /

dt[, mycols] # ---> Raises an error
```

mycols . dt mycols .
: data.table data.table . . dt . dt[, 2:3] dt[, mycols] 2:3 . mycols .

data.frame data.table

```
data.frame data.table . data.frame . . .
1. .
2. .
3. data.table data.frame( .: print.data.frame).
4. list .
5. data.frame(   ).
6. data.table .

, [ , ] : 

A[1:10, ]
A[A$var > 17, ] # A[var > 17, ] just works for data.table

. $ [[ ]] .
```

```
A$var
colname <- 'var'
A[[colname]]
A[[1]]
```

```
B <- `[.data.frame`(A, 2:4)

# We can give it a better name
select <- `[.data.frame`
B <- select(A, 2:4)
C <- select(A, c('foo', 'bar'))
```

```
" data.frame row.names , data.table key . row.names data.table .
B <- A[A$var != 0, ]
# or...
B <- with(A, A[var != 0, ]) # data.table will silently index A by var before subsetting

stuff <- c('a', 'c', 'f')
C <- A[match(stuff, A$name), ] # really worse than: setkey(A); A[stuff, ]
```

```
1 . . :
```

```
B <- select(A, 2)     #--> a table with just the second column
C <- unlist(A[1, ])  #--> the first row as a vector (coerced if necessary)
```

data.table

```
, 1.9.6 .
```

(1.9.6) data.table data.table, . [544 2015 9 intro vignette 5 .] 'setkey' 'key =' .

```
library(data.table)
DT <- data.table(
  x = letters[1:5],
  y = 5:1,
  z = (1:5) > 3
)

#> DT
#   x y     z
#1: a 5 FALSE
#2: b 4 FALSE
#3: c 3 FALSE
#4: d 2  TRUE
#5: e 1  TRUE
```

setkey . . .

```
setkey(DT, y)
```

()

```
tables()

> tables()
      NAME  NROW  NCOL  MB COLS KEY
[1,] DT      5     3  1 x,y,z y
Total: 1MB
```

```
#> DT
#   x y     z
#1: e 1  TRUE
#2: d 2  TRUE
#3: c 3 FALSE
#4: b 4 FALSE
#5: a 5 FALSE
```

v1.9.6 , . data.table "on=" . SO .

2017 1 "on"

?

key , setindex(DT, key.col) setindexv(DT, "key.col.string") . DT data.table. setindex(DT, NULL)

indices(DT) .

2 ?

"on" . 2 . . .

y DT.

```
DT
# x y      z
# 1: e 1   TRUE
# 2: d 2   TRUE
# 3: c 3 FALSE
# 4: b 4 FALSE
# 5: a 5 FALSE

# Let us set x as index
setindex(DT, x)

# Use indices to see what has been set
indices(DT)
# [1] "x"

# fast subset using index and not keyed column
DT["c", on ="x"]
#x y      z
#1: c 3 FALSE

# old way would have been rekeying DT from y to x, doing subset and
# perhaps keying back to y (now we save two sorts)
# This is a toy example above but would have been more valuable with big data sets
```

data.table : <https://riptutorial.com/ko/r/topic/849/data-table>

6: data.table

- `melt(DT, id.vars=c(..), variable.name="CategoryLabel", value.name="Value")`
- `dcast(DT, LHS ~ RHS, value.var="Value", fun.aggregate=sum) dcast(DT, LHS ~ RHS, value.var="Value", fun.aggregate=sum)`

id.vars	melt	
variable.name	melt	
value.name	melt	
value.var	dcast .	
	(LHS)	(RHS)
.	.	dcast

```
data.table . data.table .
```

Examples

```
data.table - |
```

```
data USArrests datasets data USArrests .
```

```
data("USArrests")
head(USArrests)

    Murder Assault UrbanPop Rape
Alabama     13.2     236      58 21.2
Alaska      10.0     263      48 44.5
Arizona      8.1     294      80 31.0
Arkansas     8.8     190      50 19.5
California   9.0     276      91 40.6
Colorado     7.9     204      78 38.7
```

```
?USArrests . data.table data.table . data.frame .
```

```
library(data.table)
DT <- as.data.table(USArrests, keep.rownames=TRUE)
```

```
. . . . . USArrests .
```

```
      State Crime Rate
1: Alabama Murder 13.2
2: Alaska  Murder 10.0
3: Arizona Murder  8.1
```

```
4:      Arkansas Murder 8.8
5:      California Murder 9.0
---
196:      Virginia Rape 20.7
197:      Washington Rape 26.2
198: West Virginia Rape 9.3
199:      Wisconsin Rape 10.8
200:      Wyoming Rape 15.6
```

melt .

```
DTm <- melt(DT)
names(DTm) <- c("State", "Crime", "Rate")
```

melt . USArests UrbanPop . Murder , Assault , Rape , 10 . UrbanPop . id.vars id.vars .

```
DTmu <- melt(DT, id.vars=c("rn", "UrbanPop" ),
               variable.name='Crime', value.name = "Rate")
names(DTmu)[1] <- "State"
```

(Murder, Assault) variable.name value.name value.name . .

```
          State UrbanPop Crime Rate
1:      Alabama      58 Murder 13.2
2:      Alaska       48 Murder 10.0
3:      Arizona      80 Murder  8.1
4:      Arkansas      50 Murder  8.8
5:      California     91 Murder  9.0
```

split-apply-combine . , ?

```
DTmu[, .(ViolentCrime = sum(Rate)), by=State]
```

:

```
          State ViolentCrime
1:      Alabama      270.4
2:      Alaska       317.5
3:      Arizona      333.1
4:      Arkansas      218.3
5:      California     325.6
6:      Colorado      250.6
```

data.table - II

dcast dcast .

```
DTc <- dcast(DTmu, State + UrbanPop ~ Crime)
```

```

      State UrbanPop Murder Assault Rape
1:   Alabama      58  13.2    236 21.2
2:   Alaska       48  10.0    263 44.5
3:  Arizona       80   8.1    294 31.0
4: Arkansas      50   8.8    190 19.5
5: California    91   9.0    276 40.6

```

(LHS) (RHS) . ? dcast . value.var .

dcast fun.aggregate . . Decile .

```

DTmu[, Decile := cut(UrbanPop, quantile(UrbanPop, probs = seq(0, 1, by=0.1)))]
levels(DTmu$Decile) <- paste0(1:10, "D")

```

Decile ~ Crime . fun.aggregate . .

```
dcast(DTmu, Decile ~ Crime, value.var="Rate", fun.aggregate=sum)
```

:

```
dcast(DTmu, Decile ~ Crime, value.var="Rate", fun.aggregate=mean)
```

```

      State UrbanPop Crime Rate Decile
1:   Alabama      58 Murder 13.2     4D
2:   Alaska       48 Murder 10.0     2D
3:  Arizona       80 Murder  8.1     8D
4: Arkansas      50 Murder  8.8     2D
5: California    91 Murder  9.0    10D

```

. fun.aggregate .

```
dcast(DTmu, Decile ~ Crime, value.var="Rate", fun.aggregate=sum)
```

```

      Decile Murder Assault Rape
1:    1D    39.4     808  62.6
2:    2D    35.3     815  94.3
3:    3D    22.6     451  67.7
4:    4D    54.9     898 106.0
5:    5D    42.4     758 107.6

```

data.table : <https://riptutorial.com/ko/r/topic/6934/data-table--->

7: Date

- Date : 1970-01-01 **UNIX**
- (,).
- .
- POSIXct POSIXlt POSIXlt .
- sys.Date() Date .

- lubridate ymd , mdy Date as.Date . lubridate .
- data.table IDate Date Date double .

Examples

```
Dates format(date, format = "%Y-%m-%d") POSIXct ( as.POSIXct () ) POSIXlt ( as.POSIXlt () )
```

```
d = as.Date("2016-07-21") # Current Date Time Stamp

format(d, "%a")           # Abbreviated Weekday
## [1] "Thu"

format(d, "%A")           # Full Weekday
## [1] "Thursday"

format(d, "%b")           # Abbreviated Month
## [1] "Jul"

format(d, "%B")           # Full Month
## [1] "July"

format(d, "%m")           # 00-12 Month Format
## [1] "07"

format(d, "%d")           # 00-31 Day Format
## [1] "21"

format(d, "%e")           # 0-31 Day Format
## [1] "21"

format(d, "%y")           # 00-99 Year
## [1] "16"

format(d, "%Y")           # Year with Century
## [1] "2016"
```

```
?strptime .
```

```
as.Date()
```

```
> x <- as.Date("2016-8-23")
> x
[1] "2016-08-23"
> class(x)
[1] "Date"
```

as.Date() . %Y-%m-%d , - - .

```
> as.Date("23-8-2016", format="%d-%m-%Y") # To read in an European-style date
[1] "2016-08-23"
```

. H H . "dm-yy" "dm-YYYY" "md-yy" "md-YYYY" "YYYY-md" "YYYY-dm" . "-" "/" . 1986
 11 6 1986 11 6 1986 11 6 1986 11 6 . **as.Date()** "YYYY-md" .

"%d-%m-%Y" "9-6-1962" .

```
#  

# It tries to interprets the string as YYYY-m-d  

#  

> as.Date("9-6-1962")
[1] "0009-06-19" #interprets as "%Y-%m-%d"  

>  

as.Date("9/6/1962")
[1] "0009-06-19" #again interprets as "%Y-%m-%d"  

>  

# It has no problem in understanding, if the date is in form YYYY-m-d or YYYY/m/d  

#  

> as.Date("1962-6-9")
[1] "1962-06-09" # no problem  

> as.Date("1962/6/9")
[1] "1962-06-09" # no problem  

>
```

. **as.Date()** .

%d	
%m	
%Y	2
%Y	4
%b	3
%B	

format .

```
> as.Date("9-6-1962", format="%d-%m-%Y")
```

```
[1] "1962-06-09"  
>
```

```
> as.Date("9-6-1962", "%d-%m-%Y")  
[1] "1962-06-09"  
>
```

· %b ·

```
> as.Date("6Nov1962", "%d%b%Y")  
[1] "1962-11-06"  
>
```

'-' '-' / ' '

```
> as.Date("6 Nov, 1962", "%d %b, %Y")  
[1] "1962-11-06"  
>
```

.. NA . %B ..

```
> as.Date("October 12, 2016", "%B %d, %Y")  
[1] "2016-10-12"  
>  
> as.Date("12 October, 2016", "%d %B, %Y")  
[1] "2016-10-12"  
>
```

%y . **origin tz ()**.

R as.Date() Date ISO 8601 YYYY-MM-DD strftime -style .

```
as.Date('2016-08-01')    # in ISO format, so does not require formatting string  
## [1] "2016-08-01"  
  
as.Date('05/23/16', format = '%m/%d/%y')  
## [1] "2016-05-23"  
  
as.Date('March 23rd, 2016', '%B %drd, %Y')    # add separators and literals to format  
## [1] "2016-03-23"  
  
as.Date(' 2016-08-01 foo')    # leading whitespace and all trailing characters are ignored  
## [1] "2016-08-01"  
  
as.Date(c('2016-01-01', '2016-01-02'))  
# [1] "2016-01-01" "2016-01-02"
```

Date : <https://riptutorial.com/ko/r/topic/9015/date->

8: devtools

devtools R .

1. R
2. roxygen2
3. devtools

Examples

R . . .
./ R .

LaTeX .

roxygen .

```
install.packages("devtools")
library("devtools")
install.packages("roxygen2")
library("roxygen2")
```

roxygen . doxygen .

roxygen .

```
#' Increment a variable.
#'
#' Note that the behavior of this function
#' is undefined if `x` is not of class `numeric`.
#'
#' @export
#' @author another guy
#' @name Increment Function
#' @title increment
#'
#' @param x    Variable to increment
#' @return     `x` incremented of 1
#'
#' @seealso   `other_function`
#'
#' @examples
#' increment(3)
#' > 4
increment <- function(x) {
  return (x+1)
}
```

(vignettes) .

./script1.R ./script2.R .

```
package.skeleton(name="MyPackage", code_files=c("script1.R", "script2.R"))
```

./MyPackage/man/ . . .

```
roxygenize("MyPackage")
```

./ R CMD Rd2pdf MyPackage .

1.

./MyPackage/DESCRIPTION . Package , Version , License , Description , Title , Author Maintainer .

Depends (R version <3.2.0) Imports (R version> 3.2.0) .

2.

./MyPackage/ R/ man/ **subfolders** . , : .

- data/ : . . RData `data()` `load()` `load()` .
 - tests/ : . . .
 - src/ : C / C ++ / Fortran (`Rcpp` ...).
 - exec/ : .
 - misc/ : .
-

./MyPackage/Read-and-delete-me .

devtools::install("MyPackage") .

./ R CMD build MyPackage

Github

```
myPackage      MyPackage/ . . .
```

```
devtools github  :
```

```
install_package("MyPackage", "your_github_username")
```

CRAN

```
CRAN  .  :  ( ). R CMD check .
```

```
. . .  
(vignette) . . . , . (vignette) . . .
```

```
Vignettes markdown .
```

-
- **Rmarkdown**: `install.packages("rmarkdown")`
-

```
devtools::use_vignette("MyVignette", "MyPackage")
```

```
./vignettes/MyVignette.Rmd .
```

```
Markdown .
```

```
Markdown R     Markdown .
```

```
```{r}  
Add two numbers together
add <- function(a, b) a + b
add(10, 20)
```
```

```
# Add two numbers together  
add <- function(a, b) a + b  
add(10, 20)  
## [1] 30
```

```
,      ./DESCRIPTION .
```

```
devtools  : https://riptutorial.com/ko/r/topic/10884/devtools--
```

9: ggplot2

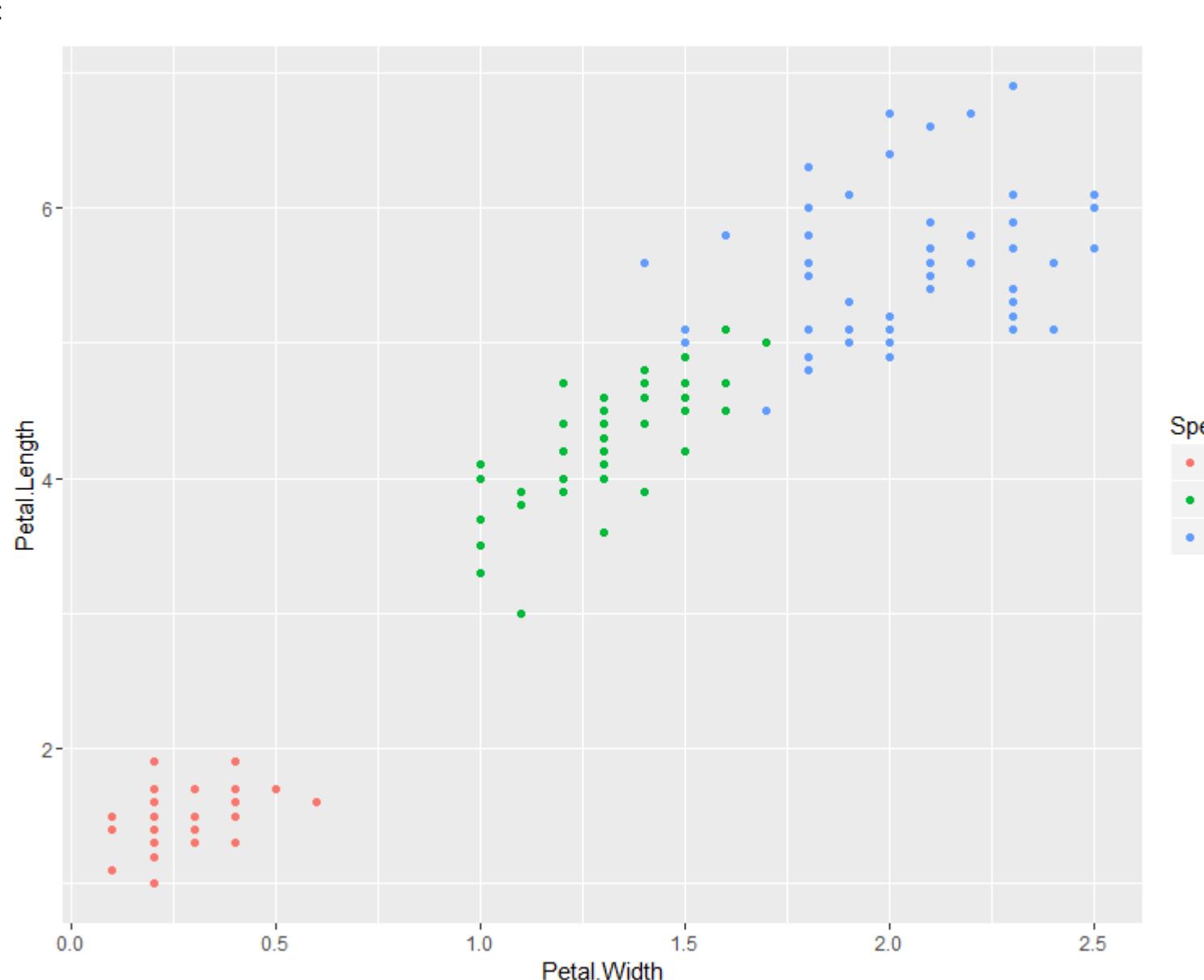
```
ggplot2   http://ggplot2.tidyverse.org/ .
```

```
( : data.frame ) .
```

```
RStudio    "ggplot2" .
```

Examples

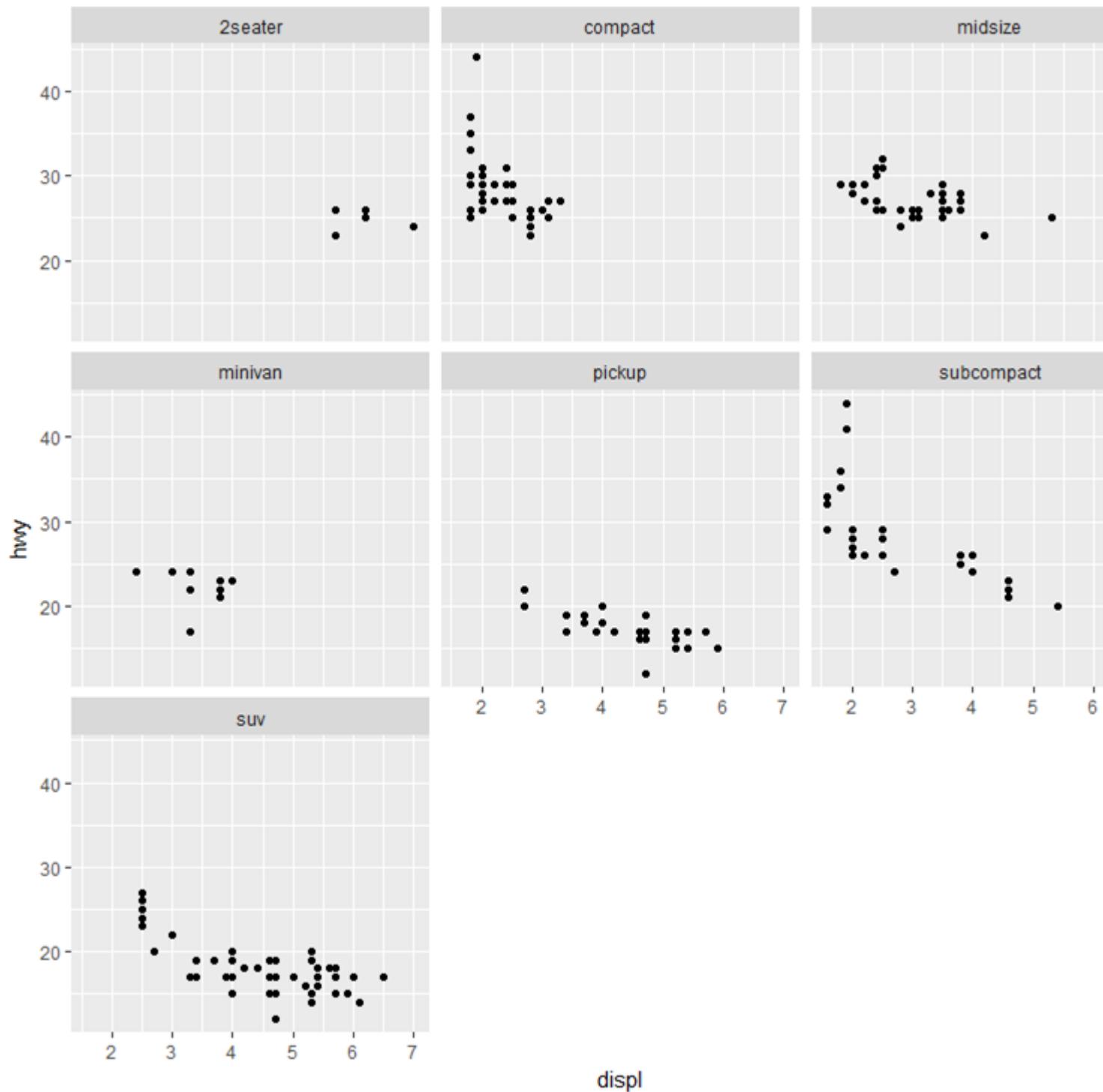
```
library(ggplot2)
ggplot(iris, aes(x = Petal.Width, y = Petal.Length, color = Species)) +
  geom_point()
```



```
facet . . . ggplot2 mpg .
```

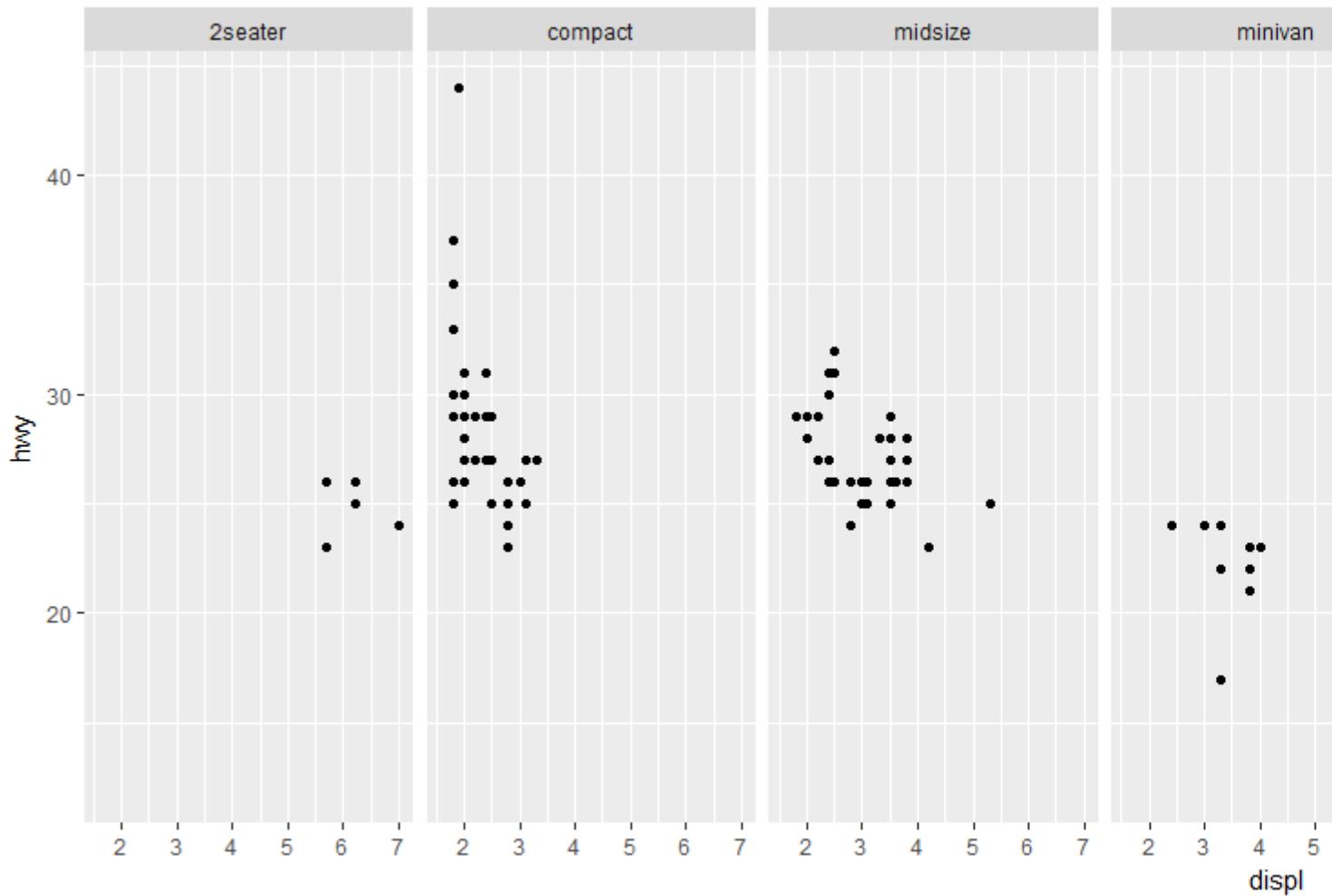
().

```
ggplot(mpg, aes(x = displ, y = hwy)) +  
  geom_point() +  
  facet_wrap(~class)
```

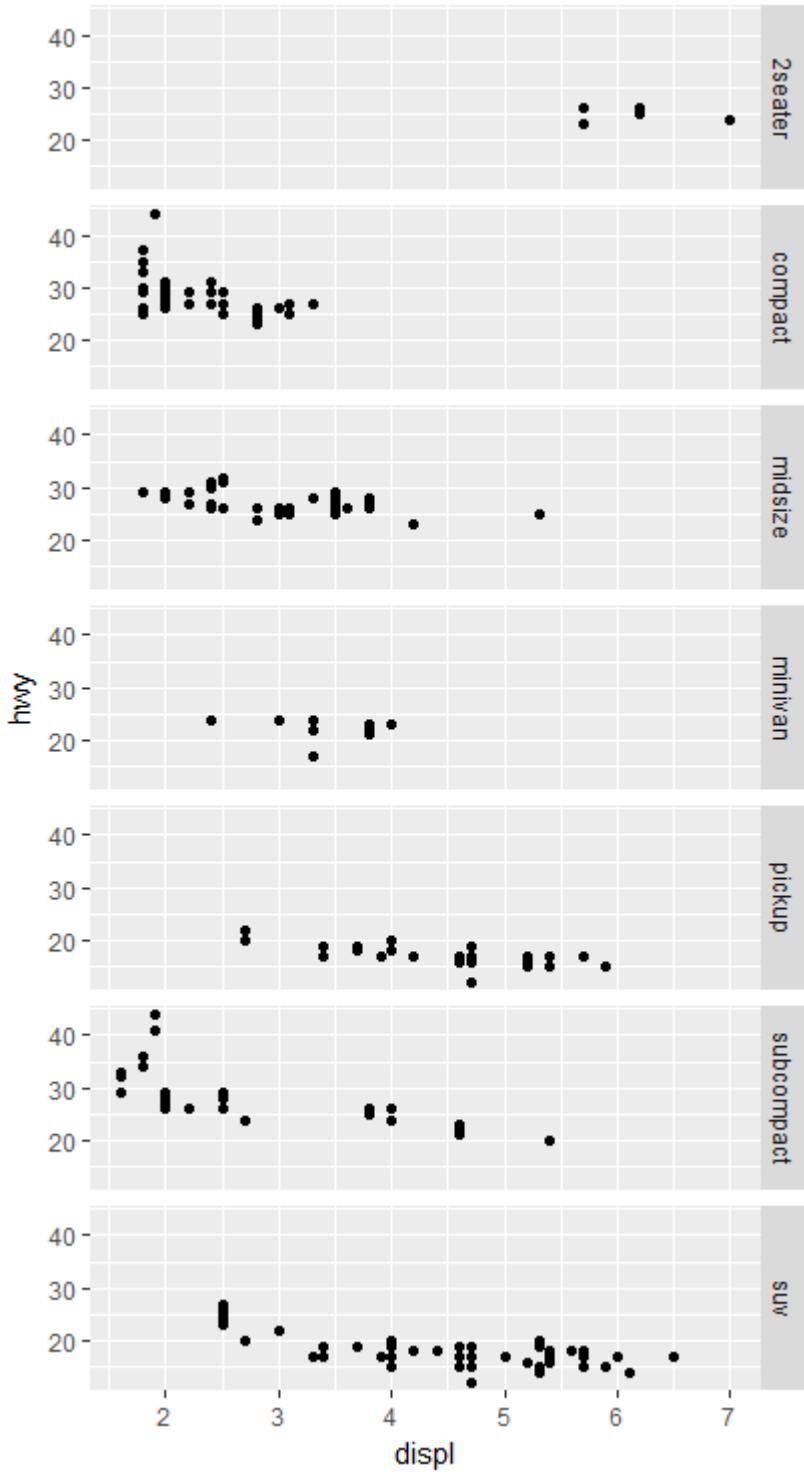


```
ggplot(mpg, aes(x = displ, y = hwy)) +  
  geom_point() +
```

```
facet_grid(.~class)
```

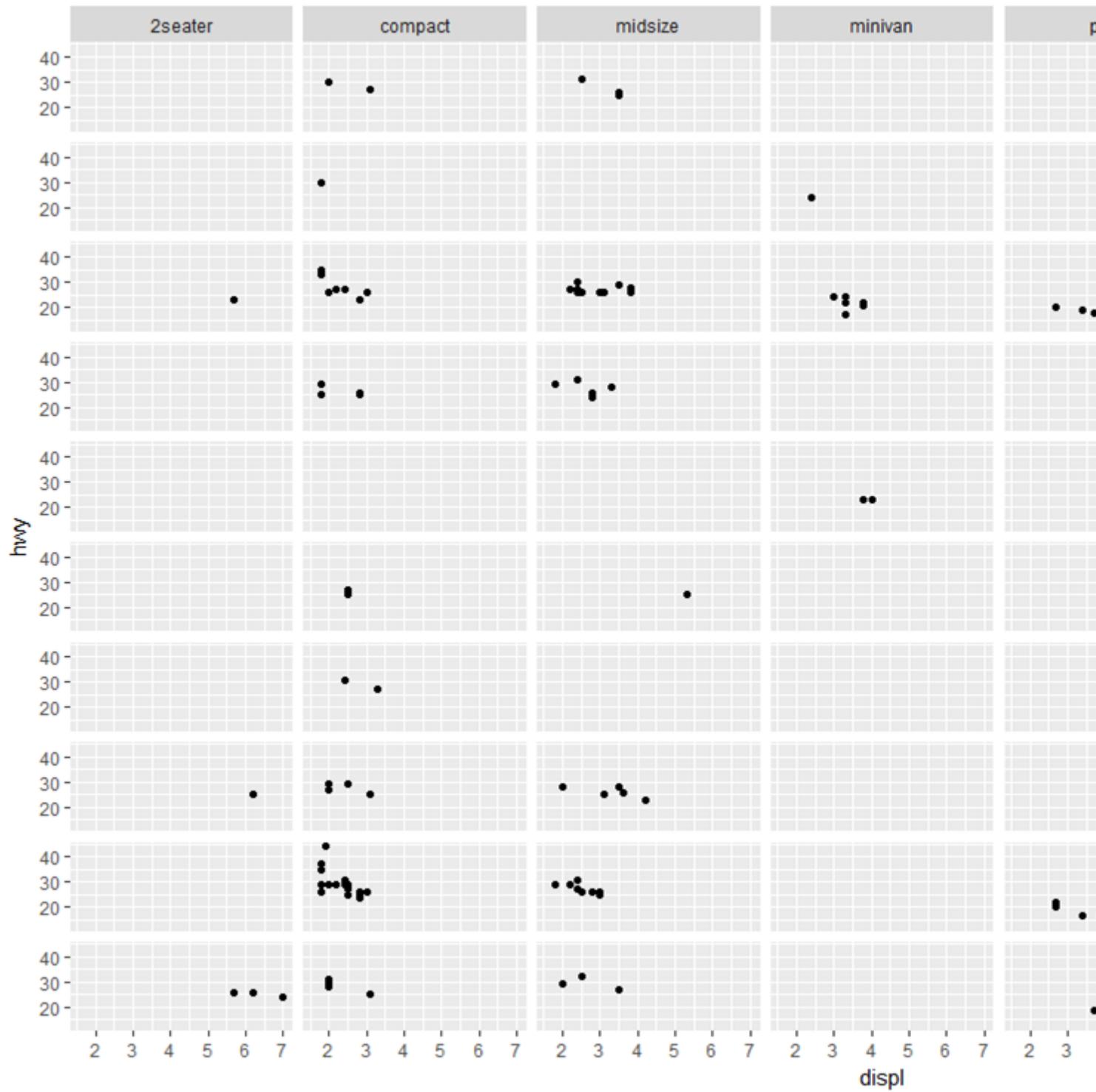


```
, :  
ggplot(mpg, aes(x = displ, y = hwy)) +  
  geom_point() +  
  facet_grid(class~.)
```



2 :

```
ggplot(mpg, aes(x = displ, y = hwy)) +  
  geom_point() +  
  facet_grid(trans~class) # "row" parameter, then "column" parameter
```



ggplot2 . 20 . .

```

set.seed(47)
sweetsWide <- data.frame(date      = 1:20,
                           chocolate = runif(20, min = 2, max = 4),
                           iceCream  = runif(20, min = 0.5, max = 1),
                           candy     = runif(20, min = 1, max = 3))

head(sweetsWide)
##   date chocolate iceCream   candy
## 1    1   3.953924  0.5890727 1.117311
## 2    2   2.747832  0.7783982 1.740851
## 3    3   3.523004  0.7578975 2.196754

```

```
## 4     4  3.644983 0.5667152 2.875028
## 5     5  3.147089 0.8446417 1.733543
## 6     6  3.382825 0.6900125 1.405674
```

```
sweetsWide  ggplot2 , R      reshape2 , data.table tidyR ()  :

# reshape from base R
sweetsLong <- reshape(sweetsWide, idvar = 'date', direction = 'long',
                      varying = list(2:4), new.row.names = NULL, times = names(sweetsWide)[-1])

# melt from 'reshape2'
library(reshape2)
sweetsLong <- melt(sweetsWide, id.vars = 'date')

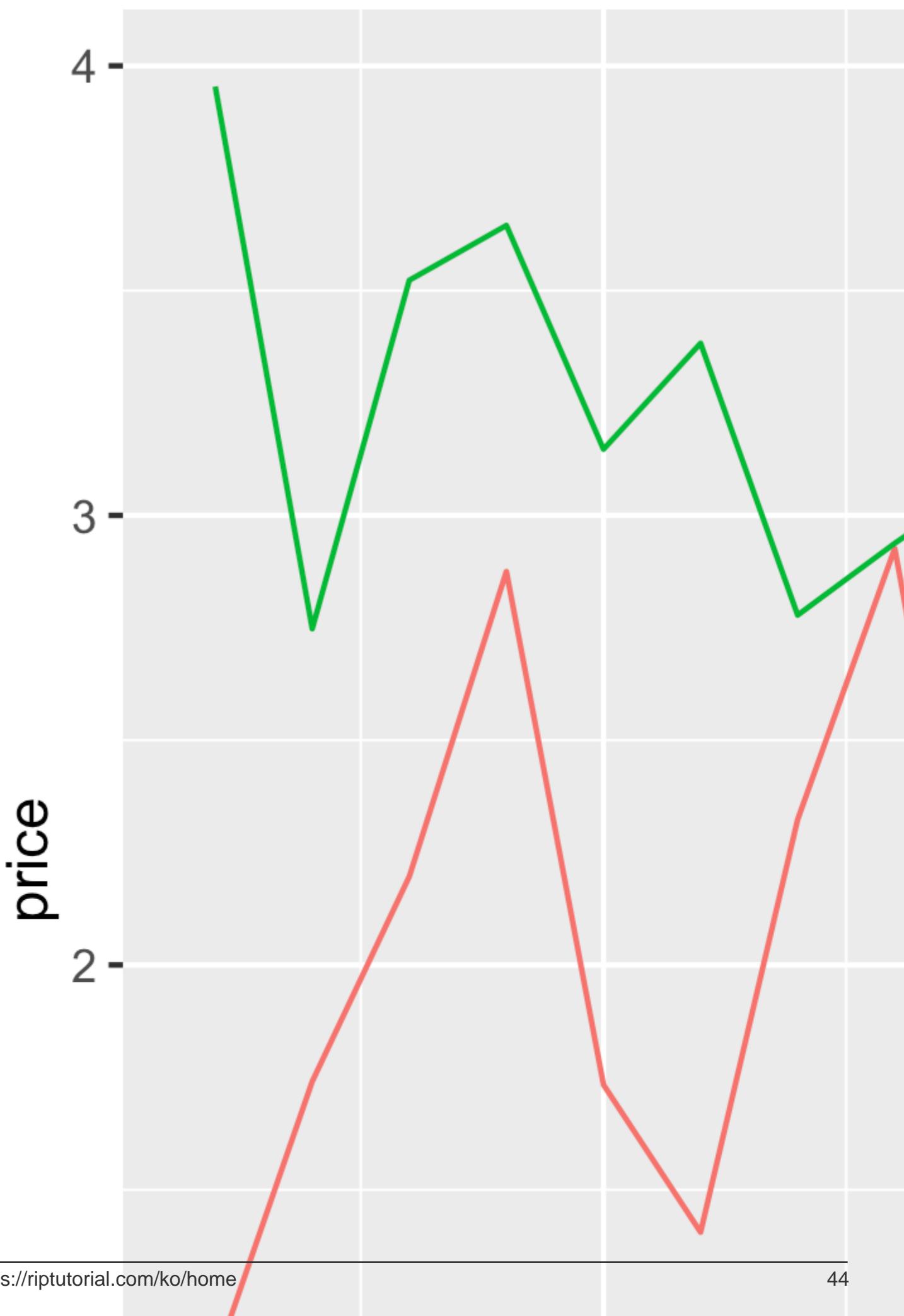
# melt from 'data.table'
# which is an optimized & extended version of 'melt' from 'reshape2'
library(data.table)
sweetsLong <- melt(setDT(sweetsWide), id.vars = 'date')

# gather from 'tidyR'
library(tidyR)
sweetsLong <- gather(sweetsWide, sweet, price, chocolate:candy)
```

```
head(sweetsLong)
##   date     sweet   price
## 1     1 chocolate 3.953924
## 2     2 chocolate 2.747832
## 3     3 chocolate 3.523004
## 4     4 chocolate 3.644983
## 5     5 chocolate 3.147089
## 6     6 chocolate 3.382825
```

```
sweetsLong
```

```
library(ggplot2)
ggplot(sweetsLong, aes(x = date, y = price, colour = sweet)) + geom_line()
```



10: GPU

GPU ". CUDA OpenCL. NVIDIA CUDA NVIDIA GPU . (: NVIDIA, AMD, Intel) (CPU GPU) SDK () . R GPU .

CUDA Toolkit OpenCL SDK R . R GPU CUDA NVIDIA GPU . .

1. [gputools](#)
- 2.
3. [HiPLARM](#)
- 4.

OpenCL .

1. [OpenCL](#) - R OpenCL
2. [gpuR](#) -

- GPU .

Examples

gpuR gpuMatrix

```
library(gpuR)

# gpuMatrix objects
X <- gpuMatrix(rnorm(100), 10, 10)
Y <- gpuMatrix(rnorm(100), 10, 10)

# transfer data to GPU when operation called
# automatically copied back to CPU
Z <- X %*% Y
```

gpuR vclMatrix

```
library(gpuR)

# vclMatrix objects
X <- vclMatrix(rnorm(100), 10, 10)
Y <- vclMatrix(rnorm(100), 10, 10)

# data always on GPU
# no data transfer
Z <- X %*% Y
```

GPU : <https://riptutorial.com/ko/r/topic/4680/gpu-->

11: hclust

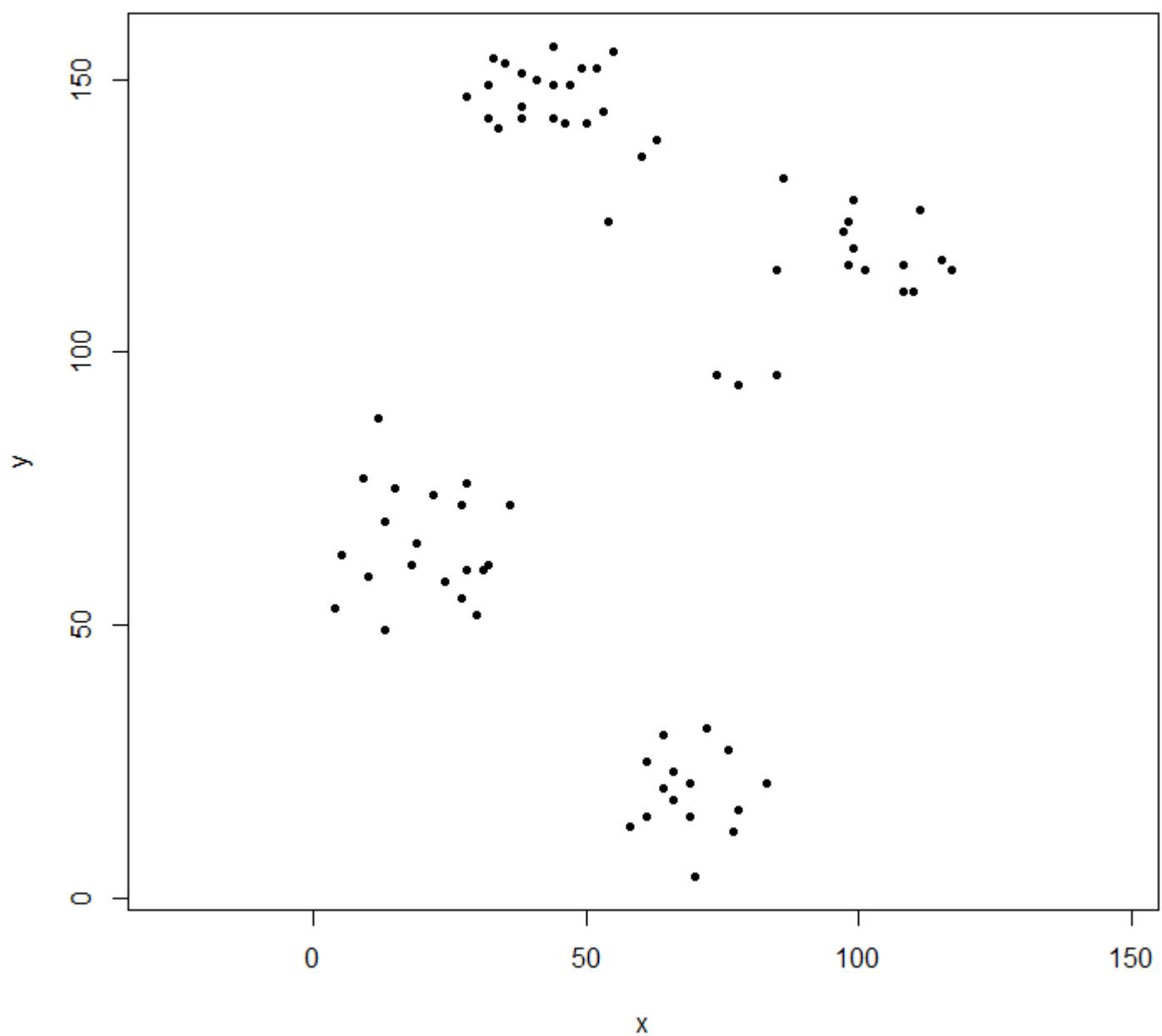
```
stats hclust .  
hclust . CRAN .
```

Examples

```
1 - hclust , end ,
```

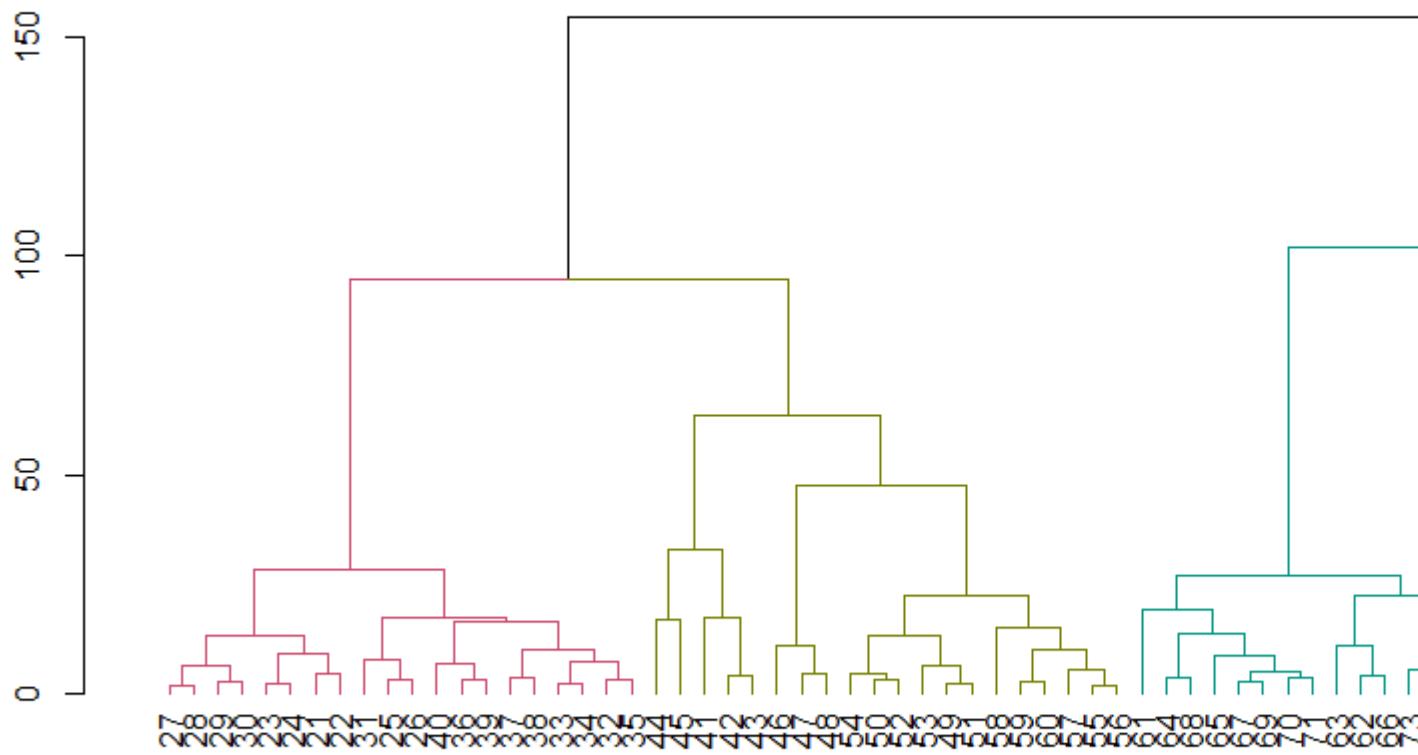
```
ruspini .
```

```
library(cluster) ## to get the ruspini data  
plot(ruspini, asp=1, pch=20) ## take a look at the data
```



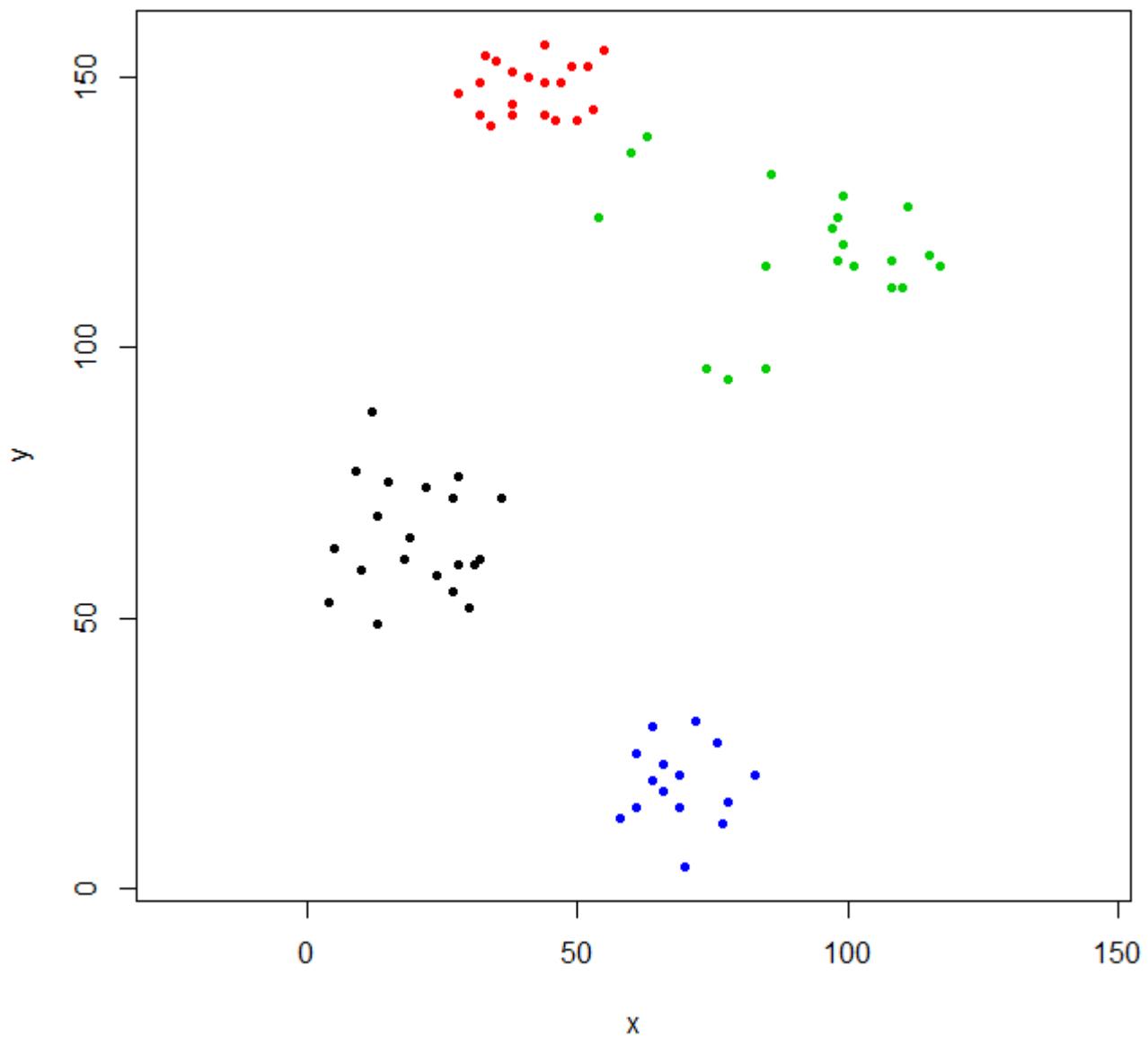
hclust . . . hang .

```
ruspini_hc_defaults <- hclust(dist(ruspini))
dend <- as.dendrogram(ruspini_hc_defaults)
if(!require(dendextend)) install.packages("dendextend"); library(dendextend)
dend <- color_branches(dend, k = 4)
plot(dend)
```

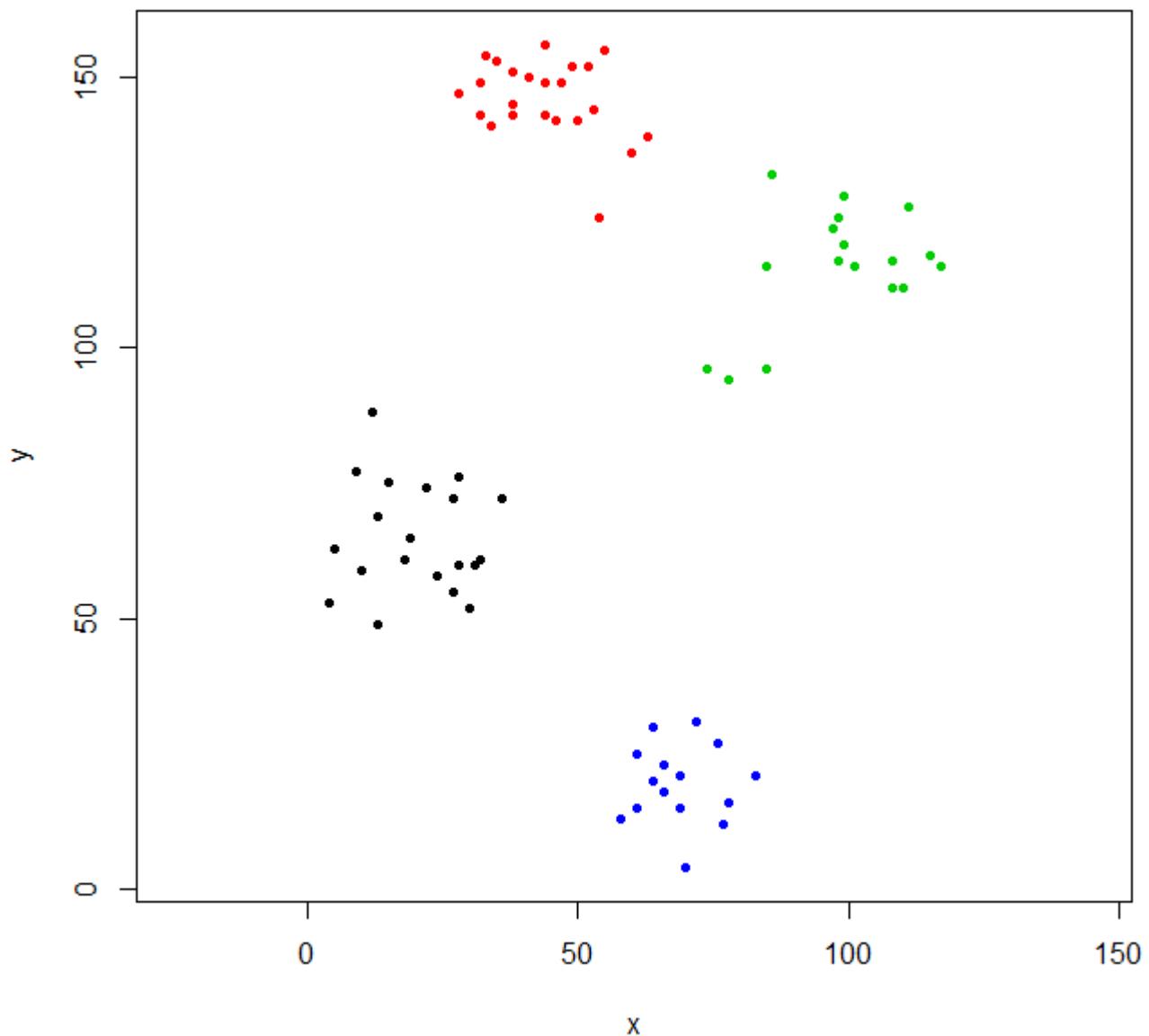


.k .

```
rhc_def_4 = cutree(ruspinis_hc_defaults,k=4)
plot(ruspinis, pch=20, asp=1, col=rhc_def_4)
```



```
scaled_ruspini_hc_defaults = hclust(dist(scale(ruspini)))
srhc_def_4 = cutree(scaled_ruspini_hc_defaults,4)
plot(ruspini, pch=20, asp=1, col=srhc_def_4)
```



". method

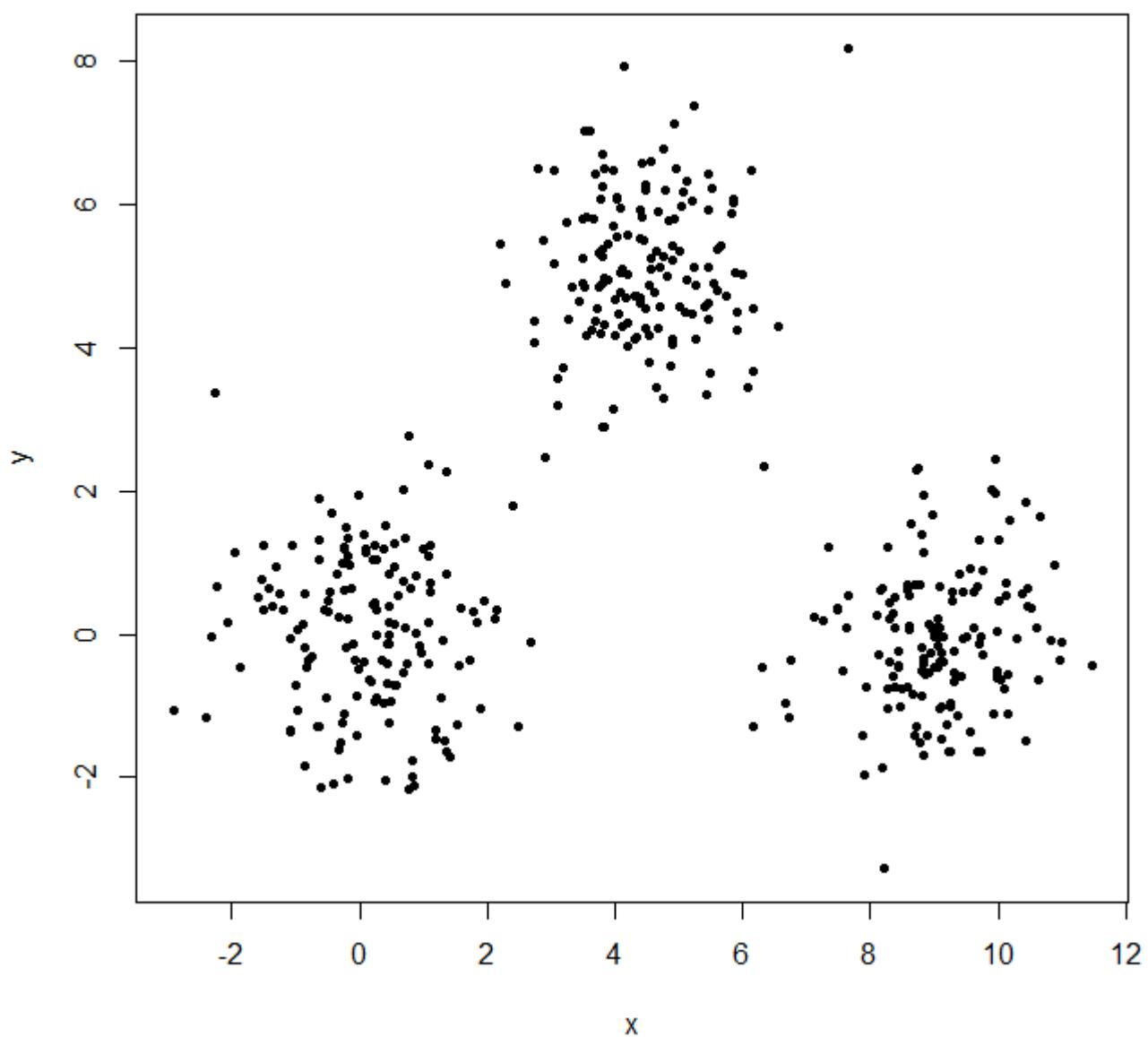
```
ruspini_hc_single = hclust(dist(ruspini), method="single")
```

2 - hclust

1 .

3 .

```
set.seed(656)
x = c(rnorm(150, 0, 1), rnorm(150, 9, 1), rnorm(150, 4.5, 1))
y = c(rnorm(150, 0, 1), rnorm(150, 0, 1), rnorm(150, 5, 1))
XYdf = data.frame(x,y)
plot(XYdf, pch=20)
```



```

XY_sing = hclust(dist(XYdf), method="single")
XYs3 = cutree(XY_sing, k=3)
table(XYs3)
XYs3
  1   2   3
448   1   1

```

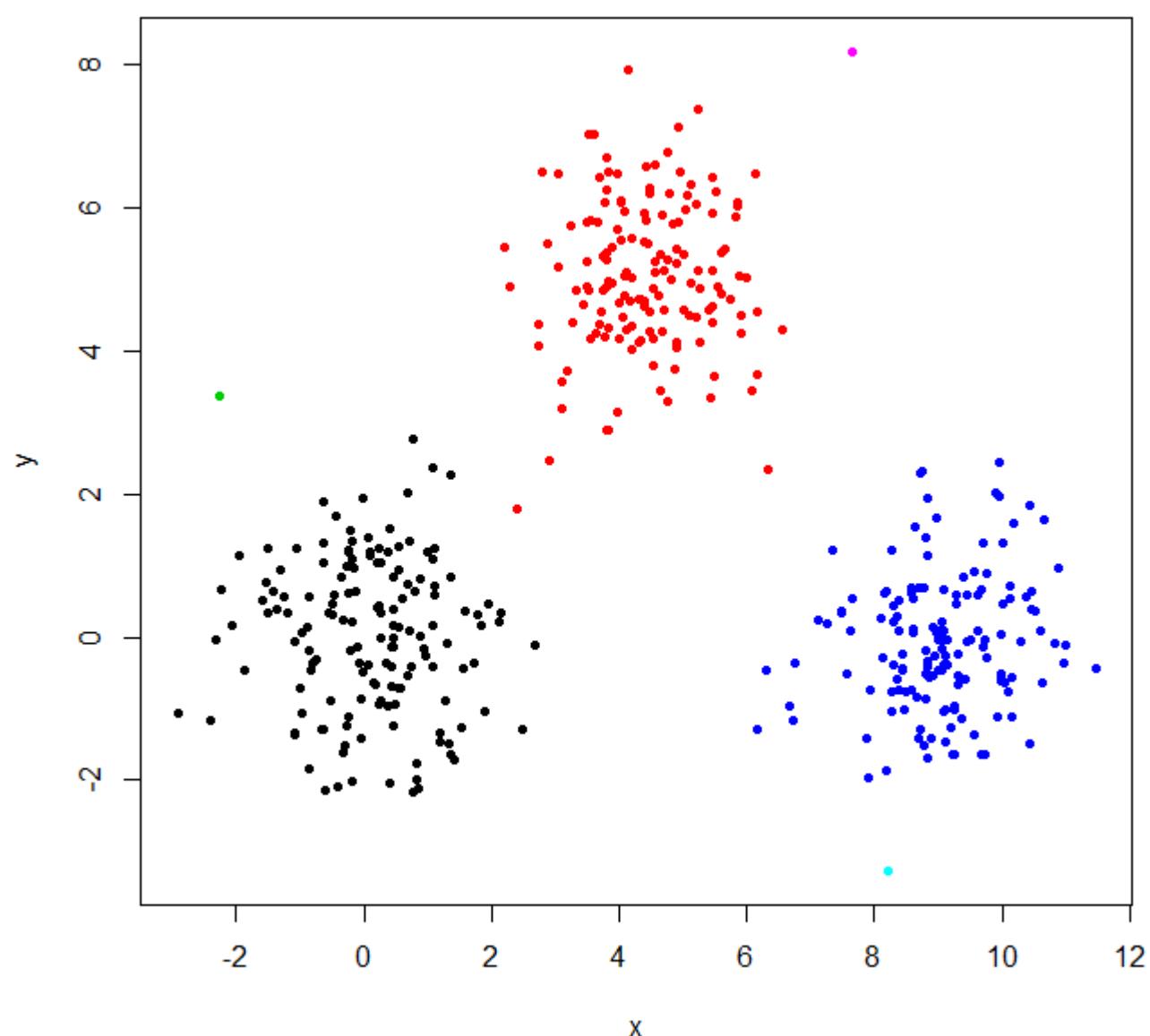
hclust . "" k .

```

XYs6 = cutree(XY_sing, k=6)
table(XYs6)
XYs6
  1   2   3   4   5   6
148 150   1 149   1   1

```

```
plot(XYdf, pch=20, col=XYs6)
```



StackOverflow

hclust : <https://riptutorial.com/ko/r/topic/8084/hclust---->

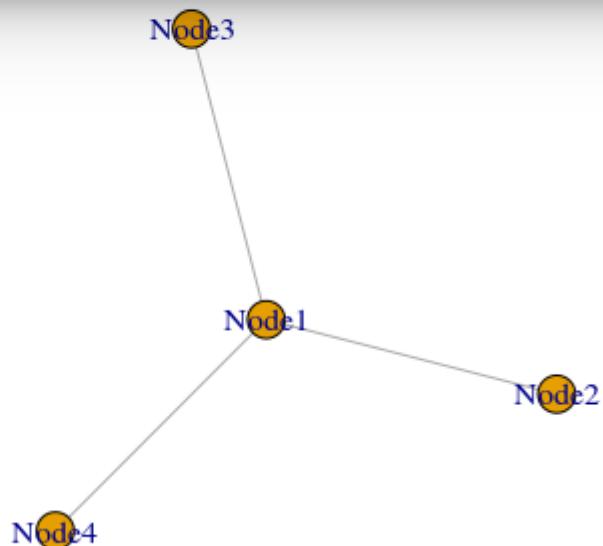
12: igraph

Examples

R igraph . R v.3.2.3 igraph .

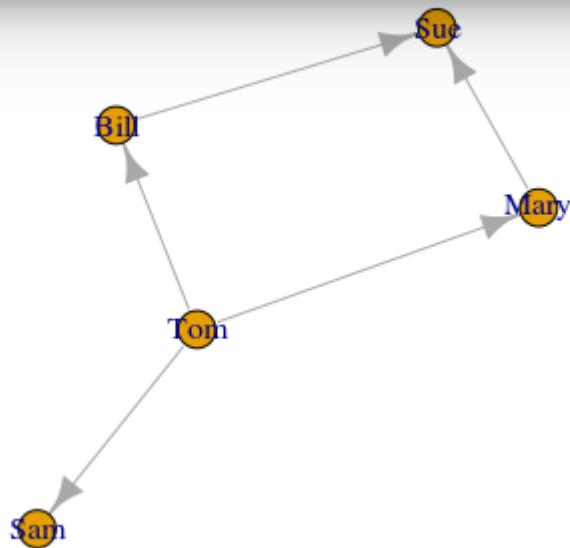
```
g<-graph.formula (Node1-Node2, Node1-Node3, Node4-Node1)
plot(g)
```

```
> g<-graph.formula(Node1-Node2, Node1-Node3, Node4-Node1)
> plot(g)
>
```



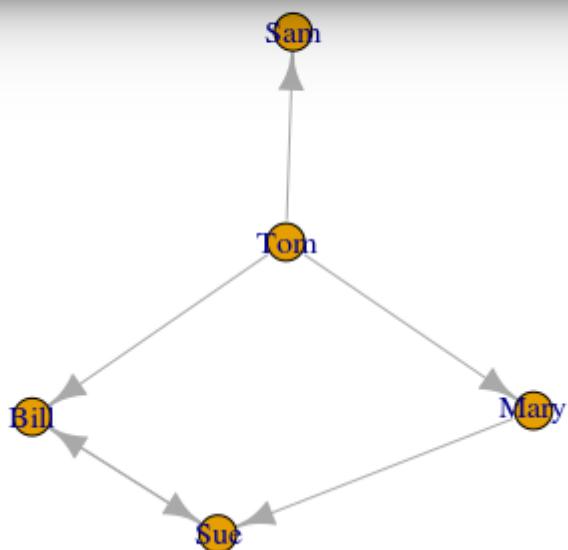
```
dg<-graph.formula(Tom-->Mary, Tom-->Bill, Tom-->Sam, Sue-->Mary, Bill-->Sue)
plot(dg)
```

```
> dg<-graph.formula(Tom-->Mary, Tom-->Bill, Tom-->Sam, Sue-->Mary, Bill-->Sue)
> plot(dg)
>
```



```
dg<-graph.formula(Tom-->Mary, Tom-->Bill, Tom-->Sam, Sue-->Mary, Bill++Sue)
plot(dg)
```

```
> dg<-graph.formula(Tom-->Mary, Tom-->Bill, Tom-->Sam, Sue-->Mary, Bill++Sue)
> plot(dg)
>
```



igraph : <https://riptutorial.com/ko/r/topic/4851/igraph--->

13: JSON

Examples

JSON R /

```
jsonlite      JSON . JSON      fromJSON() toJSON() respectively    vectors , matrices
data.frames  JSON .
```

JSON ()

```
library(jsonlite)

## vector to JSON
toJSON(c(1,2,3))
# [1] 1 2 3

fromJSON('[1,2,3]')
# [1] 1 2 3
```

JSON .

```
toJSON(list(myVec = c(1,2,3)))
# {"myVec": [1,2,3]}

fromJSON('{"myVec": [1,2,3]}')
# $myVec
# [1] 1 2 3
```

```
## list structures
lst <- list(a = c(1,2,3),
            b = list(letters[1:6]))

toJSON(lst)
# {"a": [1,2,3], "b": [{"a": "a", "b": "b", "c": "c", "d": "d", "e": "e", "f": "f"}]}

fromJSON('{"a": [1,2,3], "b": [{"a": "a", "b": "b", "c": "c", "d": "d", "e": "e", "f": "f"}]} ')
# $a
# [1] 1 2 3
#
# $b
# [,1] [,2] [,3] [,4] [,5] [,6]
# [1,] "a"   "b"   "c"   "d"   "e"   "f"
```

data.frame JSON ()

```
## converting a data.frame to JSON
df <- data.frame(id = seq_along(1:10),
                  val = letters[1:10])

toJSON(df)
```

```

#
[{"id":1,"val":"a"}, {"id":2,"val":"b"}, {"id":3,"val":"c"}, {"id":4,"val":"d"}, {"id":5,"val":"e"}, {"id":6,"val":"f"}, {"id":7,"val":"g"}, {"id":8,"val":"h"}, {"id":9,"val":"i"}, {"id":10,"val":"j"}]

## reading a JSON string
fromJSON('[{"id":1,"val":"a"}, {"id":2,"val":"b"}, {"id":3,"val":"c"}, {"id":4,"val":"d"}, {"id":5,"val":"e"}, {"id":6,"val":"f"}, {"id":7,"val":"g"}, {"id":8,"val":"h"}, {"id":9,"val":"i"}, {"id":10,"val":"j"}]

#      id val
# 1    1   a
# 2    2   b
# 3    3   c
# 4    4   d
# 5    5   e
# 6    6   f
# 7    7   g
# 8    8   h
# 9    9   i
# 10  10  j

```

JSON .

```

## Reading JSON from URL
googleway_issues <- fromJSON("https://api.github.com/repos/SymbolixAU/googleway/issues")

googleway_issues$url
# [1] "https://api.github.com/repos/SymbolixAU/googleway/issues/20"
"https://api.github.com/repos/SymbolixAU/googleway/issues/19"
# [3] "https://api.github.com/repos/SymbolixAU/googleway/issues/14"
"https://api.github.com/repos/SymbolixAU/googleway/issues/11"
# [5] "https://api.github.com/repos/SymbolixAU/googleway/issues/9"
"https://api.github.com/repos/SymbolixAU/googleway/issues/5"
# [7] "https://api.github.com/repos/SymbolixAU/googleway/issues/2"

```

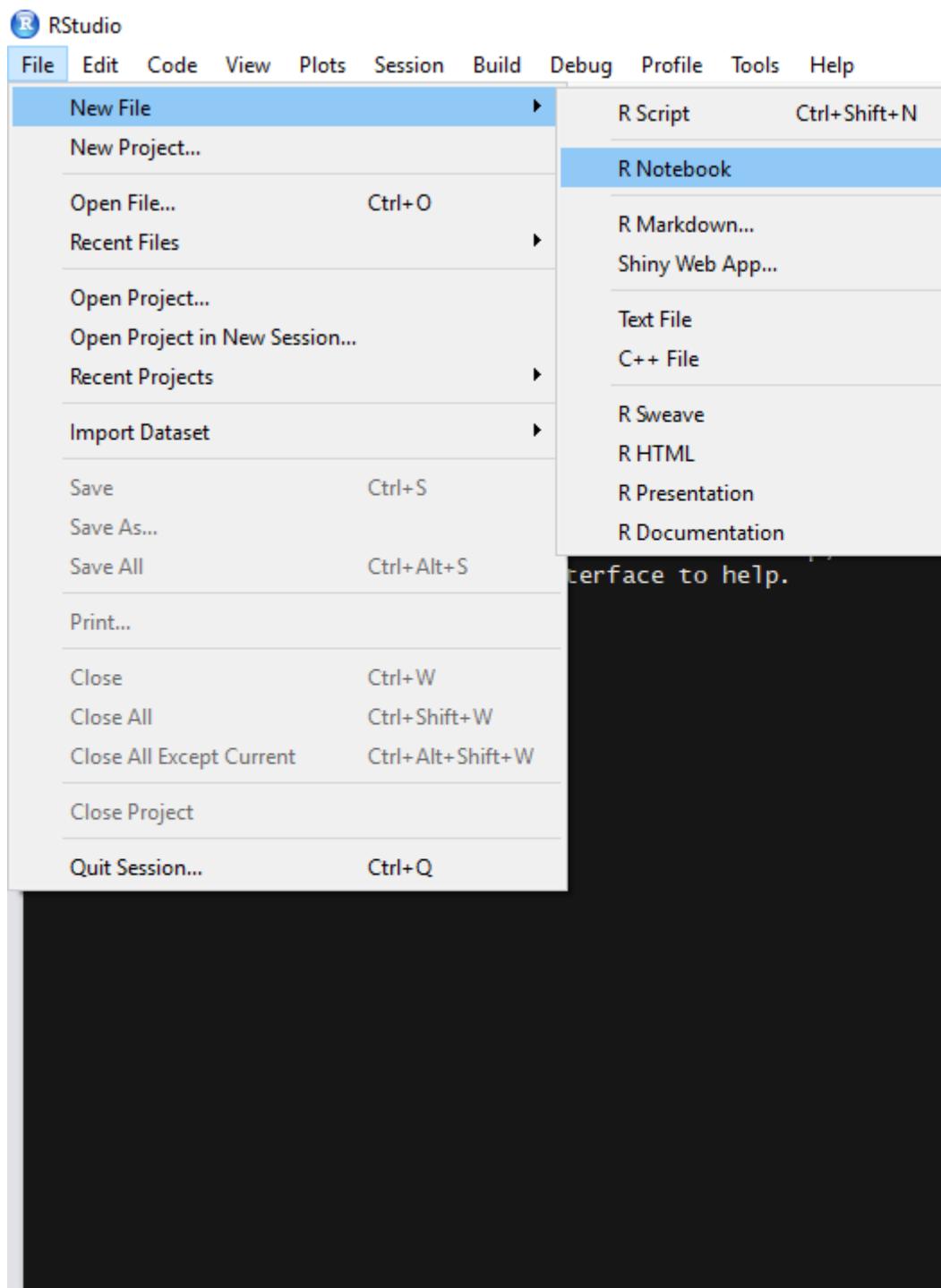
JSON : <https://riptutorial.com/ko/r/topic/2460/json>

14: R Markdown (RStudio)

R Notebook R Markdown , . R / R Markdown . : R RStudio RStudio 1.0 .

Examples

-> -> R RStudio
R Notebook RStudio . RStudio .



(R) Ctrl + Alt + I (OS X : Cmd + Option + I) .

RStudio

File Edit Code View Plots Session Build Debug Profile Tools Help

Go to file/function Addins

Untitled1

ABC Preview

1 ---
2 title: "R Notebook"
3 output: html_notebook
4 ---
5
6 This is an [R Markdown](http://rmarkdown.rstudio.com) Notebook.
notebook, the results appear beneath the code.
7
8 Try executing this chunk by clicking the *Run* button within the
inside it and pressing *Ctrl+Shift+Enter*.
9
10 ``{r}
11 plot(cars)
12 ``
13
14 Add a new chunk by clicking the *Insert chunk* button on the toolbar or by pressing *
15
16 when you save the notebook, an HTML file containing the code and output will be saved
(click the *Preview* button or press *Ctrl+Shift+K* to preview the HTML file).
17

17:1 (Top Level) ▾

Console

The screenshot shows the RStudio interface with a dark theme. A context menu is open from the 'Insert' button in the toolbar. The 'Insert' button has a red arrow pointing to it. The menu items are: R (selected), Python, Rcpp, SQL, Stan. A tooltip says 'Insert a new R chunk'. Other menu items like 'Execute code' and 'Preview' are visible but not selected.

() . . . **Ctrl + Shift + Enter (OS X : Cmd + Shift + Enter)** .

Ctrl + Alt + I (OS X : Cmd + Option + I) .

```
1 ---  
2 title: "R Notebook"  
3 output: html_notebook  
4 ---  
5  
6 This is an [R Markdown](http://rmarkdown.rstudio.com) Notebook. When you execute code  
notebook, the results appear beneath the code.  
7  
8 Try executing this chunk by clicking the *Run* button within the chunk or by placing  
inside it and pressing *Ctrl+Shift+Enter*.  
9  
10 ````{r}  
11 plot(cars)  
12 ````  
13  
14 Add a new chunk by clicking the *Insert chunk* button on the toolbar or by pressing *C  
15  
16 when you save the notebook, an HTML file containing the code and output will be saved  
(click the *Preview* button or press *Ctrl+Shift+K* to preview the HTML file).  
17
```

. R . . R .

Run **Ctrl + Alt + R (OS X : Cmd + Option + R).**

R Restart R () .

RStudio

File Edit Code View Plots Session Build Debug Profile Tools Help

Untitled1*

Go to file/function Addins

```
14 data("iris")
15 head(iris,5)
16

Sepal.Length Sepal.Width Petal.Length Petal.Width
1           5.1          3.5         1.4        0.2
2           4.9          3.0         1.4        0.2
3           4.7          3.2         1.3        0.2
4           4.6          3.1         1.5        0.2
5           5.0          3.6         1.4        0.2

5 rows

17
18 Divide Iris data to x (contain the all features) and y (only the classes)
19 ````{r}
20 x <- subset(iris, select=-species)
21 y <- iris$Species
22
23
24 Create SVM Model and show summary
25 ````{r}
26 svm_model <- svm(x,y)
27
28 summary(svm_model)
29
30
31 Run Prediction
32 ````{r}
33 pred <- predict(svm_model,x)
34
35
36 you can time taken by using system.time
```

Run All:

"pdf_document" "html_notebook"

RStudio

File Edit Code View Plots Session Build Debug Profile Tools Help

Go to file/function Addins

Untitled1 x Preview

```
1 ---  
2 title: "R Notebook"  
3 output: html_notebook  
4 ---  
5  
6 This is an [R Markdown](http://rmarkdown.rstudio.com) Notebook. When you execute code  
notebook, the results appear beneath the code.  
7  
8 Try executing this chunk by clicking the *Run* button within the chunk or by placing  
inside it and pressing *Ctrl+Shift+Enter*.  
9  
10 ``{r}  
11 plot(cars)  
12 ``  
13  
14 Add a new chunk by clicking the *Insert chunk* button on the toolbar or by pressing *C  
15  
16 when you save the notebook, an HTML file containing the code and output will be saved  
(click the *Preview* button or press *Ctrl+Shift+K* to preview the HTML file).  
17
```

4:1 | # R Notebook

Console

.Rmd .nb.html . () .Rmd HTML .

RStudio .

R Markdown (RStudio) : <https://riptutorial.com/ko/r/topic/10728/r-markdown---rstudio-->

15: R

Examples

Windows

```
R Windows () R .  
, R . .  
R R . ( c:\stats\R ). , R .() (tm) :  
• .  
• miniCRAN .  
  
packageStatus .  
  
pkgs <- packageStatus() # choose mirror  
upgrade(pkgs)  
  
installr . GUI . gui Rgui RStudio .  
  
install.packages("installr") # install  
setInternet2(TRUE) # only for R versions older than 3.3.0  
installr::updateR() # updating R.  
  
https://www.r-statistics.com/tag/installr/ Windows . https://www.r-statistics.com/2015/06/ --  
- Windows on /  
. R .  
R : https://riptutorial.com/ko/r/topic/4088/r----
```

16: R

R R

Examples

R

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

installr R

installr R R

R (RStudio RStudio).

```
install.packages("installr")
library("installr")
updateR()
```



R Console

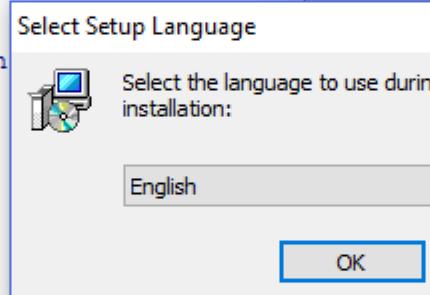
```
> library(installr)
Loading required package: stringr

Welcome to installr version 0.19.0

More information is available on the installr project website:
https://github.com/talgalili/installr/

Contact: <tal.galili@gmail.com>
Suggestions and bug-reports can be submitted at: https://github.com/talgalili/i$  
To suppress this message use:
suppressPackageStartupMessages(library(installr))

Warning message:
package 'installr' was built under R version 3.4.1
> updateR()
Installing the newest version of R,
please wait for the installer file to be download and executed.
Be sure to click 'next' as needed...
trying URL 'https://cran.rstudio.com/bin/windows/base/R-3.4.1-win.exe'
Content type 'application/x-msdos-program' length 78086510 bytes (74.5 MB)
downloaded 74.5 MB
```





R Console

```
> library(installr)
Loading required package: stringr

Welcome to installr version 0.19.0

More information is available on the installr project website:
https://github.com/talgalili/installr/

Contact: <tal.galili@gmail.com>
Suggestions and bug-reports can be submitted at: https://github.com/talgalili/i$
```

To suppress this message use:
suppressPackageStartupMessages

Warning message:
package 'installr' was built under R version 3.4.1
> updateR()
Installing the newest version of R,
please wait for the installer file to be download and
Be sure to click 'next' as needed...
trying URL 'https://cran.rstudio.com/bin/windows/base/R-3.5.1.exe'
Content type 'application/x-msdos-program' length 78086
downloaded 74.5 MB

Question



Do you wish to copy your packages from the
newer version of R?



R Console

```
> library(installr)
Loading required package: stringr

Welcome to installr version 0.19.0

More information is available on the installr project website:
https://github.com/talgalili/installr/

Contact: <tal.galili@gmail.com>
Suggestions and bug-reports can be submitted at: https://github.com/talgalili/i$
```

To suppress this message use:
suppressPackageStartupMessages

Question



Once your packages are copied to the new R
do you wish to KEEP the packages from the
installation?
(if you choose 'NO' - you will erase your pac

```
Warning message:
package 'installr' was built under R version 3.4.1
> updateR()
Installing the newest version of R,
please wait for the installer file to be download and
Be sure to click 'next' as needed...
trying URL 'https://cran.rstudio.com/bin/windows/base/'
Content type 'application/x-msdos-program' length 7808
downloaded 74.5 MB
```



R Console

```
> library(installr)
Loading required package: stringr

Welcome to installr version 0.19.0

More information is available on the installr project website:
https://github.com/talgalili/installr/

Contact: <tal.galili@gmail.com>
Suggestions and bug-reports can be submitted at: https://github.com/talgalili/i$
```

To suppress this message use:
suppressPackageStartupMessages()

Warning message:
package 'installr' was built under R version 3.4.1
> updateR()
Installing the newest version of R,
please wait for the installer file to be download and
Be sure to click 'next' as needed...
trying URL 'https://cran.rstudio.com/bin/windows/base/R-3.4.1.exe'
Content type 'application/x-msdos-program' length 78086
downloaded 74.5 MB

Question



Do you wish to copy your 'Rprofile.site' from
the newer version of R?



R Console

```
> library(installr)
Loading required package: stringr

Welcome to installr version 0.19.0

More information is available on the installr project website:
https://github.com/talgalili/installr/

Contact: <tal.galili@gmail.com>
Suggestions and bug-reports can be submitted at: https://github.com/talgalili/i$  
To suppress this message use:
suppressPackageStartupMessages(lis  
Warning message:
package 'installr' was built under R version 3.4.1
> updateR()
Installing the newest version of R,
please wait for the installer file to be download and ex
Be sure to click 'next' as needed...
trying URL 'https://cran.rstudio.com/bin/windows/base/R-3
Content type 'application/x-msdos-program' length 7808651
downloaded 74.5 MB
```

Question



Do you wish to update your packages in

Yes

R .

R

R .

version

R : <https://riptutorial.com/ko/r/topic/10729/r--->

17: R

Examples

R R "" .

, " " .

- ?
- R ?
- ?
- apply ? , R ?

, apply , R .

/ .

```
apply(mtcars, 1, mean)
      Mazda RX4      Mazda RX4 Wag      Datsun 710      Hornet 4 Drive      Hornet
Sportabout          Valiant        Duster 360
      29.90727      29.98136      23.59818      38.73955
53.66455          35.04909      59.72000
      Merc 240D      Merc 230      Merc 280      Merc 280C      Merc
450SE            Merc 450SL      Merc 450SLC
      24.63455      27.23364      31.86000      31.78727
46.43091          46.50000      46.35000
      Cadillac Fleetwood Lincoln Continental Chrysler Imperial
Civic              Toyota Corolla      Toyota Corona
      66.23273      66.05855      65.97227      19.44091
17.74227          18.81409      24.88864
      Dodge Challenger      AMC Javelin      Camaro Z28      Pontiac Firebird      Fiat
X1-9              Porsche 914-2      Lotus Europa
      47.24091      46.00773      58.75273      57.37955
18.92864          24.77909      24.88027
      Ford Pantera L      Ferrari Dino      Maserati Bora
      60.97182      34.50818      63.15545      Volvo 142E
                                         26.26273
```

? .

1. data.frame matrix .(apply .) .matrix . . . apply(iris, 2, class) str(iris)
sapply(iris, class) apply(iris, 2, class) .
2. , . , R nrow(mtcars) . mean R ?? :
3. R , ?

: .

```
rowMeans(mtcars)
      Mazda RX4      Mazda RX4 Wag      Datsun 710      Hornet 4 Drive      Hornet
Sportabout          Valiant        Duster 360
      29.90727      29.98136      23.59818      38.73955
53.66455          35.04909      59.72000
      Merc 240D      Merc 230      Merc 280      Merc 280C      Merc
```

| | | | | | |
|--------------------|---------------------|-------------------|------------------|----------|-------|
| 450SE | Merc 450SL | Merc 450SLC | | | |
| | 24.63455 | 27.23364 | 31.86000 | 31.78727 | |
| 46.43091 | 46.50000 | 46.35000 | | | |
| Cadillac Fleetwood | Lincoln Continental | Chrysler Imperial | | Fiat 128 | Honda |
| Civic | Toyota Corolla | Toyota Corona | | | |
| | 66.23273 | 66.05855 | 65.97227 | 19.44091 | |
| 17.74227 | 18.81409 | 24.88864 | | | |
| Dodge Challenger | AMC Javelin | Camaro Z28 | Pontiac Firebird | | Fiat |
| X1-9 | Porsche 914-2 | Lotus Europa | | | |
| | 47.24091 | 46.00773 | 58.75273 | 57.37955 | |
| 18.92864 | 24.77909 | 24.88027 | | | |
| Ford Pantera L | Ferrari Dino | Maserati Bora | Volvo 142E | | |
| | 60.97182 | 34.50818 | 63.15545 | 26.26273 | |

R . data.frame matrix . rowMeans

```
rowMeans(iris)
Error in rowMeans(iris) : 'x' must be numeric
```

? , mtcars (A data.frame list list A vector).

```
Reduce(`+`, mtcars)/ncol(mtcars)
[1] 29.90727 29.98136 23.59818 38.73955 53.66455 35.04909 59.72000 24.63455 27.23364 31.86000
31.78727 46.43091 46.50000 46.35000 66.23273 66.05855
[17] 65.97227 19.44091 17.74227 18.81409 24.88864 47.24091 46.00773 58.75273 57.37955 18.92864
24.77909 24.88027 60.97182 34.50818 63.15545 26.26273
```

(NA).

R .

```
aggregate(. ~ cyl, mtcars, mean)
cyl      mpg      disp       hp      drat       wt      qsec       vs       am      gear
carb
1   4 26.66364 105.1364  82.63636 4.070909 2.285727 19.13727 0.9090909 0.7272727 4.090909
1.545455
2   6 19.74286 183.3143 122.28571 3.585714 3.117143 17.97714 0.5714286 0.4285714 3.857143
3.428571
3   8 15.10000 353.1000 209.21429 3.229286 3.999214 16.77214 0.0000000 0.1428571 3.285714
3.500000
```

R C (C R).

? R rowsum rowsum , .

```
rowsum(mtcars[-2], mtcars$cyl)/table(mtcars$cyl)
mpg      disp       hp      drat       wt      qsec       vs       am      gear      carb
4 26.66364 105.1364  82.63636 4.070909 2.285727 19.13727 0.9090909 0.7272727 4.090909 1.545455
6 19.74286 183.3143 122.28571 3.585714 3.117143 17.97714 0.5714286 0.4285714 3.857143 3.428571
8 15.10000 353.1000 209.21429 3.229286 3.999214 16.77214 0.0000000 0.1428571 3.285714 3.500000
```

```
. data.frame ? matrix ?
```

```
( ). .
```

```
m .
```

```
set.seed(100)
m <- matrix(sample(1e2), 10)
m
 [,1] [,2] [,3] [,4] [,5] [,6] [,7] [,8] [,9] [,10]
[1,]    8   33   39   86   71  100   81   68   89   84
[2,]   12   16   57   80   32   82   69   11   41   92
[3,]   62   91   53   13   42   31   60   70   98   79
[4,]   66   94   29   67   45   59   20   96   64    1
[5,]   36   63   76    6   10   48   85   75   99    2
[6,]   18     4   27   19   44   56   37   95   26   40
[7,]    3   24   21   25   52   51   83   28   49   17
[8,]   46     5   22   43   47   74   35   97   77   65
[9,]   55   54   78   34   50   90   30   61   14   58
[10,]  88   73   38   15    9   72    7   93   23   87
```

```
:
```

```
apply(m, 1, var)
[1] 871.6556 957.5111 699.2111 941.4333 1237.3333 641.8222 539.7889 759.4333 500.4889
1255.6111
```

```
,
```

```
RowVar <- function(x) {
  rowSums((x - rowMeans(x))^2) / (dim(x)[2] - 1)
}
RowVar(m)
[1] 871.6556 957.5111 699.2111 941.4333 1237.3333 641.8222 539.7889 759.4333 500.4889
1255.6111
```

R : <https://riptutorial.com/ko/r/topic/3327/r--->

18: Rcpp

Examples

```
Rcpp  R : cppFunction() evalCpp()      .  sourceCpp()      C ++  cppFunction() .
```

```
R C ++ .  "" .
```

```
# Note - This is R code.  
# cppFunction in Rcpp allows for rapid testing.  
require(Rcpp)  
  
# Creates a function that multiples each element in a vector  
# Returns the modified vector.  
cppFunction(  
  NumericVector exfun(NumericVector x, int i){  
    x = x*i;  
    return x;  
  }")  
  
# Calling function in R  
exfun(1:5, 3)
```

```
C ++ .
```

```
# Use evalCpp to evaluate C++ expressions  
evalCpp("std::numeric_limits<double>::max()")  
## [1] 1.797693e+308
```

Rcpp

```
Rcpp R C ++ . .
```

```
// [[Rcpp::attribute]]
```

```
// [[Rcpp::export]]
```

```
sourceCpp() C ++ .
```

```
C ++ .
```

```
// Add code below into C++ file Rcpp_example.cpp  
  
#include <Rcpp.h>  
using namespace Rcpp;  
  
// Place the export tag right above function declaration.  
// [[Rcpp::export]]
```

```

double muRcpp(NumericVector x){

    int n = x.size(); // Size of vector
    double sum = 0; // Sum value

    // For loop, note cpp index shift to 0
    for(int i = 0; i < n; i++){
        // Shorthand for sum = sum + x[i]
        sum += x[i];
    }

    return sum/n; // Obtain and return the Mean
}

// Place dependent functions above call or
// declare the function definition with:
double muRcpp(NumericVector x);

// [[Rcpp::export]]
double varRcpp(NumericVector x, bool bias = true){

    // Calculate the mean using C++ function
    double mean = muRcpp(x);
    double sum = 0;

    int n = x.size();

    for(int i = 0; i < n; i++){
        sum += pow(x[i] - mean, 2.0); // Square
    }

    return sum/(n-bias); // Return variance
}

```

R C++ .

```

require(Rcpp)

# Compile File
sourceCpp("path/to/file/Rcpp_example.cpp")

# Make some sample data
x = 1:5

all.equal(muRcpp(x), mean(x))
## TRUE

all.equal(varRcpp(x), var(x))
## TRUE

```

Rcpp

C ++ .

```
// [[Rcpp::plugins(name)]]
```

:

```

// built-in C++11 plugin
// [[Rcpp::plugins cpp11]]]

// built-in C++11 plugin for older g++ compiler
// [[Rcpp::plugins cpp0x]]]

// built-in C++14 plugin for C++14 standard
// [[Rcpp::plugins cpp14]]]

// built-in C++1y plugin for C++14 and C++17 standard under development
// [[Rcpp::plugins cpp1y]]]

// built-in OpenMP++11 plugin
// [[Rcpp::plugins openmp]]]

```

Rcpp Rcpp.h Rcpp<PACKAGE>.h(:**RcppArmadillo**) .

```
// [[Rcpp::depends(Rcpp<PACKAGE>)]]
```

:

```

// Use the RcppArmadillo package
// Requires different header file from Rcpp.h
#include <RcppArmadillo.h>
// [[Rcpp::depends(RcppArmadillo)]]]

// Use the RcppEigen package
// Requires different header file from Rcpp.h
#include <RcppEigen.h>
// [[Rcpp::depends(RcppEigen)]]
```

Rcpp : <https://riptutorial.com/ko/r/topic/1404/rcpp>

19: RESTful R

OpenCPU R . .

Examples

opencpu

. <https://www.opencpu.org/apps.html>

R .

```
library(opencpu)
opencpu$start(port = 5936)
```

URL R . XML, html, JSON .

, R cURL .

```
#curl uses http post method for -X POST or -d "arg=value"
curl http://localhost:5936/opcpu/library/MASS/scripts/ch01.R -X POST
curl http://localhost:5936/opcpu/library/stats/R/rnorm -d "n=10&mean=5"
```

. R ().

/ocpu/tmp/ .

:

```
curl https://public.opencpu.org/opcpu/library/stats/R/rnorm -d n=5
/ocpu/tmp/x009f9e7630/R/.val
/ocpu/tmp/x009f9e7630/stdout
/ocpu/tmp/x009f9e7630/source
/ocpu/tmp/x009f9e7630/console
/ocpu/tmp/x009f9e7630/info
```

x009f9e7630 .

```
/ocpu/tmp/x009f9e7630/R/.val /ocpu/tmp/x009f9e7630/R/.val rnorm(5)
/ocpu/tmp/x009f9e7630/R/console rnorm(5) .
```

RESTful R : <https://riptutorial.com/ko/r/topic/8323/restful-r->

20: RMarkdown knitr

- :
 - YAML , .

```
|  
|  
|  
: "r format(Sys.time(), '%d %B, %Y')"  
|  
: 10 .html html_output .PDF , pdf_document , ..
```

—
—
—

| | | HTML | pdf | ODT | RTF | MD | | | | | | |
|--------------------------|--|------|-----|-----|-----|----|--|--|--|--|--|--|
| citation_package | , natbib, biblatex none
LaTeX | | | | | | | | | | | |
| | R "none", "hide" "show" . | | | | | | | | | | | |
| CSS | CSS | | | | | | | | | | | |
| dev | (: "png") | | | | | | | | | | | |
| | () | | | | | | | | | | | |
| fig_caption | ? | | | | | | | | | | | |
| fig_height,
fig_width | () | | | | | | | | | | | |
| | : "tango", "pygments",
"kate", "zenburn",
"textmate" | | | | | | | | | | | |
| | (in_header, before_body,
after_body) | | | | | | | | | | | |
| | ()? | | | | | | | | | | | |
| keep_md | knitr .md | | | | | | | | | | | |
| keep_tex | knitr .tex . | | | | | | | | | | | |

| | | HTML | pdf | ODT | RTF | MD | | | |
|-----------------|---------------------------------------|------|-----|-----|-----|----|--|--|--|
| latex_engine | ""pdflatex ","xelatex ",
lualatex" | | | | | | | | |
| lib_dir | (, MathJax) | | | | | | | | |
| mathjax | URL MathJax / URL . | | | | | | | | |
| md_extensions | R Markdown Markdown | | | | | | | | |
| number_sections | | | | | | | | | |
| pandoc_args | Pandoc | | | | | | | | |
| preserve_yaml | YAML ? | | | | | | | | |
| reference_docx | docx docx | | | | | | | | |
| self_contained | | | | | | | | | |
| slide_level | | | | | | | | | |
| | ? | | | | | | | | |
| | , em , | | | | | | | | |
| | Pandoc | | | | | | | | |
| | Bootswatch Beamer | | | | | | | | |
| toc | | | | | | | | | |
| toc_depth | | | | | | | | | |
| toc_float | . | | | | | | | | |

Examples

Rstudio

.Rmd , .R r .

Rstudio render .

```
---
title: "Rstudio exemple of a rmd file"
author: 'stack user'
date: "22 July 2016"
```

```

output: html_document
---

The header is used to define the general parameters and the metadata.

## R Markdown

This is an R Markdown document.
It is a script written in markdown with the possibility to insert chunk of R code in it.
To insert R code, it needs to be encapsulated into inverted quote.

Like that for a long piece of code:

```{r cars}
summary(cars)
```

And like ``r cat("that")`` for small piece of code.

## Including Plots

You can also embed plots, for example:

```{r echo=FALSE}
plot(pressure)
```

```

ioslides

. jQuery CSS knitr ioslides . jQuery . .

```
<script src="https://ajax.googleapis.com/ajax/libs/jquery/1.12.2/jquery.min.js"></script>
```

jQuery DOM() . , HTML . (\$(document).ready(function() { ... })), .backdrop.title-slide , .backdrop .segue " <footer></footer> (: </slide>). label .

CSS .

```
<footer> ( footer::after ) footer::after :
```

- label .
- 12
- (20 , 60)

```

---
title: "Adding a footer to presentation slides"
author: "Martin Schmelzer"
date: "26 Juli 2016"
output: ioslides_presentation
---

```{r setup, include=FALSE}
knitr::opts_chunk$set(echo = FALSE)

```

```
```

<script src="https://ajax.googleapis.com/ajax/libs/jquery/1.12.2/jquery.min.js"></script>

<script>
$(document).ready(function() {
    $('slide:not(.title-slide, .backdrop, .segue)').append('<footer label="My amazing
footer!"></footer>');
})
</script>

<style>
footer:after {
    content: attr(label);
    font-size: 12pt;
    position: absolute;
    bottom: 20px;
    left: 60px;
    line-height: 1.9;
}
</style>

## Slide 1

This is slide 1.

## Slide 2

This is slide 2

# Test

## Slide 3

And slide 3.
```

Slide 1

This is slide 1.

My amazing footer

RMarkdown knitr : <https://riptutorial.com/ko/r/topic/2999/rmarkdown--knitr->

21: RMarkdown

Examples

HTML LaTeX

-
- **xtable**
-

HTML

```
---
```

```
title: "Printing Tables"
author: "Martin Schmelzer"
date: "29 Juli 2016"
output: html_document
---
```

```
```{r setup, include=FALSE}
knitr::opts_chunk$set(echo = TRUE)
library(knitr)
library(xtable)
library(pander)
df <- mtcars[1:4,1:4]
```

# Print tables using `kable`
```{r, 'kable'}
kable(df)
```

# Print tables using `xtable`
```{r, 'xtable', results='asis'}
print(xtable(df), type="html")
```

# Print tables using `pander`
```{r, 'pander'}
pander(df)
```
```

Printing Tables

Martin Schmelzer

29 Juli 2016

Print tables using kable

```
kable(df)
```

| | mpg | cyl | disp | hp |
|----------------|------|-----|------|-----|
| Mazda RX4 | 21.0 | 6 | 160 | 110 |
| Mazda RX4 Wag | 21.0 | 6 | 160 | 110 |
| Datsun 710 | 22.8 | 4 | 108 | 93 |
| Hornet 4 Drive | 21.4 | 6 | 258 | 110 |

Print tables using xtable

```
print(xtable(df), type="html")
```

| | mpg cyl disp hp |
|----------------|-----------------|
| Mazda RX4 | 21.0 6 160 110 |
| Mazda RX4 Wag | 21.0 6 160 110 |
| Datsun 710 | 22.8 4 108 93 |
| Hornet 4 Drive | 21.4 6 258 110 |

Print tables using pander

```
pander(df)
```

| | mpg | cyl | disp | hp |
|----------------|------|-----|------|-----|
| Mazda RX4 | 21 | 6 | 160 | 110 |
| Mazda RX4 Wag | 21 | 6 | 160 | 110 |
| Datsun 710 | 22.8 | 4 | 108 | 93 |
| Hornet 4 Drive | 21.4 | 6 | 258 | 110 |

PDF

```
---
```

```
title: "Printing Tables"
author: "Martin Schmelzer"
date: "29 Juli 2016"
output: pdf_document
---
```

```
```{r setup, include=FALSE}
knitr::opts_chunk$set(echo = TRUE)
library(knitr)
library(xtable)
library(pander)
df <- mtcars[1:4,1:4]
```

# Print tables using `kable`
```{r, 'kable'}
kable(df)
```

# Print tables using `xtable`
```{r, 'xtable', results='asis'}
print(xtable(df, caption="My Table"))
```

# Print tables using `pander`
```{r, 'pander'}
pander(df)
```
```

Printing Tables

Martin Schwalzer
29.Juli.2016

Print tables using `kable`

```
kable(df)
```

| | mpg | cyl | disp | hp |
|----------------|------|-----|------|-----|
| Mazda RX4 | 21.0 | 6 | 160 | 110 |
| Mazda RX4 Wag | 21.0 | 6 | 160 | 110 |
| Datsun 710 | 22.8 | 4 | 108 | 93 |
| Hornet 4 Drive | 21.0 | 6 | 158 | 110 |

Print tables using `xtable`

```
print(xtable(df), caption = "My Table")
```

% latex table generated in R 3.3.1 by xtable 1.8-2 package % Fri Jul 29 10:18:41 2016

| | mpg | cyl | disp | hp |
|----------------|------|------|--------|--------|
| Mazda RX4 | 21.0 | 6.00 | 160.00 | 110.00 |
| Mazda RX4 Wag | 21.0 | 6.00 | 160.00 | 110.00 |
| Datsun 710 | 22.8 | 4.00 | 108.00 | 93.00 |
| Hornet 4 Drive | 21.0 | 6.00 | 158.00 | 110.00 |

Table 2: My Table

Print tables using `pander`

```
pander(df)
```

| | mpg | cyl | disp | hp |
|----------------|------|-----|------|-----|
| Mazda RX4 | 21 | 6 | 160 | 110 |
| Mazda RX4 Wag | 21 | 6 | 160 | 110 |
| Datsun 710 | 22.8 | 4 | 108 | 93 |
| Hornet 4 Drive | 21.0 | 6 | 158 | 110 |

xtable ?

```
options(xtable.comment = FALSE)
```

LaTeX preamble

LaTeX (`:\\usepackage`) RMarkdown .

1. YAML header-includes :

```
---
```

```
title: "Including LaTeX Preamble Commands in RMarkdown"
```

```
header-includes:
```

- \renewcommand{\familydefault}{\cmss}
- \usepackage[cm, slantedGreek]{sfmath}
- \usepackage[T1]{fontenc}

```
output: pdf_document
```

```
---
```

```
```{r setup, include=FALSE}
```

```
knitr:::opts_chunk$set(echo = TRUE, external=T)
```

```
```
```

```
# Section 1
```

```
As you can see, this text uses the Computer Modern Font!
```

Including LaTeX Preamble Commands in RMarkdown

Section 1

As you can see, this text uses the Computer Modern Font!

2. `includes`, `in_header` `includes`

```
---
```

```
title: "Including LaTeX Preamble Commands in RMarkdown"
output:
  pdf_document:
    includes:
      in_header: includes.tex
---
```

```
```{r setup, include=FALSE}
knitr::opts_chunk$set(echo = TRUE, external=T)
```

# Section 1

As you can see, this text uses the Computer Modern Font!
```

`include.tex` header-includes . . .

```
template . . .
```

```
---
```

```
title: "My Template"
author: "Martin Schmelzer"
output:
  pdf_document:
    template: myTemplate.tex
---
```

`bibtex` `cna` `YAML` `bibliography`: . . . `biblio-style`: . . .

```
---
```

```
title: "Including Bibliography"
author: "John Doe"
output: pdf_document
bibliography: references.bib
---
```

```
# Abstract

@R_Core_Team_2016

# References
```

Including Bibliography
John Doe

Abstract

R Core Team (2016)

References

R Core Team. 2016. *R: A Language and Environment for Statistical Computing*. Vienna, Austria: R Foundation for Statistical Computing. <http://www.R-project.org/>.

R-markdown

R-

R-markdown R . R

```
`r 2*2`
```

```
```{r name, echo=TRUE, include=TRUE, ...}
```

```
2*2
```

```
````
```

().

- **echo** (boolean)
- ()
- **fig.width** (numeric)
- **fig.height** ()
- **fig.cap** (character)

tag=value .

R-markdown

R-markdown . R-markdown R .

```
# Title #
```

```
This is **plain markdown** text.
```

```
```{r code, include=FALSE, echo=FALSE}
```

```
Just declare variables
```

```
income <- 1000
taxes <- 125
```

```
````
```

```
My income is: `r income ` dollars and I payed `r taxes ` dollars in taxes.
```

```
Below is the sum of money I will have left:
```

```
```{r gain, include=TRUE, echo=FALSE}
```

```
gain <- income-taxes
```

```
gain
~~~  
```{r plotOutput, include=TRUE, echo=FALSE, fig.width=6, fig.height=6}  
pie(c(income,taxes), label=c("income", "taxes"))  
~~~
```

## R-markdown

R knitr R-markdown R markdown .

R-markdown pdf / html .

1. knitr R-markdown markdown .
2. markdown *pandoc* pdf / html .

knitr knit2html() knit2pdf() . knit2pdf() knit2pdf()

income.Rmd R pdf .

```
library(knitr)
knit2pdf("income.Rmd", "income.pdf")
```

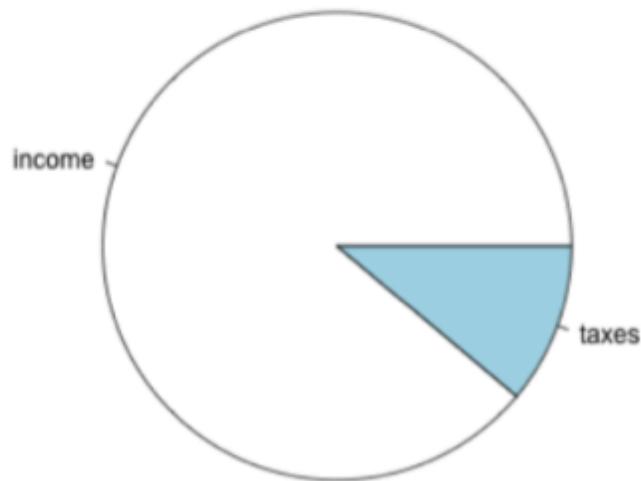
## Title

This is plain **markdown** text.

My income is: 1000 dollars and I payed 125 dollars in taxes.

Below is the sum of money I will have left:

```
[1] 875
```



RMarkdown : <https://riptutorial.com/ko/r/topic/4572/rmarkdown--->

## 22: RMD

YAML	
toc	
number_sections	
bibliography	
csl	

- RMD .
- install.packages("rmarkdown") R rmarkdown .
- Rmarkdown . YAML . link-citations: true
- .

MODS .mods	
BibLaTeX	.
BibTeX	.bibtex
RIS	
EndNote	.enl
EndNote XML	.xml
ISI	.wos
MEDLINE	.medline
	.coop
JSON citeproc	.json

## Examples

RMD YAML . PDF , (toc) .

```

```

```
title: "Writing an academic paper in R"
author: "Author"
date: "Date"
```

```
output:
 pdf_document:
 number_sections: yes
 toc: yes
 bibliography: bibliography.bib

```

bibliography.bib :

```
@ARTICLE{Meyer2000,
 AUTHOR="Bernd Meyer",
 TITLE="A constraint-based framework for diagrammatic reasoning",
 JOURNAL="Applied Artificial Intelligence",
 VOLUME= "14",
 ISSUE = "4",
 PAGES= "327--344",
 YEAR=2000
}
```

.bib @ bibkey (:Meyer2000 .

```
Introduction

'@Meyer2000` results in @Meyer2000.

'@Meyer2000 [p. 328]` results in @Meyer2000 [p. 328]

'[@Meyer2000]` results in [@Meyer2000]

'[-@Meyer2000]` results in [-@Meyer2000]

Summary

References
```

RStudio (Ctrl + Shift + K) rmarkdown::render("<path-to-your-RMD-file">) RMD  
rmarkdown::render("<path-to-your-RMD-file">) .

# Writing an academic paper in

*Author*

*Date*

## Contents

**1 Introduction**

**2 Summary**

**References**

## 1 Introduction

@Meyer2000 results in Meyer (2000).

@Meyer2000 [p. 328] results in Meyer (2000, 328)

[@Meyer2000] results in (Meyer 2000)

[-@Meyer2000] results in (2000)

## 2 Summary

## References

Meyer, Bernd. 2000. “A Constraint-Based Framework for Diagrammatic Reasoning.” *Intelligence* 14 (4): 327–44.

```

```

```
title: "Writing an academic paper in R"
author: "Author"
date: "Date"
output:
 pdf_document:
 number_sections: yes
 toc: yes
 bibliography: bibliography.bib
 csl: elsevier-harvard.csl

```

```
Introduction
```

```
`@Meyer2000` results in @Meyer2000.
```

```
`@Meyer2000 [p. 328]` results in @Meyer2000 [p. 328]
```

```
`[@Meyer2000]` results in [@Meyer2000]
```

```
`[-@Meyer2000]` results in [-@Meyer2000]
```

```
Summary
```

```
Reference
```

# Writing an academic paper in R

*Author*

*Date*

## Contents

1 Introduction	1
2 Summary	1
Reference	1

## 1 Introduction

- @Meyer2000 results in Meyer (2000).
- @Meyer2000 [p. 328] results in Meyer (2000, p. 328)
- [@Meyer2000] results in (Meyer, 2000)
- [-@Meyer2000] results in (2000)

## 2 Summary

## Reference

Meyer, B., 2000. A constraint-based framework for diagrammatic reasoning. *Applied Artificial Intelligence* 14, 327–344.

# 23: RODBC

## Examples

### RODBC Excel

RODBC R RDMS Windows Excel SQL Excel .

```
require(RODBC)
con = odbcConnectExcel("myfile.xlsx") # open a connection to the Excel file
sqlTables(con)$TABLE_NAME # show all sheets
df = sqlFetch(con, "Sheet1") # read a sheet
df = sqlQuery(con, "select * from [Sheet1 $]") # read a sheet (alternative SQL syntax)
close(con) # close the connection to the file
```

### SQL Server

RODBC SQL Server . 'Driver', SQL Server, "Atilla" sqlQuery .

```
library(RODBC)
cn <- odbcDriverConnect(connection="Driver={SQL
Server};server=localhost;database=Atilla;trusted_connection=yes;")
tbl <- sqlQuery(cn, 'select top 10 * from table_1')
```

```
library(RODBC)
con <- odbcDriverConnect("driver={Sql Server};server=servername;trusted connection=true")
dat <- sqlQuery(con, "select * from table");
close(con)
```

SQL Server . [connectionstrings.com](http://connectionstrings.com) .

databasename.schema.objectname .

RODBC : <https://riptutorial.com/ko/r/topic/2471/rodbc>

# **24: R**

## **Examples**

**R PDF PMF**

### **PMF BINOMIAL**

10 . 2 6 ?

dbinom .

```
> dbinom(2, 10, 1/6)
[1] 0.29071
```

### **POISSON PMF**

20 . 18 ?

dpois .

```
> dpois(18, 20)
[1] 0.08439355
```

### **PDF**

5 2 x = 2.5 pdf .

```
> dnorm(2.5, mean=5, sd=2)
[1] 0.09132454
```

**R :** <https://riptutorial.com/ko/r/topic/4333/r-->

# 25: R

## Examples

R , , Base . . .

. . . :

```
one <- function() { 1 }
one()
[1] 1

two <- function() { 1 + 1 }
two()
[1] 2
```

({ } . . . , . . .  
. ( vec ) ( 6) . . .

```
vec <- 4:9
subtract.length <- function(x) { x - length(x) }
subtract.length(vec)
[1] -2 -1 0 1 2 3
```

length() (, Base) . . .

```
vec2 <- (4:7)/2

msdf <- function(x, multiplier=4) {
 mult <- x * multiplier
 subl <- subtract.length(x)
 data.frame(mult, subl)
}

msdf(vec2, 5)
 mult subl
1 10.0 -2.0
2 12.5 -1.5
3 15.0 -1.0
4 17.5 -0.5
```

multiplier=4 multiplier 4 . 4  
, ( one , two , subtract.length )  
. . . \*apply Base .

data.frame .

```
df <- data.frame(first=5:9, second=(0:4)^2, third=-1:3)
```

```
apply(df, 2, function(x) { sqrt(sum(x^2)) })
 first second third
15.968719 18.814888 3.872983
```

```
x <- sample(1:6, 12, replace=TRUE)
mat <- matrix(x, nrow=3)

apply(mat, 1, function(x) { seq(min(x), max(x)) })
```

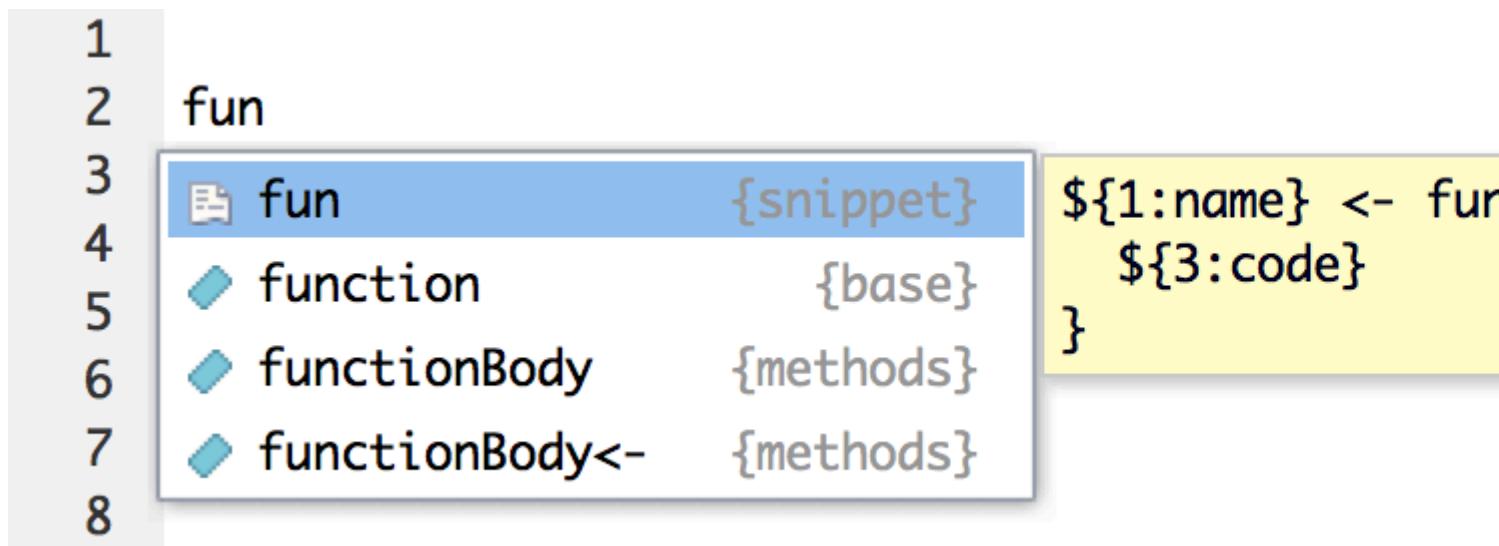
```
(function() { 1 })()
[1] 1
```

~

```
f <- function() { 1 }
f()
[1] 1
```

## RStudio

""RStudio IDE Tab .



```
name <- function(variables) {
}
```

```
name <- function(df, x, y) {
 require(tidyverse)
 out <-
 return(out)
}
```

Global Options -> Code Edit Snippets .

[[ .R ]]

```
basic.stats <- function(dset, vars){
 for(i in 1:length(vars)){
 print(vars[i])
 print(summary(dset[[vars[i]]]))
 }
}

basic.stats(iris, c("Sepal.Length", "Petal.Width"))

, (1,,,3)R . dset[[vars[i]]] vars i dset . , iris[["Sepal.Length"]] iris
Sepal.Length .
```

R : <https://riptutorial.com/ko/r/topic/7937/r-->

## 26: R ODE

- `ode (y, times, func, parms, method, ...)`

```
|
| () :ODE ()
| ; .
ODE
| () :func
:lsoda
```

- . "The Lorenz model" "Lorenz"

```
return(list(c(dX, dY, dZ)))
```

```
yini <- c(X = 1, Y = 1, Z = 1)
```

## Examples

Lorenz X, Y Z . .

$$\frac{dX}{dt} = a \cdot X + Y \cdot Z$$

$$\frac{dY}{dt} = b \cdot (Y - Z)$$

$$\frac{dZ}{dT} = -X \cdot Y + c \cdot Y - Z$$

$$X(0) = Y(0) = Z(0) = 1$$

a, b c

$$a = -8/3$$

$$b = -10$$

$$c = 28$$

```
library(deSolve)
```

```

Define R-function

Lorenz <- function (t, y, parms) {
 with(as.list(c(y, parms)), {
 dX <- a * X + Y * Z
 dY <- b * (Y - Z)
 dZ <- -X * Y + c * Y - Z

 return(list(c(dX, dY, dZ)))
 })
}

Define parameters and variables

parms <- c(a = -8/3, b = -10, c = 28)
yini <- c(X = 1, Y = 1, Z = 1)
times <- seq(from = 0, to = 100, by = 0.01)

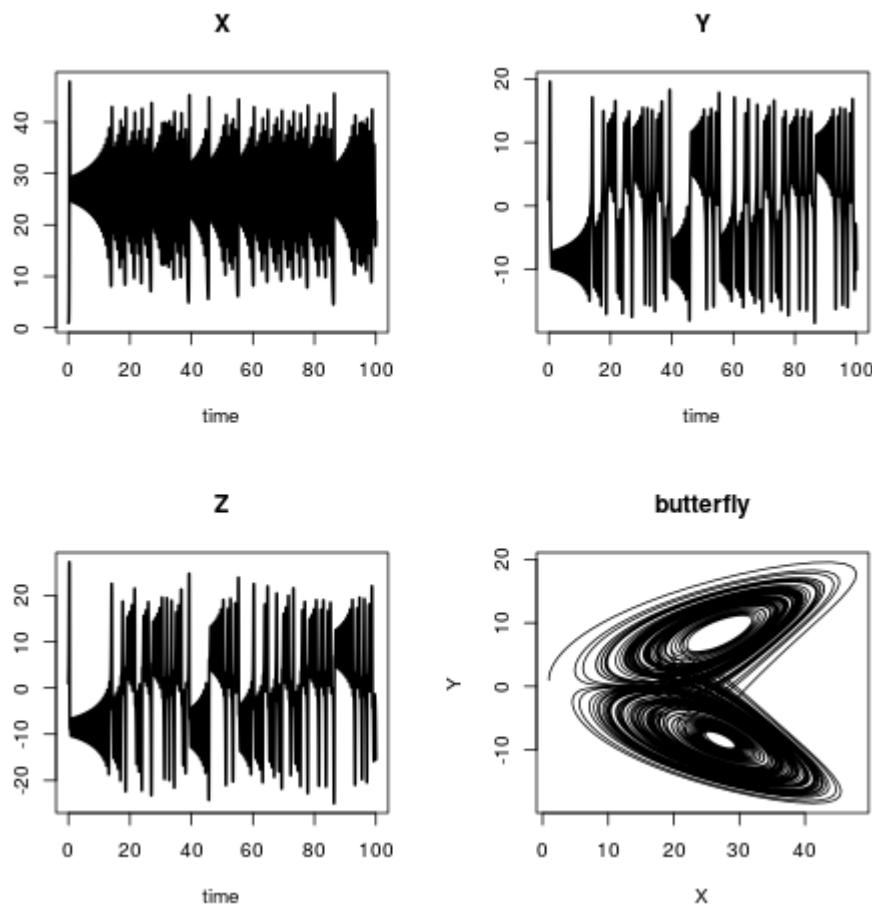
Solve the ODEs

out <- ode(y = yini, times = times, func = Lorenz, parms = parms)

Plot the results

plot(out, lwd = 2)
plot(out[, "X"], out[, "Y"],
 type = "l", xlab = "X",
 ylab = "Y", main = "butterfly")

```



## Lotka-Volterra :

```

library(deSolve)

Define R-function

LV <- function(t, y, parms) {
 with(as.list(c(y, parms)), {
 dP <- rG * P * (1 - P/K) - rI * P * C
 dC <- rI * P * C * AE - rM * C

 return(list(c(dP, dC), sum = C+P))
 })
}

Define parameters and variables

parms <- c(rI = 0.2, rG = 1.0, rM = 0.2, AE = 0.5, K = 10)
yini <- c(P = 1, C = 2)
times <- seq(from = 0, to = 200, by = 1)

Solve the ODEs

```

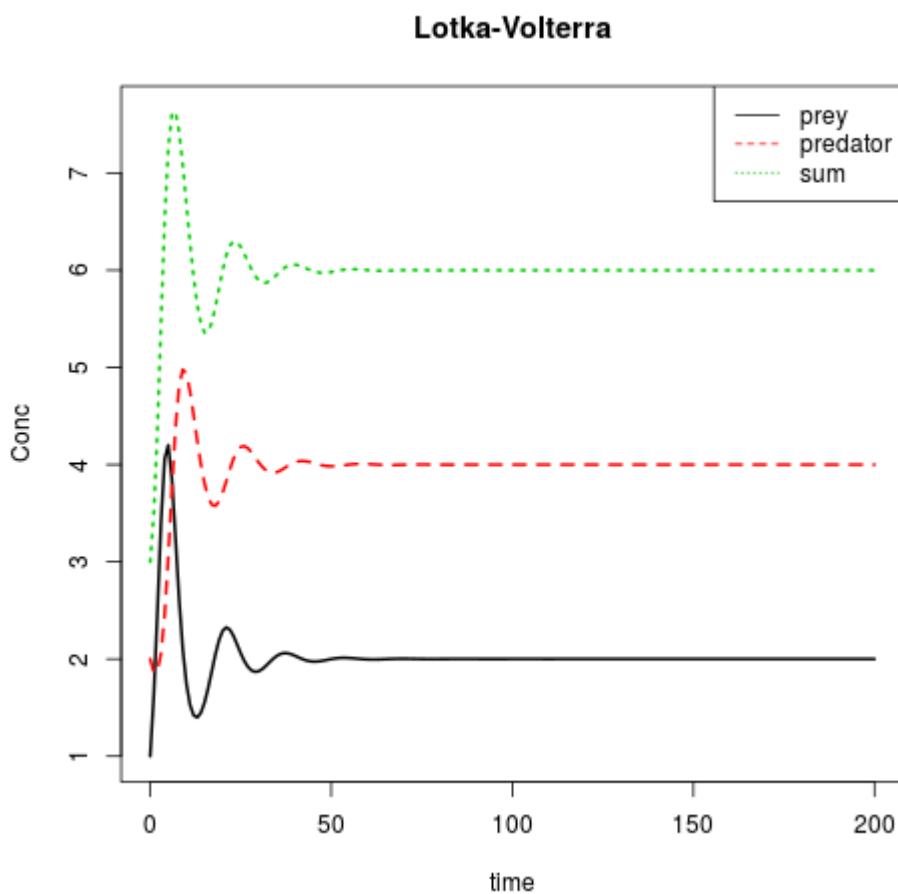
```

out <- ode(y = yini, times = times, func = LV, parms = parms)

Plot the results

```

`matplot(out[,1], out[,2:4], type = "l", xlab = "time", ylab = "Conc",
 main = "Lotka-Volterra", lwd = 2)
legend("topright", c("prey", "predator", "sum"), col = 1:3, lty = 1:3)`



## ODE - R

```

library(deSolve)

Define parameters and variables

```

`eps <- 0.01;
M <- 10
k <- M * eps^2/2
L <- 1
L0 <- 0.5
r <- 0.1
w <- 10
g <- 1`

```

parameter <- c(eps = eps, M = M, k = k, L = L, L0 = L0, r = r, w = w, g = g)

yini <- c(xl = 0, yl = L0, xr = L, yr = L0,
 ul = -L0/L, vl = 0,
 ur = -L0/L, vr = 0,
 lam1 = 0, lam2 = 0)

times <- seq(from = 0, to = 3, by = 0.01)

Define R-function

caraxis_R <- function(t, y, parms) {
 with(as.list(c(y, parms)), {
 yb <- r * sin(w * t)
 xb <- sqrt(L * L - yb * yb)
 Ll <- sqrt(xl^2 + yl^2)
 Lr <- sqrt((xr - xb)^2 + (yr - yb)^2)

 dxl <- ul; dyl <- vl; dxr <- ur; dyr <- vr

 dul <- (L0-Ll) * xl/Ll + 2 * lam2 * (xl-xr) + lam1*xb
 dvl <- (L0-Ll) * yl/Ll + 2 * lam2 * (yl-yr) + lam1*yb - k * g

 dur <- (L0-Lr) * (xr-xb)/Lr - 2 * lam2 * (xl-xr)
 dvr <- (L0-Lr) * (yr-yb)/Lr - 2 * lam2 * (yl-yr) - k * g

 c1 <- xb * xl + yb * yl
 c2 <- (xl - xr)^2 + (yl - yr)^2 - L * L

 return(list(c(dxl, dyl, dxr, dyr, dul, dvl, dur, dvr, c1, c2)))
 })
}

```

## ODE - C

```

sink("caraxis_C.c")
cat("
/* suitable names for parameters and state variables */

#include <R.h>
#include <math.h>
static double parms[8];

#define eps parms[0]
#define m parms[1]
#define k parms[2]
#define L parms[3]
#define L0 parms[4]
#define r parms[5]
#define w parms[6]
#define g parms[7]

/*-----
initialising the parameter common block
-----*/

```

```

*/
void init_C(void (* daeparms)(int *, double *)) {
 int N = 8;
 daeparms (&N, parms);
}
/* Compartments */

#define xl y[0]
#define yl y[1]
#define xr y[2]
#define yr y[3]
#define lam1 y[8]
#define lam2 y[9]

/*-----
the residual function
-----*/
void caraxis_C (int *neq, double *t, double *y, double *ydot,
 double *yout, int* ip)
{
 double yb, xb, Lr, Ll;

 yb = r * sin(w * *t) ;
 xb = sqrt(L * L - yb * yb);
 Ll = sqrt(xl * xl + yl * yl) ;
 Lr = sqrt((xr-xb)*(xr-xb) + (yr-yb)*(yr-yb)) ;

 ydot[0] = y[4];
 ydot[1] = y[5];
 ydot[2] = y[6];
 ydot[3] = y[7];

 ydot[4] = (L0-Ll) * xl/Ll + lam1*xb + 2*lam2*(xl-xr) ;
 ydot[5] = (L0-Ll) * yl/Ll + lam1*yb + 2*lam2*(yl-yr) - k*g;
 ydot[6] = (L0-Lr) * (xr-xb)/Lr - 2*lam2*(xl-xr) ;
 ydot[7] = (L0-Lr) * (yr-yb)/Lr - 2*lam2*(yl-yr) - k*g ;

 ydot[8] = xb * xl + yb * yl;
 ydot[9] = (xl-xr) * (xl-xr) + (yl-yr) * (yl-yr) - L*L;

}

", fill = TRUE)
sink()
system("R CMD SHLIB caraxis_C.c")
dyn.load(paste("caraxis_C", .Platform$dynlib.ext, sep = ""))
dllname_C <- dyn.load(paste("caraxis_C", .Platform$dynlib.ext, sep = ""))[[1]]

```

## ODE - Fortran

```

sink("caraxis_fortran.f")
cat("
c-----
c Initialiser for parameter common block
c-----
 subroutine init_fortran(daeparms)

 external daeparms
 integer, parameter :: N = 8

```

```

double precision parms(N)
common /myparms/parms

call daeparms(N, parms)
return
end

c-----
c rate of change
c-----
subroutine caraxis_fortran(neq, t, y, ydot, out, ip)
implicit none
integer neq, IP(*)
double precision t, y(neq), ydot(neq), out(*)
double precision eps, M, k, L, L0, r, w, g
common /myparms/ eps, M, k, L, L0, r, w, g

double precision xl, yl, xr, yr, ul, vl, ur, vr, lam1, lam2
double precision yb, xb, Ll, Lr, dxl, dyl, dxr, dyr
double precision dul, dvl, dur, dvr, c1, c2

c expand state variables
xl = y(1)
yl = y(2)
xr = y(3)
yr = y(4)
ul = y(5)
vl = y(6)
ur = y(7)
vr = y(8)
lam1 = y(9)
lam2 = y(10)

yb = r * sin(w * t)
xb = sqrt(L * L - yb * yb)
Ll = sqrt(xl**2 + yl**2)
Lr = sqrt((xr - xb)**2 + (yr - yb)**2)

dxl = ul
dyl = vl
dxr = ur
dyr = vr

dul = (L0-Ll) * xl/Ll + 2 * lam2 * (xl-xr) + lam1*xb
dvl = (L0-Ll) * yl/Ll + 2 * lam2 * (yl-yr) + lam1*yb - k*g
dur = (L0-Lr) * (xr-xb)/Lr - 2 * lam2 * (xl-xr)
dvr = (L0-Lr) * (yr-yb)/Lr - 2 * lam2 * (yl-yr) - k*g

c1 = xb * xl + yb * yl
c2 = (xl - xr)**2 + (yl - yr)**2 - L * L

c function values in ydot
ydot(1) = dxl
ydot(2) = dyl
ydot(3) = dxr
ydot(4) = dyr
ydot(5) = dul
ydot(6) = dvl
ydot(7) = dur
ydot(8) = dvr
ydot(9) = c1

```

```

ydot(10) = c2
return
end
", fill = TRUE)

sink()
system("R CMD SHLIB caraxis_fortran.f")
dyn.load(paste("caraxis_fortran", .Platform$dynlib.ext, sep = ""))
dllname_fortran <- dyn.load(paste("caraxis_fortran", .Platform$dynlib.ext, sep = ""))[[1]]

```

## ODE -

### ODE - R , ODE - C ODE - Fortran - .

```

library(microbenchmark)

R <- function(){
 out <- ode(y = yini, times = times, func = caraxis_R,
 parms = parameter)
}

C <- function(){
 out <- ode(y = yini, times = times, func = "caraxis_C",
 initfunc = "init_C", parms = parameter,
 dllname = dllname_C)
}

fortran <- function(){
 out <- ode(y = yini, times = times, func = "caraxis_fortran",
 initfunc = "init_fortran", parms = parameter,
 dllname = dllname_fortran)
}

```

```

all.equal(tail(R()), tail(fortran()))
all.equal(R()[,2], fortran()[,2])
all.equal(R()[,2], C()[,2])

```

(: ):

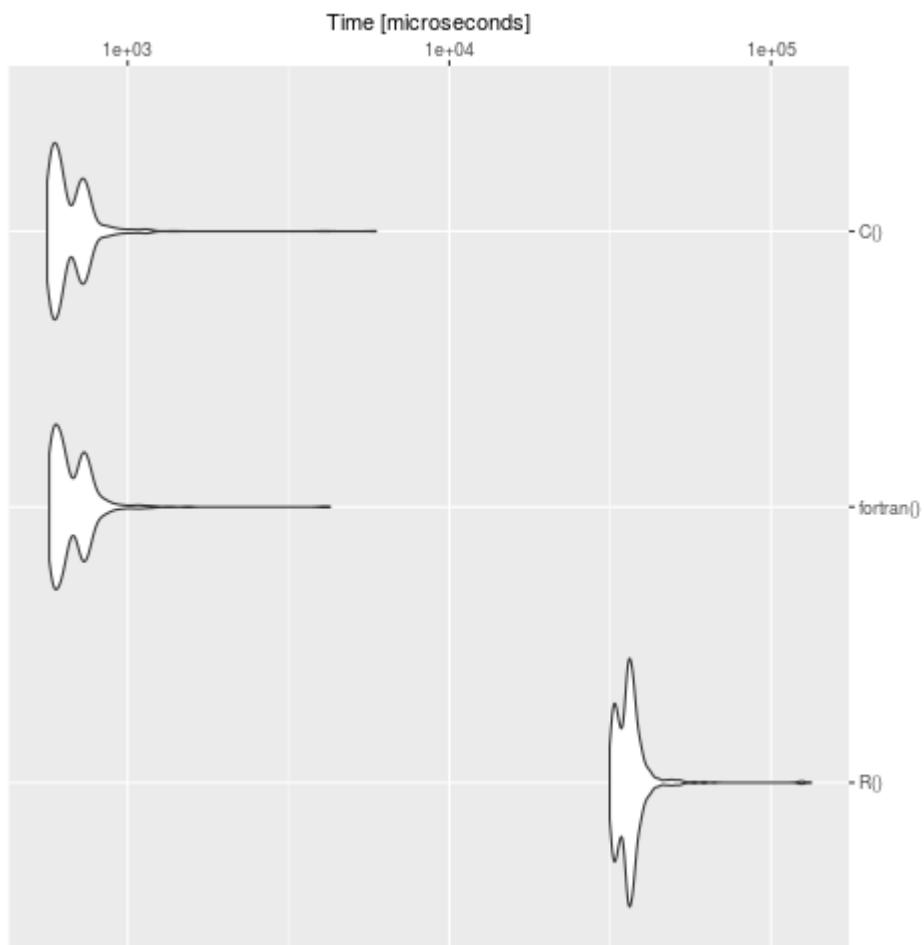
```

bench <- microbenchmark::microbenchmark(
 R(),
 fortran(),
 C(),
 times = 1000
)

summary(bench)

 expr min lq mean median uq max neval cld
 R() 31508.928 33651.541 36747.8733 36062.2475 37546.8025 132996.564 1000 b
 fortran() 570.674 596.700 686.1084 637.4605 730.1775 4256.555 1000 a
 C() 562.163 590.377 673.6124 625.0700 723.8460 5914.347 1000 a

```



C R . . . code ODE R C

R ODE : <https://riptutorial.com/ko/r/topic/7448/r-ode-->

# 27: R

R 4 . . .

S3, S4, Reference Classes S6.

## Examples

S3

S3 R OO .

S3 . class get (got?) .

```
> class(3)
[1] "numeric"
```

class .

```
> bicycle <- 2
> class(bicycle) <- 'vehicle'
> class(bicycle)
[1] "vehicle"
```

attr .

```
> velocipede <- 2
> attr(velocipede, 'class') <- 'vehicle'
> class(velocipede)
[1] "vehicle"
```

```
> class(x = bicycle) <- c('human-powered vehicle', class(x = bicycle))
> class(x = bicycle)
[1] "human-powered vehicle" "vehicle"
```

R . .

:

```
> summary.vehicle <- function(object, ...) {
+ message('this is a vehicle')
+ }
> summary(object = my_bike)
this is a vehicle
```

summary.bicycle .

```
> summary.bicycle <- function(object, ...) {
+ message('this is a bicycle')
+ }
> summary(object = my_bike)
this is a bicycle
```

R : <https://riptutorial.com/ko/r/topic/9723/r--->

## 28: R -

### Examples

0 0

0 .

```
data("GermanCredit")
variances<-apply(GermanCredit, 2, var)
variances[which(variances<=0.0025)]
```

"nearZeroVar (, ) . . . "

```
library(caret)
names(GermanCredit) [nearZeroVar(GermanCredit)]
```

NA

```
library(VIM)
data(sleep)
colMeans(is.na(sleep))

 BodyWgt BrainWgt NonD Dream Sleep Span Gest
0.00000000 0.00000000 0.22580645 0.19354839 0.06451613 0.06451613 0.06451613
 Pred Exp Danger
0.00000000 0.00000000 0.00000000
```

NonD Dream . 20 % ( )

```
library(purrr) # in order to use keep()

select correlatable vars
toCorrelate<-mtcars %>% keep(is.numeric)

calculate correlation matrix
correlationMatrix <- cor(toCorrelate)

pick only one out of each highly correlated pair's mirror image
correlationMatrix[upper.tri(correlationMatrix)]<-0

and I don't remove the highly-correlated-with-itself group
```

```
diag(correlationMatrix)<-0

find features that are highly correlated with another feature at the +- 0.85 level
apply(correlationMatrix, 2, function(x) any(abs(x)>=0.85))

mpg cyl disp hp drat wt qsec vs am gear carb
TRUE TRUE TRUE FALSE FALSE FALSE FALSE FALSE FALSE FALSE FALSE
```

MPG , . cyl disp . . .

R - : <https://riptutorial.com/ko/r/topic/7561/r----->

# 29: R I/O

## Examples

### Rds RData (Rda)

```
.rds .Rdata (.rda) R R . (:write.table .
```

- R .
  - R ( :, ) .
- 

```
saveRDS / readRDS R .
```

```
, .rds iris :
```

```
saveRDS(object = iris, file = "my_data_frame.rds")
```

```
:
```

```
iris2 <- readRDS(file = "my_data_frame.rds")
```

---

```
save() save() .Rdata .
```

```
:2 :
```

```
save(iris, cars, file = "myIrisAndCarsData.Rdata")
```

```
:
```

```
load("myIrisAndCarsData.Rdata")
```

```
save load :
```

```
save(iris, cars, file = "myIrisAndCarsData.Rdata", envir = foo <- new.env())
load("myIrisAndCarsData.Rdata", envir = foo)
foo$cars
```

```
save(iris, cars, file = "myIrisAndCarsData.Rdata", envir = foo <- new.env())
load("myIrisAndCarsData.Rdata", envir = foo)
foo$cars
```

R I/O : <https://riptutorial.com/ko/r/topic/5540/r----i---o>

# 30: R

## Examples

### RCurl

imdb .

```
R> library(RCurl)
R> library(XML)
R> url <- "http://www.imdb.com/chart/top"
R> top <- getURL(url)
R> parsed_top <- htmlParse(top, encoding = "UTF-8")
R> top_table <- readHTMLTable(parsed_top) [[1]]
R> head(top_table[1:10, 1:3])

Rank & Title IMDb Rating
1 1. The Shawshank Redemption (1994) 9.2
2 2. The Godfather (1972) 9.2
3 3. The Godfather: Part II (1974) 9.0
4 4. The Dark Knight (2008) 8.9
5 5. Pulp Fiction (1994) 8.9
6 6. The Good, the Bad and the Ugly (1966) 8.9
7 7. Schindler's List (1993) 8.9
8 8. 12 Angry Men (1957) 8.9
9 9. The Lord of the Rings: The Return of the King (2003) 8.9
10 10. Fight Club (1999) 8.8
```

R : <https://riptutorial.com/ko/r/topic/4336/r-->

# 31: R

R . R ?regex . ICU .

## Examples

`grep`

```
General syntax:
grep(<pattern>, <character vector>)

mystring <- c('The number 5',
 'The number 8',
 '1 is the loneliest number',
 'Company, 3 is',
 'Git SSH tag is git@github.com',
 'My personal site is www.personal.org',
 'path/to/my/file')

grep('5', mystring)
[1] 1
grep('@', mystring)
[1] 5
grep('number', mystring)
[1] 1 2 3
```

x|y "x" "y" .

```
grep('5|8', mystring)
[1] 1 2
grep('com|org', mystring)
[1] 5 6
```

. Regex . " "

```
grep('The number .', mystring)
[1] 1 2
```

!

```
tricky <- c('www.personal.org', 'My friend is a cyborg')
grep('.org', tricky)
[1] 1 2
```

(\). R ( ).

```
grep('\.org', tricky)
Error: '\.' is an unrecognized escape in character string starting "'\.'
grep('\\.org', tricky)
[1] 1
```

( [ ] ) .

```
grep('[13]', mystring)
[1] 3 4
grep('[@/]', mystring)
[1] 5 7
```

. [0-4] 0, 1, 2, 3 4 [AZ] [Az] [A-z0-9] ( )

```
grep('[0-4]', mystring)
[1] 3 4
grep('[A-Z]', mystring)
[1] 1 2 4 5 6
```

R . [:lower:] az, [:upper:] AZ, [:alpha:] Az, [:digit:] 0-9, [:alnum:] A-z0-9 .  
[[:digit:]] ( ). [@[:digit:]/] @, / 0-9 .

```
grep('[[[:digit:]]]', mystring)
[1] 1 2 3 4
grep('@[:digit:]/]', mystring)
[1] 1 2 3 4 5 7
```

( ^ ) . , [^5] "5" .

```
grep('The number [^5]', mystring)
[1] 2
```

R : <https://riptutorial.com/ko/r/topic/9743/r--->

# 32: S4

GOF Java (OOP) ("").

R , S4 Object System .

## Examples

R S4 Class State Machine .

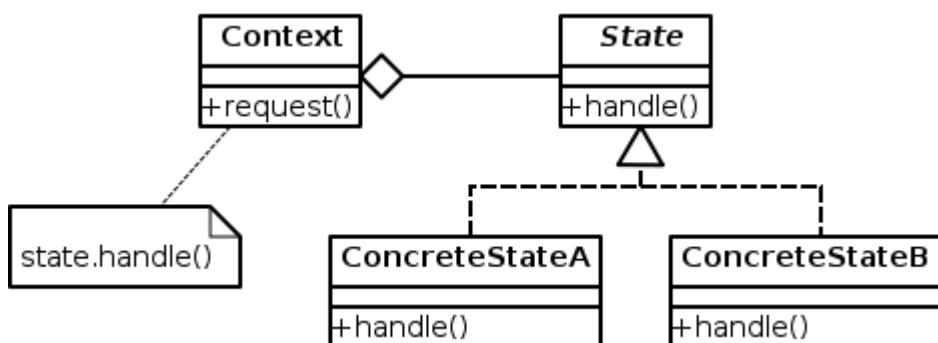
( ";" ) . . . : Name; [Address;] Phone . . . , . . .

```
GREGORY BROWN; 25 NE 25TH; +1-786-987-6543
DAVID SMITH; 786-123-4567
ALAN PEREZ; 25 SE 50TH; +1-786-987-5553
```

- : "^( [AZ] ' ? \s+ ) \* \* [AZ] + ( \s+ [AZ] {1,2} \. ? , ? + ) \* [AZ] + ( (-| \s+) [AZ] + ) \* \$ " . : RAFAEL REAL,  
DAVID R. SMITH, ERNESTO PEREZ GONZALEZ, O' CONNOR BROWN, LUIS PEREZ-MENA
- : "^\s[0-9]{1,4} (\s+[AZ]{1,2}[0-9]{1,2}[AZ]{1,2}|[AZ]\s[0-9]+)\$" . : 11020 LE JEUNE ROAD , 87  
SW 27TH . , , .
- : "^\s\*(\s+(-| \s+)) \* [0-9]{3} (-| \s+) [0-9]{3} (-| \s+) [0-9]{4} \$" . : 305-123-4567, 305 123  
4567, +1-786-123-4567 .

- , .
- R "\s".
- [regex101.com](http://regex101.com) .

. State () (State Pattern ).



- Context : , . ( handle() ), ( handle() State ) . . . Context .
  - : state
  - : handle(), ...

- State : . . .
  - : name, pattern
  - : doAction(), isState(pattern), ...
- Concrete States ( ) : Context State . InitState, NameState, AddressState, PhoneState .

:action, handle(), doAction() goNext() . doAction() Context handle() ( State Context ) .

## S4 Person .

```
setClass(Class = "Person",
 slots = c(name = "character", address = "character", phone = "character")
)
```

. setClass prototype, representation "initialize" .

```
setMethod("initialize", "Person",
 definition = function(.Object, name = NA_character_,
 address = NA_character_, phone = NA_character_) {
 .Object@name <- name
 .Object@address <- address
 .Object@phone <- phone
 .Object
 }
)
```

## initialize methods . R .

```
> initialize
```

```
function (.Object, ...) {...}
```

setMethod exactly (.Object) .

show , Java toString() .

```
setMethod("show", signature = "Person",
 definition = function(object) {
 info <- sprintf("%s@[name='%s', address='%s', phone='%s']",
 class(object), object@name, object@address, object@phone)
 cat(info)
 invisible(NULL)
 }
)
```

: toString() Java .

(Person) R (: ). ( ).

```

setGeneric(name = "as.list", signature = c('x'),
 def = function(x) standardGeneric("as.list"))

Suggestion taken from here:
http://stackoverflow.com/questions/30386009/how-to-extend-as-list-in-a-canonical-way-to-s4-
objects
setMethod("as.list", signature = "Person",
 definition = function(x) {
 mapply(function(y) {
 #apply as.list if the slot is again an user-defined object
 #therefore, as.list gets applied recursively
 if (inherits(slot(x,y),"Person")) {
 as.list(slot(x,y))
 } else {
 #otherwise just return the slot
 slot(x,y)
 }
 },
 slotNames(class(x)),
 SIMPLIFY=FALSE)
 }
)

```

**R OO . Statisticians . . 1) ( setGeneric ) 2) ( setMethod ) . .**

**S4 , State .**

```

setClass(Class = "State", slots = c(name = "character", pattern = "character"))

setMethod("initialize", "State",
 definition = function(.Object, name = NA_character_, pattern = NA_character_) {
 .Object@name <- name
 .Object@pattern <- pattern
 .Object
 }
)

setMethod("show", signature = "State",
 definition = function(object) {
 info <- sprintf("%s@[name='%s', pattern='%s']", class(object),
 object@name, object@pattern)
 cat(info)
 invisible(NULL)
 }
)

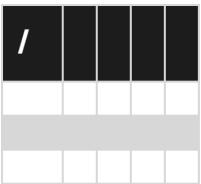
setGeneric(name = "isState", signature = c('obj', 'input'),
 def = function(obj, input) standardGeneric("isState"))

setGeneric(name = "doAction", signature = c('obj', 'input', 'context'),
 def = function(obj, input, context) standardGeneric("doAction"))

```

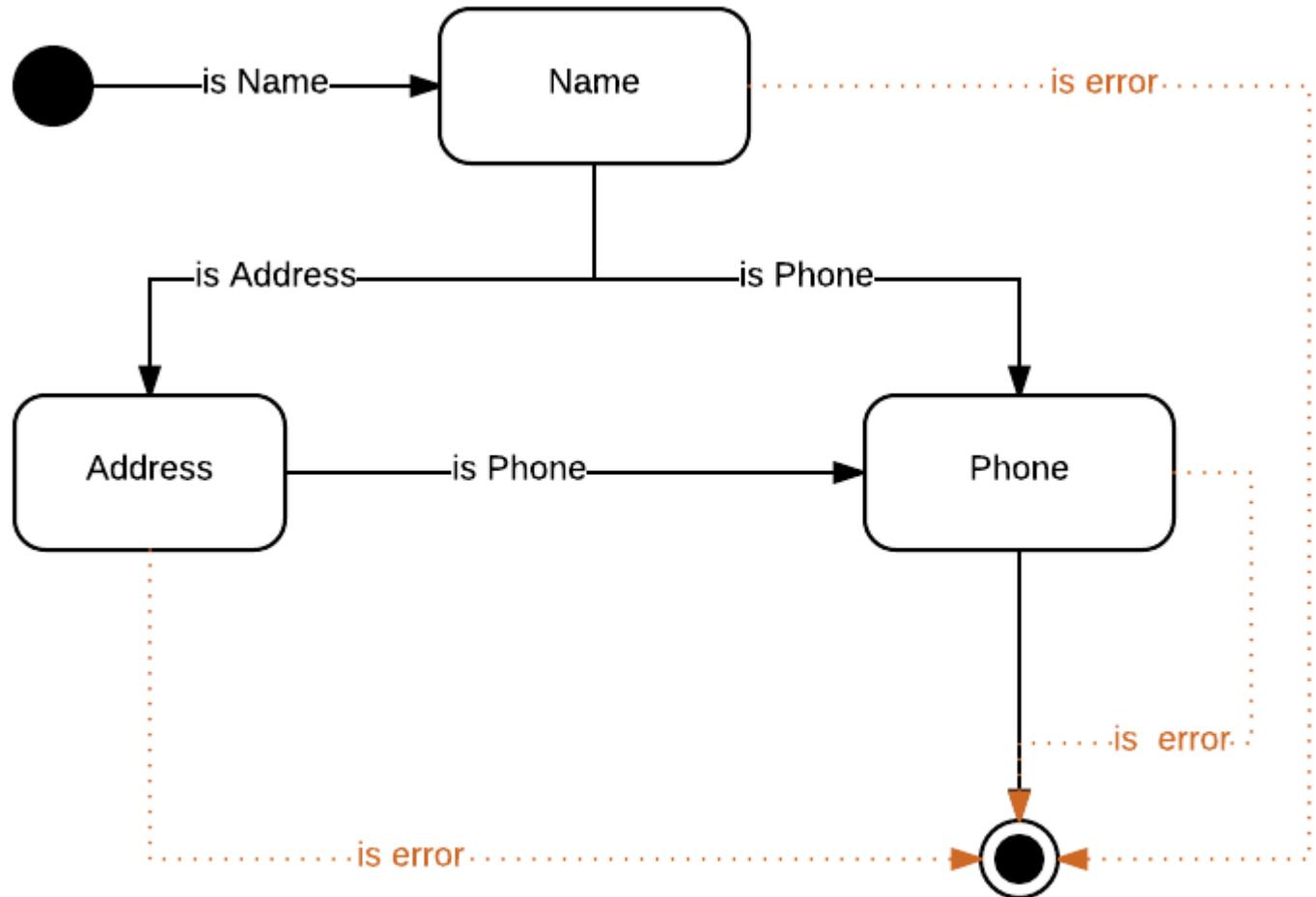
**State name pattern ( isState() ) ( doAction() doAction() ).**





: [row, col]=[i,j] i j .

Name ( ) . UML State Machine .



**is error:** when the input argument has an invalid pattern

State

:

```

setClass("InitState", contains = "State")

setMethod("initialize", "InitState",
 definition = function(.Object, name = "init", pattern = NA_character_) {
 .Object@name <- name
 .Object@pattern <- pattern
 .Object
 }
)

```

```

 }
)

setMethod("show", signature = "InitState",
 definition = function(object) {
 callNextMethod()
 }
)

```

R contains .

show callNextMethod() ).

. NA .

State .

```

setMethod(f = "isState", signature = "InitState",
 definition = function(obj, input) {
 nameState <- new("NameState")
 result <- isState(nameState, input)
 return(result)
 }
)

```

( pattern ), name .

```

setMethod(f = "doAction", signature = "InitState",
 definition = function(obj, input, context) {
 nameState <- new("NameState")
 if (isState(nameState, input)) {
 person <- context@person
 person@name <- trimws(input)
 context@person <- person
 context@state <- nameState
 } else {
 msg <- sprintf("The input argument: '%s' cannot be identified", input)
 stop(msg)
 }
 return(context)
 }
)

```

doAction . @-operator . get/set get/set (OO : ), get-set .

doAction .

```

setClass ("NameState", contains = "State")

setMethod("initialize","NameState",
 definition=function(.Object, name="name",
 pattern = "^([A-Z]'?\\s+)* *[A-Z]+(\\s+[A-Z]{1,2}\\.?,* +)* [A-Z]+((-|\\s+) [A-Z]+)*$"
{
 .Object@pattern <- pattern
}

```

```

 .Object@name <- name
 .Object
}
)

setMethod("show", signature = "NameState",
definition = function(object) {
 callNextMethod()
}
)

```

grepl .

```

setMethod(f="isState", signature="NameState",
definition=function(obj, input) {
 result <- grepl(obj@pattern, input, perl=TRUE)
 return(result)
}
)

```

```

setMethod(f = "doAction", signature = "NameState",
definition=function(obj, input, context) {
 addressState <- new("AddressState")
 phoneState <- new("PhoneState")
 person <- context@person
 if (isState(addressState, input)) {
 person@address <- trimws(input)
 context@person <- person
 context@state <- addressState
 } else if (isState(phoneState, input)) {
 person@phone <- trimws(input)
 context@person <- person
 context@state <- phoneState
 } else {
 msg <- sprintf("The input argument: '%s' cannot be identified", input)
 stop(msg)
 }
 return(context)
}
)
```

- person : address phone .
- state

isState() . ( addressState, phoneState ) .

( ) .

```

setClass("AddressState", contains = "State")

setMethod("initialize", "AddressState",
definition = function(.Object, name="address",

```

```

pattern = "^\s[0-9]{1,4}(\s+[A-Z]{1,2}[0-9]{1,2}[A-Z]{1,2}|[A-Z\s0-9]+)$") {
 .Object@pattern <- pattern
 .Object@name <- name
 .Object
}
)

setMethod("show", signature = "AddressState",
definition = function(object) {
 callNextMethod()
}
)

setMethod(f="isState", signature="AddressState",
definition=function(obj, input) {
 result <- grepl(obj@pattern, input, perl=TRUE)
 return(result)
}
)

setMethod(f = "doAction", "AddressState",
definition=function(obj, input, context) {
 phoneState <- new("PhoneState")
 if (isState(phoneState, input)) {
 person <- context@person
 person@phone <- trimws(input)
 context@person <- person
 context@state <- phoneState
 } else {
 msg <- sprintf("The input argument: '%s' cannot be identified", input)
 stop(msg)
 }
 return(context)
}
)

```

```

setClass("PhoneState", contains = "State")

setMethod("initialize", "PhoneState",
definition = function(.Object, name = "phone",
 pattern = "^\s*(\+|-|\s+)*[0-9]{3}(-|\s+)[0-9]{3}(-|\s+)[0-9]{4}$") {
 .Object@pattern <- pattern
 .Object@name <- name
 .Object
}
)

setMethod("show", signature = "PhoneState",
definition = function(object) {
 callNextMethod()
}
)

setMethod(f = "isState", signature = "PhoneState",
definition = function(obj, input) {
 result <- grepl(obj@pattern, input, perl = TRUE)
 return(result)
}
)
```

```

context persons .

setMethod(f = "doAction", "PhoneState",
 definition = function(obj, input, context) {
 context <- addPerson(context, context@person)
 context@state <- new("InitState")
 return(context)
 }
)

```

Context . . .

```

setClass(Class = "Context",
 slots = c(state = "State", persons = "list", person = "Person")
)

```

- state :
- person : .
- persons : .

: name name .

```

setMethod(f="initialize", signature="Context",
 definition = function(.Object) {
 .Object@state <- new("InitState")
 .Object@persons <- list()
 .Object@person <- new("Person")
 return(.Object)
 }
)

setMethod("show", signature = "Context",
 definition = function(object) {
 cat("An object of class ", class(object), "\n", sep = "")
 info <- sprintf("[state='%s', persons='%s', person='%s']", object@state,
 toString(object@persons), object@person)
 cat(info)
 invisible(NULL)
 }
)

setGeneric(name = "handle", signature = c('obj', 'input', 'context'),
 def = function(obj, input, context) standardGeneric("handle"))

setGeneric(name = "addPerson", signature = c('obj', 'person'),
 def = function(obj, person) standardGeneric("addPerson"))

setGeneric(name = "parseLine", signature = c('obj', 's'),
 def = function(obj, s) standardGeneric("parseLine"))

setGeneric(name = "parseLines", signature = c('obj', 's'),
 def = function(obj, s) standardGeneric("parseLines"))

setGeneric(name = "as.df", signature = c('obj'),
 def = function(obj) standardGeneric("as.df"))

```

- handle() : state doAction() .
- addPerson : persons person .
- parseLine() : .
- parseLines() : ()
- as.df() : persons .

```
handle() context state doAction() .
```

```
setMethod(f = "handle", signature = "Context",
 definition = function(obj, input) {
 obj <- doAction(obj@state, input, obj)
 return(obj)
 }
)

setMethod(f = "addPerson", signature = "Context",
 definition = function(obj, person) {
 obj@persons <- c(obj@persons, person)
 return(obj)
 }
)
```

, R- strsplit() . handle() ( state , person , persons ) context .

```
setMethod(f = "parseLine", signature = "Context",
 definition = function(obj, s) {
 elements <- strsplit(s, ";")[[1]]
 # Adding an empty field for considering the end state.
 elements <- c(elements, "")
 n <- length(elements)
 input <- NULL
 for (i in (1:n)) {
 input <- elements[i]
 obj <- handle(obj, input)
 }
 return(obj@person)
 }
)
```

R , ( obj ) :

```
setMethod(f = "parseLines", signature = "Context",
 definition = function(obj, s) {
 n <- length(s)
 listOfPersons <- list()
 for (i in (1:n)) {
 ipersons <- parseLine(obj, s[i])
 listOfPersons[[i]] <- ipersons
 }
 obj@persons <- listOfPersons
 return(obj)
 }
)
```

```
)
```

```
persons S4 Person . R . as.list Person . persons , lapply() . lappy() ,
data.frame persons.list . rbind() ()
```

```
Sugestion taken from this post:
http://stackoverflow.com/questions/4227223/r-list-to-data-frame
setMethod(f = "as.df", signature = "Context",
definition = function(obj) {
 persons <- obj@persons
 persons.list <- lapply(persons, as.list)
 persons.ds <- do.call(rbind, lapply(persons.list, data.frame, stringsAsFactors = FALSE))
 return(persons.ds)
})
```

```
s <- c(
 "GREGORY BROWN; 25 NE 25TH; +1-786-987-6543",
 "DAVID SMITH; 786-123-4567",
 "ALAN PEREZ; 25 SE 50TH; +1-786-987-5553"
)
```

```
context .
```

```
context <- new("Context")
context <- parseLines(context, s)
```

```
df <- as.df(context)
> df
 name address phone
1 GREGORY BROWN 25 NE 25TH +1-786-987-6543
2 DAVID SMITH <NA> 786-123-4567
3 ALAN PEREZ 25 SE 50TH +1-786-987-5553
```

```
show .
```

```
> show(context@persons[[1]])
Person@[name='GREGORY BROWN', address='25 NE 25TH', phone='+1-786-987-6543']
```

```
:
```

```
>show(new("PhoneState"))
PhoneState@[name='phone', pattern='^\\s*(\\+1(-|\\s+))*[0-9]{3}(-|\\s+)[0-9]{3}(-|\\s+)[0-9]{4}$']
```

```
as.list() .
```

```
> as.list(context@persons[[1]])
$name
```

```
[1] "GREGORY BROWN"
```

```
$address
```

```
[1] "25 NE 25TH"
```

```
$phone
```

```
[1] "+1-786-987-6543"
```

```
>
```

OO R State . R OO OOP . . . : object.setID("A1") Java / C # R setID(object, "A1") . . . this ." " . . ( "Person" , "isState" ) .

S4 Java / C # . , State Pattern . . if-else , State if-else .

:

.

S4 : <https://riptutorial.com/ko/r/topic/9126/s4----->

# 33: sqldf

## Examples

```
sqldf() sqldf R. SQL SQLite .
```

```
ggplot2 "diamonds" 10 .
```

```
data("diamonds")
head(diamonds)
```

```
A tibble: 6 x 10
 carat cut color clarity depth table price x y z
 <dbl> <ord> <ord> <ord> <dbl> <dbl> <int> <dbl> <dbl> <dbl>
1 0.23 Ideal E SI2 61.5 55 326 3.95 3.98 2.43
2 0.21 Premium E SI1 59.8 61 326 3.89 3.84 2.31
3 0.23 Good E VS1 56.9 65 327 4.05 4.07 2.31
4 0.29 Premium I VS2 62.4 58 334 4.20 4.23 2.63
5 0.31 Good J SI2 63.3 58 335 4.34 4.35 2.75
6 0.24 Very Good J VVS2 62.8 57 336 3.94 3.96 2.48
```

```
require(sqldf)
sqldf("select * from diamonds limit 10")
```

```
 carat cut color clarity depth table price x y z
1 0.23 Ideal E SI2 61.5 55 326 3.95 3.98 2.43
2 0.21 Premium E SI1 59.8 61 326 3.89 3.84 2.31
3 0.23 Good E VS1 56.9 65 327 4.05 4.07 2.31
4 0.29 Premium I VS2 62.4 58 334 4.20 4.23 2.63
5 0.31 Good J SI2 63.3 58 335 4.34 4.35 2.75
6 0.24 Very Good J VVS2 62.8 57 336 3.94 3.96 2.48
7 0.24 Very Good I VVS1 62.3 57 336 3.95 3.98 2.47
8 0.26 Very Good H SI1 61.9 55 337 4.07 4.11 2.53
9 0.22 Fair E VS2 65.1 61 337 3.87 3.78 2.49
10 0.23 Very Good H VS1 59.4 61 338 4.00 4.05 2.39
```

```
"E" 10 .
```

```
sqldf("select * from diamonds where color = 'E' limit 10")
```

```
 carat cut color clarity depth table price x y z
1 0.23 Ideal E SI2 61.5 55 326 3.95 3.98 2.43
2 0.21 Premium E SI1 59.8 61 326 3.89 3.84 2.31
3 0.23 Good E VS1 56.9 65 327 4.05 4.07 2.31
4 0.22 Fair E VS2 65.1 61 337 3.87 3.78 2.49
5 0.20 Premium E SI2 60.2 62 345 3.79 3.75 2.27
6 0.32 Premium E I1 60.9 58 345 4.38 4.42 2.68
7 0.23 Very Good E VS2 63.8 55 352 3.85 3.92 2.48
8 0.23 Very Good E VS1 60.7 59 402 3.97 4.01 2.42
9 0.23 Very Good E VS1 59.5 58 402 4.01 4.06 2.40
10 0.23 Good E VS1 64.1 59 402 3.83 3.85 2.46
```

SQL      " " ( ).

1

```
sqldf("select count(*) from diamonds where carat > 1 and color = 'E'")
```

count (\*)

```
sqldf("select *, count(*) as cnt_big_E_colored_stones from diamonds where carat > 1 and color = 'E' group by clarity")
```

, cut price ?

```
sqldf("select cut, max(price) from diamonds group by cut")
```

		cut	max(price)
1		Fair	18574
2		Good	18788
3		Ideal	18806
4	Premium		18823
5	Very Good		18818

`sqldf` : <https://riptutorial.com/ko/r/topic/2100/sqldf>

# 34: stringi

```
install.packages("stringi")
```

```
:
```

```
require("stringi")
```

## Examples

```
stri_count_fixed("babab", "b")
[1] 3
stri_count_fixed("babab", "ba")
[1] 2
stri_count_fixed("babab", "bab")
[1] 1
```

```
length(gregexpr("b", "babab") [[1]])
[1] 3
length(gregexpr("ba", "babab") [[1]])
[1] 2
length(gregexpr("bab", "babab") [[1]])
[1] 1
```

```
stri_count_fixed("babab", c("b", "ba"))
[1] 3 2
stri_count_fixed(c("babab", "bbb", "bca", "abc"), c("b", "ba"))
[1] 3 0 1 0
```

R :

```
sapply(c("b", "ba"), function(x) length(gregexpr(x, "babab") [[1]]))
b ba
3 2
```

- a

- a

```
stri_count_regex("a1 b2 a3 b4 aa", "a.")
[1] 3
stri_count_regex("a1 b2 a3 b4 aa", "a\\d")
[1] 2
```

```
stri_dup("abc", 3)
[1] "abcabcabc"
```

R .

```
paste0(rep("abc", 3), collapse = "")
[1] "abcabcabc"
```

```
stri_paste(LETTERS, "-", 1:13)
[1] "A-1" "B-2" "C-3" "D-4" "E-5" "F-6" "G-7" "H-8" "I-9" "J-10" "K-11" "L-12" "M-
13"
[14] "N-1" "O-2" "P-3" "Q-4" "R-5" "S-6" "T-7" "U-8" "V-9" "W-10" "X-11" "Y-12" "Z-
13"
```

R .

```
> paste(LETTERS, 1:13, sep="-")
[1] "A-1" "B-2" "C-3" "D-4" "E-5" "F-6" "G-7" "H-8" "I-9" "J-10" "K-11" "L-12" "M-
13"
[14] "N-1" "O-2" "P-3" "Q-4" "R-5" "S-6" "T-7" "U-8" "V-9" "W-10" "X-11" "Y-12" "Z-
13"
```

```
stri_split_fixed(c("To be or not to be.", "This is very short sentence."), " ")
[[1]]
[1] "To" "be" "or" "not" "to" "be."
#
[[2]]
[1] "This" "is" "very" "short" "sentence."
```

```
stri_split_fixed("Apples, oranges and pineapllies.", c(" ", ", ", "s"))
[[1]]
[1] "Apples," "oranges" "and" "pineapllies."
#
[[2]]
[1] "Apples" "oranges and pineapllies."
#
[[3]]
[1] "Apple" ", orange" "and pineaplle" ". "
```

stringi : <https://riptutorial.com/ko/r/topic/1670/stringi--->

# 35: strsplit

- strsplit (
- 
- 
- =
- perl = FALSE
- useBytes = FALSE)

## Examples

```
strsplit . R data.frame .
```

```
strsplit : :
```

```
temp <- c("this,that,other", "hat,scarf,food", "woman,man,child")
get a list split by commas
myList <- strsplit(temp, split=", ")
print myList
myList
[[1]]
[1] "this" "that" "other"

[[2]]
[1] "hat" "scarf" "food"

[[3]]
[1] "woman" "man" "child"
```

```
split . , temp2 temp .
```

```
temp2 <- c("this, that, other", "hat,scarf ,food", "woman; man ; child")
myList2 <- strsplit(temp2, split=" ?[,;] ?")
myList2
[[1]]
[1] "this" "that" "other"

[[2]]
[1] "hat" "scarf" "food"

[[3]]
[1] "woman" "man" "child"
```

1..

2.. R , R .

strsplit : <https://riptutorial.com/ko/r/topic/2762/strsplit->

# 36: texreg

texreg ( ) . HTML .doc (MS Office Word) .

## Examples

```
models
fit1 <- lm(mpg ~ wt, data = mtcars)
fit2 <- lm(mpg ~ wt+hp, data = mtcars)
fit3 <- lm(mpg ~ wt+hp+cyl, data = mtcars)

export to html
texreg::htmlreg(list(fit1,fit2,fit3),file='models.html')

export to doc
texreg::htmlreg(list(fit1,fit2,fit3),file='models.doc')
```

	<b>Model 1</b>	<b>Model 2</b>	<b>Model 3</b>
(Intercept)	37.29*** (1.88)	37.23*** (1.60)	38.75*** (1.79)
wt	-5.34*** (0.56)	-3.88*** (0.63)	-3.17*** (0.74)
hp		-0.03** (0.01)	-0.02 (0.01)
cyl			-0.94 (0.55)
R2	0.75	0.83	0.84
Adj. R2	0.74	0.81	0.83
Num. obs.	32	32	32
RMSE	3.05	2.59	2.51

\*\*\*p < 0.001, \*\*p < 0.01, \*p < 0.05

Statistical models

texreg::htmlreg()

```
export to html
texreg::htmlreg(list(fit1,fit2,fit3),file='models.html',
 single.row = T,
 custom.model.names = LETTERS[1:3],
 leading.zero = F,
 digits = 3)
```

	<b>A</b>	<b>B</b>	<b>C</b>
(Intercept)	37.285 (1.878)***	37.227 (1.599)***	38.752 (1.787)***
wt	-5.344 (0.559)***	-3.878 (0.633)***	-3.167 (0.741)***
hp		-0.032 (0.009)**	-0.018 (0.012)
cyl			-0.942 (0.551)
R2	0.753	0.827	0.843
Adj. R2	0.745	0.815	0.826
Num. obs.	32	32	32
RMSE	3.046	2.593	2.512

\*\*\*p < 0.001, \*\*p < 0.01, \*p < 0.05

### Statistical models

**texreg** . . : <https://riptutorial.com/ko/r/topic/9037/texreg----->

# 37: tidyverse

: tidyverse gather() spread().

## Examples

### spread()

```
library(tidyverse)

example data
set.seed(123)
df <- data.frame(
 name = rep(c("firstName", "secondName"), each=4),
 numbers = rep(1:4, 2),
 value = rnorm(8)
)
df
#> #> name numbers value
#> # 1 firstName 1 -0.56047565
#> # 2 firstName 2 -0.23017749
#> # 3 firstName 3 1.55870831
#> # 4 firstName 4 0.07050839
#> # 5 secondName 1 0.12928774
#> # 6 secondName 2 1.71506499
#> # 7 secondName 3 0.46091621
#> # 8 secondName 4 -1.26506123
```

" "

```
spread(data = df,
 key = numbers,
 value = value)
#> #> name 1 2 3 4
#> # 1 firstName -0.5604756 -0.2301775 1.5587083 0.07050839
#> # 2 secondName 0.1292877 1.7150650 0.4609162 -1.26506123
```

" "

```
spread(data = df,
 key = name,
 value = value)
#> #> numbers firstName secondName
#> # 1 1 -0.56047565 0.1292877
#> # 2 2 -0.23017749 1.7150650
#> # 3 3 1.55870831 0.4609162
#> # 4 4 0.07050839 -1.2650612
```

### gather()

```
library(tidyr)

example data
df <- read.table(text ="
 numbers firstName secondName
1 1 1.5862639 0.4087477
2 2 0.1499581 0.9963923
3 3 0.4117353 0.3740009
4 4 -0.4926862 0.4437916", header = T)
df
#> #> numbers firstName secondName
#> #> 1 1 1.5862639 0.4087477
#> #> 2 2 0.1499581 0.9963923
#> #> 3 3 0.4117353 0.3740009
#> #> 4 4 -0.4926862 0.4437916
```

'numbers' .

```
gather(data = df,
 key = numbers,
 value = myValue)
#> #> numbers numbers myValue
#> #> 1 1 firstName 1.5862639
#> #> 2 2 firstName 0.1499581
#> #> 3 3 firstName 0.4117353
#> #> 4 4 firstName -0.4926862
#> #> 5 1 secondName 0.4087477
#> #> 6 2 secondName 0.9963923
#> #> 7 3 secondName 0.3740009
#> #> 8 4 secondName 0.4437916
```

tidyr : <https://riptutorial.com/ko/r/topic/9195/tidyr-->

# 38: xgboost

## Examples

### xgboost

```
library(caret) # for dummyVars
library(RCurl) # download https data
library(Metrics) # calculate errors
library(xgboost) # model

#####
Load data from UCI Machine Learning Repository (http://archive.ics.uci.edu/ml/datasets.html)
urlfile <- 'https://archive.ics.uci.edu/ml/machine-learning-databases/adult/adult.data'
x <- getURL(urlfile, ssl.verifypeer = FALSE)
adults <- read.csv(textConnection(x), header=F)

adults <-read.csv('https://archive.ics.uci.edu/ml/machine-learning-
databases/adult/adult.data', header=F)
names(adults)=c('age','workclass','fnlwgt','education','educationNum',
 'maritalStatus','occupation','relationship','race',
 'sex','capitalGain','capitalLoss','hoursWeek',
 'nativeCountry','income')
clean up data
adults$income <- ifelse(adults$income==' <=50K',0,1)
binarize all factors
library(caret)
dmy <- dummyVars(~ ., data = adults)
adultsTrsf <- data.frame(predict(dmy, newdata = adults))
#####

what we're trying to predict adults that make more than 50k
outcomeName <- c('income')
list of features
predictors <- names(adultsTrsf)[!names(adultsTrsf) %in% outcomeName]

play around with settings of xgboost - eXtreme Gradient Boosting (Tree) library
https://github.com/tqchen/xgboost/wiki/Parameters
max.depth - maximum depth of the tree
nrounds - the max number of iterations

take first 10% of the data only!
trainPortion <- floor(nrow(adultsTrsf)*0.1)

trainSet <- adultsTrsf[1:floor(trainPortion/2),]
testSet <- adultsTrsf[(floor(trainPortion/2)+1):trainPortion,]

smallestError <- 100
for (depth in seq(1,10,1)) {
 for (rounds in seq(1,20,1)) {

 # train
 bst <- xgboost(data = as.matrix(trainSet[,predictors]),
 label = trainSet[,outcomeName],
 max.depth=depth, nround=rounds,
 objective = "reg:linear", verbose=0)
 gc()
 }
}
```

```

 # predict
 predictions <- predict(bst, as.matrix(testSet[,predictors]),
outputmargin=TRUE)
 err <- rmse(as.numeric(testSet[,outcomeName]), as.numeric(predictions))

 if (err < smallestError) {
 smallestError = err
 print(paste(depth,rounds,err))
 }
 }
}

cv <- 30
trainSet <- adultsTrsf[1:trainPortion,]
cvDivider <- floor(nrow(trainSet) / (cv+1))

smallestError <- 100
for (depth in seq(1,10,1)) {
 for (rounds in seq(1,20,1)) {
 totalError <- c()
 indexCount <- 1
 for (cv in seq(1:cv)) {
 # assign chunk to data test
 dataTestIndex <- c((cv * cvDivider):(cv * cvDivider + cvDivider))
 dataTest <- trainSet[dataTestIndex,]
 # everything else to train
 dataTrain <- trainSet[-dataTestIndex,]

 bst <- xgboost(data = as.matrix(dataTrain[,predictors]),
 label = dataTrain[,outcomeName],
 max.depth=depth, nround=rounds,
 objective = "reg:linear", verbose=0)
 gc()
 predictions <- predict(bst, as.matrix(dataTest[,predictors]),
outputmargin=TRUE)

 err <- rmse(as.numeric(dataTest[,outcomeName]),
as.numeric(predictions))
 totalError <- c(totalError, err)
 }
 if (mean(totalError) < smallestError) {
 smallestError = mean(totalError)
 print(paste(depth,rounds,smallestError))
 }
 }
}

#####
Test both models out on full data set

trainSet <- adultsTrsf[1:trainPortion,]

assign everything else to test
testSet <- adultsTrsf[(trainPortion+1):nrow(adultsTrsf),]

bst <- xgboost(data = as.matrix(trainSet[,predictors]),
 label = trainSet[,outcomeName],
 max.depth=4, nround=19, objective = "reg:linear", verbose=0)
pred <- predict(bst, as.matrix(testSet[,predictors]), outputmargin=TRUE)
rmse(as.numeric(testSet[,outcomeName]), as.numeric(pred))

```

```
bst <- xgboost(data = as.matrix(trainSet[,predictors]),
 label = trainSet[,outcomeName],
 max.depth=3, nround=20, objective = "reg:linear", verbose=0)
pred <- predict(bst, as.matrix(testSet[,predictors]), outputmargin=TRUE)
rmse(as.numeric(testSet[,outcomeName]), as.numeric(pred))
```

xgboost : <https://riptutorial.com/ko/r/topic/3239/xgboost>

# 39:

( : as.numeric, as.data.frame ) R .

## Examples

R . ,

```
x = 1:3
x
[1] 1 2 3
typeof(x)
#[1] "integer"

x[2] = "hi"
x
#[1] "1" "hi" "3"
typeof(x)
#[1] "character"
```

x integer integer . x[2] = "hi" , R x character .

: <https://riptutorial.com/ko/r/topic/9793/>

# 40:

## Examples

```
: / . ! @DataTx " " . (-))
```

```
R ("" " ") nlme () lme4 () .
```

```
library(nlme)
library(lme4)
m1.nlme <- lme(Reaction~Days, random=~Days|Subject, data=sleepstudy, method="REML")
m1.lme4 <- lmer(Reaction~Days+(Days|Subject), data=sleepstudy, REML=TRUE)
all.equal(fixef(m1.nlme), fixef(m1.lme4))
[1] TRUE
```

- 
- **nlme S-PLUS Pinheiro Bates 2000 Mixed-effects** ( vignette("lmer", package="lme4")  
2015 *Journal of Statistical Software* / vignette("lmer", package="lme4") lme4 )
- lme4
- nlme (P) - lme4 lmerTest afex
- nlme ( / )

[GLMM FAQ](#) (GLMM)

: <https://riptutorial.com/ko/r/topic/3460/>--

# 41:

## Examples

XY

R .

XY . XY f} .

```
rgdal sp . R Spatial*DataFrame (* Points , Lines Polygons) .
```

OpenGeocode .

CSV . rgdal .

```
setwd("D:/GeocodeExample/")
library(rgdal)
```

CSV data.frame R.

```
xy <- read.csv("worldcities.csv", stringsAsFactors = FALSE)
```

(: , ) .

```
head(xy)
str(xy)
```

"-33.532" . SpatialPointsDataFrame() numeric . .

```
xy$latitude <- as.numeric(xy$latitude)
xy$longitude <- as.numeric(xy$longitude)
```

NA . .

```
xy <- xy[!is.na(xy$longitude),]
```

XY . (CRS) .

```
xySPoints <- SpatialPointsDataFrame(coords = c(xy[,c("longitude", "latitude")]),
proj4string = CRS("+proj=longlat +ellps=WGS84 +datum=WGS84 +no_defs"),
data = xy
)
```

```
plot(xySPoints, pch = ".")
```



(.shp)

## rgdal

ESRI shape rgdal rgdal readOGR() R .

```
library(rgdal)
shp <- readOGR(dsn = "/path/to/your/file", layer = "filename")
```

dsn / layer (: .shp ) .

shapefile raster shapefile .

```
library(raster)
shp <- shapefile("path/to/your/file.shp")
```

rgdal import .

tmap rgdal::readORG .

```
library(tmap)
sph <- read_shape("path/to/your/file.shp")
```

: <https://riptutorial.com/ko/r/topic/2093/>

# 42:

## Examples

R Wilkinson-Rogers 1 .

Linear Regressions lm formula . formula .

```
my_formula1 <- formula(mpg ~ wt)
class(my_formula1)
gives "formula"

mod1 <- lm(my_formula1, data = mtcars)
coef(mod1)
gives (Intercept) wt
37.285126 -5.344472
```

~ (LHS) (RHS) . formula formula .

```
form <- mpg ~ wt
class(form)
#[1] "formula"
```

formula ~ .

```
form_mt <- formula(mpg ~ wt, env = mtcars)
```

wt () . 0 -1 formula .

```
coef(lm(mpg ~ 0 + wt, data = mtcars))
coef(lm(mpg ~ wt -1, data = mtcars))
```

a b a:b formula :

```
coef(lm(mpg ~ wt:vs, data = mtcars))
```

formula a + b + a:b . a\*b . \*

```
coef(lm(mpg ~ wt*vs, data = mtcars))
```

\*

```
coef(lm(mpg ~ wt*vs*hp, data = mtcars))
```

7 . 3 , , .

3 . - .

```
coef(lm(mpg ~ wt*vs*hp - wt:vs:hp, data = mtcars))
```

^

```
coef(lm(mpg ~ (wt + vs + hp) ^ 2, data = mtcars))
```

.

.. data (LHS) .:

```
coef(lm(mpg ~ ., data = mtcars))
```

10 .. . . ( ?update.formula ?update.formula ).

1. GN Wilkinson CE Rogers. . C () Vol. 22, No.3 (1973), pp. 392-399

, 2 2

y ~ . . y . . y ~ var1 + var2 + var3+...+var15

y ~ . ^ 2 () 2 . y ~ var1 + var2 + ...+var15 + var1:var2 + var1:var3 + var1:var4...and so on

y ~ var1 + var2 + ...+var15 + I(var1^2) + I(var2^2) + I(var3^2)...+I(var15^2) : I(var^2) 2 .

y ~ poly(var1, degree = 2) + poly(var2, degree = 2)+...poly(var15, degree = 2)

y ~ poly(var1, var2, var3, ....var15, degree = 2) .

poly(var1, degree = 2) var1 + I(var1^2) .

3 poly() degree = 3 .

poly I(var, 2) . I(var, 2) poly() poly() .

, 2 2 .

y ~ .^2 + I(var1^2) + I(var2^2)+...I(var15^2)

4 :

```
old <- reformulate('y ~ x1+x2+x3+x4')
new <- reformulate(" y ~ .^2 + I(x1^2) + I(x2^2) + I(x3^2) + I(x4^2) ")
tmp <- .Call(stats:::C_updateform, old, new)
terms.formula(tmp, simplify = TRUE)

~y ~ x1 + x2 + x3 + x4 + I(x1^2) + I(x2^2) + I(x3^2) + I(x4^2) +
x1:x2 + x1:x3 + x1:x4 + x2:x3 + x2:x4 + x3:x4
attr(),"variables")
list(~y, x1, x2, x3, x4, I(x1^2), I(x2^2), I(x3^2), I(x4^2))
attr(),"factors")
x1 x2 x3 x4 I(x1^2) I(x2^2) I(x3^2) I(x4^2) x1:x2 x1:x3 x1:x4 x2:x3 x2:x4 x3:x4
~y 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
```

```

x1 1 0 0 0 0 0 0 1 1 1 0 0 0
x2 0 1 0 0 0 0 0 1 0 0 1 1 1 0
x3 0 0 1 0 0 0 0 0 1 0 0 1 0 1
x4 0 0 0 1 0 0 0 0 0 1 0 1 0 1
I(x1^2) 0 0 0 0 1 0 0 0 0 0 0 0 0 0
I(x2^2) 0 0 0 0 0 1 0 0 0 0 0 0 0 0
I(x3^2) 0 0 0 0 0 0 1 0 0 0 0 0 0 0
I(x4^2) 0 0 0 0 0 0 0 1 0 0 0 0 0 0
attr(),"term.labels")
[1] "x1" "x2" "x3" "x4" "I(x1^2)" "I(x2^2)" "I(x3^2)" "I(x4^2)"
[9] "x1:x2" "x1:x3" "x1:x4" "x2:x3" "x2:x4" "x3:x4"
attr(),"order")
[1] 1 1 1 1 1 1 1 2 2 2 2 2
attr(),"intercept")
[1] 1
attr(),"response")
[1] 1
attr(),".Environment")
<environment: R_GlobalEnv>

```

: <https://riptutorial.com/ko/r/topic/1061/>

# 43:

## Examples

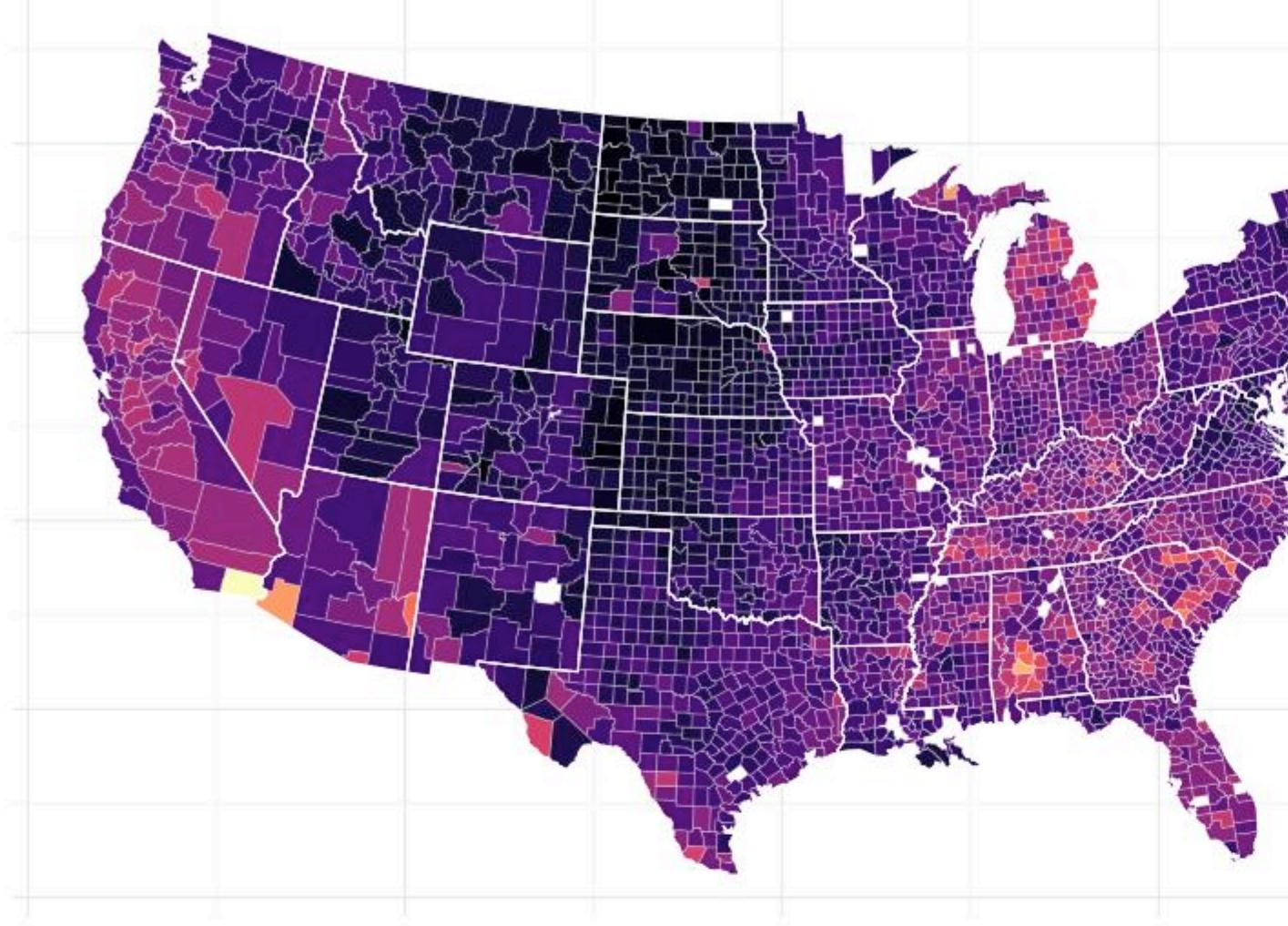
**viridis** -

Viridis ( chromis viridis fish ) Python matplotlib ( ). R .

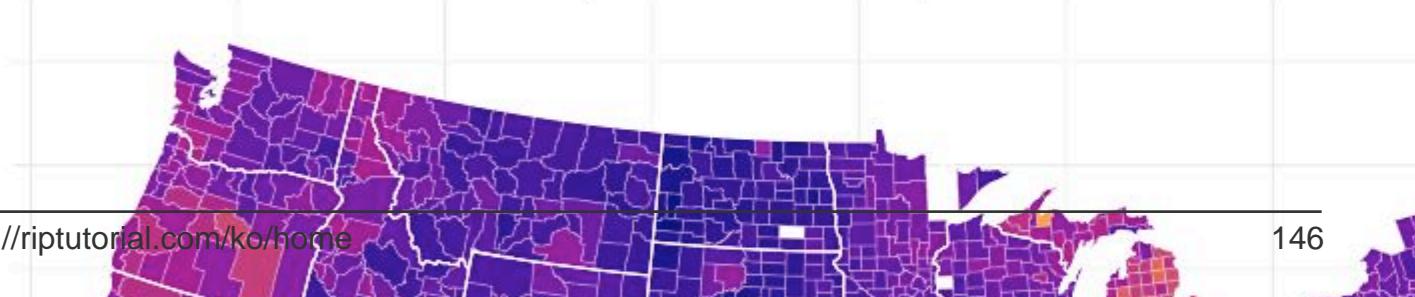
magma , plasma , inferno viridis 4 () . option A , B , C D . 4 .

U

option A aka 'magma'



option C aka 'plasma'



( souce )

---

CRAN github

---

viridis

---

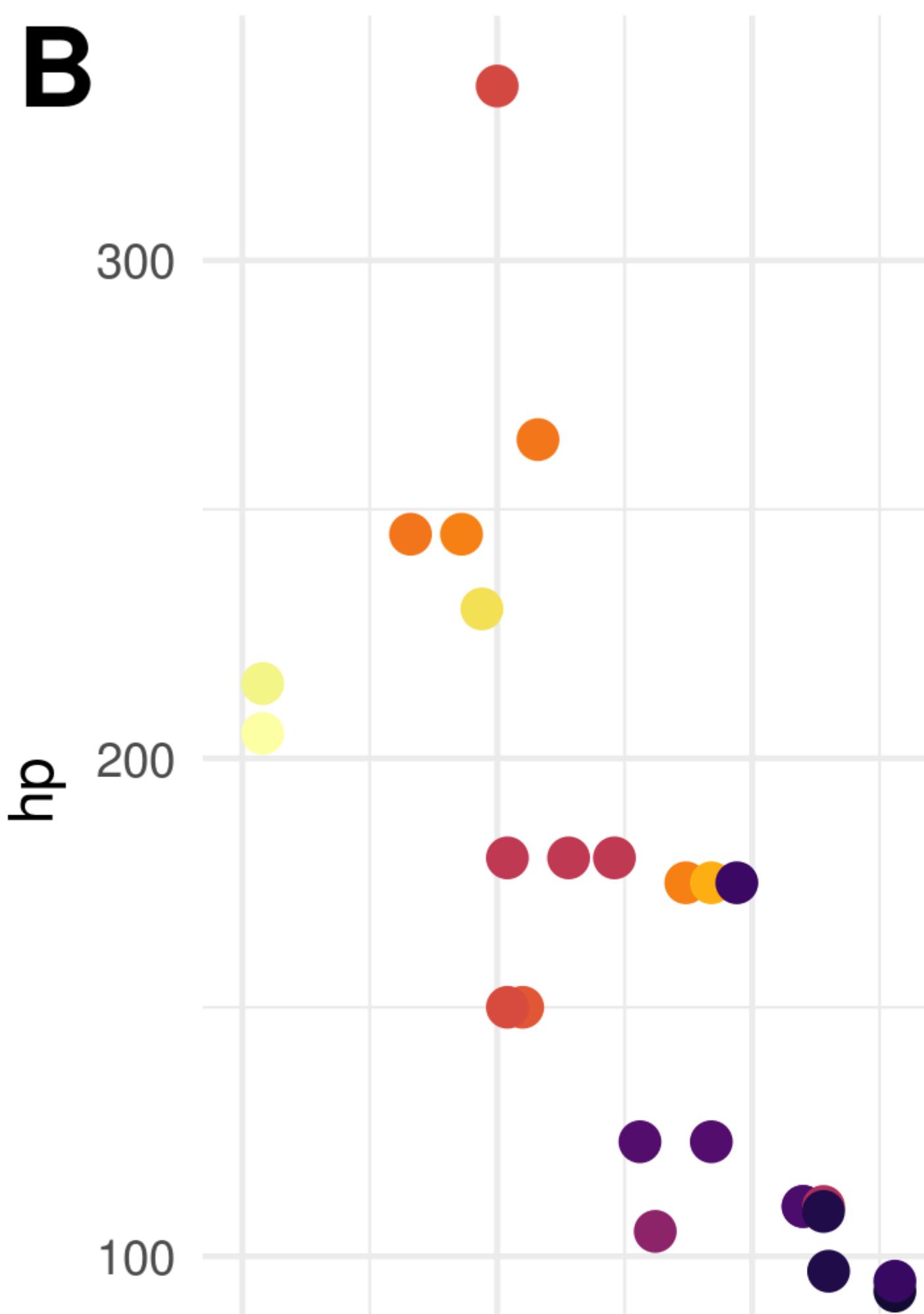
viridis ggplot2 . ggplot2 scale\_color\_viridis() scale\_fill\_viridis() . .

```
library(viridis)
library(ggplot2)

gg1 <- ggplot(mtcars) +
 geom_point(aes(x = mpg, y = hp, color = disp), size = 3) +
 scale_color_viridis(option = "B") +
 theme_minimal() +
 theme(legend.position = c(.8,.8))

gg2 <- ggplot(mtcars) +
 geom_violin(aes(x = factor(cyl), y = hp, fill = factor(cyl))) +
 scale_fill_viridis(discrete = T) +
 theme_minimal() +
 theme(legend.position = 'none')

library(cowplot)
output <- plot_grid(gg1,gg2, labels = c('B','D'),label_size = 20)
print(output)
```

**B**

## Examples

Random Forest (Breiman, L. (2001), Random Forests, *Machine Learning* 45 (5) , 5-32 ).

randomForest Breiman Fortran R .

```
iris factor R . . .
```

```
library(randomForest)

rf <- randomForest(x = iris[, 1:4],
 y = iris$Species,
 ntree = 500,
 do.trace = 100)

rf

Call:
randomForest(x = iris[, 1:4], y = iris$Species, ntree = 500, do.trace = 100)
Type of random forest: classification
Number of trees: 500
No. of variables tried at each split: 2
#
OOB estimate of error rate: 4%
Confusion matrix:
setosa versicolor virginica class.error
setosa 50 0 0 0.00
versicolor 0 47 3 0.06
virginica 0 3 47 0.06
```

	. factor .
ntree	CART
do.trace	,

: <https://riptutorial.com/ko/r/topic/8326/>

# 45:

## Examples

R Map , Reduce , Filter , Find , Position , Negate .

Map .

```
words <- list("this", "is", "an", "example")
Map(toupper, words)
```

Reduce 2 .

```
Reduce(`*`, 1:10)
```

Filter TRUE .

```
Filter(is.character, list(1,"a",2,"b",3,"c"))
```

Find TRUE .

```
Find(is.character, list(1,"a",2,"b",3,"c"))
```

Position TRUE .

```
Position(is.character, list(1,"a",2,"b",3,"c"))
```

Negate TRUE FALSE TRUE .

```
is.noncharacter <- Negate(is.character)
is.noncharacter("a")
is.noncharacter(mean)
```

: <https://riptutorial.com/ko/r/topic/5050/>

# 46:

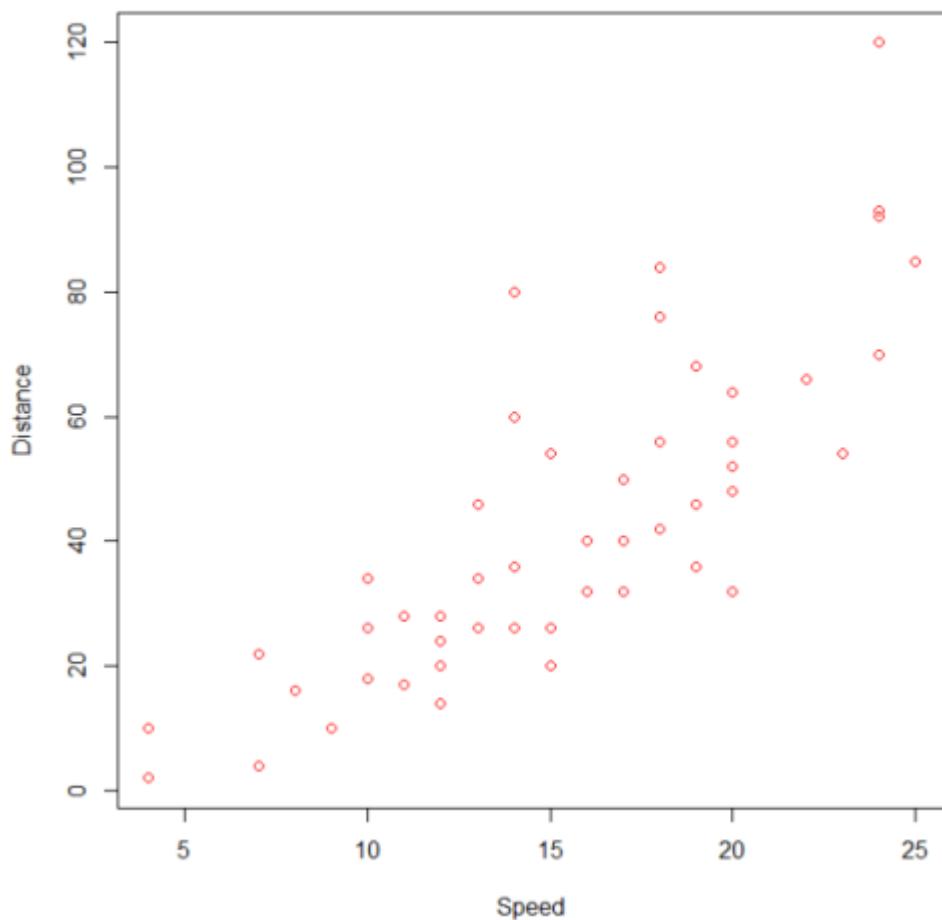
```
x X . data$variablex data[,x]
y Y . data$variabley data[,y]
main
sub
xlab X
ylab Y
pch
col
type . "p" , "l" , "b" , "c" "b" , "o" 'overplotted' "h" 'histogram' (") , "s" , "s" , "n"
" " par . par .
```

## Examples

```
plot() . cars 1920 cars .(()).
```

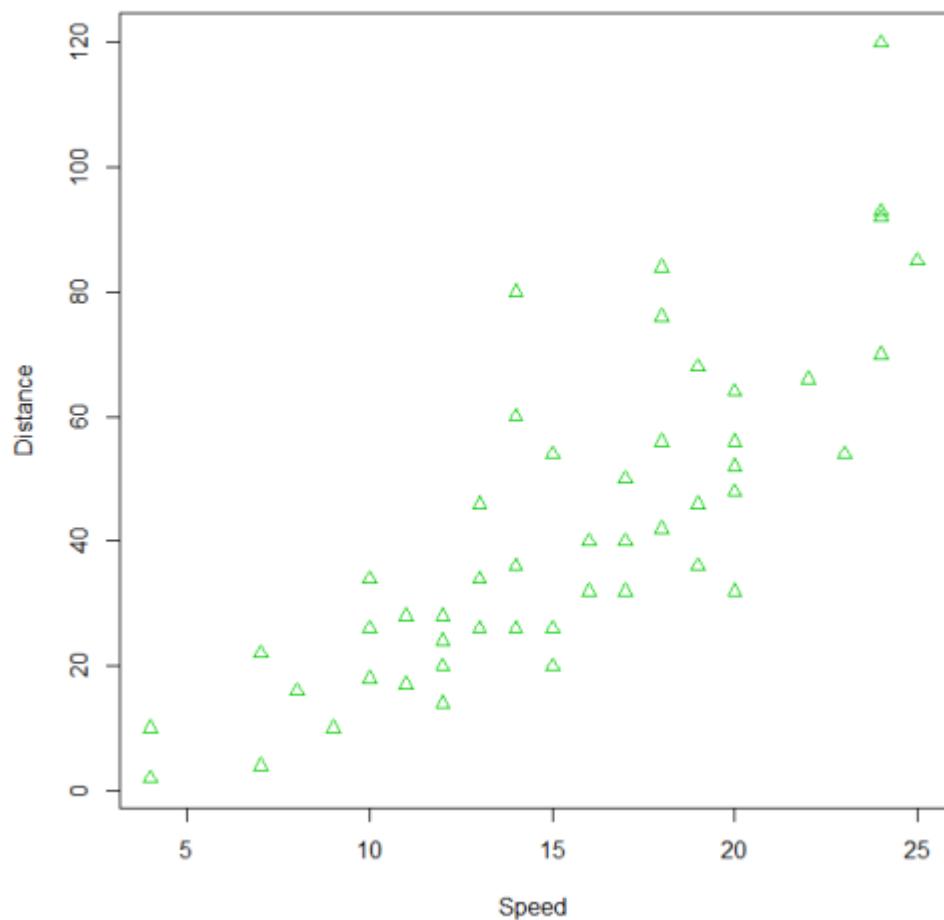
```
plot(x = cars$speed, y = cars$dist, pch = 1, col = 1,
 main = "Distance vs Speed of Cars",
 xlab = "Speed", ylab = "Distance")
```

**Distance to stop vs Speed of Cars**



```
with(cars, plot(dist~speed, pch = 2, col = 3,
 main = "Distance to stop vs Speed of Cars",
 xlab = "Speed", ylab = "Distance"))
```

### Distance to stop vs Speed of Cars

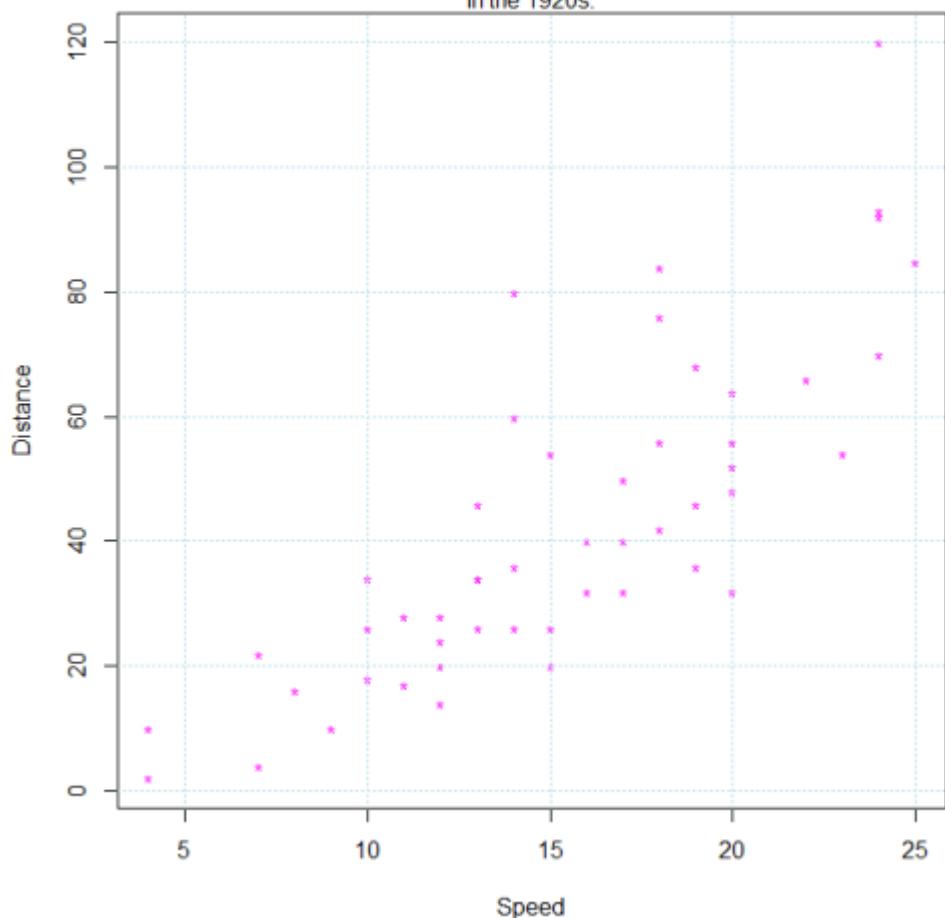


```
points() , text() , mtext() , lines() , grid() .

plot(dist~speed, pch = "*", col = "magenta", data=cars,
 main = "Distance to stop vs Speed of Cars",
 xlab = "Speed", ylab = "Distance")
mtext("In the 1920s.")
grid(col="lightblue")
```

### Distance to stop vs Speed of Cars

In the 1920s.



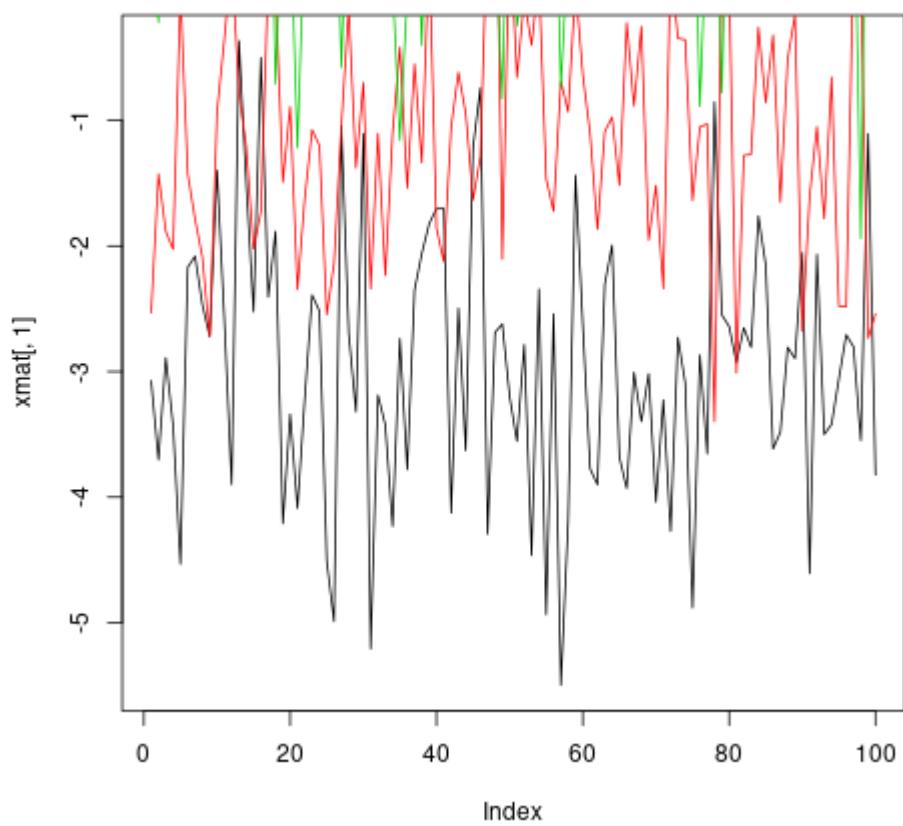
## Matplot

```
matplot () .
```

```
xmat <- cbind(rnorm(100, -3), rnorm(100, -1), rnorm(100, 1), rnorm(100, 3))
head(xmat)
[,1] [,2] [,3] [,4]
[1,] -3.072793 -2.53111494 0.6168063 3.780465
[2,] -3.702545 -1.42789347 -0.2197196 2.478416
[3,] -2.890698 -1.88476126 1.9586467 5.268474
[4,] -3.431133 -2.02626870 1.1153643 3.170689
[5,] -4.532925 0.02164187 0.9783948 3.162121
[6,] -2.169391 -1.42699116 0.3214854 4.480305
```

```
plot points lines .
```

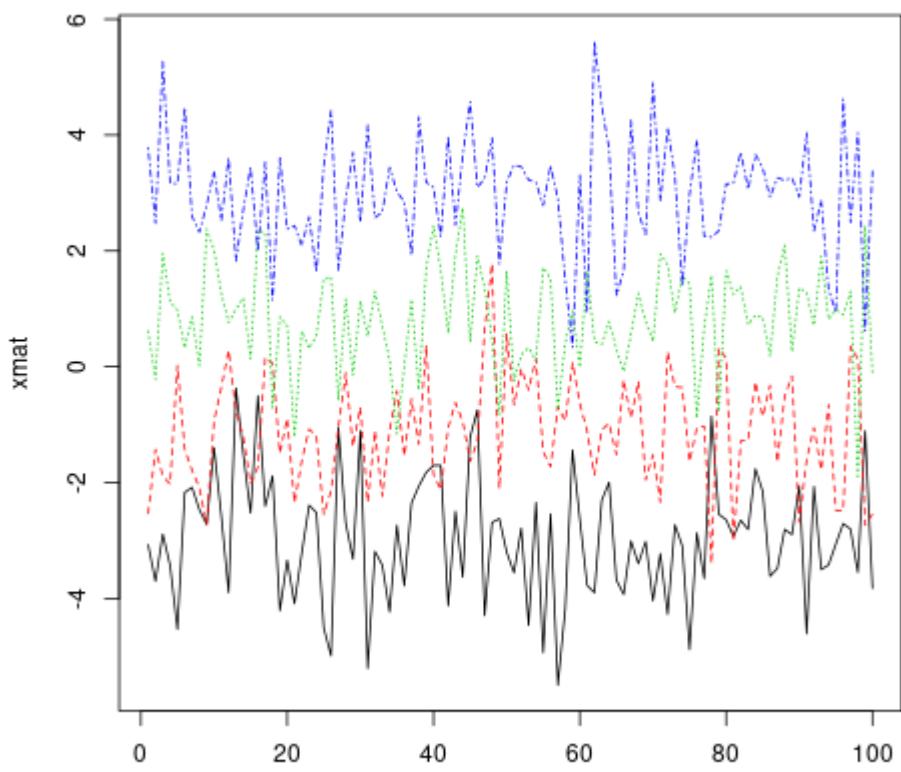
```
plot(xmat[,1], type = 'l')
lines(xmat[,2], col = 'red')
lines(xmat[,3], col = 'green')
lines(xmat[,4], col = 'blue')
```



```
plot
```

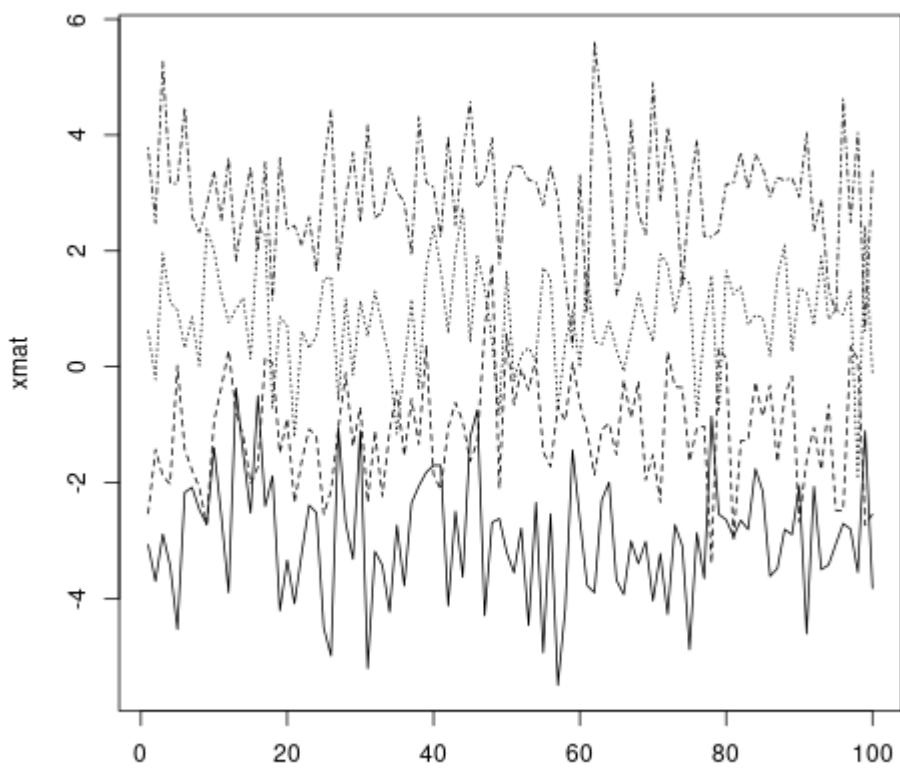
```
matplotlib
```

```
matplot(xmat, type = 'l')
```



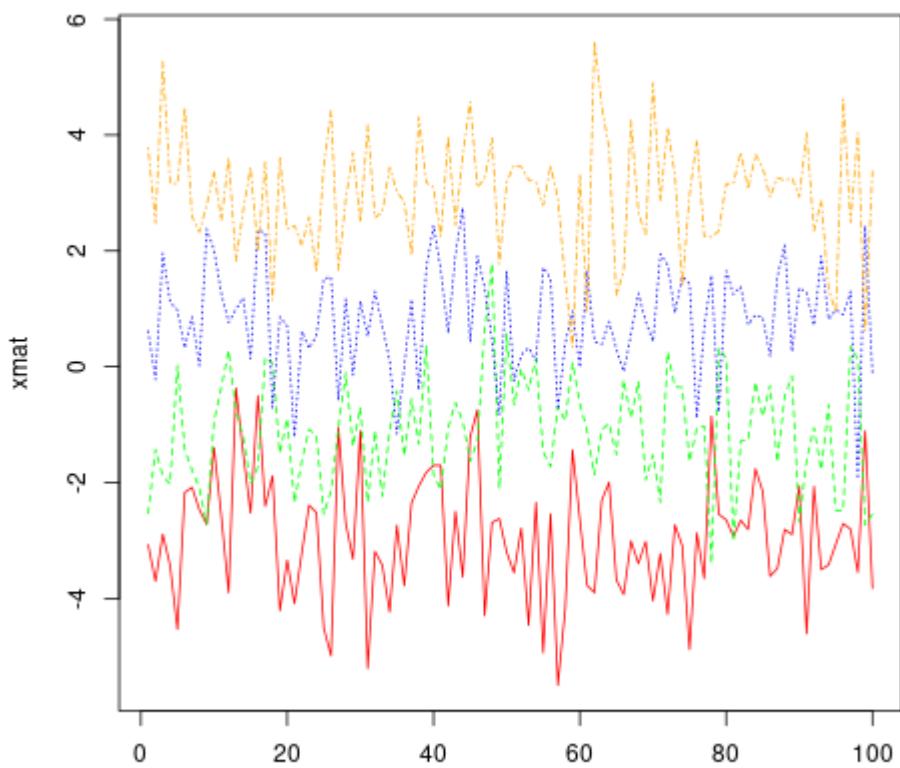
```
matplot (col) (lty) ., () ...
```

```
matplot(xmat, type = 'l', col = 'black')
```



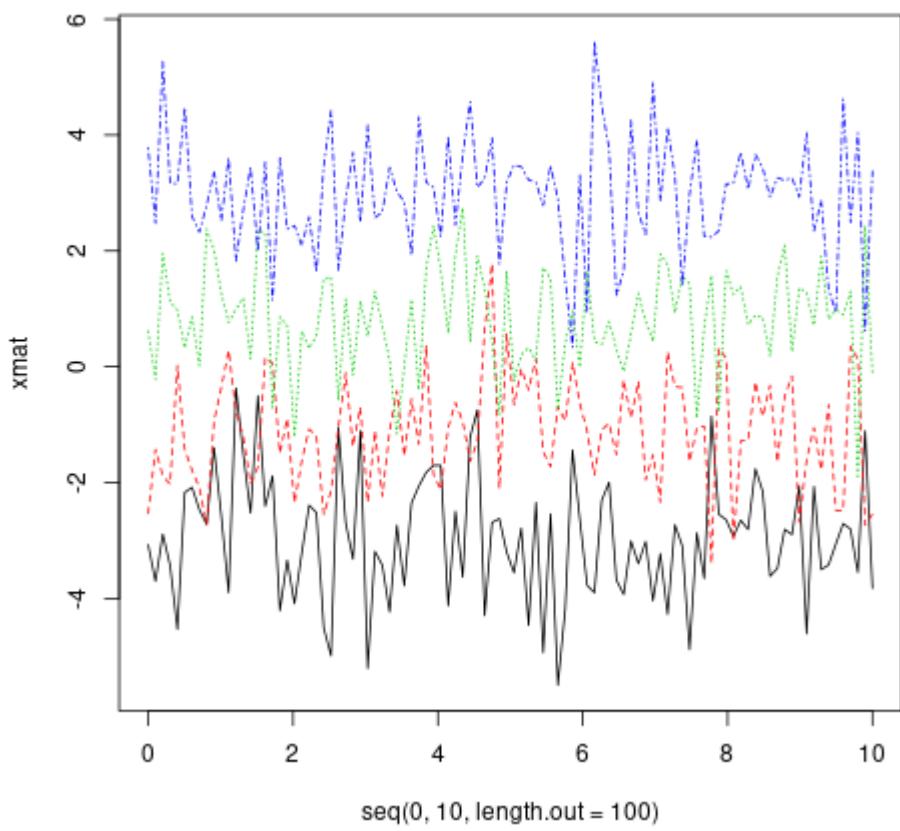
... ( R ).

```
matplot(xmat, type = 'l', col = c('red', 'green', 'blue', 'orange'))
```



```
main , xlab , xmin plot . ?par .
plot , , matplot y x . x y .

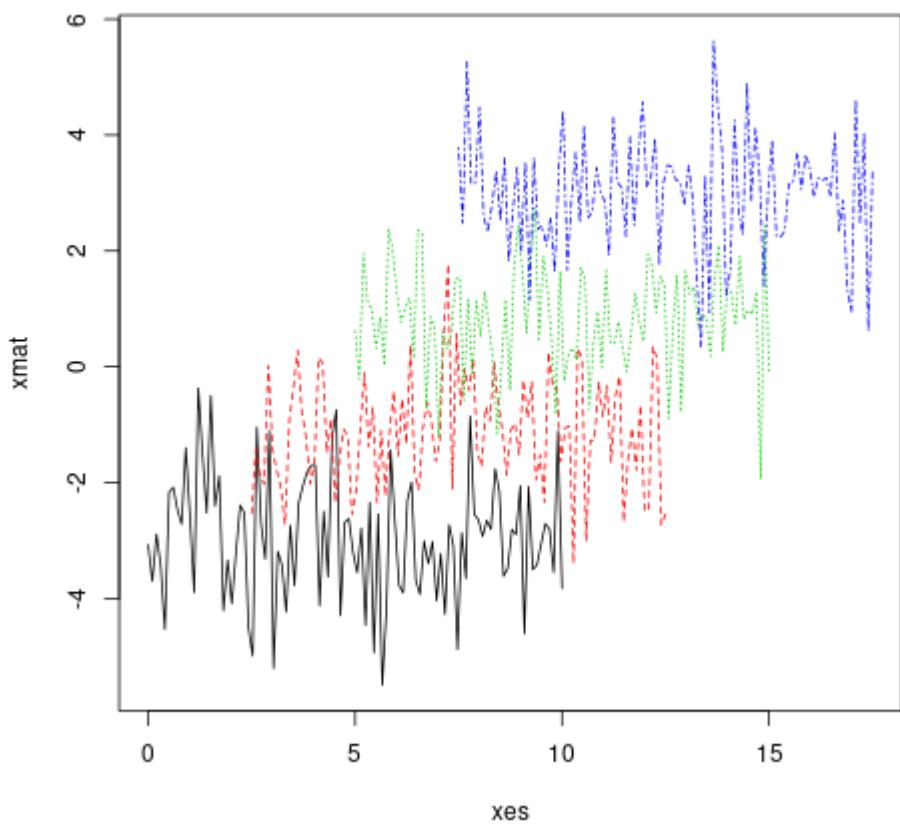
matplot(x = seq(0, 10, length.out = 100), y = xmat, type='l')
```



seq(0, 10, length.out = 100)

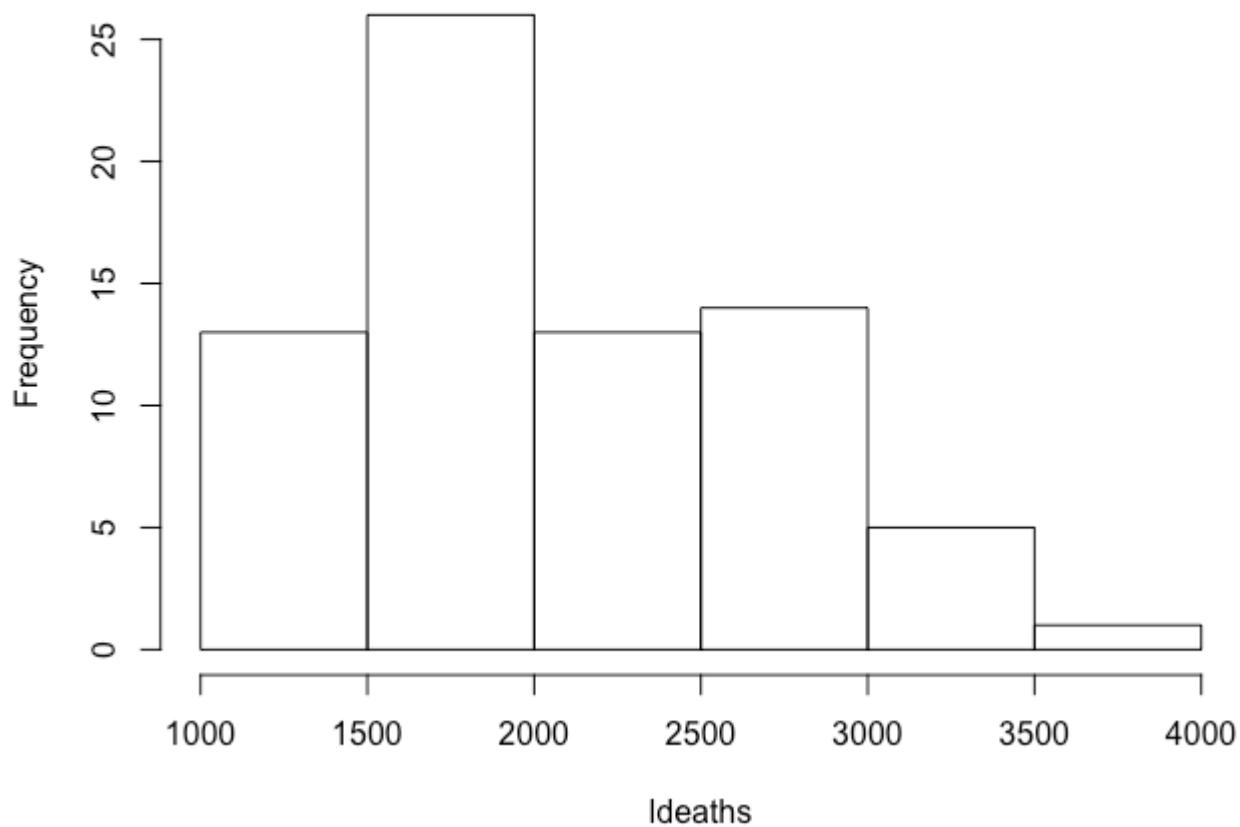
, x y .

```
xes <- cbind(seq(0, 10, length.out = 100),
 seq(2.5, 12.5, length.out = 100),
 seq(5, 15, length.out = 100),
 seq(7.5, 17.5, length.out = 100))
matplot(x = xes, y = xmat, type = 'l')
```



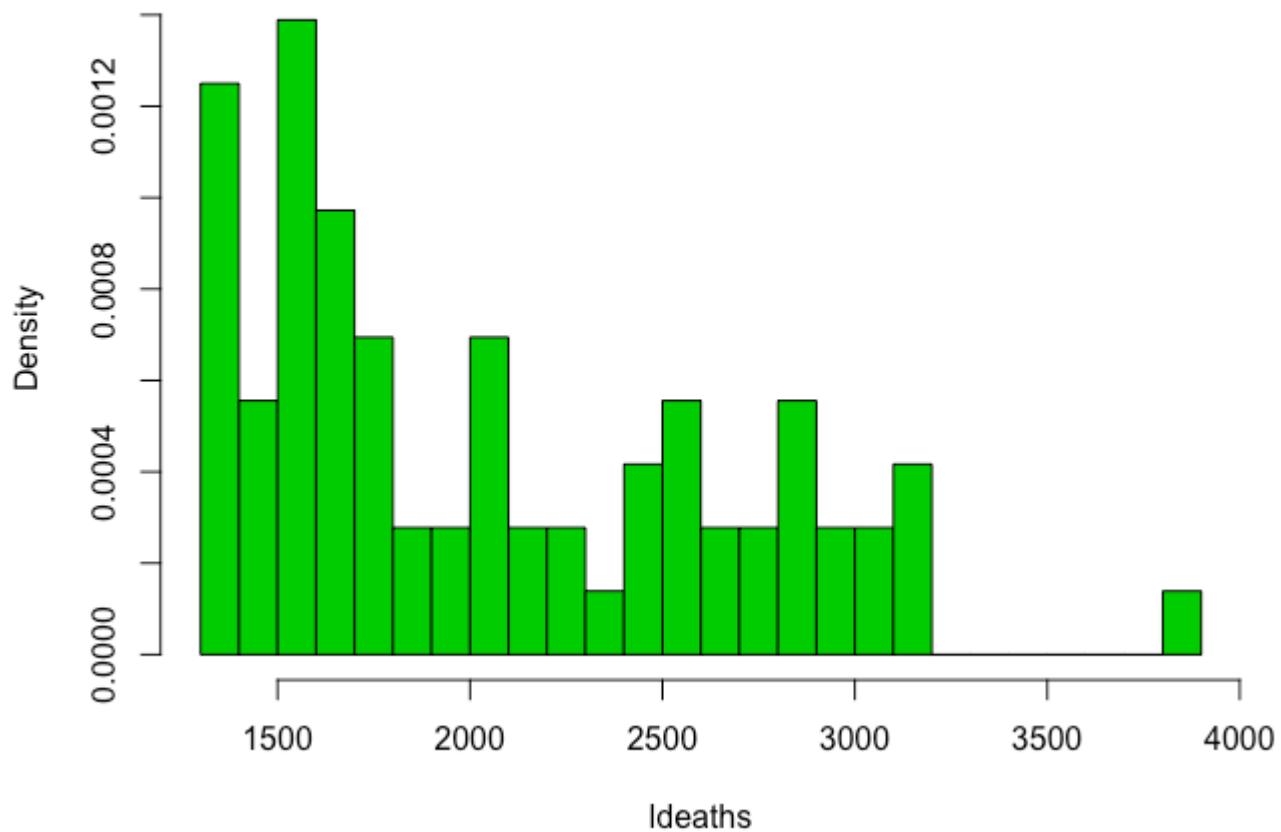
```
hist(ldeaths)
```

## Histogram of ldeaths



```
hist(ldeaths, breaks = 20, freq = F, col = 3)
```

## Histogram of Deaths

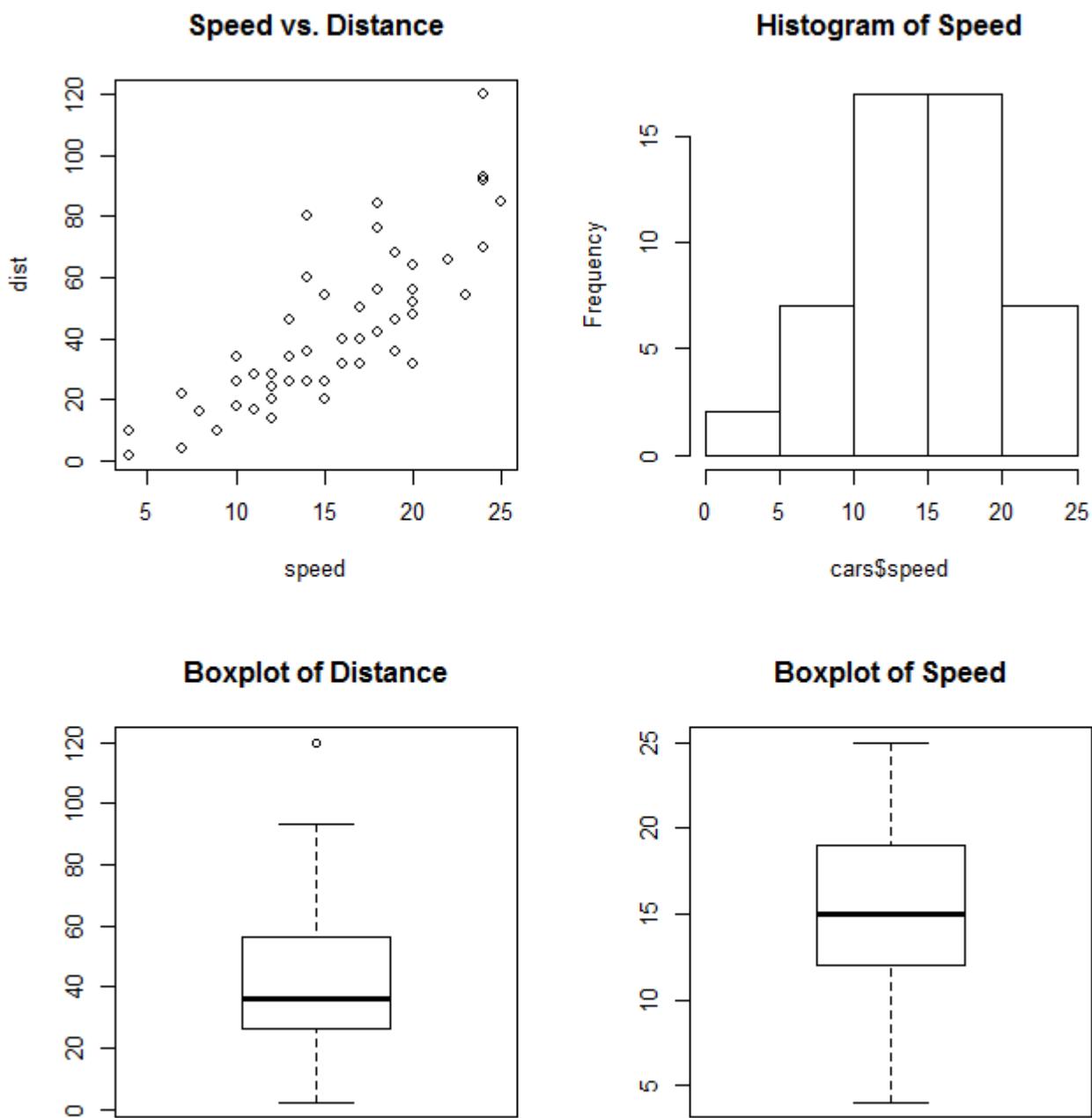


```
(: Scatterplot Barplot) . R par() layout() .
```

```
par()
```

```
par mfrrow mfcoll nrowss ncolsc(nrows, ncols) .
```

```
par(mfrrow=c(2,2))
plot(cars, main="Speed vs. Distance")
hist(cars$speed, main="Histogram of Speed")
boxplot(cars$dist, main="Boxplot of Distance")
boxplot(cars$speed, main="Boxplot of Speed")
```

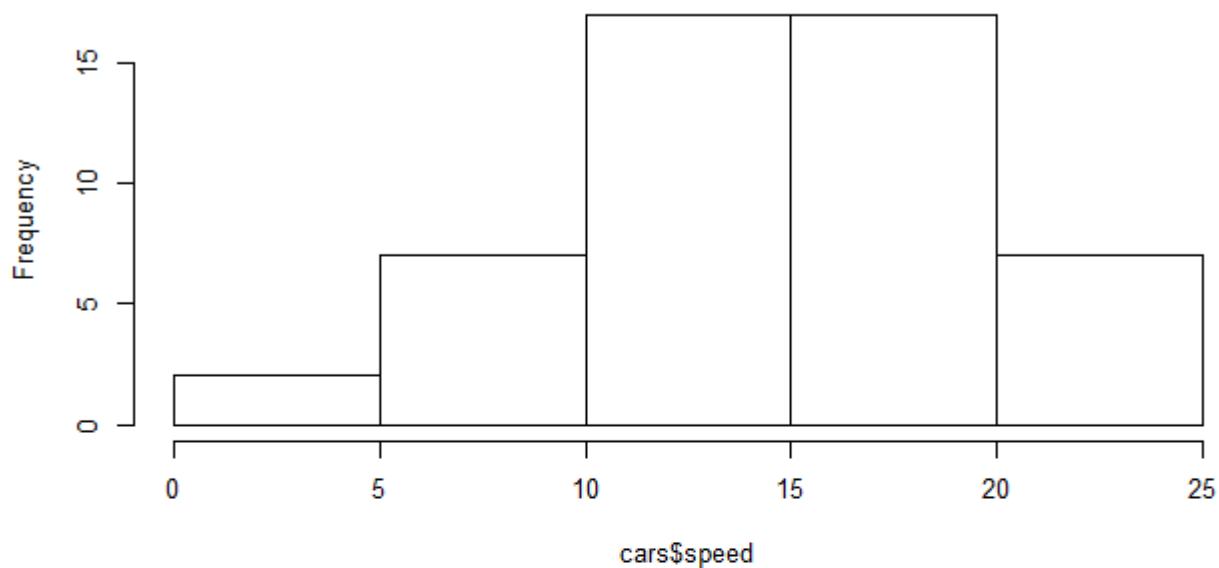


```
layout()
```

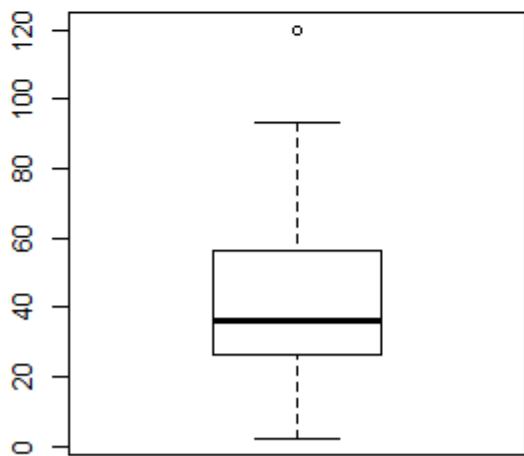
```
layout()
```

```
layout(matrix(c(1,1,2,3), 2,2, byrow=T))
hist(cars$speed, main="Histogram of Speed")
boxplot(cars$dist, main="Boxplot of Distance")
boxplot(cars$speed, main="Boxplot of Speed")
```

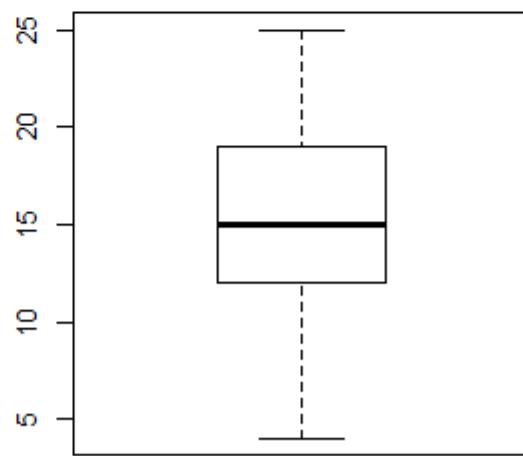
### Histogram of Speed



### Boxplot of Distance



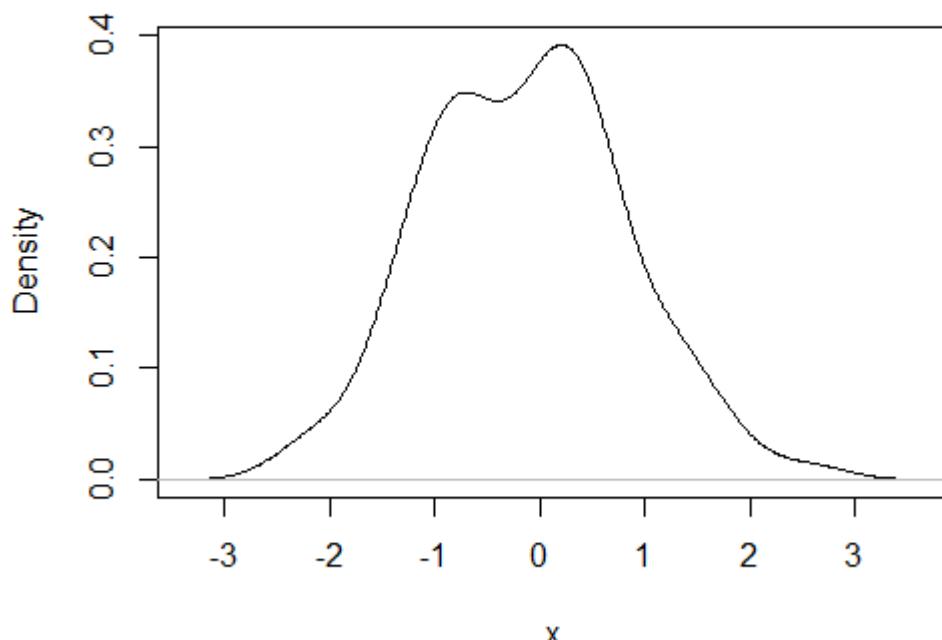
### Boxplot of Speed



```
plot(density(rnorm(100)),main="Normal density",xlab="x")
```

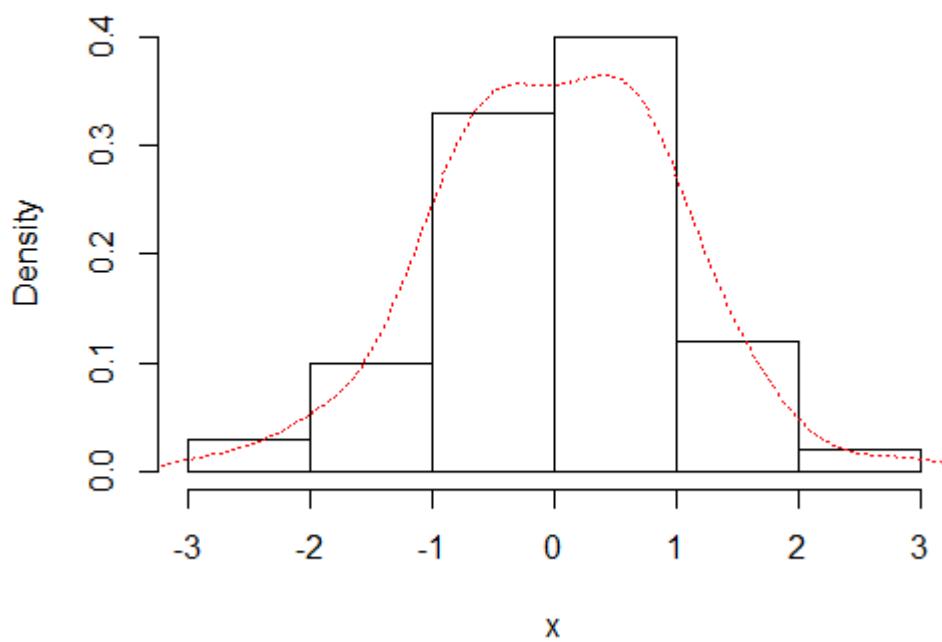
~

## Normal density



```
x=rnorm(100)
hist(x,prob=TRUE,main="Normal density + histogram")
lines(density(x),lty="dotted",col="red")
```

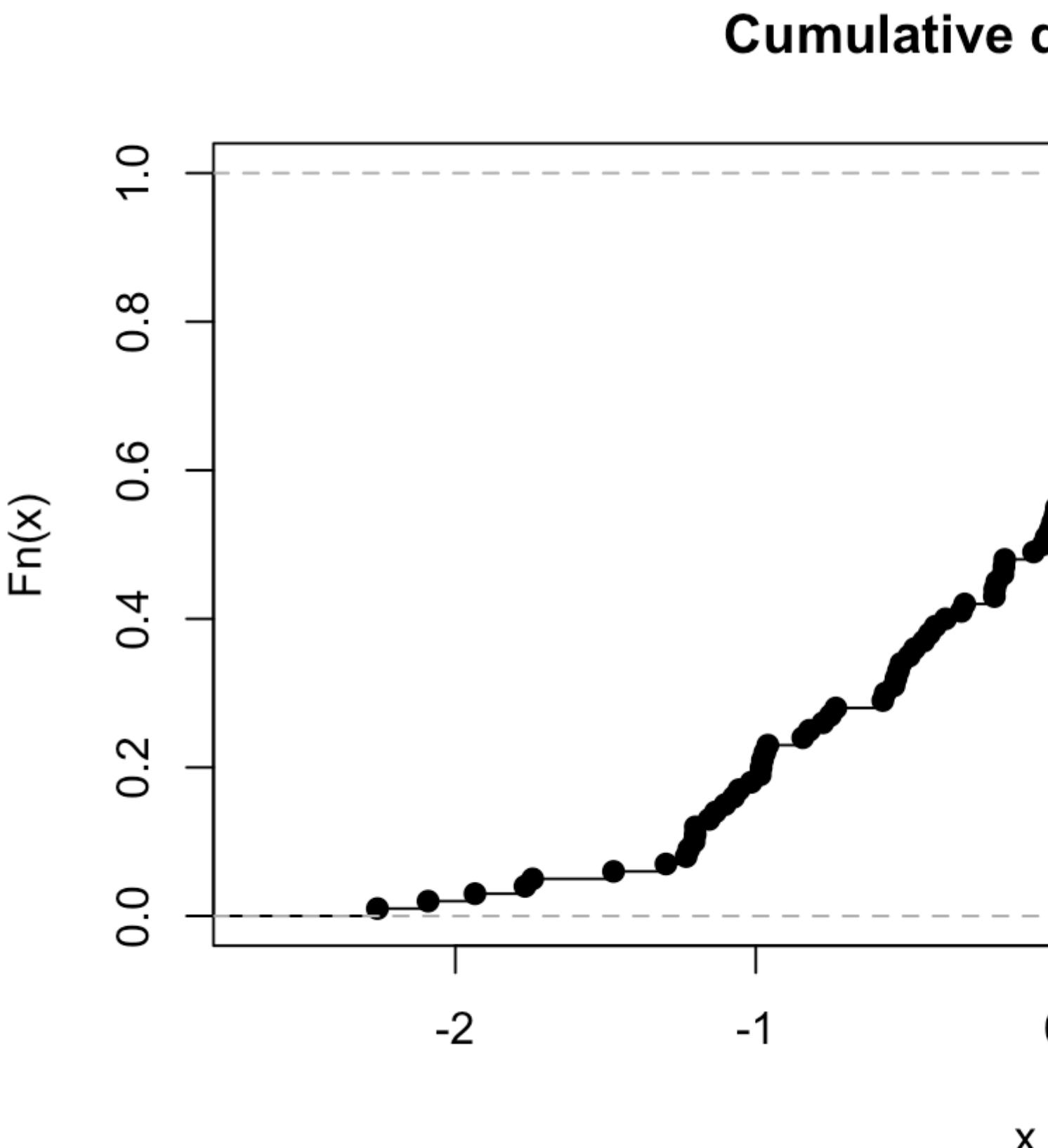
## Normal density + histogram



```
. ecdf()
```

```
plot(ecdf(rnorm(100)),main="Cumulative distribution",xlab="x")
```

~



## R\_Plots

```
x_values <- rnorm(n = 20 , mean = 5 , sd = 8) #20 values generated from Normal(5,8)
y_values <- rbeta(n = 20 , shape1 = 500 , shape2 = 10) #20 values generated from Beta(500,10)
```

```
x_values y_values
```

```
plot(x = x_values, y = y_values, type = "p") #standard scatter-plot
plot(x = x_values, y = y_values, type = "l") # plot with lines
plot(x = x_values, y = y_values, type = "n") # empty plot
```

```
?plot()
```

```
#boxplot is an easy way to see if we have some outliers in the data.
```

```
z<- rbeta(20 , 500 , 10) #generating values from beta distribution
z[c(19 , 20)] <- c(0.97 , 1.05) # replace the two last values with outliers
boxplot(z) # the two points are the outliers of variable z.
```

```
hist(x = x_values) # Histogram for x vector
hist(x = x_values, breaks = 3) #use breaks to set the numbers of bars you want
```

- **Pie\_charts**

```
P <- c(rep('A' , 3) , rep('B' , 10) , rep('C' , 7))
t <- table(P) # this is a frequency matrix of variable P
pie(t) # And this is a visual version of the matrix above
```

: <https://riptutorial.com/ko/r/topic/1377/>

## Examples

```
. : , . ().
```

```
exampleList1 <- list('a', 'b')
exampleList2 <- list(1, 2)
exampleList3 <- list('a', 1, 2)
```

```
str .
```

```
str(exampleList1)
str(exampleList2)
str(exampleList3)
```

```
, , , . [.
```

```
Returns List
exampleList3[1]
exampleList3[1:2]
```

```
[[.
```

```
Returns Character
exampleList3[[1]]
```

```
exampleList4 <- list(
 num = 1:3,
 numeric = 0.5,
 char = c('a', 'b')
)
```

```
exampleList4[['char']]
```

```
$.
```

```
exampleList4$num
```

```
. $.
```

```

exampleList5 <- exampleList4[2:3]

exampleList4$num
c(1, 2, 3)

exampleList5$num
0.5

exampleList5[['num']]
NULL

```

```

Numeric vector
exampleVector1 <- c(12, 13, 14)
Character vector
exampleVector2 <- c("a", "b", "c", "d", "e", "f")
Matrix
exampleMatrix1 <- matrix(rnorm(4), ncol = 2, nrow = 2)
List
exampleList3 <- list('a', 1, 2)

exampleList6 <- list(
 num = exampleVector1,
 char = exampleVector2,
 mat = exampleMatrix1,
 list = exampleList3
)
exampleList6
#$num
#[1] 12 13 14
#
#$char
#[1] "a" "b" "c" "d" "e" "f"
#
#$mat
[,1] [,2]
#[1,] 0.5013050 -1.88801542
#[2,] 0.4295266 0.09751379
#
#$list
#$list[[1]]
#[1] "a"
#
#$list[[2]]
#[1] 1
#
#$list[[3]]
#[1] 2

```

(: ) .list list .

```

l1 <- list(c(1, 2, 3), c("a", "b", "c"))
l1
[[1]]
[1] 1 2 3
##
[[2]]

```

```

[1] "a" "b" "c"

. . . names . .

names(l1)
NULL
names(l1) <- c("vector1", "vector2")
l1
$vector1
[1] 1 2 3
##
$vector2
[1] "a" "b" "c"

```

```

l2 <- list(vec = c(1, 3, 5, 7, 9),
 mat = matrix(data = c(1, 2, 3), nrow = 3))
l2
$vec
[1] 1 3 5 7 9
##
$mat
[,1]
[1,] 1
[2,] 2
[3,] 3
names(l2)
[1] "vec" "mat"

```

"vec" "mat"

R . . . ( )

```

Function example which returns a single element numeric vector
exampleFunction1 <- function(num1, num2){
 result <- num1 + num2
 return(result)
}

Using example function 1
exampleFunction1(1, 2)

Function example which returns a simple numeric vector
exampleFunction2 <- function(num1, num2, multiplier){
 tempResult1 <- num1 + num2
 tempResult2 <- tempResult1 * multiplier
 result <- c(tempResult1, tempResult2)
 return(result)
}

Using example function 2
exampleFunction2(1, 2, 4)

```

R 1 ( ). () ()

```
We will be using mtcars dataset here
Function which returns a result that is supposed to contain multiple type of results
This can be solved by putting the results into a list
exampleFunction3 <- function(dataframe, removeColumn, sumColumn) {
 resultDataFrame <- dataframe[, -removeColumn]
 resultSum <- sum(dataframe[, sumColumn])
 resultList <- list(resultDataFrame, resultSum)
 return(resultList)
}

Using example function 3
exampleResult <- exampleFunction3(mtcars, 2, 4)
exampleResult[[1]]
exampleResult[[2]]
```

vector data.frame , .

. (:n mxn , data.frame data.table ) . . .

```
res <- list(character(0), c("Luzhuang", "Laisu", "Peihui"), character(0),
 c("Anjiangping", "Xinzhai", "Yongfeng"), character(0), character(0),
 c("Puji", "Gaotun", "Banjingcun"), character(0), character(0),
 character(0))
res
```

```
[[1]]
character(0)

[[2]]
[1] "Luzhuang" "Laisu" "Peihui"

[[3]]
character(0)

[[4]]
[1] "Anjiangping" "Xinzhai" "Yongfeng"

[[5]]
character(0)

[[6]]
character(0)

[[7]]
[1] "Puji" "Gaotun" "Banjingcun"

[[8]]
character(0)

[[9]]
character(0)

[[10]]
```

```
character(0)
```

```
res <- sapply(res, function(s) if (length(s) == 0) NA_character_ else paste(s, collapse = ""))
res
```

```
[1] NA "Luzhuang Laisu Peihui" NA
"Anjiangping Xinzhai Yongfeng" NA

[6] NA "Puji Gaotun Banjingcun" NA
NA
```

```
:
```

. , Azure ML R . . .

```
> df
 name height team fun_index title age desc Y
1 Andrea 195 Lazio 97 6 33 eccellente 1
2 Paja 165 Fiorentina 87 6 31 deciso 1
3 Roro 190 Lazio 65 6 28 strano 0
4 Gioele 70 Lazio 100 0 2 simpatico 1
5 Cacio 170 Juventus 81 3 33 duro 0
6 Edola 171 Lazio 72 5 32 svampito 1
7 Salami 175 Inter 75 3 30 doppiopasso 1
8 Braugo 180 Inter 79 5 32 gjn 0
9 Benna 158 Juventus 80 6 28 esaurito 0
10 Riggio 182 Lazio 92 5 31 certezza 1
11 Giordano 185 Roma 79 5 29 buono 1

> number <- "42"
```

```
> paste(df$name[4],"is a",df3$team[4], "supporter.")
[1] "Gioele is a Lazio supporter."
> paste("The answer to THE question is", number)
[1] "The answer to THE question is 42"
```

```
l <- list(df,number)
dataframe_container <- data.frame(out2 = as.integer(serialized(l, connection=NULL)))
```

```
#----- unserialize -----
unser_obj <- unserialize(as.raw(dataframe_container$out2))
#----- taking back the elements-----
df_mod <- unser_obj[1][[1]]
number_mod <- unser_obj[2][[1]]
```

```
> paste(df_mod$name[4],"is a",df_mod$team[4], "supporter.")
[1] "Gioele is a Lazio supporter."
> paste("The answer to THE question is", number_mod)
[1] "The answer to THE question is 42"
```

: <https://riptutorial.com/ko/r/topic/1365/>

# 48:

R . f} .

- `data.table` ,
- `tidyverse`
- 

## Examples

reshape R reshape . ?reshape .

```
create unbalanced longitudinal (panel) data set
set.seed(1234)
df <- data.frame(identifier=rep(1:5, each=3),
 location=rep(c("up", "down", "left", "up", "center"), each=3),
 period=rep(1:3, 5), counts=sample(35, 15, replace=TRUE),
 values=rnorm(15, 5, 10))[-c(4,8,11),]
df

 identifier location period counts values
1 1 up 1 4 9.186478
2 1 up 2 22 6.431116
3 1 up 3 22 6.334104
5 2 down 2 31 6.161130
6 2 down 3 23 6.583062
7 3 left 1 1 6.513467
9 3 left 3 24 5.199980
10 4 up 1 18 6.093998
12 4 up 3 20 7.628488
13 5 center 1 10 9.573291
14 5 center 2 33 9.156725
15 5 center 3 11 5.228851
```

data.frame . , 2 , 3 4 . . : , .

data.frame ,

```
reshape wide on time variable
df.wide <- reshape(df, idvar="identifier", timevar="period",
 v.names=c("values", "counts"), direction="wide")
df.wide

 identifier location values.1 counts.1 values.2 counts.2 values.3 counts.3
1 1 up 9.186478 4 6.431116 22 6.334104 22
5 2 down NA NA 6.161130 31 6.583062 23
7 3 left 6.513467 1 NA NA 5.199980 24
10 4 up 6.093998 18 NA NA 7.628488 20
13 5 center 9.573291 10 9.156725 33 5.228851 11
```

NAs .

```
"v.names" . "drop" . data.frame non-varying / non-id v.names .
```

```
reshape(df, idvar="identifier", timevar="period", direction="wide",
 drop="location")
```

```
df.wide .
```

```
reshape(df.wide, direction="long")
```

```
remove "." separator in df.wide names for counts and values
names(df.wide)[grep("\\.", names(df.wide))] <-
 gsub("\\.", "", names(df.wide)[grep("\\.", names(df.wide))])
```

```
reshape long reshape "varying" . . .
```

```
reshape(df.wide, idvar="identifier",
 varying=list(c(3,5,7), c(4,6,8)), direction="long")
```

```
long reshaping "v.names" .
```

```
"9" "" reshape time .
```

	[cm]	[yr]
178	20	
174		45
182	31	

[cm]	178	
[cm]	174	
[cm]	182	
[yr]	20	
[yr]	45	

	[yr]	31
--	------	----

Base R . mtcars . . .

```
mtcars # shows the dataset
data <- data.frame(observation=row.names(mtcars), mtcars)
```

## R

R long : stack() unstack() .

```
long <- stack(data)
long # this shows the long format
wide <- unstack(long)
wide # this shows the wide format
```

## tidyR

gather() wide long spread() long wide .

```
library(tidyR)
long <- gather(data, variable, value, 2:12) # where variable is the name of the
variable column, value indicates the name of the value column and 2:12 refers to
the columns to be converted.
long # shows the long result
wide <- spread(long, variable, value)
wide # shows the wide result (~data)
```

## data.table

data.table reshape2 melt() , dcast() .

```
library(data.table)
long <- melt(data, 'observation', 2:12, 'variable', 'value')
long # shows the long result
wide <- dcast(long, observation ~ variable)
wide # shows the wide result (~data)
```

: <https://riptutorial.com/ko/r/topic/2904/>-----

# 49:

## Examples

### tbl\_df

```
tbl_df (tibble diff) . tibble .
```

```
as_data_frame tbl_df .
```

```
library(tibble)
mtcars_tbl <- as_data_frame(mtcars)
```

```
data.frames tbl_dfs .
```

```
A tibble: 32 x 11
... with 22 more rows
 mpg cyl disp hp drat wt qsec vs am gear carb
 * <dbl>
1 21.0 6 160.0 110 3.90 2.620 16.46 0 1 4 4
2 21.0 6 160.0 110 3.90 2.875 17.02 0 1 4 4
3 22.8 4 108.0 93 3.85 2.320 18.61 1 1 4 1
4 21.4 6 258.0 110 3.08 3.215 19.44 1 0 3 1
5 18.7 8 360.0 175 3.15 3.440 17.02 0 0 3 2
6 18.1 6 225.0 105 2.76 3.460 20.22 1 0 3 1
7 14.3 8 360.0 245 3.21 3.570 15.84 0 0 3 4
8 24.4 4 146.7 62 3.69 3.190 20.00 1 0 4 2
9 22.8 4 140.8 95 3.92 3.150 22.90 1 0 4 2
10 19.2 6 167.6 123 3.92 3.440 18.30 1 0 4 4
... with 22 more rows
```

- ( 32 x 11 )
- ( dbl )
- .( options(tibble.print\_max = [number]) options(tibble.print\_max = [number]) ).

```
dplyr group_by() tbl_dfs .
```

:

**tidyverse** ?

```
tidyverse R Hadley / Rstudio . tidyverse . .
```

```
tidyverse API . tidyverse .
```

```
tidyverse R for Data Science. tidyverse tidyverse .
```

```
())
```

?

R

```
install.package("tidyverse")
library("tidyverse")
```

/ . /

—  
?

:

- `ggplot2` : SO\_doc
- `dplyr` : ( Rcpp ) SO\_doc
- `tidyr` : SO\_doc
- `readr` : .
- `purrr` : JS underscore.js, lodash lazy.js R
- `tibble` : .
- `magrittr` : SO\_doc

:

- `hms` :
- `stringr` :
- `lubridate` : / SO\_doc
- `forcats` : .

:

- `DBI` : R (DBMS) .
- : SPSS, SAS Stata . SO\_doc
- `httr` : httr API .
- `jsonlite` : JSON
- `readxl` : read.xls .xlsx SO\_doc
- `rvest` : rvest SO\_doc .
- `xml2` : XML

:

- `modelr` :
- : .

tidyverse .

- `knitr` : API . SO\_docs : 1 , 2

- rmarkdown : Rstudio . SO\_docs : [1](#) , [2](#) , [3](#) , [4](#)

: <https://riptutorial.com/ko/r/topic/1395/>

## Examples

**R** ,

```
sample(5)
[1] 4 5 3 1 2
```

:

```
sample(10:15)
[1] 11 15 12 10 14 13
```

pracma .

```
randperm(a, k)
Generates one random permutation of k of the elements a, if a is a vector,
or of 1:a if a is a single integer.
a: integer or numeric vector of some length n.
k: integer, smaller as a or length(a).

Examples
library(pracma)
randperm(1:10, 3)
[1] 3 7 9

randperm(10, 10)
[1] 4 5 10 8 2 7 6 9 3 1

randperm(seq(2, 10, by=2))
[1] 6 4 10 2 8
```

**R** set.seed() . , ( ).

```
> sample(1:10,5)
[1] 6 9 2 7 10
> sample(1:10,5)
[1] 7 6 1 2 10
```

```
> rnorm(5)
[1] 0.4874291 0.7383247 0.5757814 -0.3053884 1.5117812
> rnorm(5)
[1] 0.38984324 -0.62124058 -2.21469989 1.12493092 -0.04493361
```

( 1 ) .

```
> set.seed(1)
```

```
> sample(letters,2)
[1] "g" "j"
> set.seed(1)
> sample(letters,2)
[1] "g" "j"
```

`rexp()` :

```
> set.seed(1)
> rexp(5)
[1] 0.7551818 1.1816428 0.1457067 0.1397953 0.4360686
> set.seed(1)
> rexp(5)
[1] 0.7551818 1.1816428 0.1457067 0.1397953 0.4360686
```

**5 .**

**0 10**

```
runif(5, min=0, max=10)
[1] 2.1724399 8.9209930 6.1969249 9.3303321 2.4054102
```

**0 1**

```
rnorm(5, mean=0, sd=1)
[1] -0.97414402 -0.85722281 -0.08555494 -0.37444299 1.20032409
```

**10 0.5**

```
rbinom(5, size=10, prob=0.5)
[1] 4 3 5 2 3
```

**0.2**

```
rgeom(5, prob=0.2)
[1] 14 8 11 1 3
```

**3 , 10 5**

```
rhyper(5, m=3, n=10, k=5)
[1] 2 0 1 1 1
```

**10 0.8**

```
rnbnom(5, size=10, prob=0.8)
```

```
[1] 3 1 3 4 2
```

## ( ) 2

```
rpois(5, lambda=2)
[1] 2 1 2 3 4
```

## 1.5

```
rexp(5, rate=1.5)
[1] 1.8993303 0.4799358 0.5578280 1.5630711 0.6228000
```

## 0 1

```
rlogis(5, location=0, scale=1)
[1] 0.9498992 -1.0287433 -0.4192311 0.7028510 -1.2095458
```

## 15

```
rchisq(5, df=15)
[1] 14.89209 19.36947 10.27745 19.48376 23.32898
```

## a = 1 b = 0.5

```
rbeta(5, shape1=1, shape2=0.5)
[1] 0.1670306 0.5321586 0.9869520 0.9548993 0.9999737
```

## 3 = 0.5

```
rgamma(5, shape=3, scale=0.5)
[1] 2.2445984 0.7934152 3.2366673 2.2897537 0.8573059
```

## 0 1

```
rcauchy(5, location=0, scale=1)
[1] -0.01285116 -0.38918446 8.71016696 10.60293284 -0.68017185
```

## 0 1 ( )

```
rlnorm(5, meanlog=0, sdlog=1)
[1] 0.8725009 2.9433779 0.3329107 2.5976206 2.8171894
```

## 0.5 1

```
rweibull(5, shape=0.5, scale=1)
[1] 0.337599112 1.307774557 7.233985075 5.840429942 0.005751181
```

## Wilcoxon 10 , 20 .

```
rwilcox(5, 10, 20)
[1] 111 88 93 100 124
```

## 5 3

```
rmultinom(5, size=5, prob=c(0.1,0.1,0.8))
 [,1] [,2] [,3] [,4] [,5]
 [1,] 0 0 1 1 0
 [2,] 2 0 1 1 0
 [3,] 3 5 3 3 5
```

: <https://riptutorial.com/ko/r/topic/1578/>

# 51: - (POSIXct POSIXlt)

R - POSIXct POSIXlt (?DateTimeClasses).

POSIXct .

•

•

## Examples

```
test date-time object
options(digits.secs = 3)
d = as.POSIXct("2016-08-30 14:18:30.58", tz = "UTC")

format(d,"%S") # 00-61 Second as integer
[1] "30"

format(d,"%OS") # 00-60.99... Second as fractional
[1] "30.579"

format(d,"%M") # 00-59 Minute
[1] "18"

format(d,"%H") # 00-23 Hours
[1] "14"

format(d,"%I") # 01-12 Hours
[1] "02"

format(d,"%p") # AM/PM Indicator
[1] "PM"

format(d,"%z") # Signed offset
[1] "+0000"

format(d,"%Z") # Time Zone Abbreviation
[1] "UTC"
```

?strptime .

/

POSIXct POSIXlt - . " " .

```
as.POSIXct("11:38", # time string
 format = "%H:%M") # formatting string
```

```
[1] "2016-07-21 11:38:00 CDT"
strptime("11:38",
 format = "%H:%M") # identical, but makes a POSIXlt object
[1] "2016-07-21 11:38:00 CDT"

as.POSIXct("11 AM",
 format = "%I %p")
[1] "2016-07-21 11:00:00 CDT"
```

```
as.POSIXct("11:38:22", # time string without timezone
 format = "%H:%M:%S",
 tz = "America/New_York") # set time zone
[1] "2016-07-21 11:38:22 EDT"

as.POSIXct("2016-07-21 00:00:00",
 format = "%F %T") # shortcut tokens for "%Y-%m-%d" and "%H:%M:%S"
```

?strptime .

---

- .
- **time** , . 0.
- **string** **tz** .
  
- **tz** .
  - CST "CST6CDT" "America/Chicago" .
- .
  - R : OlsonNames()
  - R : system("cat \$R\_HOME/share/zoneinfo/zone.tab")
- **IANA (Internet Assigned Numbers Authority)**
  - **tz** ([Wikipedia](#))
  - **IANA TZ (2016e)**

**POSIXct** . **POSIXct** . .

```
adding/subtracting times - 60 seconds
as.POSIXct("2016-01-01") + 60
[1] "2016-01-01 00:01:00 AEDT"

adding 3 hours, 14 minutes, 15 seconds
as.POSIXct("2016-01-01") + ((3 * 60 * 60) + (14 * 60) + 15)
[1] "2016-01-01 03:14:15 AEDT"
```

, as.difftime **datetime** . . :

```
as.POSIXct("2016-01-01") +
```

```
as.difftime(3, units="hours") +
as.difftime(14, units="mins") +
as.difftime(15, units="secs")
[1] "2016-01-01 03:14:15 AEDT"
```

/ difftime() , , , .

```
using POSIXct objects
difftime(
 as.POSIXct("2016-01-01 12:00:00"),
 as.POSIXct("2016-01-01 11:59:59"),
 unit = "secs")
Time difference of 1 secs
```

- seq.POSIXt() seq .

- (POSIXct POSIXlt) : <https://riptutorial.com/ko/r/topic/9027/-----posixct--posixlt->

# 52:

R, . ?Dates ", " ?DateTimeClasses ", " ?difftime " " . : - .

- **POSIXct**

date-time POSIXct 1970-01-01 00:00:00 UTC **UNIX epoch** . Sys.Time() Sys.Time() .

- **POSIXlt**

, , , , , . strptime .

- **Date**

## /

POSIXct tidyverse UNIX . POSIXlt .

```
origin = as.POSIXct("1970-01-01 00:00:00", format ="%Y-%m-%d %H:%M:%S", tz = "UTC")

origin
[1] "1970-01-01 UTC"

origin + 47
[1] "1970-01-01 00:00:47 UTC"

as.numeric(origin) # At epoch
0

as.numeric(Sys.time()) # Right now (output as of July 21, 2016 at 11:47:37 EDT)
1469116057

posixlt = as.POSIXlt(Sys.time(), format ="%Y-%m-%d %H:%M:%S", tz = "America/Chicago")

Conversion to POISXct
posixct = as.POSIXct(posixlt)
posixct

Accessing components
posixlt$sec # Seconds 0-61
posixlt$min # Minutes 0-59
posixlt$hour # Hour 0-23
posixlt$mdy # Day of the Month 1-31
posixlt$mon # Months after the first of the year 0-11
posixlt$year # Years since 1900.

ct = as.POSIXct("2015-05-25")
lt = as.POSIXlt("2015-05-25")

object.size(ct)
520 bytes
object.size(lt)
```

```
1816 bytes
```

- 
- **data.table** **IDate** **ITime**
- 
- 
- 

## Examples

R , .

```
Sys.Date() # Returns date as a Date object
[1] "2016-07-21"

Sys.time() # Returns date & time at current locale as a POSIXct object
[1] "2016-07-21 10:04:39 CDT"

as.numeric(Sys.time()) # Seconds from UNIX Epoch (1970-01-01 00:00:00 UTC)
[1] 1469113479

Sys.timezone() # Time zone at current location
[1] "Australia/Melbourne"
```

OlsonNames() Olson / IANA .

```
str(OlsonNames())
#> chr [1:589] "Africa/Abidjan" "Africa/Accra" "Africa/Addis_Ababa" "Africa/Algiers"
"Africa/Asmara" "Africa/Asmera" "Africa/Bamako" ...
```

```
eom <- function(x, p=as.POSIXlt(x)) as.Date(modifyList(p, list(mon=p$mon + 1, mday=0)))
```

```
x <- seq(as.POSIXct("2000-12-10"), as.POSIXct("2001-05-10"), by="months")
> data.frame(before=x, after=eom(x))
 before after
1 2000-12-10 2000-12-31
2 2001-01-10 2001-01-31
3 2001-02-10 2001-02-28
4 2001-03-10 2001-03-31
5 2001-04-10 2001-04-30
6 2001-05-10 2001-05-31
```

```
>
```

```
:

> eom('2000-01-01')
[1] "2000-01-31"
```

```
:

date <- as.Date("2017-01-20")

> as.POSIXlt(cut(date, "month"))
[1] "2017-01-01 EST"
```

```
num . mondate .
```

```
:

moveNumOfMonths <- function(date, num) {
 as.Date(mondate(date) + num)
}
```

```
1 :

:
```

```
> moveNumOfMonths("2017-10-30",-1)
[1] "2017-09-30"
```

```
2 :

:
```

```
> moveNumOfMonths("2017-10-30",-2)
[1] "2017-08-30"
```

```
2 :

:
```

```
> moveNumOfMonths("2017-02-28", 2)
[1] "2017-04-30"
```

```
2 2 , 4 .

:
```

```
> moveNumOfMonths("2016-11-30", 2)
[1] "2017-01-31"
> moveNumOfMonths("2017-01-31", -2)
[1] "2016-11-30"
```

```
11 30 .
```

```
> moveNumOfMonths("2017-01-30", -2)
[1] "2016-11-30"
> moveNumOfMonths("2016-11-30", 2)
[1] "2017-01-31"
```

1 31 11 2 1 .

: <https://riptutorial.com/ko/r/topic/1157/>

## 53: /

```
expr "try part" tryCatch . (,) readLines . explicitly return " "() insided
// . AFAIU, (simpleCondition) tryCatch .
"try part" . . finally = <expression> ("try part") .
```

tryCatch

```
tryCatch expr . . , (, return(NA) () warning error ?tryCatch). tryCatch ().
```

"try part" NA y NA . NULL lapply NULL " / " y 3 2 NULL . return , NULL (, ).

""

```
urls , (readLines) :
```

```
Warning message:
In file(con, "r") : cannot open file 'I'm no URL': No such file or directory
```

```
. readLines warn = FALSE , .
```

```
suppressWarnings(readLines(con = url))
```

```
readLines(con = url, warn = FALSE)
```

## Examples

### tryCatch ()

```
URL HTML . () () .
```

tryCatch

```
readUrl <- function(url) {
 out <- tryCatch(
 #####
 # Try part: define the expression(s) you want to "try" #
 #####
 {
 # Just to highlight:
 # If you want to use more than one R expression in the "try part"
 }
)
}
```

```

then you'll have to use curly brackets.
Otherwise, just write the single expression you want to try and

message("This is the 'try' part")
readLines(con = url, warn = FALSE)
},

#####
Condition handler part: define how you want conditions to be handled
#####

Handler when a warning occurs:
warning = function(cond) {
 message(paste("Reading the URL caused a warning:", url))
 message("Here's the original warning message:")
 message(cond)

 # Choose a return value when such a type of condition occurs
 return(NULL)
}

Handler when an error occurs:
error = function(cond) {
 message(paste("This seems to be an invalid URL:", url))
 message("Here's the original error message:")
 message(cond)

 # Choose a return value when such a type of condition occurs
 return(NA)
}

#####

finally = {
 message(paste("Processed URL:", url))
 message("Some message at the end\n")
}

}

return(out)
}

```

## URL URL

```

urls <- c(
 "http://stat.ethz.ch/R-manual/R-devel/library/base/html/connections.html",
 "http://en.wikipedia.org/wiki/Xz",
 "I'm no URL"
)

y <- lapply(urls, readUrl)
Processed URL: http://stat.ethz.ch/R-manual/R-devel/library/base/html/connections.html
Some message at the end
#

```

```
Processed URL: http://en.wikipedia.org/wiki/Xz
Some message at the end
#
URL does not seem to exist: I'm no URL
Here's the original error message:
cannot open the connection
Processed URL: I'm no URL
Some message at the end
#
Warning message:
In file(con, "r") : cannot open file 'I'm no URL': No such file or directory
```

```
length(y)
[1] 3

head(y[[1]])
[1] "<!DOCTYPE html PUBLIC "-//W3C//DTD HTML 4.01 Transitional//EN">"
[2] "<html><head><title>R: Functions to Manipulate Connections</title>"
[3] "<meta http-equiv=\"Content-Type\" content=\"text/html; charset=utf-8\">"
[4] "<link rel=\"stylesheet\" type=\"text/css\" href=\"R.css\">"
[5] "</head><body>"
[6] ""

y[[3]]
[1] NA
```

/ : <https://riptutorial.com/ko/r/topic/4060/>----

# 54:

## Examples

```
R
```

```
a
```

```
a <- matrix(1:9, 3, 3)
```

```
?
```

```
> class(a)
[1] "matrix"
```

```
:
```

```
> a %*% t(a)
 [,1] [,2] [,3]
[1,] 66 78 90
[2,] 78 93 108
[3,] 90 108 126
```

```
a ?
```

```
> dim(a)
[1] 3 3
> nrow(a)
[1] 3
> ncol(a)
[2] 3
```

```
head , tail str .
```

```
> head(a, 1)
 [,1] [,2] [,3]
[1,] 1 4 7
> tail(a, 1)
 [,1] [,2] [,3]
[3,] 3 6 9
> str(a)
int [1:3, 1:3] 1 2 3 4 5 6 7 8 9
```

```
(:) . str . : a
```

```
a <- c(a)
```

```
?
```

```
> class(a)
[1] "integer"
```

, a . .

```
> dim(a)
NULL
```

, : .

```
> length(a)
[1] 9
```

: .

```
> class(a * 1.0)
[1] "numeric"
```

data.frames .

```
a <- as.data.frame(a)
names(a) <- c("var1", "var2", "var3")
```

: .

```
> names(a)
[1] "var1" "var2" "var3"
```

R .

: <https://riptutorial.com/ko/r/topic/3565/>

# 55:

( ).

```
TRUE , FALSE NA . . T F R TRUE FALSE T F
```

## Examples

```
. (: ! , | , & , xor()) (&& , ||) . if cond .
```

Logical Operators		
!		!
&	()	x y
&&	( )	x && y
	()	x
	( )	x
xor	() OR	xor (x, y)

```
|| TRUE . . && FALSE FALSE .
```

```
> x <- 5
> x > 6 || stop("X is too small")
Error: X is too small
> x > 3 || stop("X is too small")
[1] TRUE
```

```
is.logical() .
```

```
as.logical() .
```

```
> x <- 2
> z <- x > 4
> z
[1] FALSE
> class(x)
[1] "numeric"
> as.logical(2)
[1] TRUE
```

```
as.numeric() double . NA NA NA .
```

## NAs

```
> TRUE & NA
[1] NA
> FALSE & NA
[1] FALSE
> TRUE || NA
[1] TRUE
> FALSE || NA
[1] NA
```

: <https://riptutorial.com/ko/r/topic/9016/>

# 56:

```
NA .
```

```
NA (NA) . (: sqrt(-1)) NaN () .
```

## Examples

```
anyNA . is.na .
```

```
vec <- c(1, 2, 3, NA, 5)

anyNA(vec)
[1] TRUE
is.na(vec)
[1] FALSE FALSE FALSE TRUE FALSE
```

```
is.na (FALSE = 0, TRUE = 1).
```

```
sum(is.na(vec))
[1] 1
```

```
colSums colSums colSums is.na .
```

```
colSums(is.na(airquality))
Ozone Solar.R Wind Temp Month Day
37 7 0 0 0 0
```

```
naniar (github CRAN)
```

## NA

```
read.* read.* R "NA" . NA . (.), (-) (:empty) NA . read.* na.strings R / NA
na.strings .
```

```
read.csv("name_of_csv_file.csv", na.strings = "-")
```

```
NA .
```

```
read.csv('missing.csv', na.strings = c('.','-'))
```

```
NA write.csv na .
```

## NAs

```
NA logical .
```

```
class(NA)
#[1] "logical"
```

NA .

```
x <- c(1, NA, 1)
class(x[2])
#[1] "numeric"
```

NA NA\_character\_ , NA\_integer\_ , NA\_real\_ NA\_complex\_ . NA\_integer\_ NA\_integer\_ . Date :

```
class(Sys.Date() [NA_integer_])
[1] "Date"
```

## TRUE / FALSE / NA

NA NA NA . , NA OR TRUE TRUE TRUE NA OR FALSE NA NA TRUE FALSE

```
NA | TRUE
[1] TRUE
TRUE | TRUE is TRUE and FALSE | TRUE is also TRUE.
```

```
NA | FALSE
[1] NA
TRUE | FALSE is TRUE but FALSE | FALSE is FALSE.
```

```
NA & TRUE
[1] NA
TRUE & TRUE is TRUE but FALSE & TRUE is FALSE.
```

```
NA & FALSE
[1] FALSE
TRUE & FALSE is FALSE and FALSE & FALSE is also FALSE.
```

NA .

```
df <- data.frame(v1=0:9,
 v2=c(rep(1:2, each=4), NA, NA),
 v3=c(NA, letters[2:10]))
```

```
df[df$v2 == 1 & !is.na(df$v2),]
```

```
v1 v2 v3
```

```
#1 0 1 <NA>
```

```
#2 1 1 b
```

```
#3 2 1 c
```

```
#4 3 1 d
```

```
df[df$v2 == 1,]
```

```
v1 v2 v3
```

```
#1 0 1 <NA>
```

```
#2 1 1 b
```

```
#3 2 1 c
```

```
#4 3 1 d
```

```
#NA NA NA <NA>
```

```
#NA.1 NA NA <NA>
```

NA . SPSS 99 .

```
num.vec <- c(1, 2, 3, 99, 5)
num.vec
[1] 1 2 3 99 5
```

NA .

```
num.vec[num.vec == 99] <- NA
```

```
is.na<- . (?is.na) .
```

```
is.na<- . . .
```

```
is.na(num.vec) <- num.vec == 99
```

```
num.vec
[1] 1 2 3 NA 5
```

```
num.vec[!is.na(num.vec)]
num.vec[complete.cases(num.vec)]
na.omit(num.vec)
[1] 1 2 3 5
```

```
mean(num.vec) # returns: [1] NA
```

na.rm NA .

```
mean(num.vec, na.rm = TRUE) # returns: [1] 2.75
an alternative to using 'na.rm = TRUE':
mean(num.vec[!is.na(num.vec)]) # returns: [1] 2.75
```

lm R na.action . na.omit options(na.action = 'na.exclude') R .

na.action na.action . :

```
lm(y2 ~ y1, data = anscombe, na.action = 'na.exclude')
```

: <https://riptutorial.com/ko/r/topic/3388/>

# 57:

```
sub gsub .
```

## Examples

```
pattern . replacement .
```

```
", forename" "forename surname" .
```

```
library(randomNames)
set.seed(1)

strings <- randomNames(5)
strings
[1] "Sigg, Zachary" "Holt, Jake" "Ortega, Sandra" "De La Torre,
Nichole"
[5] "Perkins, Donovan"

sub("^(.+),\\s(.+)$", "\\2 \\1", strings)
[1] "Zachary Sigg" "Jake Holt" "Sandra Ortega" "Nichole De La Torre"
[5] "Donovan Perkins"
```

```
sub("^(.+),\\s(.+)", "\\1", strings)
[1] "Sigg" "Holt" "Ortega" "De La Torre" "Perkins"
```

```
(). :
```

```
2,14,14,14,19
```

```
:
```

```
2,14,19
```

```
gsub .
```

```
gsub("(\\d+) (,\\1)+", "\\1", "2,14,14,14,19")
[1] "2,14,19"
```

```
,
```

```
> gsub("(\\d+) (,\\1)+", "\\1", "2,14,14,14,19,20,21")
[1] "2,14,19,20,21"
```

1. (\d+) : (1) ( ) 1. ( \" \') ( \\" ) . \d\ [0-9] .

2. , : :, ( )  
3. \\1: 1 , .

one,two,two,three,four,four,five,six

\d \w . \w , . [a-zA-Z0-9\_] .

```
> gsub("(\\w+)(,\\1)+", "\\1", "one,two,two,three,four,four,five,six")
[1] "one,two,three,four,five,six"
>
```

: <https://riptutorial.com/ko/r/topic/9219/>--

# 58:

R . , . . . - - -

## Examples

Vector1 .

```
set.seed(123)
Vector1 <- rnorm(20)
```

```
set.seed(123)
Vector1[sample(1:length(Vector1), 5)] <- NA
```

is.na Vector .

```
Vector1 <- Vector1[!is.na(Vector1)]
```

Vector1 NA .

NA . . complete.cases .

6 NAs

```
x <- head(airquality)
```

Solar.R NAs 2 .

```
x_no_NA <- x[complete.cases(x),]
```

x\_no\_NA NAs

: <https://riptutorial.com/ko/r/topic/8165/>

# 59:

R . R . R .

## Examples

R

<https://vincentarelbundock.github.io/Rdatasets/datasets.html>

(1888) .

```
library(tidyverse)

swiss %>%
 ggplot(aes(x = Agriculture, y = Fertility,
 color = Catholic > 50)) +
 geom_point() +
 stat_ellipse()
```

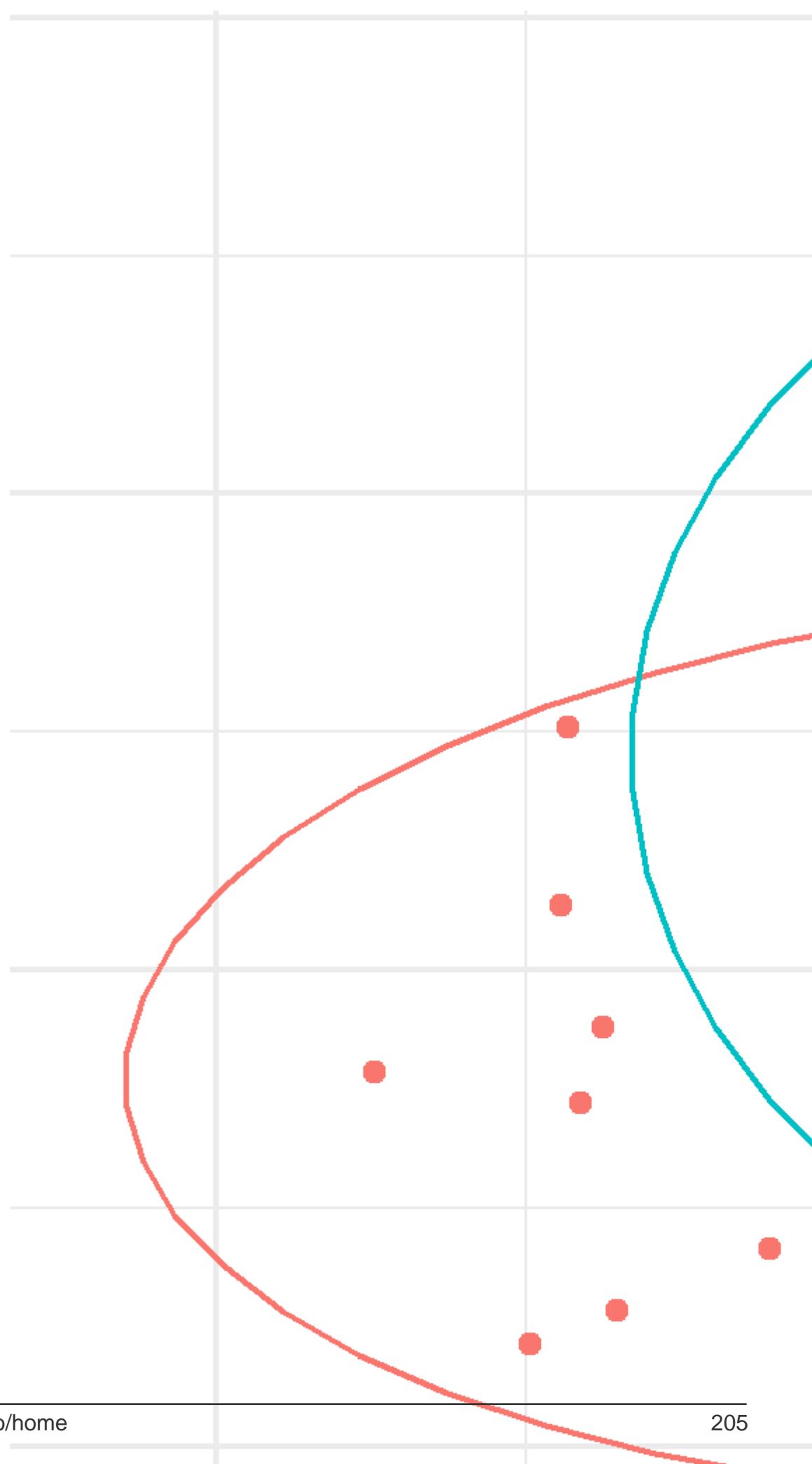
Fertility

110

90

70

50



# 60:

- `data.frame(..., row.names = NULL, check.rows = FALSE, check.names = TRUE, stringsAsFactors = default.stringsAsFactors ())`
- `as.data.frame(x, row.names = NULL, = FALSE, ...)` #
- `as.data.frame(x, ..., stringsAsFactors = default.stringsAsFactors ()) " # S3`
- `as.data.frame(x, row.names = NULL, = FALSE, ..., stringsAsFactors = default.stringsAsFactors ()) "`
- `is.data.frame(x)`

## Examples

### `data.frame`

```
data.frame : . () " ". ().
```

```
data.frame .
```

```
> structure(list(character()), class = "data.frame")
NULL
<0 rows> (or 0-length row.names)
```

```
. data.frame . 3 2 data.frame (a b).
```

```
> structure(list(a = 1:3, b = letters[1:3]), class = "data.frame")
[1] a b
<0 rows> (or 0-length row.names)
```

```
data.frame . 1:3 .
```

```
> structure(list(a = 1:3, b = letters[1:3]), class = "data.frame", row.names = 1:3)
 a b
1 1 a
2 2 b
3 3 c
```

```
3 2 data.frame . nrow() , ncol() dim() .
```

```
> x <- structure(list(a = numeric(3), b = character(3)), class = "data.frame", row.names = 1:3)
> nrow(x)
[1] 3
> ncol(x)
[1] 2
> dim(x)
```

```
[1] 3 2
```

```
R data.frame (structure()) . data.frame() . . . , data.frame()
```

```
> str(data.frame("a a a" = numeric(3), "b-b-b" = character(3)))
'data.frame': 3 obs. of 2 variables:
$ a.a.a: num 0 0 0
$ b.b.b: Factor w/ 1 level ":" 1 1 1
```

```
as.data.frame() . data.frame data.frame . , .
```

```
> m <- matrix(letters[1:9], nrow = 3)
> m
 [,1] [,2] [,3]
[1,] "a" "d" "g"
[2,] "b" "e" "h"
[3,] "c" "f" "i"
```

```
:
```

```
> as.data.frame(m)
V1 V2 V3
1 a d g
2 b e h
3 c f i
> str(as.data.frame(m))
'data.frame': 3 obs. of 3 variables:
$ V1: Factor w/ 3 levels "a","b","c": 1 2 3
$ V2: Factor w/ 3 levels "d","e","f": 1 2 3
$ V3: Factor w/ 3 levels "g","h","i": 1 2 3
```

---

■ [ , ][ , ] \$

- data[rows, columns] matrix
  - ()
- list :
  - data[columns]
  - data[[one\_column]]
- \$ data\$column\_name

```
mtcars .
```

■ data[rows, columns]

```
mtcars [] . .
```

```
get the first row
```

```
mtcars[1,]
get the first five rows
mtcars[1:5,]
```

20

```
get the first column
mtcars[, 1]
get the first, third and fifth columns:
mtcars[, c(1, 3, 5)]
```

```
. mtcars[1,]
```

( )

. data.frame      . numeric    character    .

```
get the mpg column
mtcars[, "mpg"]
get the mpg, cyl, and disp columns
mtcars[, c("mpg", "cyl", "disp")]
```

```
first four rows of the mpg column
mtcars[1:4, "mpg"]

2nd and 5th row of the mpg, cyl, and disp columns
mtcars[c(2, 5), c("mpg", "cyl", "disp")]
```

1

```
multiple columns returns a data frame
class(mtcars[, c("mpg", "cyl")])
[1] "data.frame"
single column returns a vector
class(mtcars[, "mpg"])
[1] "numeric"
```

( ) . drop = FALSE . R " " .

```
class(mtcars[, "mpg", drop = FALSE])
[1] "data.frame"
```

```
 , drop = FALSE

list ., (). [] [[[[

data[columns]
```

```
mtcars["mpg"]
mtcars[c("mpg", "cyl", "disp")]
my_columns <- c("mpg", "cyl", "hp")
mtcars[my_columns]
```

```
data[columns] data[, columns] data.frame list data.frame . matrix data.frame
data.frame .
```

```
When selecting a single column
like a list will return a data frame
class(mtcars["mpg"])
[1] "data.frame"
like a matrix will return a vector
class(mtcars[, "mpg"])
[1] "numeric"
```

```
data[[one_column]]
```

```
data.frame list [[.
```

```
extract a single column by name as a vector
mtcars[["mpg"]]

extract a single column by name as a data frame (as above)
mtcars[["mpg"]]
```

5

5

```
get the column "mpg"
mtcars$mpg
```

5

5

\$ . ( : RStudio) . \$ . . ,

```
my_column <- "mpg"
the below will not work
mtcars$my_column
but these will work
```

```
mtcars[, my_column] # vector
mtcars[my_column] # one-column data frame
mtcars[[my_column]] # vector
```

```
R $. (:) $.
() $
```

```
give you the values of "mpg" column
as "mtcars" has only one column having name starting with "m"
mtcars$m
will give you "NULL"
as "mtcars" has more than one columns having name starting with "d"
mtcars$d
```

```

:
()
```

```
mtcars[1,] # first row
mtcars[-1,] # everything but the first row
mtcars[-(1:10),] # everything except the first 10 rows
```

```
< .
```

```
logical vector indicating TRUE when a row has mpg less than 15
FALSE when a row has mpg >= 15
test <- mtcars$mpg < 15

extract these rows from the data frame
mtcars[test,]
```

```
extract all columns for rows where the value of cyl is 4.
mtcars[mtcars$cyl == 4,]
extract the cyl, mpg, and hp columns where the value of cyl is 4
mtcars[mtcars$cyl == 4, c("cyl", "mpg", "hp")]
```

## data.frames

```
data.frames subset() , transform() , with() within() .
subset() data.frame ().
```

```
subset(mtcars, subset = cyl == 6, select = c("mpg", "hp"))
 mpg hp
Mazda RX4 21.0 110
Mazda RX4 Wag 21.0 110
Hornet 4 Drive 21.4 110
Valiant 18.1 105
Merc 280 19.2 123
Merc 280C 17.8 123
Ferrari Dino 19.7 175
```

```
cyl == 6 mpg hp . [] .
```

```
mtcars[mtcars$cyl == 6, c("mpg", "hp")]
```

```
transform() data.frame . . . mpg2 mpg^2 mtcars data.frame :
```

```
mtcars <- transform(mtcars, mpg2 = mpg^2)
```

~

```
with() within() data.frame $ [] .
```

```
, / airquality data.frame :
```

```
aq <- within(airquality, {
 lOzone <- log(Ozone) # creates new column
 Month <- factor(month.abb[Month]) # changes Month Column
 cTemp <- round((Temp - 32) * 5/9, 1) # creates new column
 S.cT <- Solar.R / cTemp # creates new column
 rm(Day, Temp) # removes columns
})
```

```
. . data.frame . . . : 1 3 .
```

```
df1 <- data.frame(x = 1:3, y = c("a", "b", "c"))
df1
x y
1 1 a
2 2 b
3 3 c
class(df1)
[1] "data.frame"
```

```
df2 <- data.frame(x = c("1", "2", "3"), y = c("a", "b", "c"))
df2
x y
1 1 a
2 2 b
3 3 c
```

```

"X" df1 df2 . str .

str(df1)
'data.frame': 3 obs. of 2 variables:
$ x: int 1 2 3
$ y: Factor w/ 3 levels "a","b","c": 1 2 3
str(df2)
'data.frame': 3 obs. of 2 variables:
$ x: Factor w/ 3 levels "1","2","3": 1 2 3
$ y: Factor w/ 3 levels "a","b","c": 1 2 3

df1 data.frame "x" "y" 3 . "x" () "y" () . stringsAsFactors .

df3 <- data.frame(x = 1:3, y = c("a", "b", "c"), stringsAsFactors = FALSE)
str(df3)
'data.frame': 3 obs. of 2 variables:
$ x: int 1 2 3
$ y: chr "a" "b" "c"

"y" . "" . data.frame .(data.frame(x = 1:3, y = 1:4) .)

, R . :
```

```

mydataframe <- iris
str(mydataframe)
```

## do.call

```

do.call . . .

dataList <- list(1:3, 4:6, 7:9)
dataList
[[1]]
[1] 1 2 3
#
[[2]]
[1] 4 5 6
#
[[3]]
[1] 7 8 9

dataframe <- data.frame(do.call(rbind, dataList))
dataframe
X1 X2 X3
1 1 2 3
2 4 5 6
3 7 8 9
```

```

dataframeList <- list(data.frame(a = 1:2, b = 1:2, c = 1:2),
 data.frame(a = 3:4, b = 3:4, c = 3:4))
dataframeList
[[1]]
```

```

a b c
1 1 1 1
2 2 2 2

[[2]]
a b c
1 3 3 3
2 4 4 4

dataframe <- do.call(rbind, dataList)
dataframe
a b c
1 1 1 1
2 2 2 2
3 3 3 3
4 4 4 4

```

## data.frame

```

. RDBMS data.frame . . .
.
.
.
```

```

bob <- data.frame(jobs = c("scientist", "analyst"),
 pay = c(160000, 100000), age = c(30, 25))
str(bob)

```

```

'data.frame': 2 obs. of 3 variables:
 $ jobs: Factor w/ 2 levels "analyst","scientist": 2 1
 $ pay : num 160000 100000
 $ age : num 30 25

```

```

Convert *all columns* to character
bob[] <- lapply(bob, as.character)
str(bob)

```

```

'data.frame': 2 obs. of 3 variables:
 $ jobs: chr "scientist" "analyst"
 $ pay : chr "160000" "1e+05"
 $ age : chr "30" "25"

```

```

Convert only factor columns to character
bob[] <- lapply(bob, function(x) {
 if (is.factor(x)) x <- as.character(x)
 return(x)
})

```

```
columns rows .
```

```

df <- data.frame(item = c(1:10),
 price_Elasticity = c(-0.57667, 0.03205, -0.04904, 0.10342, 0.04029,

```

```
 0.0742, 0.1669, 0.0313, 0.22204, 0.06158),
totalMargin = c(-145062, 98671, 20576, -56382, 207623, 43463, 1235,
 34521, 146553, -74516))
```

```
price_Elasticity > 0 rows :
```

```
df[df$price_Elasticity > 0,]
```

item	price_Elasticity	totalMargin
2	2	0.03205
4	4	0.10342
5	5	0.04029
6	6	0.07420
7	7	0.16690
8	8	0.03130
9	9	0.22204
10	10	0.06158

```
price_Elasticity > 0 totalMargin > 0 :
```

```
df[df$price_Elasticity > 0 & df$totalMargin > 0,]
```

item	price_Elasticity	totalMargin
2	2	0.03205
5	5	0.04029
6	6	0.07420
7	7	0.16690
8	8	0.03130
9	9	0.22204

: <https://riptutorial.com/ko/r/topic/438/>

R . R ..

## Examples

R

aggregate .

```
aggregate(formula,function,data)
```

:

```
df = data.frame(group=c("Group 1","Group 1","Group 2","Group 2","Group 2"), subgroup =
c("A","A","A","A","B"),value = c(2,2.5,1,2,1.5))

sum, grouping by one column
aggregate(value~group, FUN=sum, data=df)

mean, grouping by one column
aggregate(value~group, FUN=mean, data=df)

sum, grouping by multiple columns
aggregate(value~group+subgroup,FUN=sum,data=df)

custom function, grouping by one column
in this example we want the sum of all values larger than 2 per group.
aggregate(value~group, FUN=function(x) sum(x[x>2]), data=df)
```

:

```
> df = data.frame(group=c("Group 1","Group 1","Group 2","Group 2","Group 2"), subgroup =
c("A","A","A","A","B"),value = c(2,2.5,1,2,1.5))
> print(df)
 group subgroup value
1 Group 1 A 2.0
2 Group 1 A 2.5
3 Group 2 A 1.0
4 Group 2 A 2.0
5 Group 2 B 1.5
>
> # sum, grouping by one column
> aggregate(value~group, FUN=sum, data=df)
 group value
1 Group 1 4.5
2 Group 2 4.5
>
> # mean, grouping by one column
> aggregate(value~group, FUN=mean, data=df)
 group value
```

```

1 Group 1 2.25
2 Group 2 1.50
>
> # sum, grouping by multiple columns
> aggregate(value~group+subgroup,FUN=sum,data=df)
 group subgroup value
1 Group 1 A 4.5
2 Group 2 A 3.0
3 Group 2 B 1.5
>
> # custom function, grouping by one column
> # in this example we want the sum of all values larger than 2 per group.
> aggregate(value~group, FUN=function(x) sum(x[x>2]), data=df)
 group value
1 Group 1 2.5
2 Group 2 0.0

```

## dplyr

dplyr ! group\_by () summarize () . . .

:

```

Aggregating with dplyr
library(dplyr)

df = data.frame(group=c("Group 1","Group 1","Group 2","Group 2","Group 2"),
 subgroup = c("A","A","A","B"),value = c(2,2.5,1,2,1.5))
print(df)

sum, grouping by one column
df %>% group_by(group) %>% summarize(value = sum(value)) %>% as.data.frame()

mean, grouping by one column
df %>% group_by(group) %>% summarize(value = mean(value)) %>% as.data.frame()

sum, grouping by multiple columns
df %>% group_by(group,subgroup) %>% summarize(value = sum(value)) %>% as.data.frame()

custom function, grouping by one column
in this example we want the sum of all values larger than 2 per group.
df %>% group_by(group) %>% summarize(value = sum(value[value>2])) %>% as.data.frame()

```

:

```

> library(dplyr)
>
> df = data.frame(group=c("Group 1","Group 1","Group 2","Group 2","Group 2"),
 subgroup = c("A","A","A","B"),value = c(2,2.5,1,2,1.5))
> print(df)
 group subgroup value
1 Group 1 A 2.0
2 Group 1 A 2.5
3 Group 2 A 1.0
4 Group 2 A 2.0
5 Group 2 B 1.5
>

```

```

> # sum, grouping by one column
> df %>% group_by(group) %>% summarize(value = sum(value)) %>% as.data.frame()
 group value
1 Group 1 4.5
2 Group 2 4.5
>
> # mean, grouping by one column
> df %>% group_by(group) %>% summarize(value = mean(value)) %>% as.data.frame()
 group value
1 Group 1 2.25
2 Group 2 1.50
>
> # sum, grouping by multiple columns
> df %>% group_by(group, subgroup) %>% summarize(value = sum(value)) %>% as.data.frame()
 group subgroup value
1 Group 1 A 4.5
2 Group 2 A 3.0
3 Group 2 B 1.5
>
> # custom function, grouping by one column
> # in this example we want the sum of all values larger than 2 per group.
> df %>% group_by(group) %>% summarize(value = sum(value[value>2])) %>% as.data.frame()
 group value
1 Group 1 2.5
2 Group 2 0.0

```

## data.table

**data.table**    dt[i, j, by] . " *dt* i    j . "dt ,    . list() . () . . () . () .

:

```

Aggregating with data.table
library(data.table)

dt = data.table(group=c("Group 1","Group 1","Group 2","Group 2","Group 2"), subgroup =
c("A","A","A","B","B"),value = c(2,2.5,1,2,1.5))
print(dt)

sum, grouping by one column
dt[,.(value=sum(value)),group]

mean, grouping by one column
dt[,.(value=mean(value)),group]

sum, grouping by multiple columns
dt[,.(value=sum(value)),.(group,subgroup)]

custom function, grouping by one column
in this example we want the sum of all values larger than 2 per group.
dt[,.(value=sum(value[value>2])),group]

```

:

```

> # Aggregating with data.table
> library(data.table)
>

```

```

> dt = data.table(group=c("Group 1","Group 1","Group 2","Group 2","Group 2"),
+ subgroup=c("A","A","A","B"),value = c(2,2.5,1,2,1.5))
> print(dt)
 group subgroup value
1: Group 1 A 2.0
2: Group 1 A 2.5
3: Group 2 A 1.0
4: Group 2 A 2.0
5: Group 2 B 1.5
>
> # sum, grouping by one column
> dt[,.(value=sum(value)),group]
 group value
1: Group 1 4.5
2: Group 2 4.5
>
> # mean, grouping by one column
> dt[,.(value=mean(value)),group]
 group value
1: Group 1 2.25
2: Group 2 1.50
>
> # sum, grouping by multiple columns
> dt[,.(value=sum(value)),.(group,subgroup)]
 group subgroup value
1: Group 1 A 4.5
2: Group 2 A 3.0
3: Group 2 B 1.5
>
> # custom function, grouping by one column
> # in this example we want the sum of all values larger than 2 per group.
> dt[,.(value=sum(value[value>2])),group]
 group value
1: Group 1 2.5
2: Group 2 0.0

```

: <https://riptutorial.com/ko/r/topic/10792/>--

# 62: I/O

- RMySQL
- RODBC

## Examples

### MySQL

RMySQL MariaDB MySQL R .

```
library(RMySQL)

mydb <- dbConnect(MySQL(), user='user', password='password', dbname='dbname', host='127.0.0.1')

queryString <- "SELECT * FROM table1 t1 JOIN table2 t2 on t1.id=t2.id"
query <- dbSendQuery(mydb, queryString)
data <- fetch(query, n=-1) # n=-1 to return all results
```

100,000 . SQL . . :

```
queryString <- "SELECT * FROM table1 limit 100000"
```

### MongoDB

MongoDB R MongoLite .

```
Use MongoLite library:
#install.packages("mongolite")
library(jsonlite)
library(mongolite)

Connect to the database and the desired collection as root:
db <- mongo(collection = "Tweets", db = "TweetCollector", url =
"mongodb://USERNAME:PASSWORD@HOSTNAME")

Read the desired documents i.e. Tweets inside one dataframe:
documents <- db$find(limit = 100000, skip = 0, fields = '{ "_id" : false, "Text" : true }')
```

HOSTNAME PASSWORD USERNAME TweetCollector Tweets . , Text .

. Text (: documents\$Text .

I/O : <https://riptutorial.com/ko/r/topic/5537---i---o>

## 63: R

```
R R . R source(..) R
/ file("stdin") (), stdout() () stderr() () . stdin() R stdin() ..
```

## Examples

R

R

R R (Windows R.exe R Rscript ( Rscript.exe ) ..

R Shebang . #!/usr/bin/env Rscript . Windows ..

R "hist.png" ..

```
#!/usr/bin/env Rscript

User message (\n = end the line)
cat("Input numbers, separated by space:\n")
Read user input as one string (n=1 -> Read only one line)
input <- readLines(file('stdin'), n=1)
Split the string at each space (\s == any space)
input <- strsplit(input, "\s")[[1]]
convert the obtained vector of strings to numbers
input <- as.numeric(input)

Open the output picture file
png("hist.png", width=400, height=300)
Draw the histogram
hist(input)
Close the output file
dev.off()
```

R . Shebang . cat("....\n") . file("stdin") " file("stdin") . ( scan , read.table ,
read.csv , ... ) . . . . . png(.) dev.off() . ( jpeg(.) pdf(.) ) . dev.off() . . .

R

## Linux / Mac

'''''' ..

```
chmod +x PATH/TO/SCRIPT/SCRIPTNAME.R
```

# Windows

```
"C:\Program Files\R-XXXXXXX\bin\Rscript.exe" "%~dp0\XXXXXX.R" %*
```

```
*.txt *.bat *.bat .notepad (Word) "FILENAME.bat" . "" .
```

```
XXX... :
```

- R
- .

```
: "C:\...\Rscript.exe" Windows Rscript.exe . "%~dp0\XXX.R" Rscript R . %~dp0 . %*
```

```
R . R .
```

**littler** R

**littler** ( r ) ( cran ) littler r R (Linux MacOS ).

## **littler**

R :

```
install.packages("littler")
```

```
r .
```

```
You could link to the 'r' binary installed in
'/home/*USER*/R/x86_64-pc-linux-gnu-library/3.4/littler/bin/r'
from '/usr/local/bin' in order to use 'r' for scripting.
```

r , symlink :

```
ln -s /home/*USER*/R/x86_64-pc-linux-gnu-library/3.4/littler/bin/r /usr/local/bin/r
```

## apt-get (Debian, Ubuntu) :

```
sudo apt-get install littler
```

.r littler

**R** . :

```
User message (\n = end the line)
cat("Input numbers, separated by space:\n")
Read user input as one string (n=1 -> Read only one line)
input <- readLines(file('stdin'), n=1)
Split the string at each space (\s == any space)
input <- strsplit(input, "\s")[[1]]
convert the obtained vector of strings to numbers
input <- as.numeric(input)

Open the output picture file
png("hist.png", width=400, height=300)
Draw the histogram
hist(input)
Close the output file
dev.off()
```

**shebang** .hist.r .

```
r hist.r
```

## **littler**

**shebang** littler R

```
#!/usr/bin/env r
```

**R** chmod +X /path/to/script.r .

**R** : <https://riptutorial.com/ko/r/topic/9937/-r---->

# 64:

```
dplyr R "" plyr . dplyr CRAN .
```

```
install.package("dplyr")
```

```
dplyr tbl. dplyr (0.5.0) .
```

- 
- 
- SQLite
- PostgreSQL / Redshift
- MySQL / MariaDB
- 
- MonetDB
- ()

## Examples

### dplyr

```
dplyr R .: data.frame , data.table , database . dplyr dplyr .
```

```
dplyr . (filter , arrange , select , mutate summarise) . group_by .
```

- .
- ( : \$ ).
- . , . .

```
mtcars dplyr . mtcars tbl_df (), rownames rownames_to_column tibble .
```

```
library(dplyr) # This documentation was written using version 0.5.0

mtcars_tbl <- as_data_frame(tibble::rownames_to_column(mtcars, "cars"))

examine the structure of data
head(mtcars_tbl)

A tibble: 6 × 12
cars mpg cyl disp hp drat wt qsec vs am gear carb
<chr> <dbl> <dbl>
#1 Mazda RX4 21.0 6 160 110 3.90 2.620 16.46 0 1 4 4
#2 Mazda RX4 Wag 21.0 6 160 110 3.90 2.875 17.02 0 1 4 4
#3 Datsun 710 22.8 4 108 93 3.85 2.320 18.61 1 1 4 1
#4 Hornet 4 Drive 21.4 6 258 110 3.08 3.215 19.44 1 0 3 1
#5 Hornet Sportabout 18.7 8 360 175 3.15 3.440 17.02 0 0 3 2
#6 Valiant 18.1 6 225 105 2.76 3.460 20.22 1 0 3 1
```

```
filter . data.frame () (TRUE FALSE)
```

4 - cyl:

```
filter(mtcars_tbl, cyl == 4)

A tibble: 11 x 12
cars mpg cyl disp hp drat wt qsec vs am gear carb
<chr> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl>
#1 Datsun 710 22.8 4 108.0 93 3.85 2.320 18.61 1 1 4 1
#2 Merc 240D 24.4 4 146.7 62 3.69 3.190 20.00 1 0 4 2
#3 Merc 230 22.8 4 140.8 95 3.92 3.150 22.90 1 0 4 2
#4 Fiat 128 32.4 4 78.7 66 4.08 2.200 19.47 1 1 4 1
#5 Honda Civic 30.4 4 75.7 52 4.93 1.615 18.52 1 1 4 2
... with 6 more rows
```

. 4 6 - cyl 5 - gear:

```
filter(mtcars_tbl, cyl == 4 | cyl == 6, gear == 5)

A tibble: 3 x 12
cars mpg cyl disp hp drat wt qsec vs am gear carb
<chr> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl>
#1 Porsche 914-2 26.0 4 120.3 91 4.43 2.140 16.7 0 1 5 2
#2 Lotus Europa 30.4 4 95.1 113 3.77 1.513 16.9 1 1 5 2
#3 Ferrari Dino 19.7 6 145.0 175 3.62 2.770 15.5 0 1 5 6
```

```
filter slice.slice . data.frame .
```

6 ~ 9 .

```
slice(mtcars_tbl, 6:9)

A tibble: 4 x 12
cars mpg cyl disp hp drat wt qsec vs am gear carb
<chr> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl>
#1 Valiant 18.1 6 225.0 105 2.76 3.46 20.22 1 0 3 1
#2 Duster 360 14.3 8 360.0 245 3.21 3.57 15.84 0 0 3 4
#3 Merc 240D 24.4 4 146.7 62 3.69 3.19 20.00 1 0 4 2
#4 Merc 230 22.8 4 140.8 95 3.92 3.15 22.90 1 0 4 2
```

:

```
slice(mtcars_tbl, -c(1:5, 10:n()))
```

```
slice(mtcars_tbl, 6:9) slice(mtcars_tbl, 6:9)
```

```
n()
```

---

```
arrange () . (dplyr), data.frame , . .
- hp
```

```
arrange(mtcars_tbl, hp)

A tibble: 32 x 12
cars mpg cyl disp hp drat wt qsec vs am gear carb
<chr> <dbl> <dbl>
#1 Honda Civic 30.4 4 75.7 52 4.93 1.615 18.52 1 1 4 2
#2 Merc 240D 24.4 4 146.7 62 3.69 3.190 20.00 1 0 4 2
#3 Toyota Corolla 33.9 4 71.1 65 4.22 1.835 19.90 1 1 4 1
#4 Fiat 128 32.4 4 78.7 66 4.08 2.200 19.47 1 1 4 1
#5 Fiat X1-9 27.3 4 79.0 66 4.08 1.935 18.90 1 1 4 1
#6 Porsche 914-2 26.0 4 120.3 91 4.43 2.140 16.70 0 1 5 2
... with 26 more rows
```

arrange - mpg - cyl:

```
arrange(mtcars_tbl, desc(mpg), cyl)

A tibble: 32 x 12
cars mpg cyl disp hp drat wt qsec vs am gear carb
<chr> <dbl> <dbl>
#1 Toyota Corolla 33.9 4 71.1 65 4.22 1.835 19.90 1 1 4 1
#2 Fiat 128 32.4 4 78.7 66 4.08 2.200 19.47 1 1 4 1
#3 Honda Civic 30.4 4 75.7 52 4.93 1.615 18.52 1 1 4 2
#4 Lotus Europa 30.4 4 95.1 113 3.77 1.513 16.90 1 1 5 2
#5 Fiat X1-9 27.3 4 79.0 66 4.08 1.935 18.90 1 1 4 1
#6 Porsche 914-2 26.0 4 120.3 91 4.43 2.140 16.70 0 1 5 2
... with 26 more rows
```

select . mpg, disp, wt, qsec vs mtcars\_tbl:

```
select(mtcars_tbl, mpg, disp, wt, qsec, vs)

A tibble: 32 x 5
mpg disp wt qsec vs
<dbl> <dbl> <dbl> <dbl> <dbl>
#1 21.0 160.0 2.620 16.46 0
#2 21.0 160.0 2.875 17.02 0
#3 22.8 108.0 2.320 18.61 1
#4 21.4 258.0 3.215 19.44 1
#5 18.7 360.0 3.440 17.02 0
#6 18.1 225.0 3.460 20.22 1
... with 26 more rows
```

: . cars disp vs carb:

```
select(mtcars_tbl, cars:disp, vs:carb)

A tibble: 32 x 8
cars mpg cyl disp vs am gear carb
<chr> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl>
#1 Mazda RX4 21.0 6 160.0 0 1 4 4
#2 Mazda RX4 Wag 21.0 6 160.0 0 1 4 4
#3 Datsun 710 22.8 4 108.0 1 1 4 1
#4 Hornet 4 Drive 21.4 6 258.0 1 0 3 1
#5 Hornet Sportabout 18.7 8 360.0 0 0 3 2
#6 Valiant 18.1 6 225.0 1 0 3 1
```

```
... with 26 more rows

select(mtcars_tbl, -(hp:qsec))

. , (starts_with() , ends_with() , contains() , matches() , num_range() , one_of() , everything())
select . ?select_helpers ?select ?select .

:select()
```

```
select(mtcars_tbl, cylinders = cyl, displacement = disp)

A tibble: 32 × 2
cylinders displacement
<dbl> <dbl>
#1 6 160.0
#2 6 160.0
#3 4 108.0
#4 6 258.0
#5 8 360.0
#6 6 225.0
... with 26 more rows
```

rename :

```
rename(mtcars_tbl, cylinders = cyl, displacement = disp)

A tibble: 32 × 12
cars mpg cylinders displacement hp drat wt qsec vs
<chr> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl>
#1 Mazda RX4 21.0 6 160.0 110 3.90 2.620 16.46 0
#2 Mazda RX4 Wag 21.0 6 160.0 110 3.90 2.875 17.02 0
#3 Datsun 710 22.8 4 108.0 93 3.85 2.320 18.61 1
#4 Hornet 4 Drive 21.4 6 258.0 110 3.08 3.215 19.44 1
#5 Hornet Sportabout 18.7 8 360.0 175 3.15 3.440 17.02 0
#6 Valiant 18.1 6 225.0 105 2.76 3.460 20.22 1
... with 26 more rows, and 3 more variables: am <dbl>, gear <dbl>, carb <dbl>
```

mutate . dplyr mutate . data.frame .

```
mutate(mtcars_tbl, weight_ton = wt/2, weight_pounds = weight_ton * 2000)

A tibble: 32 × 14
cars mpg cyl disp hp drat wt qsec vs am gear carb
<chr> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl>
#1 Mazda RX4 21.0 6 160.0 110 3.90 2.620 16.46 0 1 4 4
#2 Mazda RX4 Wag 21.0 6 160.0 110 3.90 2.875 17.02 0 1 4 4
#3 Datsun 710 22.8 4 108.0 93 3.85 2.320 18.61 1 1 5 4
#4 Hornet 4 Drive 21.4 6 258.0 110 3.08 3.215 19.44 1 1 5 4
#5 Hornet Sportabout 18.7 8 360.0 175 3.15 3.440 17.02 0 1 5 4
#6 Valiant 18.1 6 225.0 105 2.76 3.460 20.22 1 1 5 4
#7 Fiat 128 5.0 4 94.0 62 3.93 2.320 18.90 1 1 4 4
#8 Honda Civic 18.0 4 96.0 65 3.85 2.320 19.90 1 1 4 4
#9 Toyota Corolla 18.0 4 96.0 65 3.85 2.320 19.90 1 1 4 4
#10 Toyota Corona 18.0 4 120.0 105 3.92 3.150 22.90 1 0 4 4
#11 Dodge Challenger 14.3 8 360.0 175 3.08 3.875 17.82 0 0 3 4
#12 AMC Javelin 14.3 8 360.0 175 3.08 3.875 17.82 0 0 3 4
#13 Fiat X1-9 18.0 2 75.0 62 3.69 3.150 22.90 1 1 2 4
#14 Porsche 914-2 18.0 4 120.0 105 3.85 3.150 22.90 1 1 2 4
#15 Lotus Europa 19.2 4 108.0 95 3.70 3.150 22.90 1 1 2 4
#16 Toyota Celica 17.8 4 108.0 95 3.70 3.150 22.90 1 1 2 4
#17 Ford Mustang 15.2 8 312.0 150 3.69 3.900 18.30 0 0 3 4
#18 Volvo 142D 14.3 4 120.0 105 3.92 3.150 22.90 1 0 4 4
#19 Volvo 164P 14.3 4 120.0 105 3.92 3.150 22.90 1 0 4 4
#20 Toyota Corolla 14.3 4 96.0 65 3.92 3.150 22.90 1 0 4 4
#21 Toyota Corolla 14.3 4 96.0 65 3.92 3.150 22.90 1 0 4 4
#22 Toyota Corolla 14.3 4 96.0 65 3.92 3.150 22.90 1 0 4 4
#23 Toyota Corolla 14.3 4 96.0 65 3.92 3.150 22.90 1 0 4 4
#24 Toyota Corolla 14.3 4 96.0 65 3.92 3.150 22.90 1 0 4 4
#25 Toyota Corolla 14.3 4 96.0 65 3.92 3.150 22.90 1 0 4 4
#26 Toyota Corolla 14.3 4 96.0 65 3.92 3.150 22.90 1 0 4 4
#27 Toyota Corolla 14.3 4 96.0 65 3.92 3.150 22.90 1 0 4 4
#28 Toyota Corolla 14.3 4 96.0 65 3.92 3.150 22.90 1 0 4 4
#29 Toyota Corolla 14.3 4 96.0 65 3.92 3.150 22.90 1 0 4 4
#30 Toyota Corolla 14.3 4 96.0 65 3.92 3.150 22.90 1 0 4 4
#31 Toyota Corolla 14.3 4 96.0 65 3.92 3.150 22.90 1 0 4 4
#32 Toyota Corolla 14.3 4 96.0 65 3.92 3.150 22.90 1 0 4 4
```

```

#3 Datsun 710 22.8 4 108.0 93 3.85 2.320 18.61 1 1 4 1
1.1600 2320
#4 Hornet 4 Drive 21.4 6 258.0 110 3.08 3.215 19.44 1 0 3 1
1.6075 3215
#5 Hornet Sportabout 18.7 8 360.0 175 3.15 3.440 17.02 0 0 3 2
1.7200 3440
#6 Valiant 18.1 6 225.0 105 2.76 3.460 20.22 1 0 3 1
1.7300 3460
... with 26 more rows

```

weight\_ton weight\_pounds . R mutate .

mutate transmute .

```
transmute(mtcars_tbl, weight_ton = wt/2, weight_pounds = weight_ton * 2000)
```

```

A tibble: 32 x 2
weight_ton weight_pounds
<dbl> <dbl>
#1 1.3100 2620
#2 1.4375 2875
#3 1.1600 2320
#4 1.6075 3215
#5 1.7200 3440
#6 1.7300 3460
... with 26 more rows

```

summarise .

mpg disp .

```
summarise(mtcars_tbl, mean_mpg = mean(mpg), sd_mpg = sd(mpg),
 mean_disp = mean(disp), sd_disp = sd(disp))

A tibble: 1 x 4
mean_mpg sd_mpg mean_disp sd_disp
<dbl> <dbl> <dbl> <dbl>
#1 20.09062 6.026948 230.7219 123.9387
```

## group\_by

group\_by .

cyl mpg mean sd :

```
by_cyl <- group_by(mtcars_tbl, cyl)
summarise(by_cyl, mean_mpg = mean(mpg), sd_mpg = sd(mpg))

A tibble: 3 x 3
cyl mean_mpg sd_mpg
<dbl> <dbl> <dbl>
```

```
#1 4 26.66364 4.509828
#2 6 19.74286 1.453567
#3 8 15.10000 2.560048
```

## together

cars hp gear , cyl mpg , gear , mpg > 20 hp > (75)

```
selected <- select(mtcars_tbl, cars:hp, gear)
ordered <- arrange(selected, cyl, desc(mpg))
by_cyl <- group_by(ordered, gear)
filter(by_cyl, mpg > 20, hp > 75)
```

Source: local data frame [9 x 6]

Groups: gear [3]

```
cars mpg cyl disp hp gear
<chr> <dbl> <dbl> <dbl> <dbl> <dbl>
#1 Lotus Europa 30.4 4 95.1 113 5
#2 Porsche 914-2 26.0 4 120.3 91 5
#3 Datsun 710 22.8 4 108.0 93 4
#4 Merc 230 22.8 4 140.8 95 4
#5 Toyota Corona 21.5 4 120.1 97 3
... with 4 more rows
```

```
filter(
 group_by(
 arrange(
 select(
 mtcars_tbl, cars:hp
), cyl, desc(mpg)
), cyl
), mpg > 20, hp > 75
)
```

dplyr pipe %>%

```
mtcars_tbl %>%
 select(cars:hp) %>%
 arrange(cyl, desc(mpg)) %>%
 group_by(cyl) %>%
 filter(mpg > 20, hp > 75)
```

dplyr () summarise\_all()

```
mtcars_tbl %>%
 summarise_all(n_distinct)

A tibble: 1 x 12
cars mpg cyl disp hp drat wt qsec vs am gear carb
<int> <int> <int> <int> <int> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl>
#1 32 25 3 27 22 22 2.9 30 2 2 3 6
```

cyl .

```
mtcars_tbl %>%
 group_by(cyl) %>%
 summarise_all(n_distinct)

A tibble: 3 x 12
cyl cars mpg disp hp drat wt qsec vs am gear carb
<dbl> <int> <int> <int> <int> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl>
#1 4 11 9 11 10 10 11 11 2 2 3 2
#2 6 7 6 5 4 5 6 7 2 2 3 3
#3 8 14 12 11 9 11 13 14 1 2 2 4
```

group\_by . 3 . cyl .

summarise summarise\_at

```
mtcars_tbl %>%
 group_by(cyl) %>%
 summarise_at(c("mpg", "disp", "hp"), mean)

A tibble: 3 x 4
cyl mpg disp hp
<dbl> <dbl> <dbl> <dbl>
#1 4 26.66364 105.1364 82.63636
#2 6 19.74286 183.3143 122.28571
#3 8 15.10000 353.1000 209.21429
```

helper (?select\_helpers)

```
mtcars_tbl %>%
 group_by(cyl) %>%
 summarise_at(c("mpg", "disp", "hp"),
 c("mean", "sd"))
```

fun funs .

```
mtcars_tbl %>%
 group_by(cyl) %>%
 summarise_at(c("mpg", "disp", "hp"),
 funs(mean, sd))

A tibble: 3 x 7
cyl mpg_mean disp_mean hp_mean mpg_sd disp_sd hp_sd
<dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl>
```

```
#1 4 26.66364 105.1364 82.63636 4.509828 26.87159 20.93453
#2 6 19.74286 183.3143 122.28571 1.453567 41.56246 24.26049
#3 8 15.10000 353.1000 209.21429 2.560048 67.77132 50.97689
```

```
mtcars_tbl %>%
 group_by(cyl) %>%
 summarise_at(c("mpg", "disp", "hp"),
 c(Mean = "mean", SD = "sd"))

mtcars_tbl %>%
 group_by(cyl) %>%
 summarise_at(c("mpg", "disp", "hp"),
 funs(Mean = mean, SD = sd))

A tibble: 3 x 7
cyl mpg_Mean disp_Mean hp_Mean mpg_SD disp_SD hp_SD
<dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl>
#1 4 26.66364 105.1364 82.63636 4.509828 26.87159 20.93453
#2 6 19.74286 183.3143 122.28571 1.453567 41.56246 24.26049
#3 8 15.10000 353.1000 209.21429 2.560048 67.77132 50.97689
```

summarise\_if summarise\_if .

cyl numeric mean .

```
mtcars_tbl %>%
 group_by(cyl) %>%
 summarise_if(is.numeric, mean)

A tibble: 3 x 11
cyl mpg disp hp drat wt qsec
<dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl>
#1 4 26.66364 105.1364 82.63636 4.070909 2.285727 19.13727
#2 6 19.74286 183.3143 122.28571 3.585714 3.117143 17.97714
#3 8 15.10000 353.1000 209.21429 3.229286 3.999214 16.77214
... with 4 more variables: vs <dbl>, am <dbl>, gear <dbl>,
carb <dbl>
```

mean mean .

cyl mean :

```
mtcars_tbl %>%
 group_by(cyl) %>%
 summarise_if(function(x) is.numeric(x) & n_distinct(x) > 6, mean)

A tibble: 3 x 7
cyl mpg disp hp drat wt qsec
<dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl>
#1 4 26.66364 105.1364 82.63636 4.070909 2.285727 19.13727
#2 6 19.74286 183.3143 122.28571 3.585714 3.117143 17.97714
#3 8 15.10000 353.1000 209.21429 3.229286 3.999214 16.77214
```

(

dplyr::filter() -

```
dplyr::filter(iris, Sepal.Length > 7)
Sepal.Length Sepal.Width Petal.Length Petal.Width Species
1 7.1 3.0 5.9 2.1 virginica
2 7.6 3.0 6.6 2.1 virginica
3 7.3 2.9 6.3 1.8 virginica
4 7.2 3.6 6.1 2.5 virginica
5 7.7 3.8 6.7 2.2 virginica
6 7.7 2.6 6.9 2.3 virginica
7 7.7 2.8 6.7 2.0 virginica
8 7.2 3.2 6.0 1.8 virginica
9 7.2 3.0 5.8 1.6 virginica
10 7.4 2.8 6.1 1.9 virginica
11 7.9 3.8 6.4 2.0 virginica
12 7.7 3.0 6.1 2.3 virginica
```

dplyr::distinct() -

```
distinct(iris, Sepal.Length, .keep_all = TRUE)
Sepal.Length Sepal.Width Petal.Length Petal.Width Species
1 5.1 3.5 1.4 0.2 setosa
2 4.9 3.0 1.4 0.2 setosa
3 4.7 3.2 1.3 0.2 setosa
4 4.6 3.1 1.5 0.2 setosa
5 5.0 3.6 1.4 0.2 setosa
6 5.4 3.9 1.7 0.4 setosa
7 4.4 2.9 1.4 0.2 setosa
8 4.8 3.4 1.6 0.2 setosa
9 4.3 3.0 1.1 0.1 setosa
10 5.8 4.0 1.2 0.2 setosa
11 5.7 4.4 1.5 0.4 setosa
12 5.2 3.5 1.5 0.2 setosa
13 5.5 4.2 1.4 0.2 setosa
14 4.5 2.3 1.3 0.3 setosa
15 5.3 3.7 1.5 0.2 setosa
16 7.0 3.2 4.7 1.4 versicolor
17 6.4 3.2 4.5 1.5 versicolor
18 6.9 3.1 4.9 1.5 versicolor
19 6.5 2.8 4.6 1.5 versicolor
20 6.3 3.3 4.7 1.6 versicolor
21 6.6 2.9 4.6 1.3 versicolor
22 5.9 3.0 4.2 1.5 versicolor
23 6.0 2.2 4.0 1.0 versicolor
24 6.1 2.9 4.7 1.4 versicolor
25 5.6 2.9 3.6 1.3 versicolor
26 6.7 3.1 4.4 1.4 versicolor
27 6.2 2.2 4.5 1.5 versicolor
28 6.8 2.8 4.8 1.4 versicolor
29 7.1 3.0 5.9 2.1 virginica
30 7.6 3.0 6.6 2.1 virginica
31 7.3 2.9 6.3 1.8 virginica
32 7.2 3.6 6.1 2.5 virginica
33 7.7 3.8 6.7 2.2 virginica
```

```
34 7.4 2.8 6.1 1.9 virginica
35 7.9 3.8 6.4 2.0 virginica
```

## %> % ()

```
(%>%) dplyr . mtcars (help("mtcars")) .
```

```
library(dplyr)
library(magrittr)
df <- mtcars
df$cars <- rownames(df) #just add the cars names to the df
df <- df[,c(ncol(df),1:(ncol(df)-1))] # and place the names in the first column
```

## 1.

```
summarize . n()
```

```
df %>%
 summarize(count=n(),mean_mpg = mean(mpg, na.rm = TRUE),
 min_weight = min(wt),max_weight = max(wt))

count mean_mpg min_weight max_weight
#1 32 20.09062 1.513 5.424
```

## 2.

```
df %>%
 group_by(cyl, gear) %>%
 summarize(count=n(),mean_mpg = mean(mpg, na.rm = TRUE),
 min_weight = min(wt),max_weight = max(wt))

Source: local data frame [8 x 6]
Groups: cyl [?]
#
cyl gear count mean_mpg min_weight max_weight
<dbl> <dbl> <int> <dbl> <dbl> <dbl>
#1 4 3 1 21.500 2.465 2.465
#2 4 4 8 26.925 1.615 3.190
#3 4 5 2 28.200 1.513 2.140
#4 6 3 2 19.750 3.215 3.460
#5 6 4 4 19.750 2.620 3.440
#6 6 5 1 19.700 2.770 2.770
#7 8 3 12 15.050 3.435 5.424
#8 8 5 2 15.400 3.170 3.570
```

## dplyr NSE

```
dplyr NSE (Non-Standard Evaluation) . Shiny . select select_ .
```

```
variable1 <- "Sepal.Length"
variable2 <- "Sepal.Width"
```

```
iris %>%
select_(variable1, variable2) %>%
head(n=5)
Sepal.Length Sepal.Width
1 5.1 3.5
2 4.9 3.0
3 4.7 3.2
4 4.6 3.1
5 5.0 3.6
```

```
lazyeval interp lazeval .
```

```
variable1 <- "Sepal.Length"
variable2 <- "Sepal.Width"
variable3 <- "Species"
iris %>%
select_(variable1, variable2, variable3) %>%
group_by_(variable3) %>%
summarize_(mean1 = lazeval::interp(~mean(var), var = as.name(variable1)), mean2 =
lazyeval::interp(~mean(var), var = as.name(variable2)))
Species mean1 mean2
<fctr> <dbl> <dbl>
1 setosa 5.006 3.428
2 versicolor 5.936 2.770
3 virginica 6.588 2.974
```

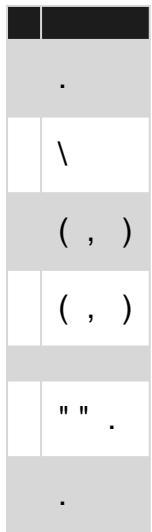
: <https://riptutorial.com/ko/r/topic/4250/>

# 65:

## Examples

```
browser . . R
```

```
browser() browser() . R
```



```
,
```

```
toDebug <- function() {
 a = 1
 b = 2

 browser()

 for(i in 1:100) {
 a = a * b
 }
}

toDebug()
```

```
Called from: toDebug
Browser[1]>
```

```
Called from: toDebug
Browser[1]> a
[1] 1
Browser[1]> b
[1] 2
Browse[1]> n
```

```
debug at #7: for (i in 1:100) {
 a = a * b
}
Browse[2]> n
debug at #8: a = a * b
Browse[2]> a
[1] 1
Browse[2]> n
debug at #8: a = a * b
Browse[2]> a
[1] 2
Browse[2]> Q
```

```
browser() .
```

```
mtcars %>% group_by(cyl) %>% {browser()}
```

```
debug debug .
```

```
debug(mean)
mean(1:3)
```

```
. undebug .
```

```
undebug(mean)
mean(1:3)
```

```
debugonce .
```

```
debugonce(mean)
mean(1:3)
mean(1:3)
```

: <https://riptutorial.com/ko/r/topic/1695/>

# 66: R with knitr

```
1. << - , ... >> =
R
@
2. \Sexpr{#R}
3. << read-external-R-file >> =
read_chunk('r-file.R')
@
<< - , ... >> =
@
```

	(TRUE / FALSE) - R
fig.width	() - R
.	() - R

```
Knitr (LaTeX) (R) knitr LaTeX (.tex) R (.R) R noweb (.Rnw) . . Rnw
LaTeX R .
```

Knitr PDF . . .

```
.Rnw PDF . . , R LaTeX ('kniting'). knitr . . knitr . .
```

```
Rscript -e "library(knitr); knit('r-noweb-file.Rnw')
```

```
.tex (r-noweb.tex) PDF . .
```

```
pdflatex r-noweb-file.tex
```

## Examples

R :

```
Knitr LaTeX R R . . R R / . . .
```

```

r-noweb-file.Rnw
\documentclass{article}

<<echo=FALSE, cache=FALSE>>=
knitr::opts_chunk$set(echo=FALSE, cache=TRUE)
knitr::read_chunk('r-file.R')
@

\begin{document}
This is an Rnw file (R noweb). It contains a combination of LaTeX and R.

One we have called the read_chunk command above we can reference sections of code in the r-file.R script.

<<Chunk1>>=
@
\end{document}

```

## R .

```

r-file.R
note the specific comment style of a single pound sign followed by four dashes

---- Chunk1 ----

print("This is R Code in an external file")

x <- seq(1:10)
y <- rev(seq(1:10))
plot(x,y)

```

## R Knitr

### Knitr LaTeX R . . . approach .

```

r-noweb-file.Rnw
\documentclass{article}
\begin{document}
This is an Rnw file (R noweb). It contains a combination of LaTeX and R.

<<my-label>>=
print("This is an R Code Chunk")
x <- seq(1:10)
@

Above is an internal code chunk.
We can access data created in any code chunk inline with our LaTeX code like this.
The length of array x is \Sexpr{length(x)}.

\end{document}

```

## Knitr LaTeX R

### Knitr LaTeX R . . . approach .

```
r-noweb-file.Rnw
\documentclass{article}
\begin{document}
This is an Rnw file (R noweb). It contains a combination of LaTeX and R.

<<code-chunk-label>>=
print("This is an R Code Chunk")
x <- seq(1:10)
y <- seq(1:10)
plot(x,y) # Brownian motion
@

\end{document}
```

R with knitr : <https://riptutorial.com/ko/r/topic/4334/-r-with-knitr>

## Examples

### GLCM

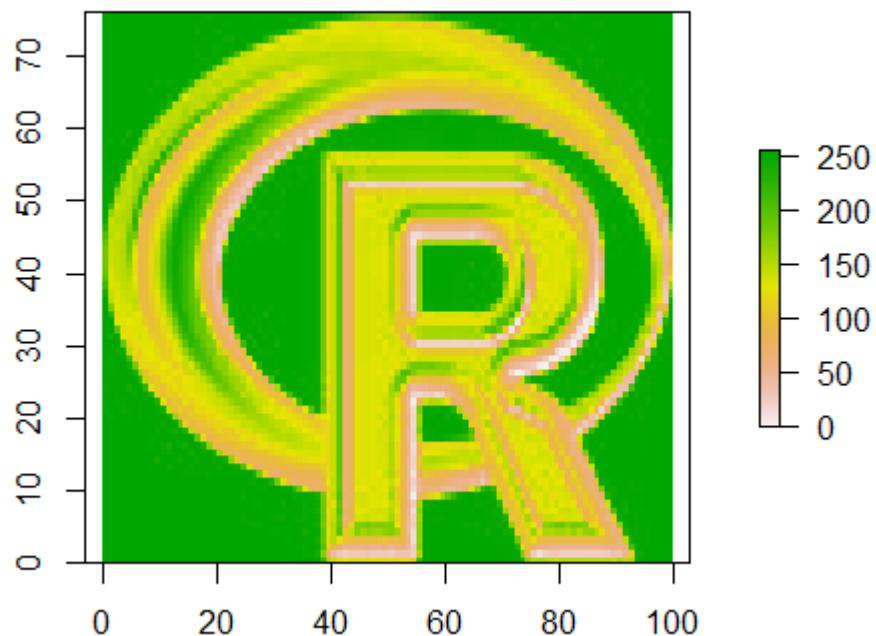
Gray Level Co-Occurrence Matrix (Haralick et al., 1973)

.glcm R RasterLayer

RasterLayer

```
library(glcm)
library(raster)

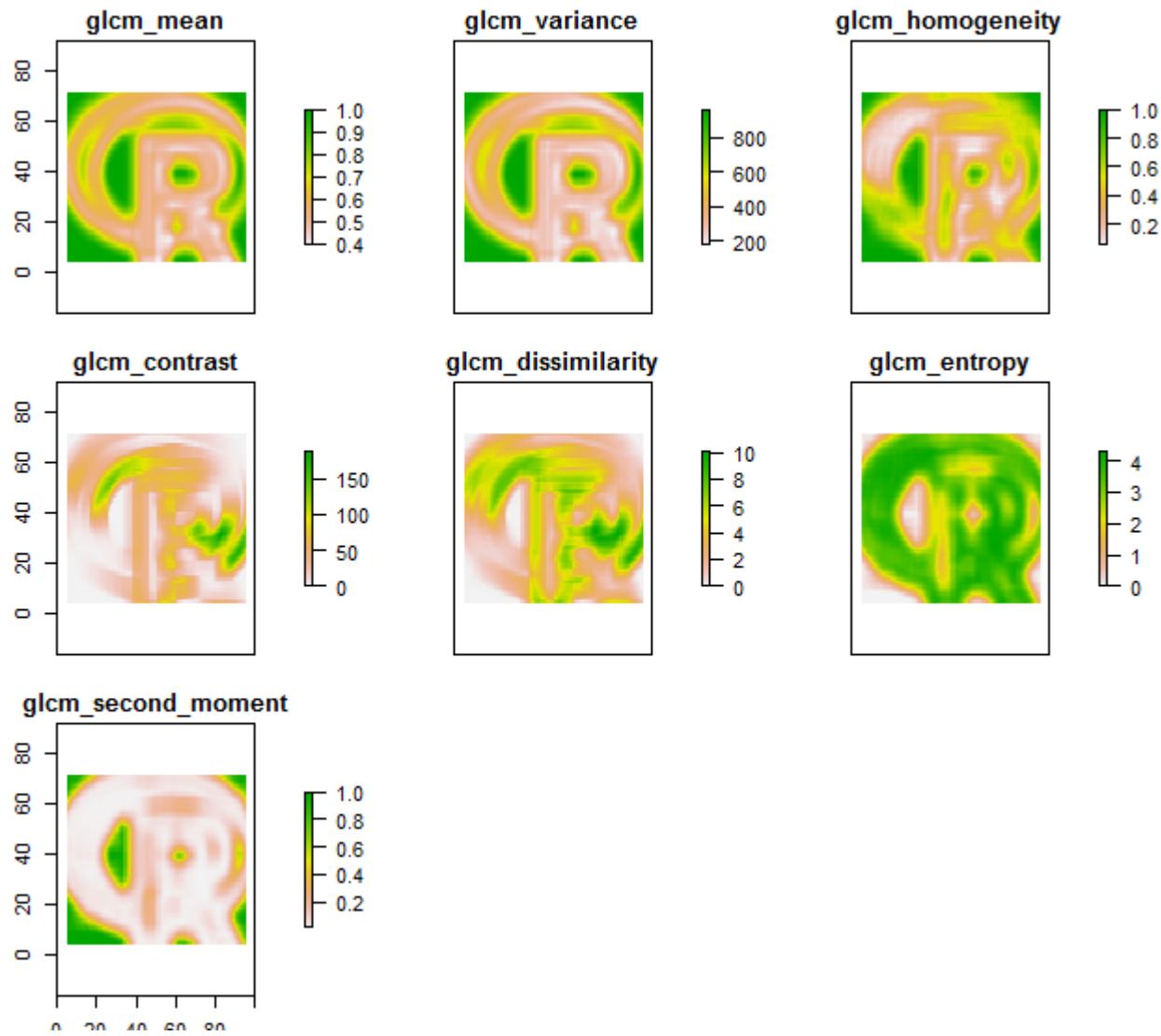
r <- raster("C:/Program Files/R/R-3.2.3/doc/html/logo.jpg")
plot(r)
```



### GLCM

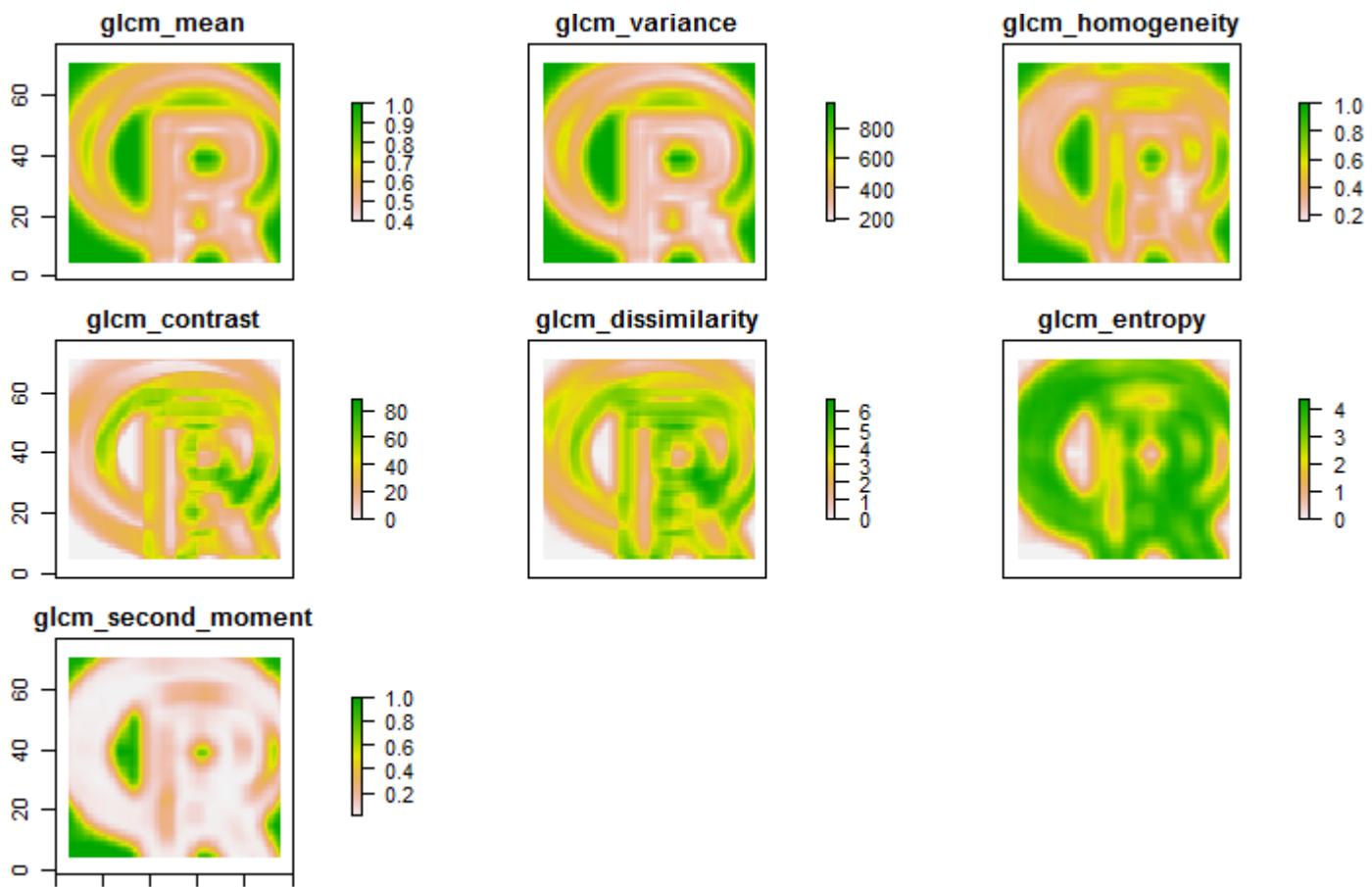
```
rglcm <- glcm(r,
 window = c(9, 9),
 shift = c(1, 1),
 statistics = c("mean", "variance", "homogeneity", "contrast",
 "dissimilarity", "entropy", "second_moment"))
)

plot(rglcm)
```



4 (0°, 45°, 90°, 135°) . shift .

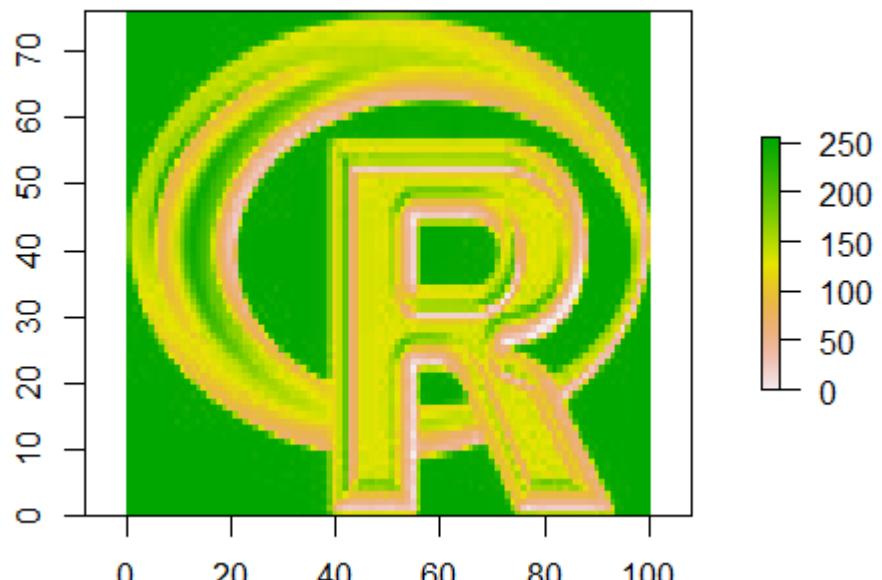
```
rglcm1 <- glcm(r,
 window = c(9,9),
 shift=list(c(0,1), c(1,1), c(1,0), c(1,-1)),
 statistics = c("mean", "variance", "homogeneity", "contrast",
 "dissimilarity", "entropy", "second_moment"))
)
plot(rglcm1)
```



mmand n    Morphologies .

```
library(raster)
library(mmand)

r <- raster("C:/Program Files/R/R-3.2.3/doc/html/logo.jpg")
plot(r)
```



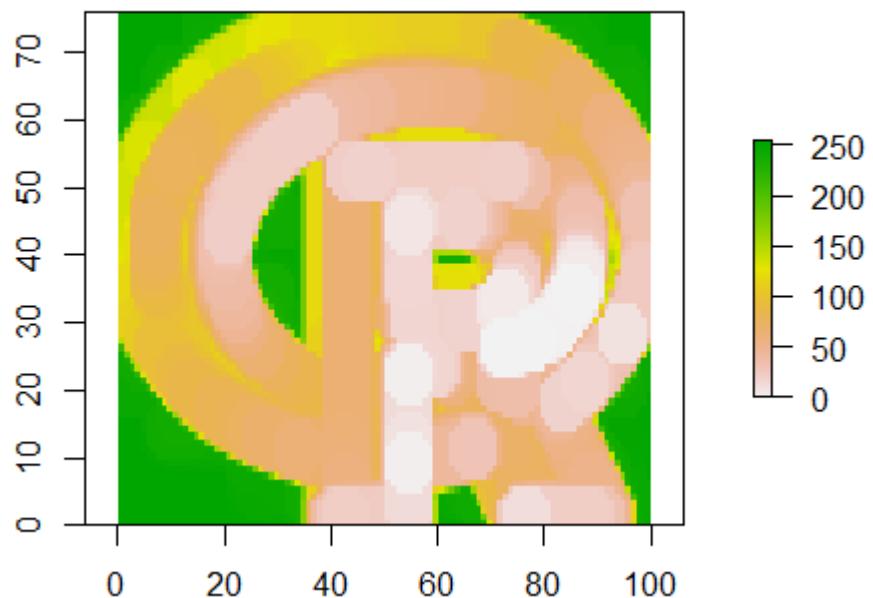
() (:9x9) (:disc, box diamond).

```
sk <- shapeKernel(c(9,9), type="disc")
, erode() . .

rArr <- as.array(r, transpose = TRUE)
rErode <- erode(rArr, sk)
rErode <- setValues(r, as.vector(aperm(rErode)))

erode() dilate() , opening() closing() . .

plot(rErode)
```



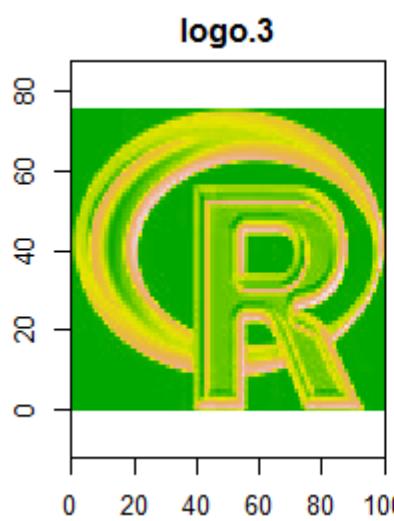
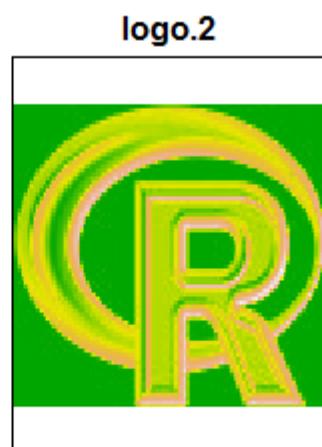
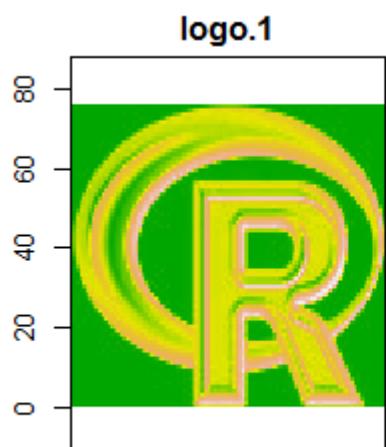
: <https://riptutorial.com/ko/r/topic/3726/>---

## 68: I/O

### Examples

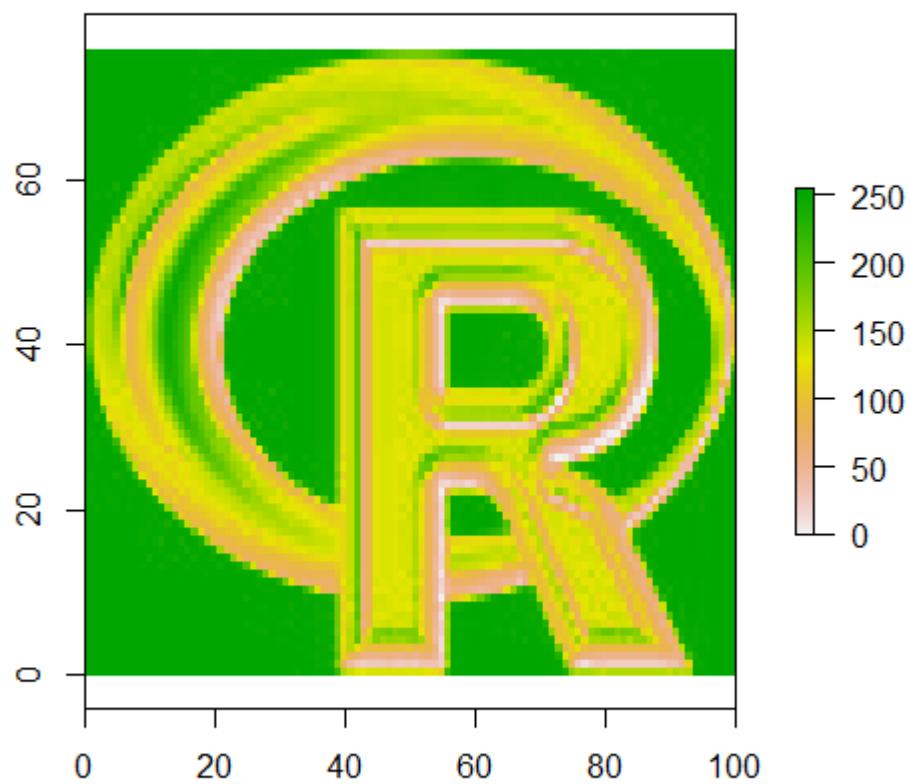
R (, , ).

```
library(raster)
r <- stack("C:/Program Files/R/R-3.2.3/doc/html/logo.jpg")
plot(r)
```



```
RasterStack [[].
```

```
plot(r[[1]])
```



I / O : <https://riptutorial.com/ko/r/topic/5539/---i---o>

## 69: -

. , R "run-length encoding" . :

```
dat <- c(1, 2, 2, 2, 3, 1, 4, 4, 1, 1)
```

1 1 ; 2 3; 3 -1 ;. R .

. R , rleid data.table ( ) .

## Examples

`rle' -

.

```
dat <- c(1, 2, 2, 2, 3, 1, 4, 4, 1, 1)
```

rle .

```
r <- rle(dat)
r
Run Length Encoding
lengths: int [1:6] 1 3 1 1 2 2
values : num [1:6] 1 2 3 1 4 1
```

r\$values r\$values .

```
r$values
[1] 1 2 3 1 4 1
```

1 2 3 1 .

r\$lengths .

```
r$lengths
[1] 1 3 1 1 2 2
```

1 1, 2 3.

R

.

```
(dat <- data.frame(x = c(1, 1, 2, 2, 2, 1), y = 1:6))
```

```
x y
1 1 1
2 1 2
3 2 3
4 2 4
5 2 5
6 1 6
```

x 2 1 3 2 1 1. y . x ( 1.5, 4 6 ).

base R, rle x run-length .

```
(r <- rle(dat$x))
Run Length Encoding
lengths: int [1:3] 2 3 1
values : num [1:3] 1 2 1
```

. length(r\$lengths) r\$lengths rep .

```
(run.id <- rep(seq_along(r$lengths), r$lengths))
[1] 1 1 2 2 2 3
```

tapply ID y .

```
data.frame(x=r$values, meanY=tapply(dat$y, run.id, mean))
x meanY
1 1 1.5
2 2 4.0
3 1 6.0
```

## data.table run

data.table . .

```
library(data.table)
(DT <- data.table(x = c(1, 1, 2, 2, 2, 1), y = 1:6))
x y
1: 1 1
2: 1 2
3: 2 3
4: 2 4
5: 2 5
6: 1 6
```

x 2 1 3 2 1 1. y . x ( 1.5, 4 6 ).

data.table rleid ID ID .

```
rleid(DT$x)
[1] 1 1 2 2 2 3
```

ID y .

```
DT[,mean(y),by=.(x, rleid(x))]
x rleid V1
1: 1 1 1.5
2: 2 2 4.0
3: 1 3 6.0
```

**long run run-length encoding ( run ) . , 1 0 1,000 :**

```
set.seed(144)
dat <- sample(rep(0:1, c(1, 1e5)), 1e7, replace=TRUE)
table(dat)
0 1
103 9999897
```

1

```
rle.df <- with(rle(dat), data.frame(values, lengths))
dim(rle.df)
[1] 207 2
head(rle.df)
values lengths
1 1 52818
2 0 1
3 1 219329
4 0 1
5 1 318306
6 0 1
```

- 52,818 1, 0, 219,329 1, 0 . run-length 207 1000 414 . rle.df write.csv .

- . rep run-length values run-length lengths .

```
decompressed <- rep(rle.df$values, rle.df$lengths)
```

```
identical(decompressed, dat)
[1] TRUE
```

**R inverse.rle rle :**

```
rle.obj <- rle(dat) # create a rle object here
class(rle.obj)
[1] "rle"

dat.inv <- inverse.rle(rle.obj) # apply the inverse.rle on the rle object
```

dat .

```
identical(dat.inv, dat)
[1] TRUE
```

- : <https://riptutorial.com/ko/r/topic/1133/>----

# 70:

```
() . (,,,) () . .
```

## Examples

**barplot()**

```
x () y . () .
```

```
barplot() R . barplot() . R heights .
```

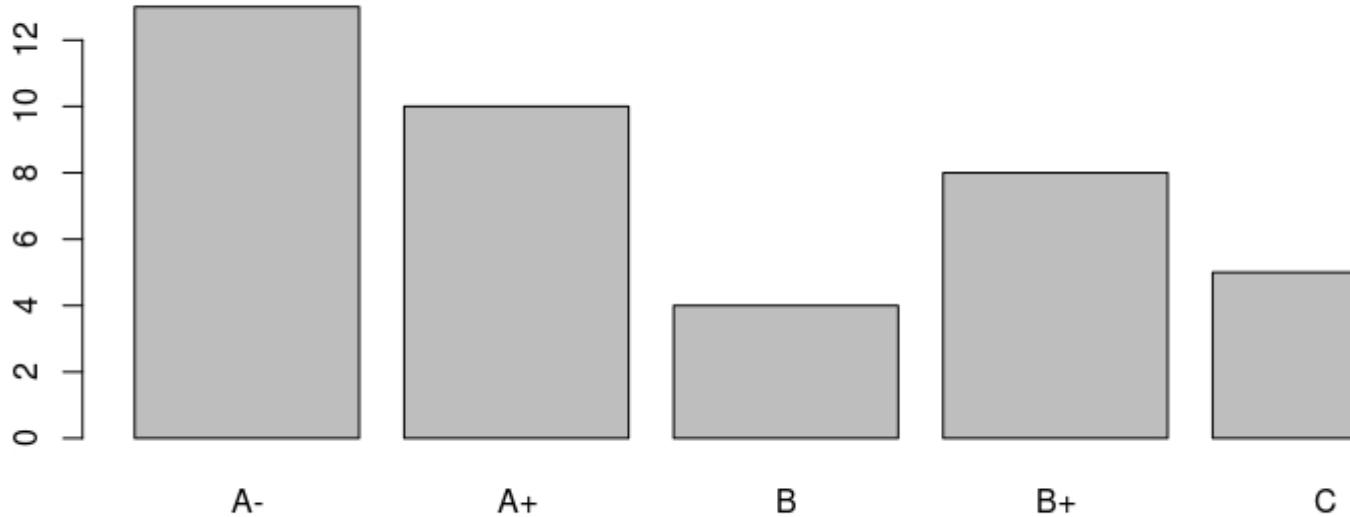
```
barplot() .
```

```
> grades<-c("A+","A-","B+","B","C")
> Marks<-sample(grades,40,replace=T,prob=c(.2,.3,.25,.15,.1))
> Marks
[1] "A+" "A-" "B+" "A-" "A+" "B" "A+" "B+" "A-" "B" "A+" "A-"
[13] "A-" "B+" "A-" "A-" "A-" "A+" "A-" "A+" "A+" "C" "C"
[25] "B" "C" "B+" "C" "B+" "B+" "B+" "A+" "B+" "A-" "A+" "A-"
[37] "A-" "B" "C" "A+"
```

```
>
```

```
> barplot(table(Marks),main="Mid-Marks in Algorithms")
```

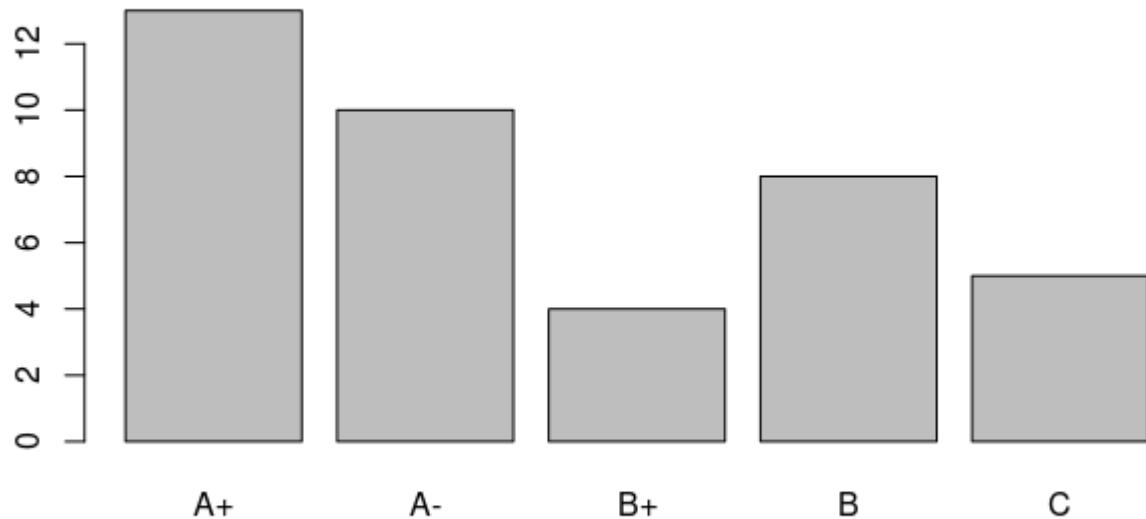
Mid-Marks in Algorithms



```
barplot() factor x lexicographical order . names.arg , .
```

```
plot to the desired horizontal axis labels
> barplot(table(Marks),names.arg=grades ,main="Mid-Marks in Algorithms")
```

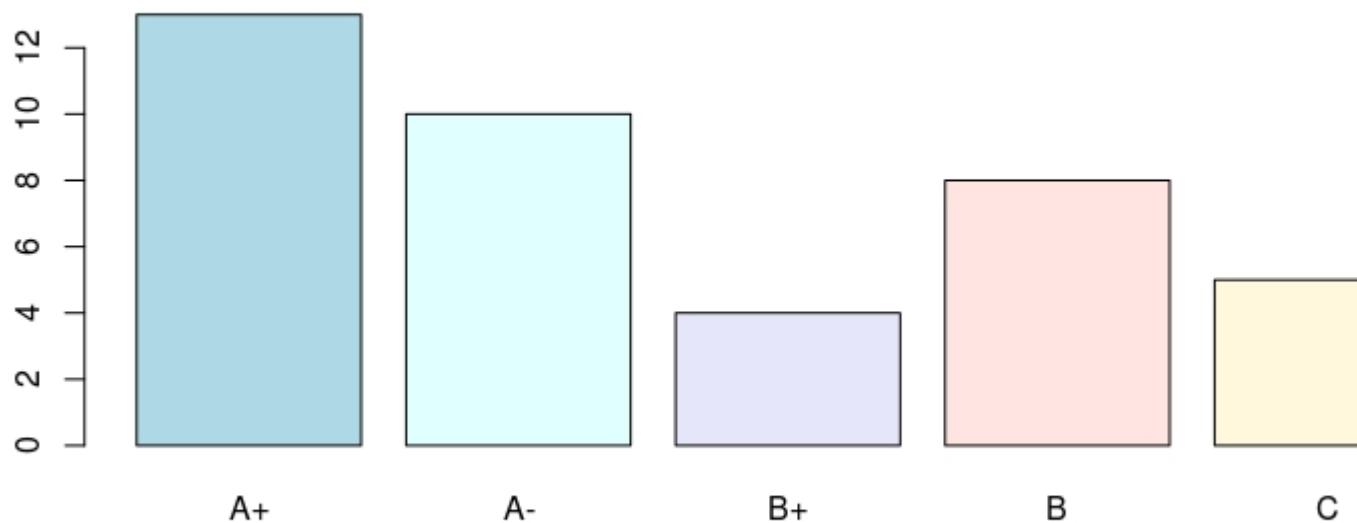
## Mid-Marks in Algorithms



```
col= .
```

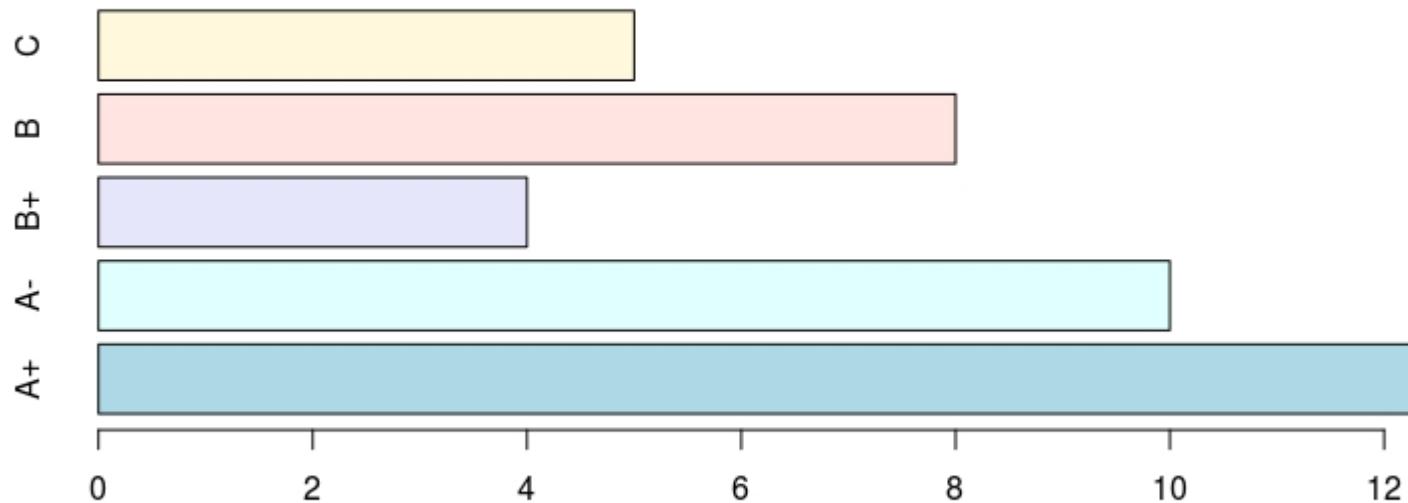
```
> barplot(table(Marks),names.arg=grades,col = c("lightblue",
"lightcyan", "lavender", "mistyrose", "cornsilk"),
main="Mid-Marks in Algorithms")
```

## Mid-Marks in Algorithms



```
> barplot(table(Marks),names.arg=grades,horiz=TRUE,col = c("lightblue",
 "lightcyan", "lavender", "mistyrose", "cornsilk"),
 main="Mid-Marks in Algorithms")
```

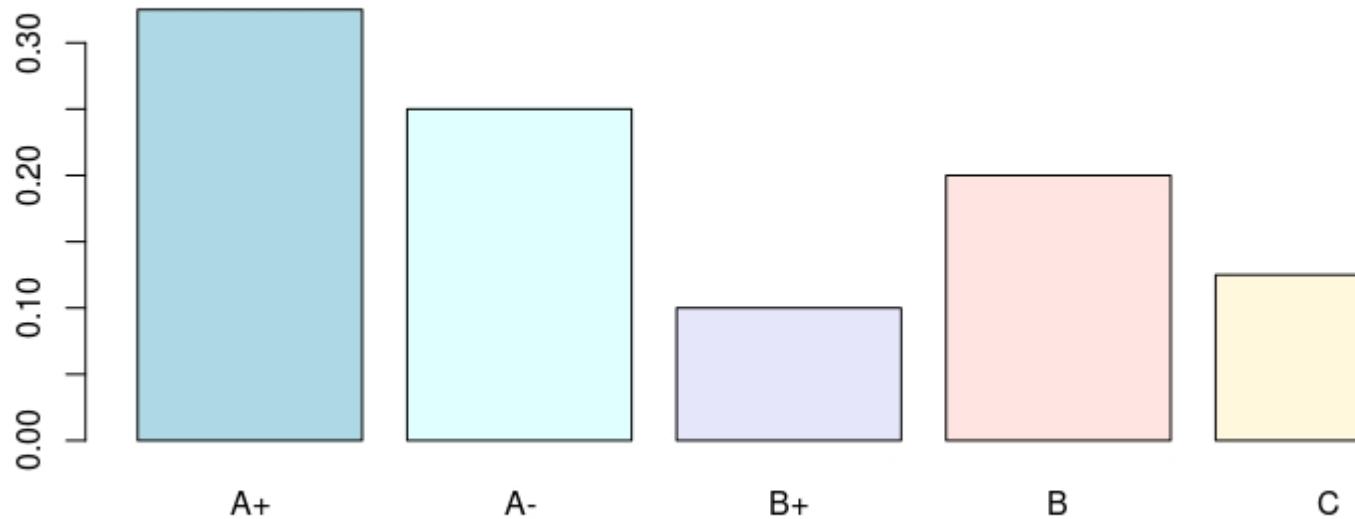
## Mid-Marks in Algorithms



y

```
> barplot(prop.table(table(Marks)),names.arg=grades,col = c("lightblue",
 "lightcyan", "lavender", "mistyrose", "cornsilk"),
 main="Mid-Marks in Algorithms")
```

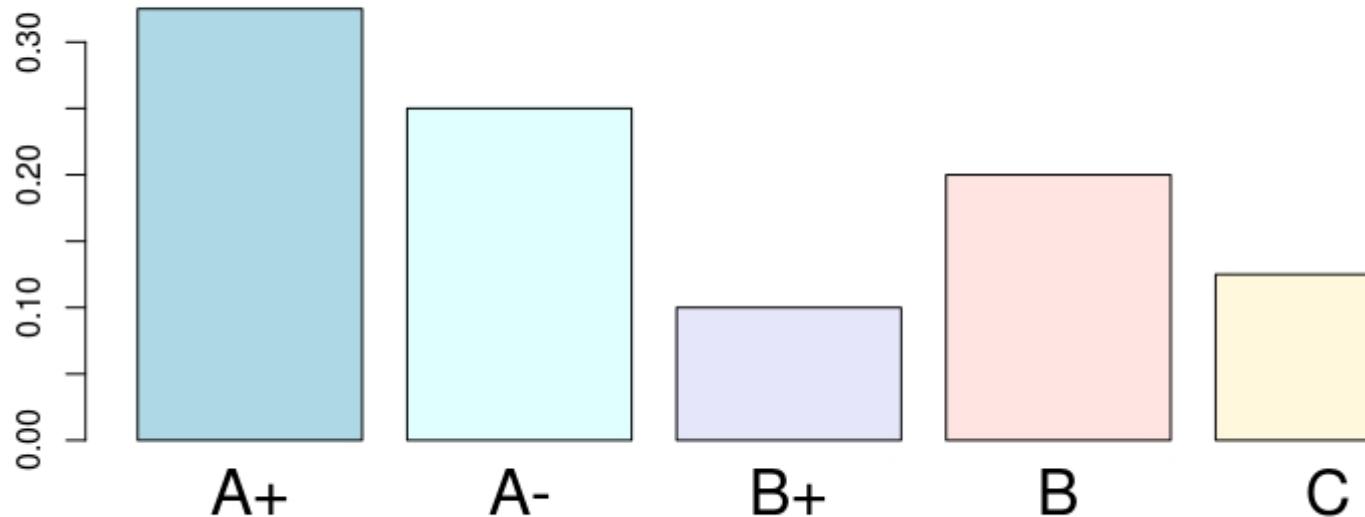
## Mid-Marks in Algorithms



```
x cex.names .
```

```
> barplot(prop.table(table(Marks)),names.arg=grades,col = c("lightblue",
 "lightcyan", "lavender", "mistyrose", "cornsilk"),
 main="Mid-Marks in Algorithms",cex.names=2)
```

## Mid-Marks in Algorithms

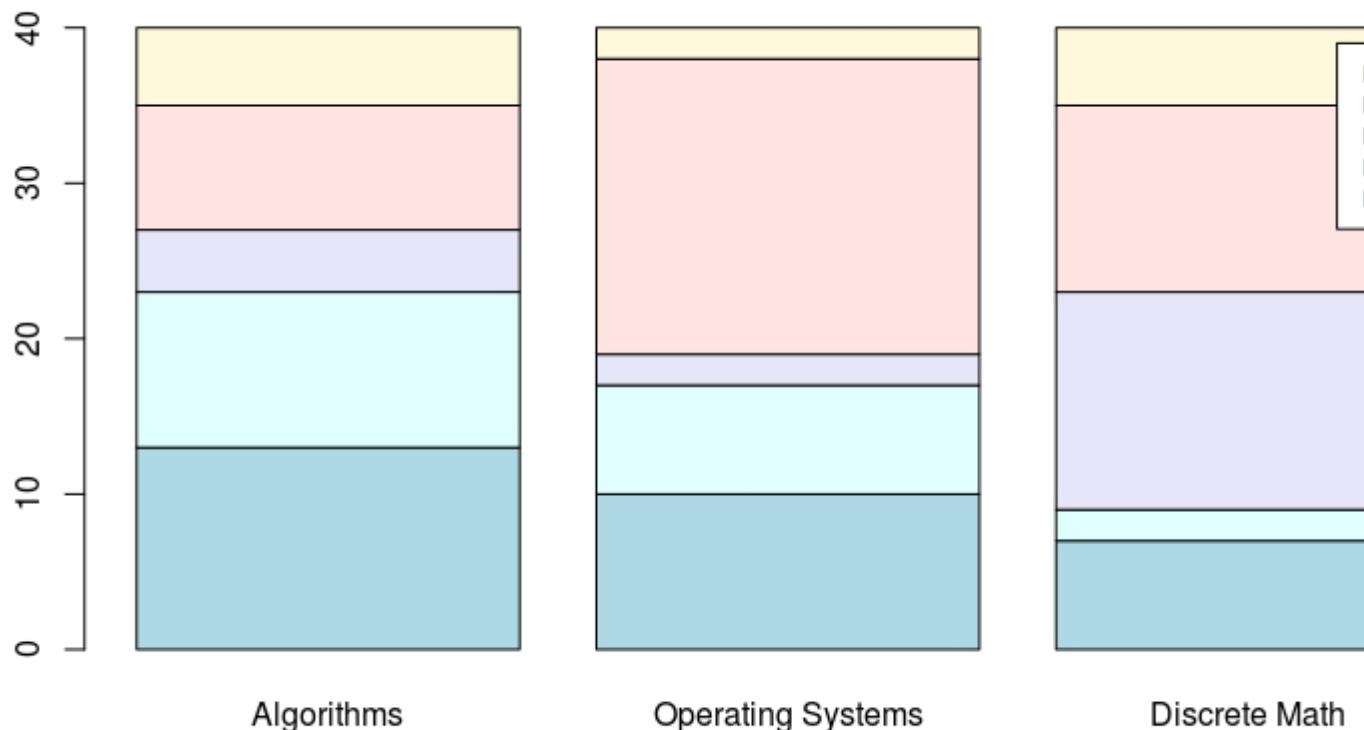


```
barplot() heights . . , . .
```

```
> gradTab
 Algorithms Operating Systems Discrete Math
A- 13 10 7
A+ 10 7 2
B 4 2 14
B+ 8 19 12
C 5 2 5
```

```
> barplot(gradTab,col = c("lightblue","lightcyan",
 "lavender", "mistyrose", "cornsilk"),legend.text = grades,
 main="Mid-Marks in Algorithms")
```

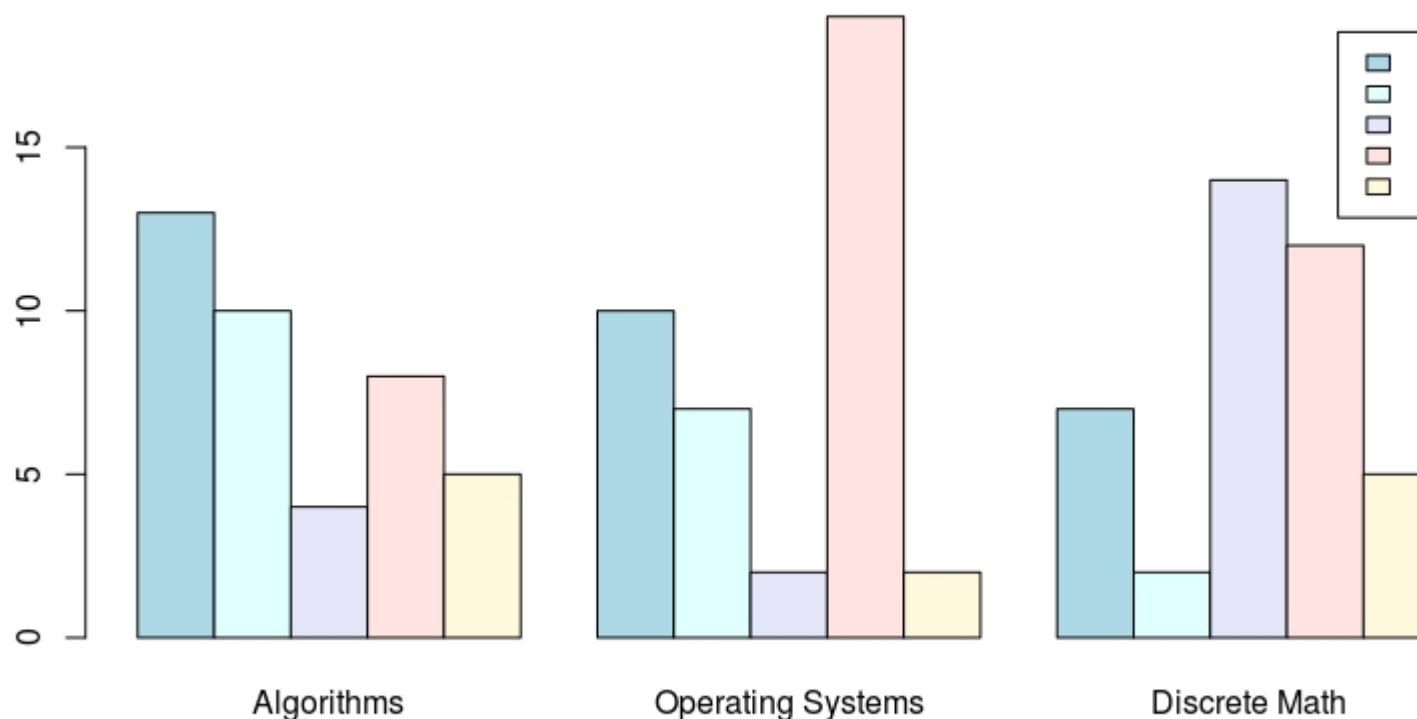
**Mid-Marks in Algorithms**



besides .

```
> barplot(gradTab,beside = T,col = c("lightblue","lightcyan",
 "lavender", "mistyrose", "cornsilk"),legend.text = grades,
 main="Mid-Marks in Algorithms")
```

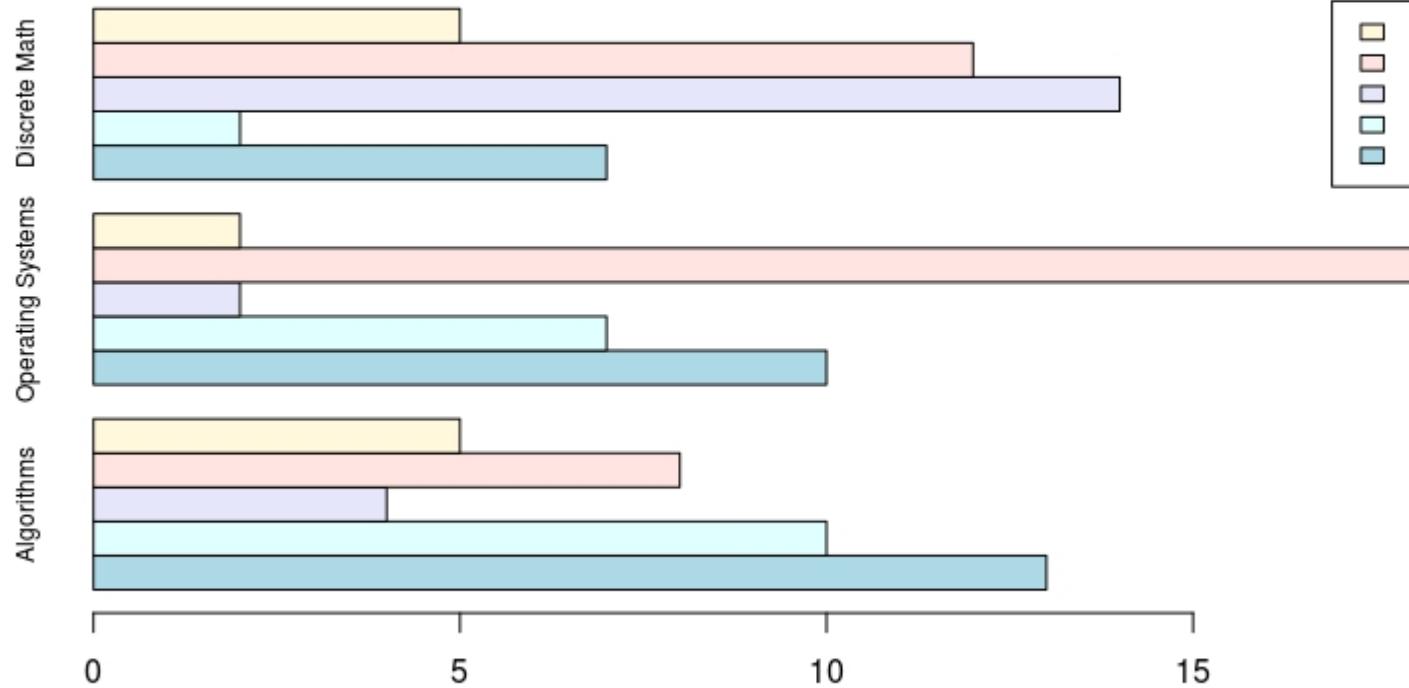
## Mid-Marks in Algorithms



```
horiz=T .
```

```
> barplot(gradTab,beside = T,horiz=T,col = c("lightblue","lightcyan",
 "lavender", "mistyrose", "cornsilk"),legend.text = grades,
 cex.names=.75,main="Mid-Marks in Algorithms")
```

## Mid-Marks in Algorithms



: <https://riptutorial.com/ko/r/topic/8091/>

# 71: :

R R .

## Examples

Q & A .

- .
- .
- .

```
library(help = "datasets")
```

- ?data.frame . SO Docs . R
- Remarks .

R> , > + . . .

- ( ).
- # ## .
- [1] .
- .

```
= <- R . , x<-1 (x <- 1 x < -1)
```

H1 .

: : <https://riptutorial.com/ko/r/topic/5410/>-----

# 72:

R

```
rand . . r* (). " " runif . " " R .
```

## Examples

```
sample . .
```

```
set.seed set.seed . sample set.seed .
```

```
sample . .
```

```
set.seed(1251)
sample(x = 10)
```

```
[1] 7 1 4 8 6 3 10 5 2 9
```

```
sample 1 x . .
```

```
library(datasets)
set.seed(1171)
iris_rand <- iris[sample(x = 1:nrow(iris)),]

> head(iris)
Sepal.Length Sepal.Width Petal.Length Petal.Width Species
1 5.1 3.5 1.4 0.2 setosa
2 4.9 3.0 1.4 0.2 setosa
3 4.7 3.2 1.3 0.2 setosa
4 4.6 3.1 1.5 0.2 setosa
5 5.0 3.6 1.4 0.2 setosa
6 5.4 3.9 1.7 0.4 setosa

> head(iris_rand)
Sepal.Length Sepal.Width Petal.Length Petal.Width Species
145 6.7 3.3 5.7 2.5 virginica
5 5.0 3.6 1.4 0.2 setosa
85 5.4 3.0 4.5 1.5 versicolor
137 6.3 3.4 5.6 2.4 virginica
128 6.1 3.0 4.9 1.8 virginica
105 6.5 3.0 5.8 2.2 virginica
```

```
sample . . () . .
```

```
set.seed(7043)
```

```
sample(x = LETTERS, size = 7)

[1] "S" "P" "J" "F" "Z" "G" "R"

size x . x .

set.seed(7305)
sample(x = letters, size = 26)

[1] "x" "z" "y" "i" "k" "f" "d" "s" "g" "v" "j" "o" "e" "c" "m" "n" "h" "u" "a" "b" "l" "r"
"w" "t" "q" "p"

sample(x = letters, size = 30)
Error in sample.int(length(x), size, replace, prob) :
 cannot take a sample larger than the population when 'replace = FALSE'
```

sample replace . replace FALSE . TRUE

```
set.seed(5062)
sample(x = c("A", "B", "C", "D"), size = 8, replace = TRUE)

[1] "D" "C" "D" "B" "A" "A" "A" "A"
```

sample . "" . 20 . , , 1/3.

```
set.seed(6472)
sample(x = c("Red", "Blue", "Green"),
 size = 20,
 replace = TRUE)
```

urn 2 , 1 , 1 . x Red . sample prob sample .

prob . 1/2 1/4.

```
set.seed(28432)
sample(x = c("Red", "Blue", "Green"),
 size = 20,
 replace = TRUE,
 prob = c(0.50, 0.25, 0.25))
```

, prob 1 . R 1 . 2 Red, 1 Blue 1 Green . . .

```
set.seed(28432)
frac_prob_example <- sample(x = c("Red", "Blue", "Green"),
 size = 200,
 replace = TRUE,
```

```
prob = c(0.50,0.25,0.25))

set.seed(28432)
numeric_prob_example <- sample(x = c("Red","Blue","Green"),
 size = 200,
 replace = TRUE,
 prob = c(2,1,1))

> identical(frac_prob_example,numeric_prob_example)
[1] TRUE
```

0 0 .

replace FALSE prob . , prob . , size 0 . :

```
set.seed(21741)
sample(x = c("Red","Blue","Green"),
 size = 2,
 replace = FALSE,
 prob = c(0.8,0.19,0.01))
```

Red ( ). 80 %, 19 %, 1 %.

Red . 20 % ( 19 %, 1 %). (19/20) 95 %, (1/20) 5 % .

set.seed randomization random seed . R set.seed .

```
set.seed(1643)
samp1 <- sample(x = 1:5,size = 200,replace = TRUE)

set.seed(1643)
samp2 <- sample(x = 1:5,size = 200,replace = TRUE)

> identical(x = samp1,y = samp2)
[1] TRUE
```

: <https://riptutorial.com/ko/r/topic/9574/>

RandomForest . Wikipedia Random Forests . R randomForest . CRAN .

## Examples

```
Used for both Classification and Regression examples
library(randomForest)
library(car) ## For the Soils data
data(Soils)

#####
RF Classification Example
set.seed(656) ## for reproducibility
S_RF_Class = randomForest(Gp ~ ., data=Soils[,c(4,6:14)])
Gp_RF = predict(S_RF_Class, Soils[,6:14])
length(which(Gp_RF != Soils$Gp)) ## No Errors

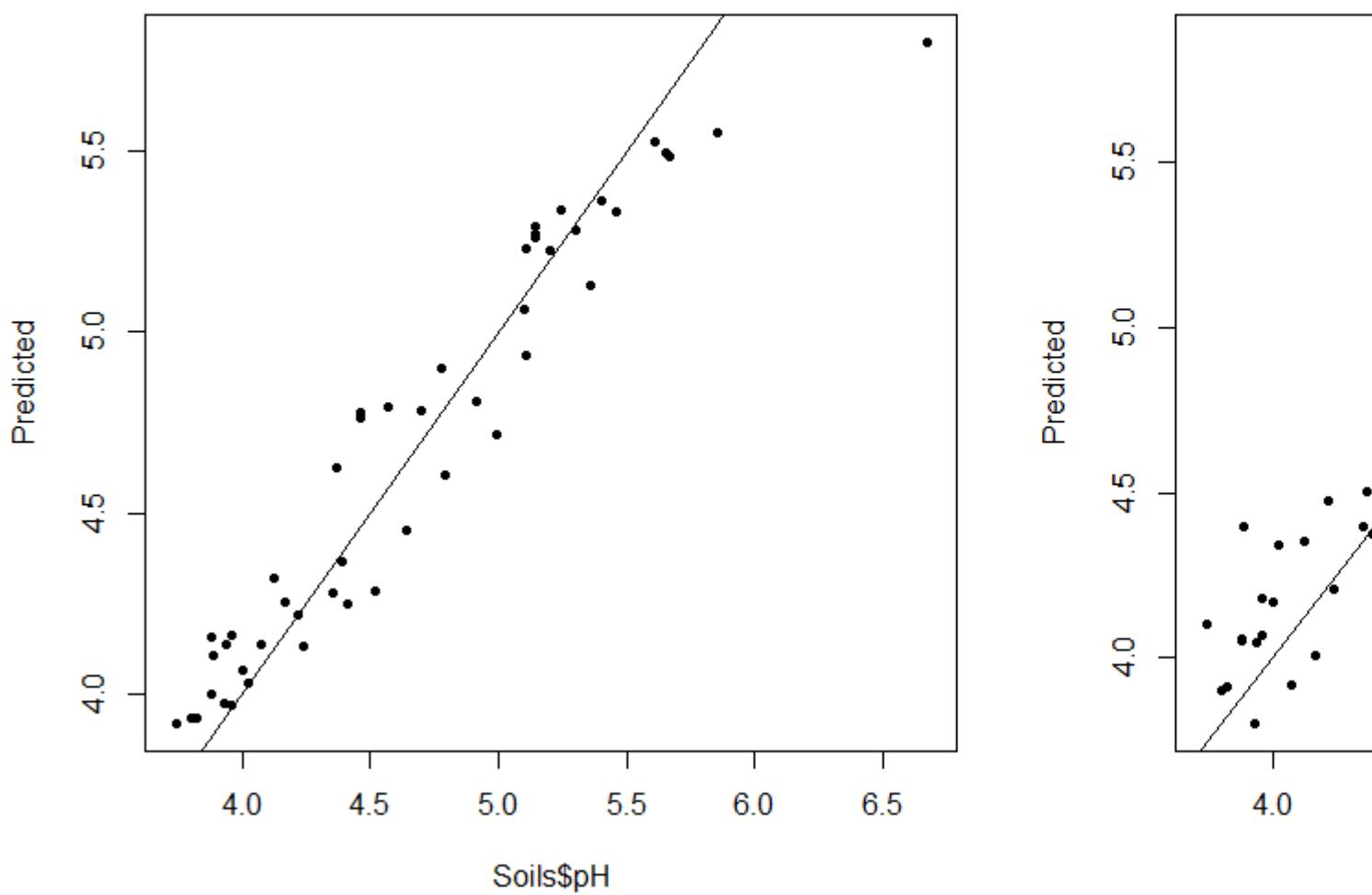
Naive Bayes for comparison
library(e1071)
S_NB = naiveBayes(Soils[,6:14], Soils[,4])
Gp_NB = predict(S_NB, Soils[,6:14], type="class")
length(which(Gp_NB != Soils$Gp)) ## 6 Errors
```

RF .

```
#####
RF Regression Example
set.seed(656) ## for reproducibility
S_RF_Reg = randomForest(pH ~ ., data=Soils[,6:14])
pH_RF = predict(S_RF_Reg, Soils[,6:14])

Compare Predictions with Actual values for RF and Linear Model
S_LM = lm(pH ~ ., data=Soils[,6:14])
pH_LM = predict(S_LM, Soils[,6:14])
par(mfrow=c(1,2))
plot(Soils$pH, pH_RF, pch=20, ylab="Predicted", main="Random Forest")
abline(0,1)
plot(Soils$pH, pH_LM, pch=20, ylab="Predicted", main="Linear Model")
abline(0,1)
```

## Random Forest



: <https://riptutorial.com/ko/r/topic/8088-->

# 74:

- ()
- 
- strsplit
- 

## Examples

```
is.character() . as.character() .
```

```
x <- "The quick brown fox jumps over the lazy dog"
class(x)
[1] "character"
is.character(x)
[1] TRUE
```

```
NA .
```

```
as.numeric("2")
[1] 2
as.numeric("fox")
[1] NA
Warning message:
NAs introduced by coercion
```

: <https://riptutorial.com/ko/r/topic/9017/>

:

•

## Examples

R print cat . R .

```
print("Hello World")
#[1] "Hello World"
cat("Hello World\n")
#Hello World
```

.( : x <- "Hello World" x print .)

cat . 1 () .

```
cat(c("hello", "world", "\n"))
#hello world
```

( \n ) .

```
cat("Hello World")
#Hello World>
```

.(RStudio 's .)

print "" . "Hello World" cat . [1] ( ).

```
print("Hello World")
#[1] "Hello World"
```

R . s print(s) print("Hello World") :

```
s <- "Hello World"
s
#[1] "Hello World"
```

```
"Hello World"
#[1] "Hello World"
```

( c() oncatenate C ), print() cat :

```
print(c("Hello World", "Here I am."))
```

```
[1] "Hello World" "Here I am."
```

```
c() .(paste .) R . . , [1] , R .
```

```
c("Hello World", "Here I am!", "This next string is really long.")
#[1] "Hello World" "Here I am!"
#[3] "This next string is really long."
```

```
print .
```

```
""" print .
```

```
print(1)
#[1] 1
print(TRUE)
#[1] TRUE
```

**Factor SO** . cat print . print() foo print(foo) invisible invisible . R  
**REPL, "read-eval-print-loop"** . cat (: ) . ( print() ) , cat() . message() warning()  
. cat , . . .

```
message("hello world")
#hello world
suppressMessages(message("hello world"))
```

. R map-reduce . . . . .

```
file() ("r").
```

```
conn <- file("/path/example.data", "r") #when file is in local system
conn1 <- file("stdin", "r") #when just standard input/output for files are available
```

```
line <- readLines(conn, n=1, warn=FALSE)
```

```
conn n=1 . (10 20) n(:10,20) . n=-1 .
```

```
; writeLines(),cat() . . . file() .
```

```
conn2 <- file("/path/result.data", "w") #when file is in local system
conn3 <- file("stdout", "w") #when just standard input/output for files are available
```

```
writeLines("text",conn2, sep = "\n")
```

**Base R** . . .

```
system("top -a -b -n 1", intern = TRUE)
system2("top", "-a -b -n 1", stdout = TRUE)
```

```
[1] "top - 08:52:03 up 70 days, 15:09, 0 users, load average: 0.00, 0.00, 0.00"
[2] "Tasks: 125 total, 1 running, 124 sleeping, 0 stopped, 0 zombie"
[3] "Cpu(s): 0.9%us, 0.3%sy, 0.0%ni, 98.7%id, 0.1%wa, 0.0%hi, 0.0%si, 0.0%st"
[4] "Mem: 12194312k total, 3613292k used, 8581020k free, 216940k buffers"
[5] "Swap: 12582908k total, 2334156k used, 10248752k free, 1682340k cached"
[6] "
[7] " PID USER PR NI VIRT RES SHR S %CPU %MEM TIME+ COMMAND "
[8] "11300 root 20 0 1278m 375m 3696 S 0.0 3.2 124:40.92 trala "
[9] " 6093 user1 20 0 1817m 269m 1888 S 0.0 2.3 12:17.96 R "
[10] " 4949 user2 20 0 1917m 214m 1888 S 0.0 1.8 11:16.73 R "
```

, UNIX top -a -b -n 1 . OS

devtools

```
devtools::system_output("top", "-a -b -n 1")
```

data.table fread read.table read.table . data.table data.frame .

```
fread("top -a -b -n 1", check.names = TRUE)
 PID USER PR NI VIRT RES SHR S X.CPU X.MEM TIME. COMMAND
1: 11300 root 20 0 1278m 375m 3696 S 0 3.2 124:40.92 trala
2: 6093 user1 20 0 1817m 269m 1888 S 0 2.3 12:18.56 R
3: 4949 user2 20 0 1917m 214m 1888 S 0 1.8 11:17.33 R
4: 7922 user3 20 0 3094m 131m 1892 S 0 1.1 21:04.95 R
```

fread 6 .

check.names = TRUE %CPU , %MEN TIME+ .

: <https://riptutorial.com/ko/r/topic/5541/>--

# 76:

- `boxplot(x, ...)` #
- `boxplot(, = NULL, ..., , na.action = NULL) ## "S3"`
- `boxplot(x, ..., range = 1.5, width = NULL, varwidth = FALSE, notch = FALSE, outline = TRUE, , = TRUE, = ("fg"), col = NULL, log = " ", = S3 (= false), == ==`

( R )	
	<code>y ~ grp . y grp () .</code>
	<code>data.frame( ).</code>
	<code>.</code>
na.action	<code>NA . . .</code>
	<code>.</code>
	<code>TRUE () boxplot .</code>
	<code>col null , . .</code>

## Examples

### `boxplot()` {graphics} box-and-whisker

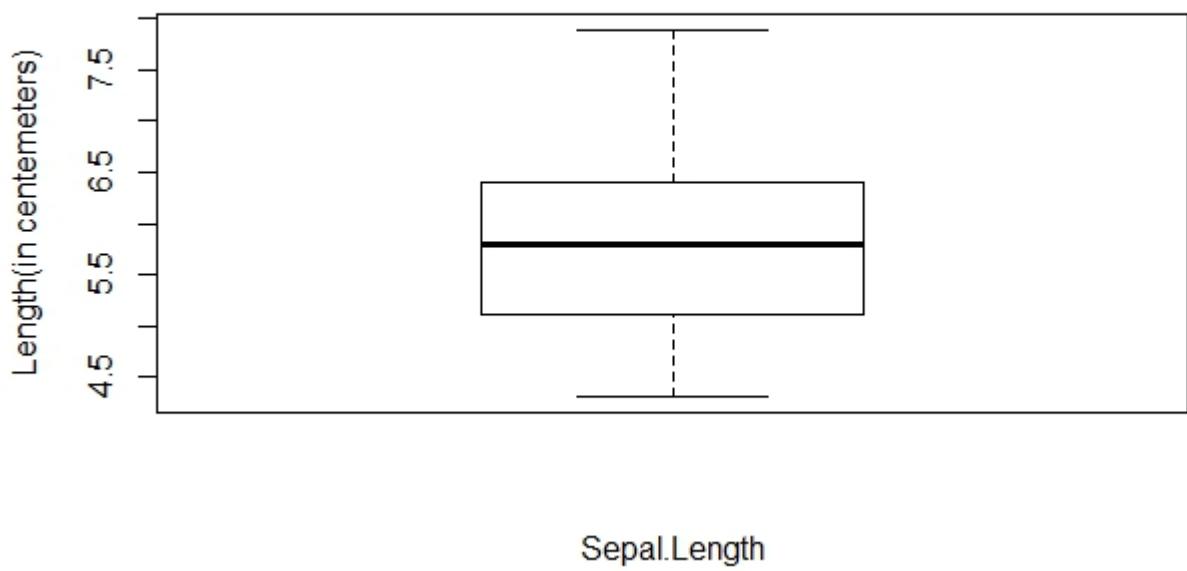
```
boxplot() iris .
```

```
> head(iris)
 Sepal.Length Sepal.Width Petal.Length Petal.Width Species
1 5.1 3.5 1.4 0.2 setosa
2 4.9 3.0 1.4 0.2 setosa
3 4.7 3.2 1.3 0.2 setosa
4 4.6 3.1 1.5 0.2 setosa
5 5.0 3.6 1.4 0.2 setosa
6 5.4 3.9 1.7 0.4 setosa
```

## boxplot (Sepal.Length)

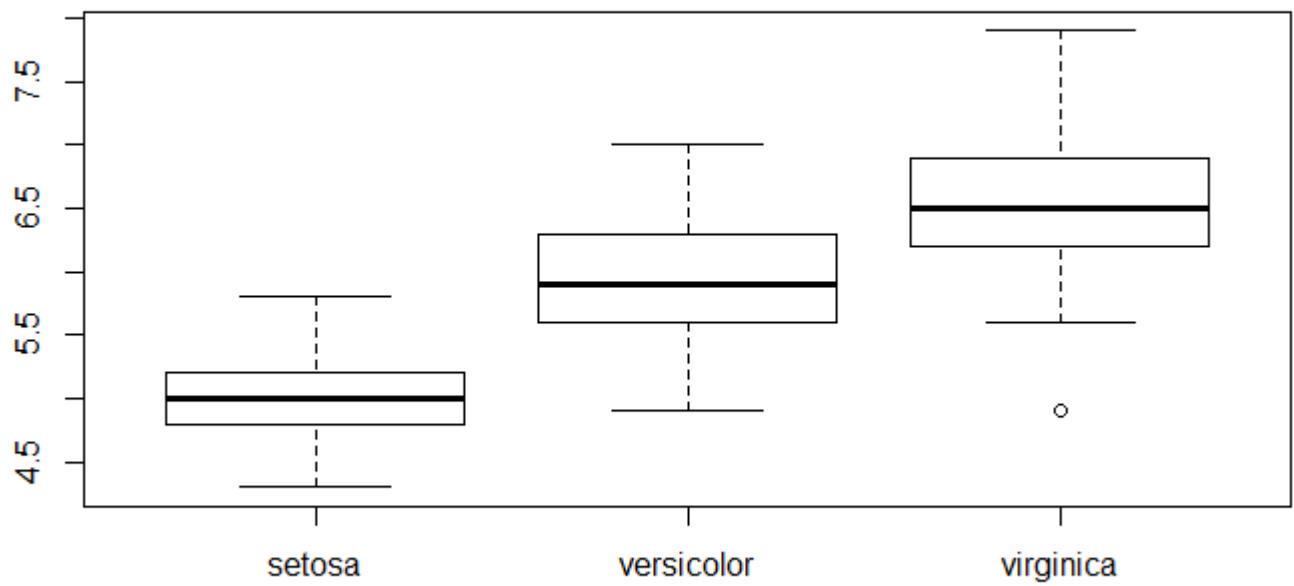
```
boxplot(iris[,1],xlab="Sepal.Length",ylab="Length(in centimeters)",
 main="Summary Charateristics of Sepal.Length(Iris Data)")
```

## Summary Characteristics of Sepal.Length(Iris Data)



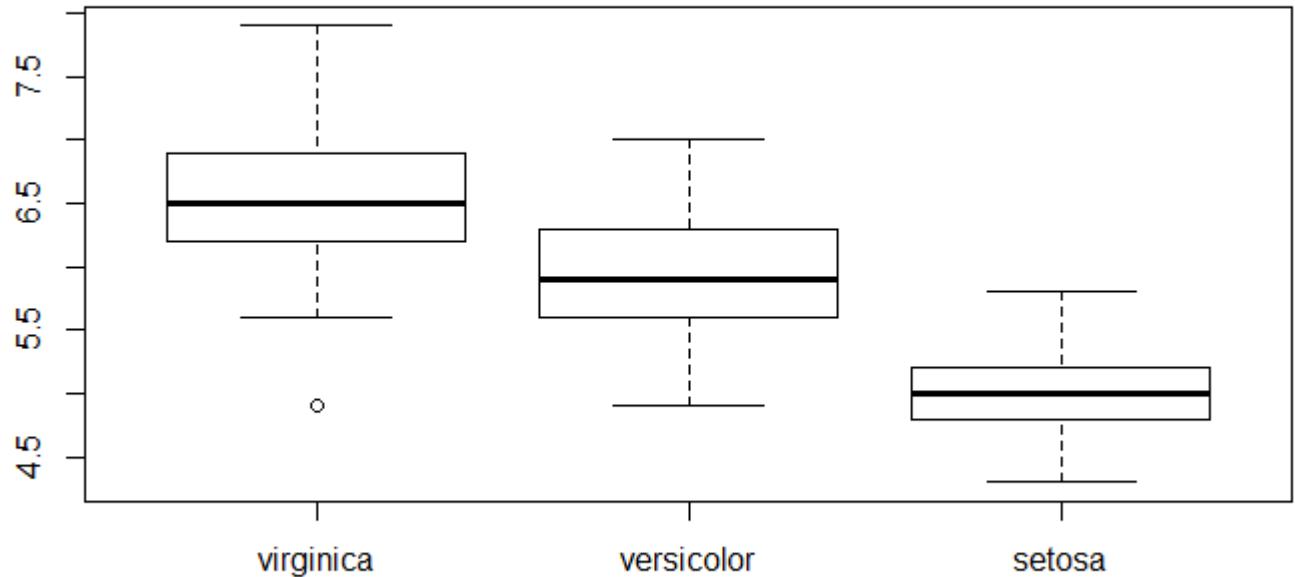
## (species)

```
boxplot(Sepal.Length~Species,data = iris)
```



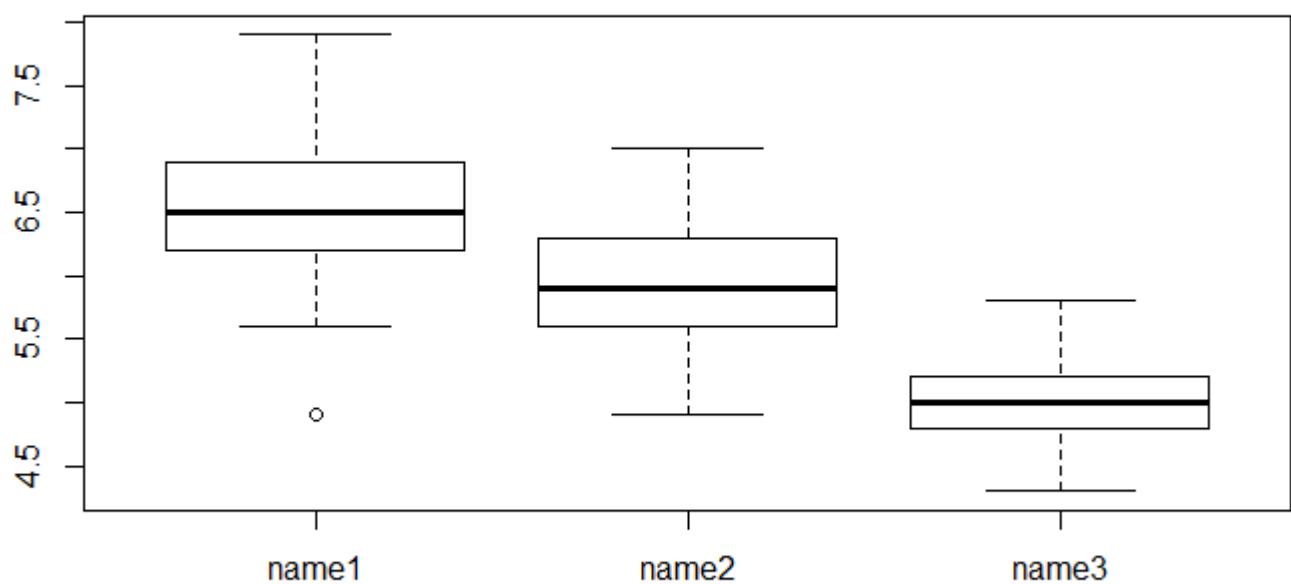
```
virginica - versicolor - setosa
```

```
newSpeciesOrder <- factor(iris$Species, levels=c("virginica","versicolor","setosa"))
boxplot(Sepal.Length~newSpeciesOrder,data = iris)
```



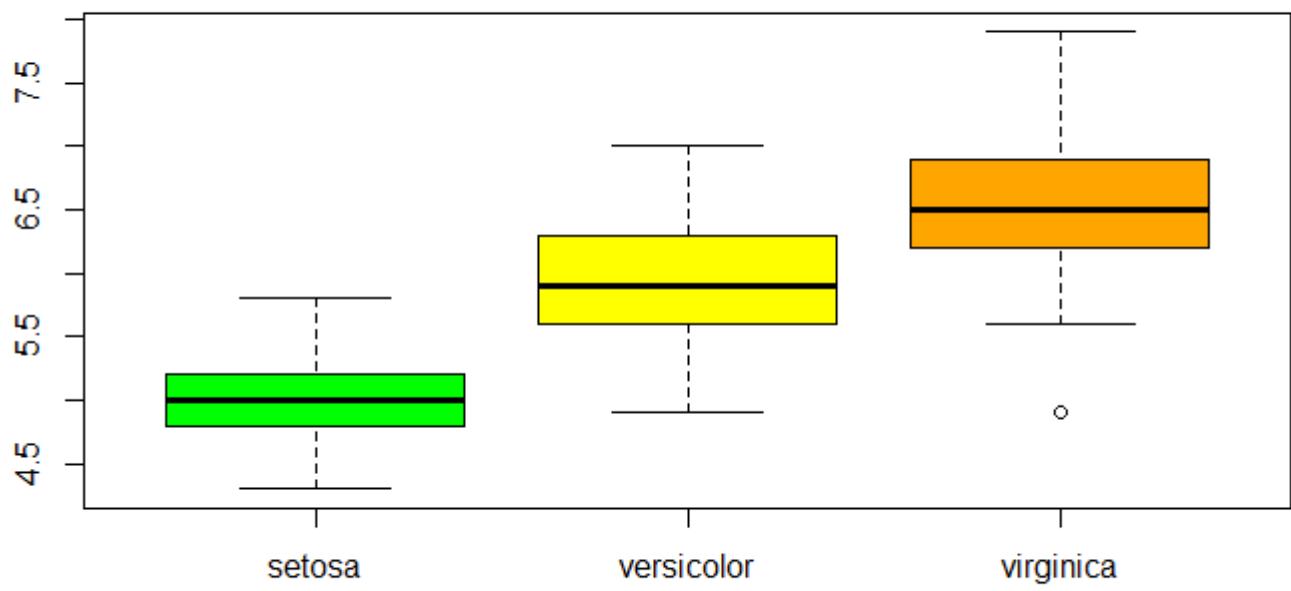
Names . . .

```
boxplot(Sepal.Length~newSpeciesOrder,data = iris,names= c("name1","name2","name3"))
```

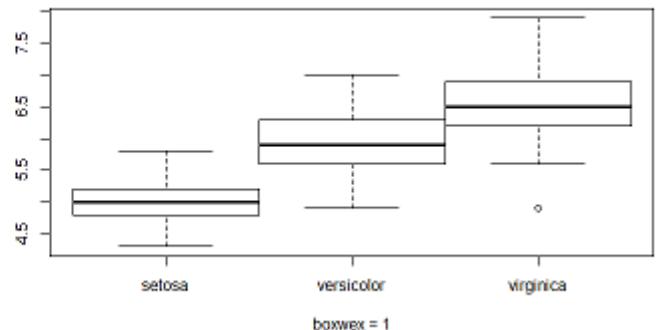
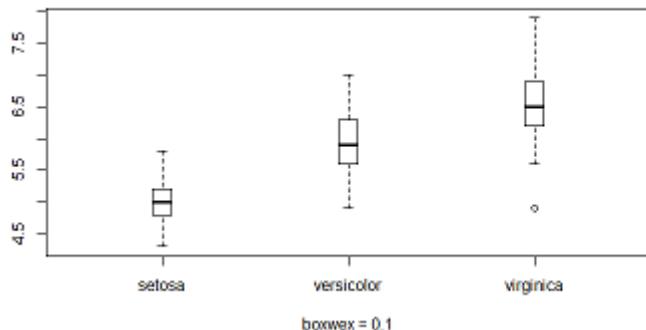


col : .

```
boxplot(Sepal.Length~Species,data = iris,col=c("green","yellow","orange"))
```



```
boxwex: .
boxplot(Sepal.Length~Species,data = iris,boxwex = 0.1)
boxplot(Sepal.Length~Species,data = iris,boxwex = 1)
```



## boxplots plot=FALSE

```
plot FALSE FALSE .
```

```
> boxplot(Sepal.Length~newSpeciesOrder,data = iris,plot=FALSE)
$stats #summary of the numerical variable for the 3 groups
 [,1] [,2] [,3]
[1,] 5.6 4.9 4.3 # extreme value
[2,] 6.2 5.6 4.8 # first quartile limit
[3,] 6.5 5.9 5.0 # median limit
[4,] 6.9 6.3 5.2 # third quartile limit
[5,] 7.9 7.0 5.8 # extreme value

$n #number of observations in each groups
[1] 50 50 50

$conf #extreme value of the notches
 [,1] [,2] [,3]
[1,] 6.343588 5.743588 4.910622
[2,] 6.656412 6.056412 5.089378

$out #extreme value
[1] 4.9

$group #group in which are the extreme value
[1] 1

$names #groups names
[1] "virginica" "versicolor" "setosa"
```

```
boxplot .
```

- **boxlty** -
- **boxlwd** -
- **boxcol** -

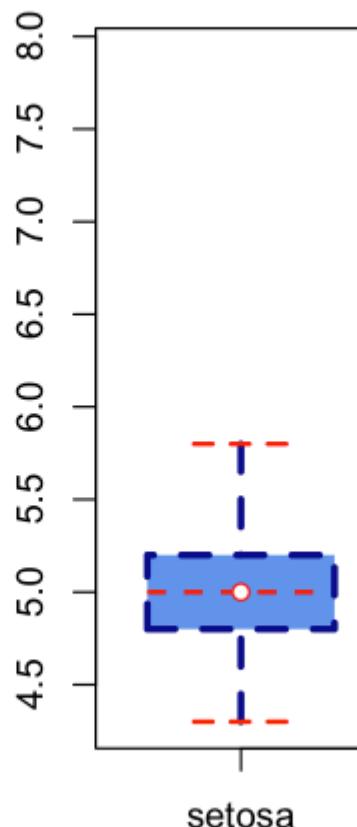
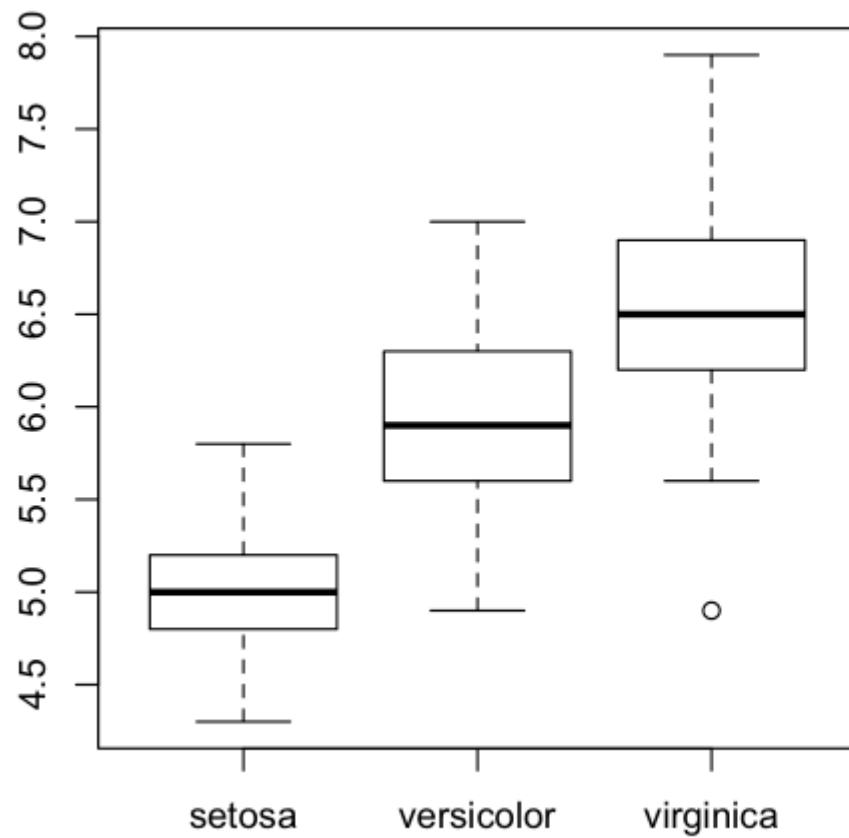
- **boxfill** -
  - **medlty** - ("blank")
  - -
  - **medcol** -
  - **medpch** - ( NA)
  - **medcex** -
  - **medbg** -
- 

- -
  - **whisklwd** -
  - **whiskcol** -
- 

- -
  - **staplelwd** -
  - **staplecol** -
- 

- **outlty** - outlier ("blank")
  - **outlwd** -
  - **outcol** -
  - **outpch** - ( NA)
  - **outcex** -
  - -
- 

```
par(mfrow=c(1,2))
Default
boxplot(Sepal.Length ~ Species, data=iris)
Modified
boxplot(Sepal.Length ~ Species, data=iris,
 boxlty=2, boxlwd=3, boxfill="cornflowerblue", boxcol="darkblue",
 medlty=2, medlwd=2, medcol="red", medpch=21, medcex=1, medbg="white",
 whisklty=2, whisklwd=3, whiskcol="darkblue",
 staplelty=2, staplelwd=2, staplecol="red",
 outlty=3, outlwd=3, outcol="grey", outpch=NA
)
```



: <https://riptutorial.com/ko/r/topic/1005/>

## Examples

:

```
x <- 1:5
x
[1] 1 2 3 4 5
```

```
10:4
[1] 10 9 8 7 6 5 4
```

```
1.25:5
[1] 1.25 2.25 3.25 4.25
```

```
-4:4
#[1] -4 -3 -2 -1 0 1 2 3 4
```

### seq ()

```
seq : 1
```

```
start (1)
```

```
(to)
```

```
seq(5)
[1] 1 2 3 4 5
```

```
seq(2, 5) # or seq(from=2, to=5)
[1] 2 3 4 5
```

```
(by)
```

```
seq(2, 5, 0.5) # or seq(from=2, to=5, by=0.5)
[1] 2.0 2.5 3.0 3.5 4.0 4.5 5.0
```

```
seq (length.out) ()
```

```
seq(2,5, length.out = 10)
[1] 2.0 2.3 2.6 2.9 3.2 3.5 3.8 4.1 4.4 4.7 5.0
```

```
, along.with length.out = length(x) .
```

```
x = 1:8
seq(2,5,along.with = x)
[1] 2.000000 2.428571 2.857143 3.285714 3.714286 4.142857 4.571429 5.000000
```

```
seq seq_along , seq_len seq.int . seq_along seq_len 1 N () . N , seq_along seq_along
seq_len .
```

```
seq_along(x)
[1] 1 2 3 4 5 6 7 8
```

```
seq_along .
```

```
counting numbers 1 through 10
seq_len(10)
[1] 1 2 3 4 5 6 7 8 9 10
indices of existing vector (or list) with seq_along
letters[1:10]
[1] "a" "b" "c" "d" "e" "f" "g" "h" "i" "j"
seq_along(letters[1:10])
[1] 1 2 3 4 5 6 7 8 9 10
```

```
seq.int seq .
```

```
sequence .
```

```
sequence(4)
[1] 1 2 3 4
sequence(c(3, 2))
[1] 1 2 3 1 2
sequence(c(3, 2, 5))
[1] 1 2 3 1 2 1 2 3 4 5
```

```
R (:) . function vector() .
```

```
vector('integer',2) # creates a vector of integers of size 2.
vector('character',2) # creates a vector of characters of size 2.
vector('logical',2) # creates a vector of logicals of size 2.
```

```
R .
```

```
integer(2) # is the same as vector('integer',2) and creates an integer vector with two
elements
character(2) # is the same as vector('integer',2) and creates an character vector with two
elements
logical(2) # is the same as vector('logical',2) and creates an logical vector with two
elements
```

```
. c() . C .
```

```
c(1, 2) # creates a integer vector of two elements: 1 and 2.
c('a', 'b') # creates a character vector of two elements: a and b.
c(T,F) # creates a logical vector of two elements: TRUE and FALSE.
```

```
R (:1) 1 . (:1.1), (:T F) (:'a') . , .
```

```
. R .
```

```
c(1,1.1,'a',T) # all types (integer, numeric, character and logical) are converted to the
'lowest' type which is character.
```

```
[(operator)] .
```

```
vec_int <- c(1,2,3)
vec_char <- c('a','b','c')
vec_int[2] # accessing the second element will return 2
vec_char[2] # accessing the second element will return 'b'
```

```
vec_int[2] <- 5 # change the second value from 2 to 5
vec_int # returns [1] 1 5 3
```

```
: (seq()) .
```

```
vec_int <- 1:10
vec_int # returns [1] 1 2 3 4 5 6 7 8 9 10
```

```
()
```

```
vec_char <- c('a','b','c','d','e')
vec_char[2:4] # returns [1] "b" "c" "d"
vec_char[c(1,3,5)] # returns [1] "a" "c" "e"
```

```
. c :
```

```
xc <- c('a' = 5, 'b' = 6, 'c' = 7, 'd' = 8)
```

```
> xc
a b c d
5 6 7 8
```

```
list :
```

```
xl <- list('a' = 5, 'b' = 6, 'c' = 7, 'd' = 8)
```

```
> xl
$a
[1] 5
```

```
$b
[1] 6
```

```
$c
[1] 7
```

```
$d
[1] 8
```

```
setNames
```

```
x <- 5:8
y <- letters[1:4]

xy <- setNames(x, y)
```

```
> xy
a b c d
5 6 7 8
```

```
, c
```

```
names
```

```
xy <- 5:8
names(xy) <- letters[1:4]
```

```
> xy["c"]
c
7
```

```
/
```

```
mydf <- data.frame(let = c('c', 'a', 'b', 'd'))
```

```
> mydf
let
1 c
2 a
3 b
4 d
```

```
xy num mydf match xy
```

```
mydf$num <- xy[match(mydf$let, names(xy))]
```

```
> mydf
 let num
1 c 7
2 a 5
3 b 6
4 d 8
```

## rep ()

rep .

```
repeat counting numbers, 1 through 5 twice
rep(1:5, 2)
[1] 1 2 3 4 5 1 2 3 4 5
```

```
repeat vector with incomplete recycling
rep(1:5, 2, length.out=7)
[1] 1 2 3 4 5 1 2
```

/ . .

```
same except repeat each integer next to each other
rep(1:5, each=2)
[1] 1 1 2 2 3 3 4 4 5 5
```

---

## rep length

```
automated length repetition
rep(1:5, 1:5)
[1] 1 2 2 3 3 3 4 4 4 4 5 5 5 5 5
hand-fed repetition length vector
rep(1:5, c(1,1,1,2,2))
[1] 1 2 3 4 4 5 5
```

rep .

---

seq rep rep\_len rep.int . rep

```
repeat counting numbers, 1 through 5 twice
rep.int(1:5, 2)
[1] 1 2 3 4 5 1 2 3 4 5
```

```
repeat vector with incomplete recycling
rep_len(1:5, length.out=7)
[1] 1 2 3 4 5 1 2
```

:

R . .

- LETTERS : 26

- letters : 26
- month.abb :
- month.name :
- pi :

1) :

```
> letters
[1] "a" "b" "c" "d" "e" "f" "g" "h" "i" "j" "k" "l" "m" "n" "o" "p" "q" "r" "s" "t" "u" "v"
"w" "x" "y" "z"

> LETTERS[7:9]
[1] "G" "H" "I"

> letters[c(1,5,3,2,4)]
[1] "a" "e" "c" "b" "d"
```

2) :

```
> month.abb
[1] "Jan" "Feb" "Mar" "Apr" "May" "Jun" "Jul" "Aug" "Sep" "Oct" "Nov" "Dec"

> month.name[1:4]
[1] "January" "February" "March" "April"

> month.abb[c(3,6,9,12)]
[1] "Mar" "Jun" "Sep" "Dec"
```

: <https://riptutorial.com/ko/r/topic/1088/>

## Examples

,

```
R . R "" .R (az , Az , 0-9), / (.) (_) .
```

- ( : 1a ), ( .11 ) - ( " ) : `1a` ` .11` . backticks :

```
list('.11' ="a")
#$` .11`
#[1] "a"
```

- ,
- . ls() -function

. foobar , foo.bar , foo\_bar , .foobar

```
R <- . = . . a<-1 (a < -1) (a <- 1) .
```

```
> foo <- 42
> fooEquals = 43
```

foo 42 . foo fooEquals 42 , fooEquals 43 .

```
> foo
[1] 42
> fooEquals
[1] 43
```

x .

```
> (x <- 5)
[1] 5
actually two function calls: first one to `<-` ; second one to the `()`-function
> is.function(``)
[1] TRUE # Often used in R help page examples for its side-effect of printing.
```

-> .

```
> 5 -> x
> x
[1] 5
>
```

R . 1 .

```
• : . : . v <- c(2, 3, 7, 10) v2 <- c("a", "b", "c") .
• : , . a <- matrix(data = c(1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12), nrow = 4, ncol = 3,
 byrow = F) . . a[1,2] , [1] 5 .
• : mylist <- list (course = 'stat', date = '04/07/2009', num_isc = 7, num_cons = 6, num_mat
 = as.character(c(45020, 45679, 46789, 43126, 42345, 47568, 45674)), results = c(30, 19, 29,
 NA, 25, 26, 27)) . () . mylist$results mylist[[6]] . : mylist[6] R .
 mylist[[6]][2] (19), mylist[6][2] .
• data.frame : . . exam <- data.frame(matr = as.character(c(45020, 45679, 46789, 43126,
 42345, 47568, 45674)), res_S = c(30, 19, 29, NA, 25, 26, 27), res_O = c(3, 3, 1, NA, 3, 2,
 NA), res_TOT = c(30, 22, 30, NA, 28, 28, 27)) . exam$matr, exam[, 'matr'] exam[1], exam[,1] .
 exam['rowname',] exam[1,] . (rownames-attribute) .

. ?Syntax . (R) . . :
```

```
> a <- 1
> b <- 2
> c <- c(2,3,4)
> d <- c(10,10,10)
> e <- c(1,2,3,4)
> f <- 1:6
> W <- cbind(1:4,5:8,9:12)
> Z <- rbind(rep(0,3),1:3,rep(10,3),c(4,7,1))
```

```
> a+b # scalar + scalar
[1] 3
> c+d # vector + vector
[1] 12 13 14
> a*b # scalar * scalar
[1] 2
> c*d # vector * vector (componentwise!)
[1] 20 30 40
> c+a # vector + scalar
[1] 3 4 5
> c^2 #
[1] 4 9 16
> exp(c)
[1] 7.389056 20.085537 54.598150
```

!

```
> c+e # warning but.. no errors, since recycling is assumed to be desired.
[1] 3 5 7 6
Warning message:
In c + e : longer object length is not a multiple of shorter object length
```

R . . . c+f # .

!

```
> Z+W # matrix + matrix #(componentwise)
> Z*W # matrix* matrix#(Standard product is always componentwise)
```

: V % \* % W

```
> W + a # matrix+ scalar is still componentwise
 [,1] [,2] [,3]
[1,] 2 6 10
[2,] 3 7 11
[3,] 4 8 12
[4,] 5 9 13

> W + c # matrix + vector... : no warnings and R does the operation in a column-wise manner
 [,1] [,2] [,3]
[1,] 3 8 13
[2,] 5 10 12
[3,] 7 9 14
[4,] 6 11 16
```

"

R

```
> foo <- 'foo'
> .foo <- 'bar'
```

ls

```
> ls()
[1] "foo"
```

all.names = TRUE 'private' .

```
> ls(all.names = TRUE)
[1] ".foo" "foo"
```

: <https://riptutorial.com/ko/r/topic/9013/>

## Examples

( ).

```
x <- 1

foo <- function(x) {
 y <- 3
 z <- x + y
 return(z)
}
```

y

: 'y' () .

```
foo <- function(x) {
 x <- 2
 y <- 3
 z <- x + y
 return(z)
}
```

```
foo(1)
x
```

5

1

```
foo <- function() {
 y <- 3
 z <- x + y
 return(z)
}

foo()
```

4

( ) .

:

```
bar <- function() {
 z <- x + y
 return(z)
}

foo <- function() {
 y <- 3
 z <- bar()
 return(z)
}

foo()
```

bar () : 'y' .

:

```
foo <- function() {

 bar <- function() {
 z <- x + y
 return(z)
 }

 y <- 3
 z <- bar()
 return(z)
}

foo()
```

4

<<- .. bar() y ..

```
bar <- function() {
 z <- x + y
 return(z)
}

foo <- function() {
 y <<- 3
 z <- bar()
 return(z)
}

foo()
```

4

R

package:base package:base ..

```
e1 <- new.env(parent = baseenv())
e2 <- new.env(parent = e1)
```

```
assign("a", 3, envir = e1)
get("a", envir = e1)
get("a", envir = e2)
```

```
e2 e1 e1 e2 a 3 . , a e2 a e1 .
```

```
assign("a", 2, envir = e2)
get("a", envir = e2)
get("a", envir = e1)
```

2

```
on.exit()
```

```
,
```

```
new_plot <- function(...) {

 old_pars <- par(mar = c(5,4,4,2) + .1, mfrow = c(1,1))
 on.exit(par(old_pars))
 plot(...)
}
```

"

```
library(plyr)
library(dplyr)
```

: 'dplyr'

'package : plyr' .

, , , , ,

'package : stats' .

,

'package : base' .

intersect, setdiff, setequal, union

```
package::function()
```

: <https://riptutorial.com/ko/r/topic/3138/>

```
. package::function() . caret , pls plyr .
Microsoft R Open (Revolution R) BLAS / LAPACK .
```

## Examples

### foreach

```
foreach R . CPU . doSNOW .
```

```
foreach 1 100000 .
```

```
library(foreach)
library(doSNOW)

cl <- makeCluster(5, type = "SOCK")
registerDoSNOW(cl)

f <- foreach(i = 1:100000, .combine = c, .inorder = F) %dopar% {
 k <- i ** 2 + sqrt(i)
 k
}
```

```
foreach .combine . . c . "+" () .
```

```
foreach-loop . k .
```

.	.	c , cbind , rbind , "+" , "*" ....
.inorder	TRUE	( i ) . FALSE . .
	base , mass , randomForest	c ("mass", "randomForest") .

```
parallel , .
```

```
localhost .
```

```
parallel::detectCores(all.tests = FALSE, logical = TRUE)
```

```
localhost .
```

```
parallelCluster <- parallel::makeCluster(parallel::detectCores())
```

```
, . mtcars . cyl mpg .
```

```

data <- mtcars
yfactor <- 'cyl'
zlevels <- sort(unique(data[[yfactor]]))
datay <- data[,1]
dataz <- data[,2]
datax <- data[,3:11]

fitmodel <- function(zlevel, datax, datay, dataz) {
 glm.fit(x = datax[dataz == zlevel,], y = datay[dataz == zlevel])
}

zlevels . , .

fitmodel <- function(zlevel, datax, datay, dataz) {
 glm.fit(x = datax[dataz == zlevel,], y = datay[dataz == zlevel])
}

for (zlevel in zlevels) {
 print("*****")
 print(zlevel)
 print(fitmodel(zlevel, datax, datay, dataz))
}

:
:

worker <- function(zlevel) {
 fitmodel(zlevel,datax, datay, dataz)
}

parallel ., parallel . . .

wrapper <- function(datax, datay, dataz) {
 # force evaluation of all paramters not supplied by parallelization apply
 force(datax)
 force(datay)
 force(dataz)
 # these variables are now in an enviroment accessible by parallel function

 # function to be applied also in the environment
 fitmodel <- function(zlevel, datax, datay, dataz) {
 glm.fit(x = datax[dataz == zlevel,], y = datay[dataz == zlevel])
 }

 # calling in this environment iterating over single parameter zlevel
 worker <- function(zlevel) {
 fitmodel(zlevel,datax, datay, dataz)
 }
 return(worker)
}

parallelcluster <- parallel::makeCluster(parallel::detectCores())
models <- parallel::parLapply(parallelcluster,zlevels,

```

```
wrapper(datax, datay, dataz))
```

```
parallel::stopCluster(parallelcluster)
```

```
parallel par apply() .
```

```
RNG . set.seed() . !
```

```
. (parallel, snow) .
```

```
s <- seed
for (i in 1:numofcores) {
 s <- nextRNGStream(s)
 # send s to worker i as .Random.seed
}
```

```
clusterSetRNGStream(cl = parallelcluster, iseed)
```

## mcparallelDo

```
mcparallelDo Unix (Linux MacOSX) R . . . future .
```

```

```

```
data(ToothGrowth)
```

```
mcparallelDo .
```

```
mcparallelDo({glm(len ~ supp * dose, data=ToothGrowth)}, "interactionPredictorModel")
```

```
binaryPredictorModel <- glm(len ~ supp, data=ToothGrowth)
gaussianPredictorModel <- glm(len ~ dose, data=ToothGrowth)
```

```
mcparallelDo targetEnvironment (: .GlobalEnv) () .
```

```
summary(interactionPredictorModel)
```

```

```

```
Example of not returning a value until we return to the top level
for (i in 1:10) {
 if (i == 1) {
 mcparallelDo({2+2}, targetValue = "output")
 }
}
```

```
if (exists("output")) print(i)
}

Example of getting a value without returning to the top level
for (i in 1:10) {
 if (i == 1) {
 mcpParallelDo({2+2}, targetValue = "output")
 }
 mcpParallelDoCheck()
 if (exists("output")) print(i)
}
```

: <https://riptutorial.com/ko/r/topic/1677/>

# 81:

R , ?Distributions .

4 .

- d -
- p -
- q - .
- r -

R ?Distributions .

## Examples

\*norm . :

```
dnorm(x, mean = 0, sd = 1, log = FALSE)
pnorm(q, mean = 0, sd = 1, lower.tail = TRUE, log.p = FALSE)
qnorm(p, mean = 0, sd = 1, lower.tail = TRUE, log.p = FALSE)
rnorm(n, mean = 0, sd = 1)
```

0

```
dnorm(0)
```

0.3989423 .

pnorm(0) .5 . , 0 .

qnorm pnorm pnorm . qnorm(.5) 0 .

, rnorm .

```
rnorm(10)
```

10 .

.

rnorm(10, mean=4, sd= 3)

rbinom dbinom, pbinom, qbinom rbinom .

dbinom() . . x quantile( x ) . defining parameters , n( ) p( ). , n = 5 , p = 0.5  
X 0,1,2,3,4,5 . , dbinom(x,n,p) x = 0, 1, 2, 3, 4, 5 P( X = x ) .

```
#Binom(n = 5, p = 0.5) probabilities
```

```

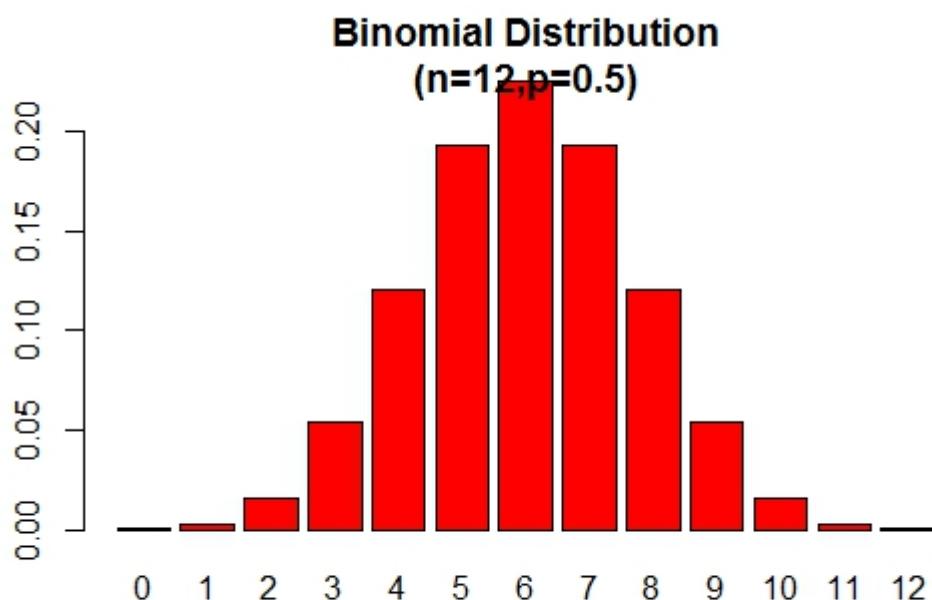
> n <- 5; p<- 0.5; x <- 0:n
> dbinom(x,n,p)
[1] 0.03125 0.15625 0.31250 0.31250 0.15625 0.03125
#To verify the total probability is 1
> sum(dbinom(x,n,p))
[1] 1
>

```

```

> x <- 0:12
> prob <- dbinom(x,12,.5)
> barplot(prob,col = "red",ylim = c(0,.2),names.arg=x,
 main="Binomial Distribution\n(n=12,p=0.5)")

```



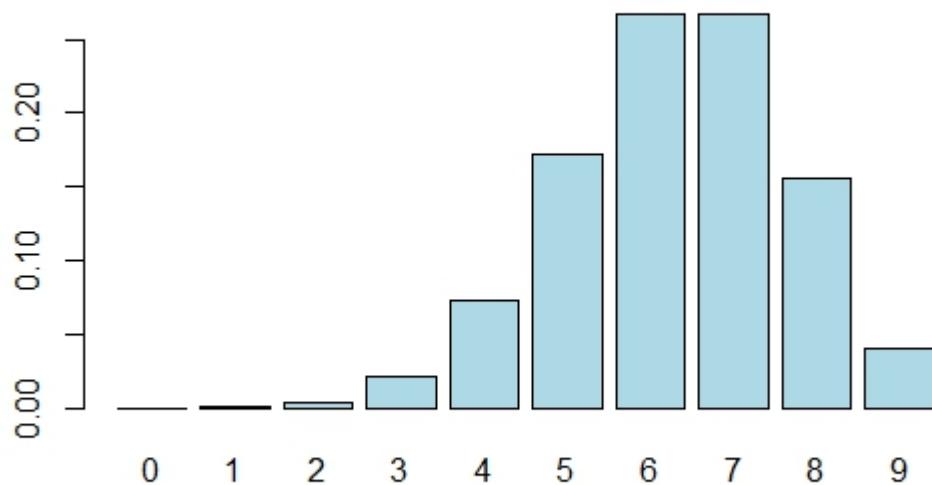
p = 0.5 . p 0.5

```

> n=9; p=.7; x=0:n; prob=dbinom(x,n,p);
> barplot(prob,names.arg = x,main="Binomial Distribution\n(n=9, p=0.7)",col="lightblue")

```

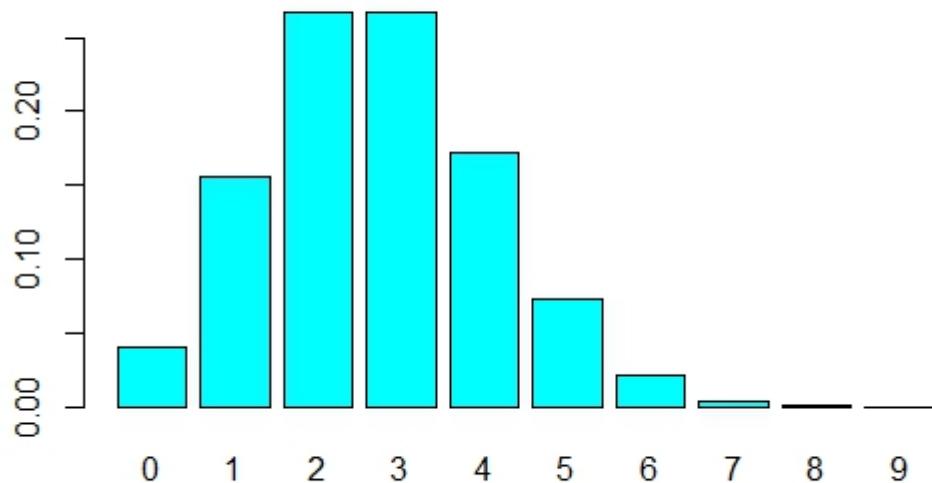
### Binomial Distribution (n=9, p=0.7)



p 0.5 .

```
> n=9; p=.3; x=0:n; prob=dbinom(x,n,p);
> barplot(prob,names.arg = x,main="Binomial Distribution\n(n=9, p=0.3)",col="cyan")
```

### Binomial Distribution (n=9, p=0.3)



pbinom() . P( X <= x ) . x quantile (x) .

```
Calculating Probabilities
P(X <= 2) in a Bin(n=5,p=0.5) distribution
> pbinom(2,5,0.5)
```

```
[1] 0.5
```

```
P(X <= 2) = P(X=0) + P(X=1) + P(X=2)
> sum(dbinom(0:2, 5, 0.5))
[1] 0.5
```

```
, : P(a <= X <= b)
```

```
P(3 <= X <= 5) = P(X=3) + P(X=4) + P(X=5) in a Bin(n=9, p=0.6) dist
> sum(dbinom(c(3, 4, 5), 9, 0.6))
[1] 0.4923556
>
```

```
:
```

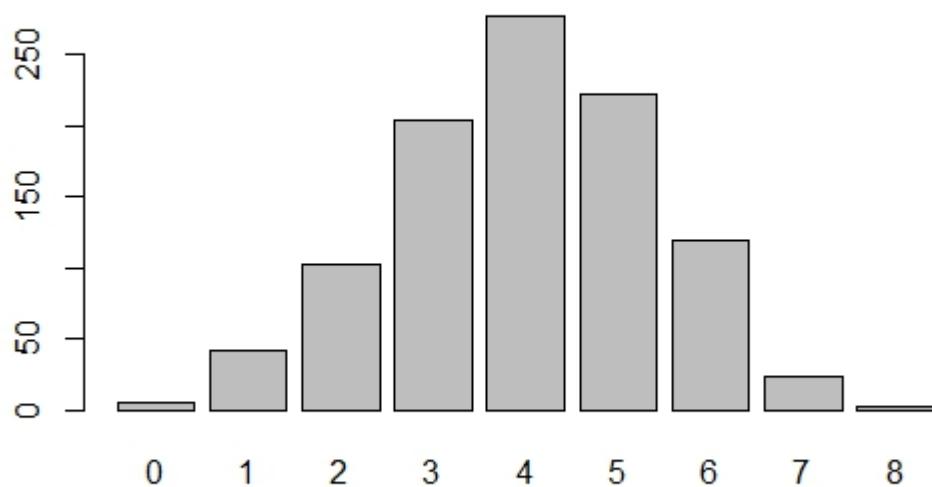
  

```
> n = 10; p = 0.4; x = 0:n;
> prob = dbinom(x, n, p)
> cdf = pbinom(x, n, p)
> distTable = cbind(x, prob, cdf)
> distTable
 x prob cdf
[1,] 0 0.0060466176 0.006046618
[2,] 1 0.0403107840 0.046357402
[3,] 2 0.1209323520 0.167289754
[4,] 3 0.2149908480 0.382280602
[5,] 4 0.2508226560 0.633103258
[6,] 5 0.2006581248 0.833761382
[7,] 6 0.1114767360 0.945238118
[8,] 7 0.0424673280 0.987705446
[9,] 8 0.0106168320 0.998322278
[10,] 9 0.0015728640 0.999895142
[11,] 10 0.0001048576 1.0000000000
>
```

```
rbinom()
```

```
Simulation
> xVal<-names(table(rbinom(1000, 8, .5)))
> barplot(as.vector(table(rbinom(1000, 8, .5))), names.arg =xVal,
 main="Simulated Binomial Distribution\n (n=8, p=0.5)")
```

### Simulated Binomial Distribution (n=8,p=0.5)



: <https://riptutorial.com/ko/r/topic/1885/>

## Examples

```
split data.frame . (for lapply / sapply).
```

```
split .
```

```
testdata <- c("e", "o", "r", "g", "a", "y", "w", "q", "i", "s", "b", "v", "x", "h", "u")
```

```
voyels consonants . , .
```

```
vowels <- c('a','e','i','o','u','y')
letter_type <- ifelse(testdata %in% vowels, "vowels", "consonants")
```

```
letter_type testdata . vowels consonants split .
```

```
split(testdata, letter_type)
#$consonants
#[1] "r" "g" "w" "q" "s" "b" "v" "x" "h"
#$vowels
#[1] "e" "o" "a" "y" "i" "u"
```

```
/ letter_type .
```

```
split data.frames .
```

```
iris .
```

```
data(iris)
```

```
split , data.frame (: Species) :
```

```
> liris <- split(iris, iris$Species)
> names(liris)
[1] "setosa" "versicolor" "virginica"
> head(liris$setosa)
 Sepal.Length Sepal.Width Petal.Length Petal.Width Species
1 5.1 3.5 1.4 0.2 setosa
2 4.9 3.0 1.4 0.2 setosa
3 4.7 3.2 1.3 0.2 setosa
4 4.6 3.1 1.5 0.2 setosa
5 5.0 3.6 1.4 0.2 setosa
6 5.4 3.9 1.7 0.4 setosa
```

(setosa ).

. lapply lapply .

```
> (lcor <- lapply(liris, FUN=function(df) cor(df[,1:4])))

$setosa
 Sepal.Length Sepal.Width Petal.Length Petal.Width
Sepal.Length 1.0000000 0.7425467 0.2671758 0.2780984
Sepal.Width 0.7425467 1.0000000 0.1777000 0.2327520
Petal.Length 0.2671758 0.1777000 1.0000000 0.3316300
Petal.Width 0.2780984 0.2327520 0.3316300 1.0000000

$versicolor
 Sepal.Length Sepal.Width Petal.Length Petal.Width
Sepal.Length 1.0000000 0.5259107 0.7540490 0.5464611
Sepal.Width 0.5259107 1.0000000 0.5605221 0.6639987
Petal.Length 0.7540490 0.5605221 1.0000000 0.7866681
Petal.Width 0.5464611 0.6639987 0.7866681 1.0000000

$virginica
 Sepal.Length Sepal.Width Petal.Length Petal.Width
Sepal.Length 1.0000000 0.4572278 0.8642247 0.2811077
Sepal.Width 0.4572278 1.0000000 0.4010446 0.5377280
Petal.Length 0.8642247 0.4010446 1.0000000 0.3221082
Petal.Width 0.2811077 0.5377280 0.3221082 1.0000000
```

( / )

```
> library(reshape)
> (topcor <- lapply(lcor, FUN=function(cormat) {
 correlations <- melt(cormat, variable_name="correlation");
 filtered <- correlations[correlations$X1 != correlations$X2,];
 filtered[which.max(filtered$correlation),]
}))

$setosa
 X1 X2 correlation
2 Sepal.Width Sepal.Length 0.7425467

$versicolor
 X1 X2 correlation
12 Petal.Width Petal.Length 0.7866681

$virginica
 X1 X2 correlation
3 Petal.Length Sepal.Length 0.8642247
```

```
> (result <- do.call("rbind", topcor))

 X1 X2 correlation
setosa Sepal.Width Sepal.Length 0.7425467
versicolor Petal.Width Petal.Length 0.7866681
virginica Petal.Length Sepal.Length 0.8642247
```

## split-apply-combine split

```
-- , .
mtcars cyl mpg . mtcars mtcars .

(spl <- split(mtcars, mtcars$cyl))
$`4`
mpg cyl disp hp drat wt qsec vs am gear carb
Datsun 710 22.8 4 108.0 93 3.85 2.320 18.61 1 1 4 1
Merc 240D 24.4 4 146.7 62 3.69 3.190 20.00 1 0 4 2
Merc 230 22.8 4 140.8 95 3.92 3.150 22.90 1 0 4 2
Fiat 128 32.4 4 78.7 66 4.08 2.200 19.47 1 1 4 1
...

$`6`
mpg cyl disp hp drat wt qsec vs am gear carb
Mazda RX4 21.0 6 160.0 110 3.90 2.620 16.46 0 1 4 4
Mazda RX4 Wag 21.0 6 160.0 110 3.90 2.875 17.02 0 1 4 4
Hornet 4 Drive 21.4 6 258.0 110 3.08 3.215 19.44 1 0 3 1
Valiant 18.1 6 225.0 105 2.76 3.460 20.22 1 0 3 1
...

$`8`
mpg cyl disp hp drat wt qsec vs am gear carb
Hornet Sportabout 18.7 8 360.0 175 3.15 3.440 17.02 0 0 3 2
Duster 360 14.3 8 360.0 245 3.21 3.570 15.84 0 0 3 4
Merc 450SE 16.4 8 275.8 180 3.07 4.070 17.40 0 0 3 3
Merc 450SL 17.3 8 275.8 180 3.07 3.730 17.60 0 0 3 3
...

. spl$`4` , spl$`6` spl$`8` (spl$"4" spl$"4" spl[["4"]]).

lapply 2 mpg .

(best2 <- lapply(spl, function(x) tail(x[order(x$mpg),], 2)))
$`4`
mpg cyl disp hp drat wt qsec vs am gear carb
Fiat 128 32.4 4 78.7 66 4.08 2.200 19.47 1 1 4 1
Toyota Corolla 33.9 4 71.1 65 4.22 1.835 19.90 1 1 4 1

$`6`
mpg cyl disp hp drat wt qsec vs am gear carb
Mazda RX4 Wag 21.0 6 160 110 3.90 2.875 17.02 0 1 4 4
Hornet 4 Drive 21.4 6 258 110 3.08 3.215 19.44 1 0 3 1

$`8`
mpg cyl disp hp drat wt qsec vs am gear carb
Hornet Sportabout 18.7 8 360 175 3.15 3.440 17.02 0 0 3 2
Pontiac Firebird 19.2 8 400 175 3.08 3.845 17.05 0 0 3 2

rbind . rbind(best2[["4"]], best2[["6"]], best2[["8"]]) , . . .

do.call(rbind, best2)
mpg cyl disp hp drat wt qsec vs am gear carb
4.Fiat 128 32.4 4 78.7 66 4.08 2.200 19.47 1 1 4 1
```

```
4.Toyota Corolla 33.9 4 71.1 65 4.22 1.835 19.90 1 1 4 1
6.Mazda RX4 Wag 21.0 6 160.0 110 3.90 2.875 17.02 0 1 4 4
6.Hornet 4 Drive 21.4 6 258.0 110 3.08 3.215 19.44 1 0 3 1
8.Hornet Sportabout 18.7 8 360.0 175 3.15 3.440 17.02 0 0 3 2
8.Pontiac Firebird 19.2 8 400.0 175 3.08 3.845 17.05 0 0 3 2
```

```
best2(2) rbind(1,) .
```

```
do.call(rbind, lapply(split(mtcars, mtcars$cyl), function(x) tail(x[order(x$mpg),], 2)))
```

```
lapply(split(x,f), FUN) ?by lapply(split(x,f), FUN) .
```

```
by(mtcars, mtcars$cyl, function(x) tail(x[order(x$mpg),], 2))
do.call(rbind, by(mtcars, mtcars$cyl, function(x) tail(x[order(x$mpg),], 2)))
```

: <https://riptutorial.com/ko/r/topic/1073/>

Drier R (SE) (NSE) 1 .

```
, summarise() summarise_()
```

lazyeval NSE .

## Examples

### dplyr

NSE . SE .

dplyr lazyeval :

```
library(dplyr)
library(lazyeval)
```

### NSE

```
filter(mtcars, cyl == 8)
filter(mtcars, cyl < 6)
filter(mtcars, cyl < 6 & vs == 1)
```

### SE ( )

```
filter_(mtcars, .dots = list(~ cyl == 8))
filter_(mtcars, .dots = list(~ cyl < 6))
filter_(mtcars, .dots = list(~ cyl < 6, ~ vs == 1))
```

### NSE

```
summarise(mtcars, mean(disp))
summarise(mtcars, mean_disp = mean(disp))
```

### SE

```
summarise_(mtcars, .dots = lazyeval::interp(~ mean(x), x = quote(disp)))
summarise_(mtcars, .dots = setNames(list(lazyeval::interp(~ mean(x), x = quote(disp))),
"mean_disp"))
summarise_(mtcars, .dots = list("mean_disp" = lazyeval::interp(~ mean(x), x = quote(disp))))
```

### NSE

```
mutate(mtcars, displ_l = disp / 61.0237)
```

### SE

```
mutate_(
 .data = mtcars,
 .dots = list(
 "displ_1" = lazyeval::interp(
 ~ x / 61.0237, x = quote(disp)
)
)
)
```

: <https://riptutorial.com/ko/r/topic/9365/>----

# 84:

## Examples

Shiny RStudio R R

Shiny

- .R
- : ui.R server.R

Shiny

- **ui** :
- **(server)** :

```
library(shiny)

Create the UI
ui <- shinyUI(fluidPage(
 # Application title
 titlePanel("Hello World!"))
))

Create the server function
server <- shinyServer(function(input, output){})

Run the app
shinyApp(ui = ui, server = server)
```

**ui.R**

```
library(shiny)

Define UI for application
shinyUI(fluidPage(
 # Application title
 titlePanel("Hello World!"))
))
```

**server.R**

```
library(shiny)

Define server logic
shinyServer(function(input, output){})
```

- : ( 0))
- :
- 

## HTML .

```
library(shiny)

ui <- fluidPage(
 radioButtons("radio",
 label = HTML('Welcome

Your favorite color is red ?'),
 choices = list("TRUE" = 1, "FALSE" = 2),
 selected = 1,
 inline = T,
 width = "100%"),
 fluidRow(column(3, textOutput("value"))))

server <- function(input, output){
 output$value <- renderPrint({
 if(input$radio == 1){return('Great !')}
 else{return("Sorry !")}})}

shinyApp(ui = ui, server = server)
```

## Welcome

Your favorite color is red ?

TRUE  FALSE

[1] "Great!"

```
library(shiny)

ui <- fluidPage(
 checkboxGroupInput("checkGroup1", label = h3("This is a Checkbox group"),
 choices = list("1" = 1, "2" = 2, "3" = 3),
 selected = 1),
 fluidRow(column(3, verbatimTextOutput("text_choice")))
)

server <- function(input, output){
 output$text_choice <- renderPrint({
 return(paste0("You have chosen the choice ", input$checkGroup1))})
}

shinyApp(ui = ui, server = server)
```

## This is a Checkbox group

- 1
- 2
- 3

```
[1] "You have chosen the choice 1"
```

- :
- :
- selected : ( NULL)
- :
- :

HTML .

```
library(shiny)

ui <- fluidPage(
 selectInput("id_selectInput",
 label = HTML('What is your favorite color ?'),
 multiple = TRUE,
 choices = list("red" = "red", "green" = "green", "blue" = "blue", "yellow" =
"yellow"),
 selected = NULL),
 br(), br(),
 fluidRow(column(3, textOutput("text_choice"))))

server <- function(input, output){
 output$text_choice <- renderPrint({
 return(input$id_selectInput)})
}

shinyApp(ui = ui, server = server)
```

What is your favorite color ?

red green blue

yellow

```
[1] "red" "green" "blue"
```

- :
- selected : ( NULL)
- multiple : TRUE FALSE
- 
- 
- selectize : TRUE FALSE (select.js )

HTML .

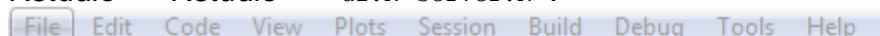
. ui.R server.R .

## 1.

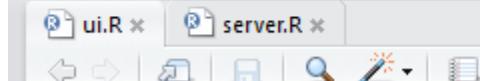
ui.R server.R . shinyApp() Shiny .

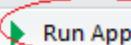
```
shinyApp("path_to_the_folder_containing_the_files")
```

Rstudio Rstudio ui.R server.R .

File Edit Code View Plots Session Build Debug Tools Help

 Go to file/function |  Addins

 ui.R \* server.R \*



Shiny App runApp() .

## 2.

R shinyApp() .

- :

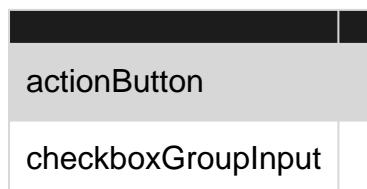
```
library(shiny)

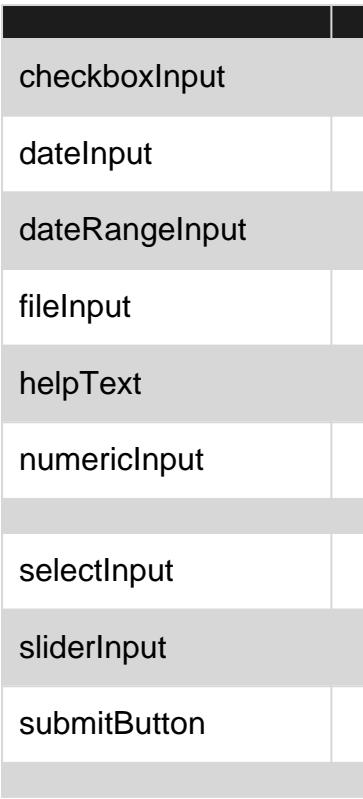
ui <- fluidPage() #Create the ui
server <- function(input, output){} #create the server

shinyApp(ui = ui, server = server) #run the App
```

- appFile Shiny .R .

```
shinyApp(appFile="path_to_my_R_file_containig_the_app")
```





```
library(shiny)

Create the UI
ui <- shinyUI(fluidPage(
 titlePanel("Basic widgets"),

 fluidRow(

 column(3,
 h3("Buttons"),
 actionButton("action", label = "Action"),
 br(),
 br(),
 submitButton("Submit")),

 column(3,
 h3("Single checkbox"),
 checkboxInput("checkbox", label = "Choice A", value = TRUE)),

 column(3,
 checkboxGroupInput("checkGroup",
 label = h3("Checkbox group"),
 choices = list("Choice 1" = 1,
 "Choice 2" = 2, "Choice 3" = 3),
 selected = 1)),

 column(3,
 dateInput("date",
 label = h3("Date input"),
 value = "2014-01-01"))

),

 fluidRow(
 column(3,
 dateRangeInput("dates", label = h3("Date range")))
)
))
```

```

column(3,
 fileInput("file", label = h3("File input")),

column(3,
 h3("Help text"),
 helpText("Note: help text isn't a true widget,",
 "but it provides an easy way to add text to",
 "accompany other widgets."),

column(3,
 numericInput("num",
 label = h3("Numeric input"),
 value = 1))

),

fluidRow(

column(3,
 radioButtons("radio", label = h3("Radio buttons"),
 choices = list("Choice 1" = 1, "Choice 2" = 2,
 "Choice 3" = 3), selected = 1)),

column(3,
 selectInput("select", label = h3("Select box"),
 choices = list("Choice 1" = 1, "Choice 2" = 2,
 "Choice 3" = 3), selected = 1)),

column(3,
 sliderInput("slider1", label = h3("Sliders"),
 min = 0, max = 100, value = 50),
 sliderInput("slider2", "",
 min = 0, max = 100, value = c(25, 75))
),

column(3,
 textInput("text", label = h3("Text input"),
 value = "Enter text..."))

)

))

Create the server function
server <- shinyServer(function(input, output){})

Run the app
shinyApp(ui = ui, server = server)

```

debug() debugonce() Shiny . browser() Shiny . : browser()

---

Showcase server.R .

- runApp("MyApp", display.mode = "showcase") runApp("MyApp", display.mode = "showcase") = "showcase" .
- Shiny app DESCRIPTION DisplayMode: Showcase DisplayMode: Showcase .

## Reactive Log Visualizer

options(shiny.reactlog=TRUE)  
F3 . Reactive Log Visualizer

. Reactive Log Visualizer R options(shiny.reactlog=TRUE)  
. Reactive Log Visualizer Windows Ctrl + F3 Mac Command +  
.

: <https://riptutorial.com/ko/r/topic/2044/>

# 85:

- `<- readline(prompt = " ")`
- `name <- readline(prompt = " ?")`

## Examples

R

R R

```
readline .
```

```
name <- readline(prompt = "What is your name?")
```

, , . :

```
result <- readline(prompt = "What is the result of 1+1?")
while(result!=2) {
 readline(prompt = "Wrong answer. What is the result of 1+1?")
}
```

. . .

```
as.numeric .

result <- as.numeric(readline(prompt = "What is the result of 1+1?"))
while(result!=2) {
 readline(prompt = "Wrong answer. What is the result of 1+1?")
}
```

: <https://riptutorial.com/ko/r/topic/5098-->

# 86:

```
R . , + function (e1, e2) .Primitive("+") function (e1, e2) .Primitive("+") e1 e2 . , +
.
:
`+` <- function(e1, e2) {e1-e2}
> 3+10
[1] -7
```

## Examples

( ).

```
3+1:5
```

:

```
[1] 4 5 6 7 8
```

: + .

- 3+1:5
- 3+c(1, 2, 3, 4, 5) .
- c(4, 5, 6, 7, 8) 3 .

R ( ) :

```
(3+1):5
```

R .

```
[1] 4 5
```

( ) .

1.

+ - :

```
> 3 + 4.5
[1] 7.5
> 3 + 4.5 + 2
```

```

[1] 9.5
> 3 + 4.5 + 2 - 3.8
[1] 5.7
> 3 + NA
#[1] NA
> NA + NA
#[1] NA
> NA - NA
#[1] NA
> NaN - NA
#[1] NaN
> NaN + NA
#[1] NaN

```

( ) .

```

> a <- 3; B <- 4.5; cc <- 2; Dd <- 3.8 ;na<-NA;nan<-NaN
> a + B
[1] 7.5
> a + B + cc
[1] 9.5
> a + B + cc - Dd
[1] 5.7
> B-nan
#[1] NaN
> a+na-na
#[1] NA
> a + na
#[1] NA
> B-nan
#[1] NaN
> a+na-na
#[1] NA

```

2.

```

> A <- c(3, 4.5, 2, -3.8);
> A
[1] 3.0 4.5 2.0 -3.8
> A + 2 # Adding a number
[1] 5.0 6.5 4.0 -1.8
> 8 - A # number less vector
[1] 5.0 3.5 6.0 11.8
> n <- length(A) #number of elements of vector A
> n
[1] 4
> A[-n] + A[n] # Add the last element to the same vector without the last element
[1] -0.8 0.7 -1.8
> A[1:2] + 3 # vector with the first two elements plus a number
[1] 6.0 7.5
> A[1:2] - A[3:4] # vector with the first two elements less the vector with elements 3 and 4
[1] 1.0 8.3

```

sum .

```
> sum(A)
[1] 5.7
> sum(-A)
[1] -5.7
> sum(A[-n]) + A[n]
[1] 5.7
```

R . . . ()

```
> B <- c(3, 5, -3, 2.7, 1.8)
> B
[1] 3.0 5.0 -3.0 2.7 1.8
> A
[1] 3.0 4.5 2.0 -3.8
> A + B # the first element of A is repeated
[1] 6.0 9.5 -1.0 -1.1 4.8
Warning message:
In A + B : longer object length is not a multiple of shorter object length
> B - A # the first element of A is repeated
[1] 0.0 0.5 -5.0 6.5 -1.2
Warning message:
In B - A : longer object length is not a multiple of shorter object length
```

```
> B[1:n] + A
[1] 6.0 9.5 -1.0 -1.1
> B[1:n] - A
[1] 0.0 0.5 -5.0 6.5
```

sum .

```
> sum(A, B)
[1] 15.2
> sum(A, -B)
[1] -3.8
> sum(A)+sum(B)
[1] 15.2
> sum(A)-sum(B)
[1] -3.8
```

: <https://riptutorial.com/ko/r/topic/4389/>

## Examples

### randomForestSRC

[CRAN](#) randomForestSRC , .

```
require(randomForestSRC)

set.seed(130948) #Other seeds give similar comparative results
x1 <- runif(1000)
y <- rnorm(1000, mean = x1, sd = .3)
data <- data.frame(x1 = x1, y = y)
head(data)
```

	x1	y
1	0.9604353	1.3549648
2	0.3771234	0.2961592
3	0.7844242	0.6942191
4	0.9860443	1.5348900
5	0.1942237	0.4629535
6	0.7442532	-0.0672639

```
(modRFSRC <- rfsrc(y ~ x1, data = data, ntree=500, nodesize = 5))
```

```
Sample size: 1000
Number of trees: 500
Minimum terminal node size: 5
Average no. of terminal nodes: 208.258
No. of variables tried at each split: 1
 Total no. of variables: 1
 Analysis: RF-R
 Family: regr
 Splitting rule: mse
 % variance explained: 32.08
 Error rate: 0.11
```

```
x1new <- runif(10000)
ynew <- rnorm(10000, mean = x1new, sd = .3)
newdata <- data.frame(x1 = x1new, y = ynew)

survival.results <- predict(modRFSRC, newdata = newdata)
survival.results
```

```
Sample size of test (predict) data: 10000
Number of grow trees: 500
Average no. of grow terminal nodes: 208.258
Total no. of grow variables: 1
```

```
Analysis: RF-R
Family: regr
% variance explained: 34.97
Test set error rate: 0.11
```

```
survival R. lung survreg() survfit() , predict .
2 2 sex .
```

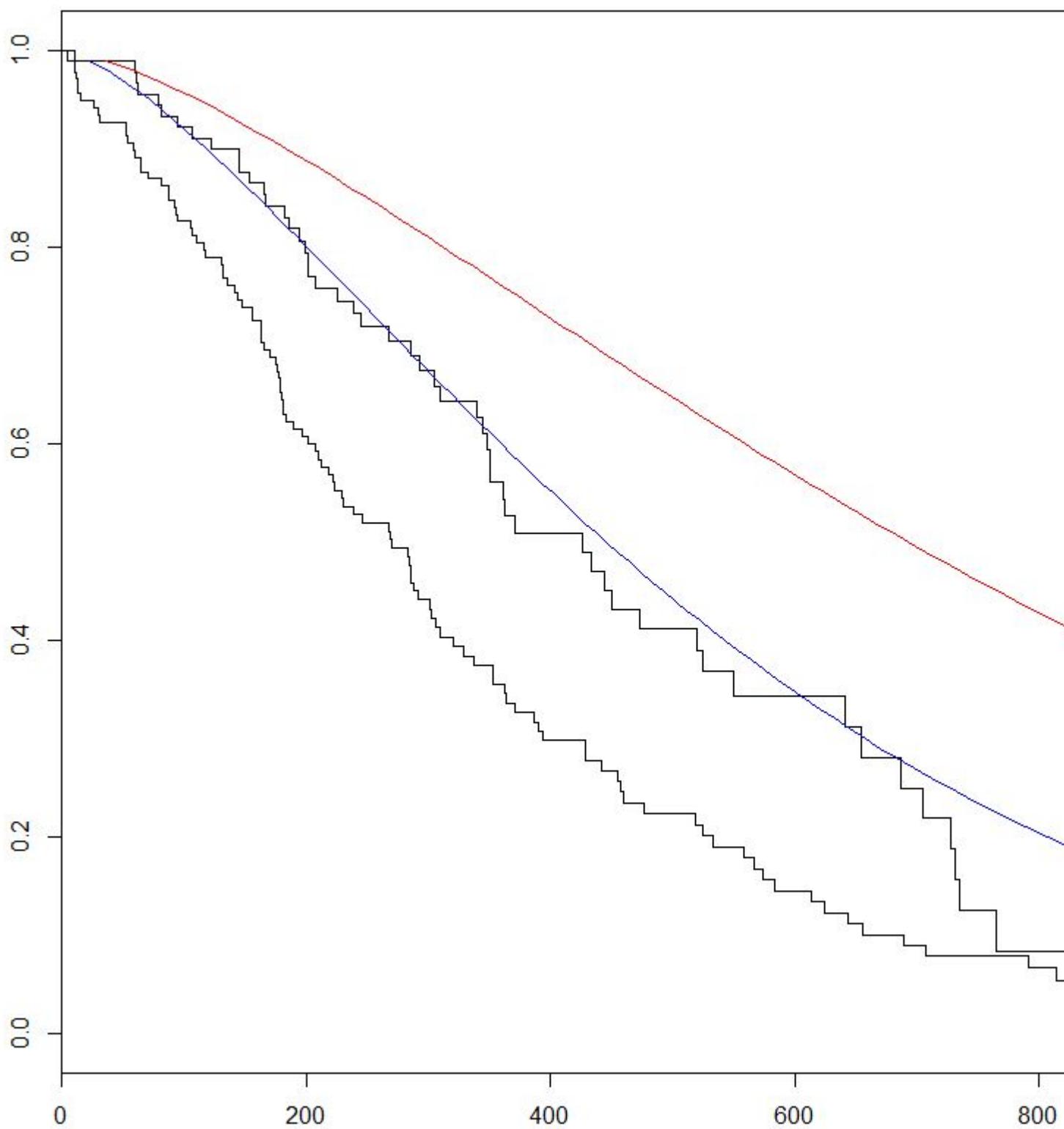
```
require(survival)
s <- with(lung, Surv(time, status))

sWei <- survreg(s ~ as.factor(sex)+age+ph.ecog+wt.loss+ph.karno, dist='weibull', data=lung)

fitKM <- survfit(s ~ sex, data=lung)
plot(fitKM)

lines(predict(sWei, newdata = list(sex = 1,
 age = 1,
 ph.ecog = 1,
 ph.karno = 90,
 wt.loss = 2),
 type = "quantile",
 p = seq(.01, .99, by = .01)),
 seq(.99, .01, by =-.01),
 col = "blue")

lines(predict(sWei, newdata = list(sex = 2,
 age = 1,
 ph.ecog = 1,
 ph.karno = 90,
 wt.loss = 2),
 type = "quantile",
 p = seq(.01, .99, by = .01)),
 seq(.99, .01, by =-.01),
 col = "red")
```



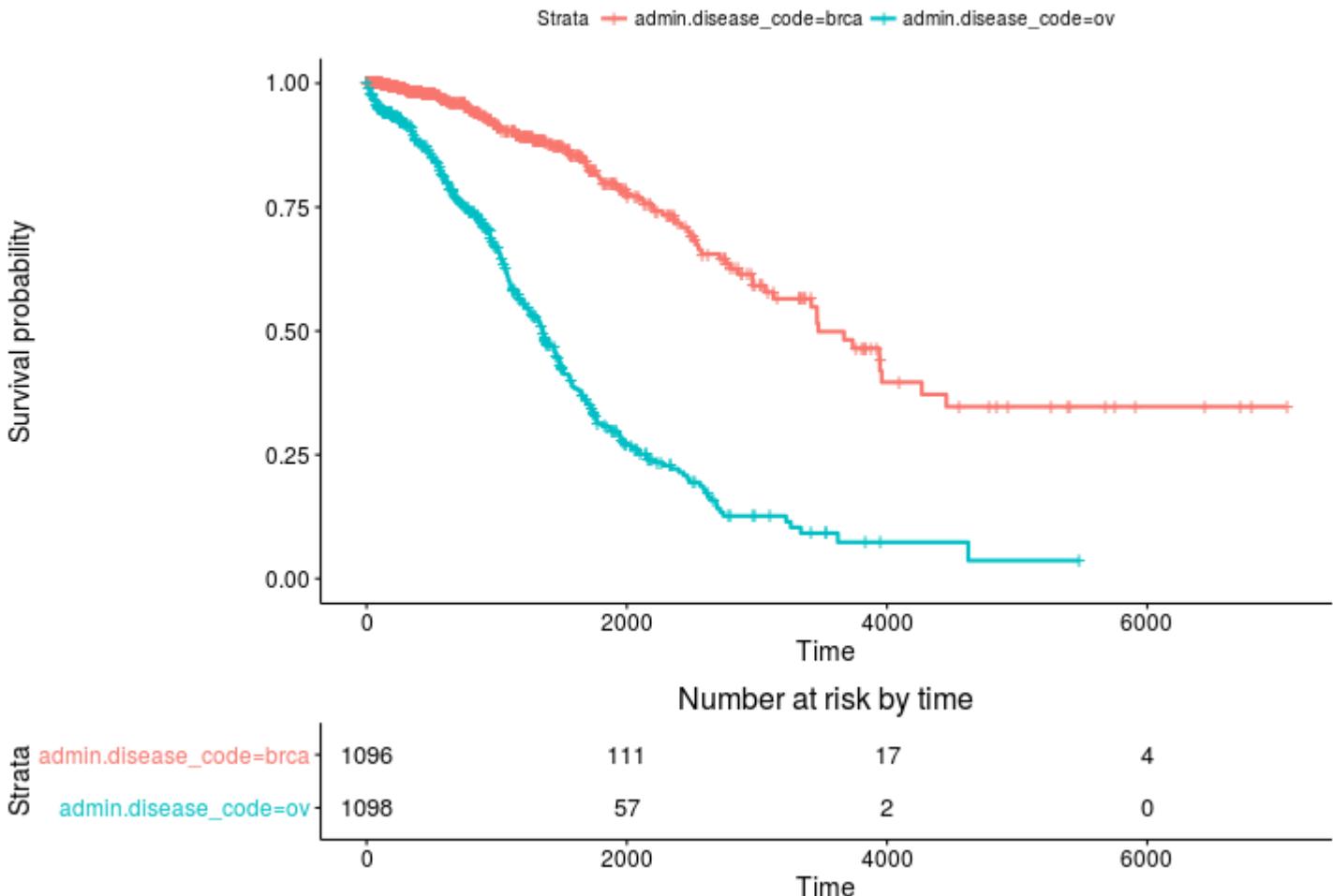
## Kaplan Meier

```
install.packages('survminer')
source("https://bioconductor.org/biocLite.R")
biocLite("RTCGA.clinical") # data for examples
library(RTCGA.clinical)
survivalTCGA(BRCA.clinical, OV.clinical,
 extract.cols = "admin.disease_code") -> BRCAOV.survInfo
```

```

library(survival)
fit <- survfit(Surv(times, patient.vital_status) ~ admin.disease_code,
 data = BRCAOV.survInfo)
library(survminer)
ggsurvplot(fit, risk.table = TRUE)

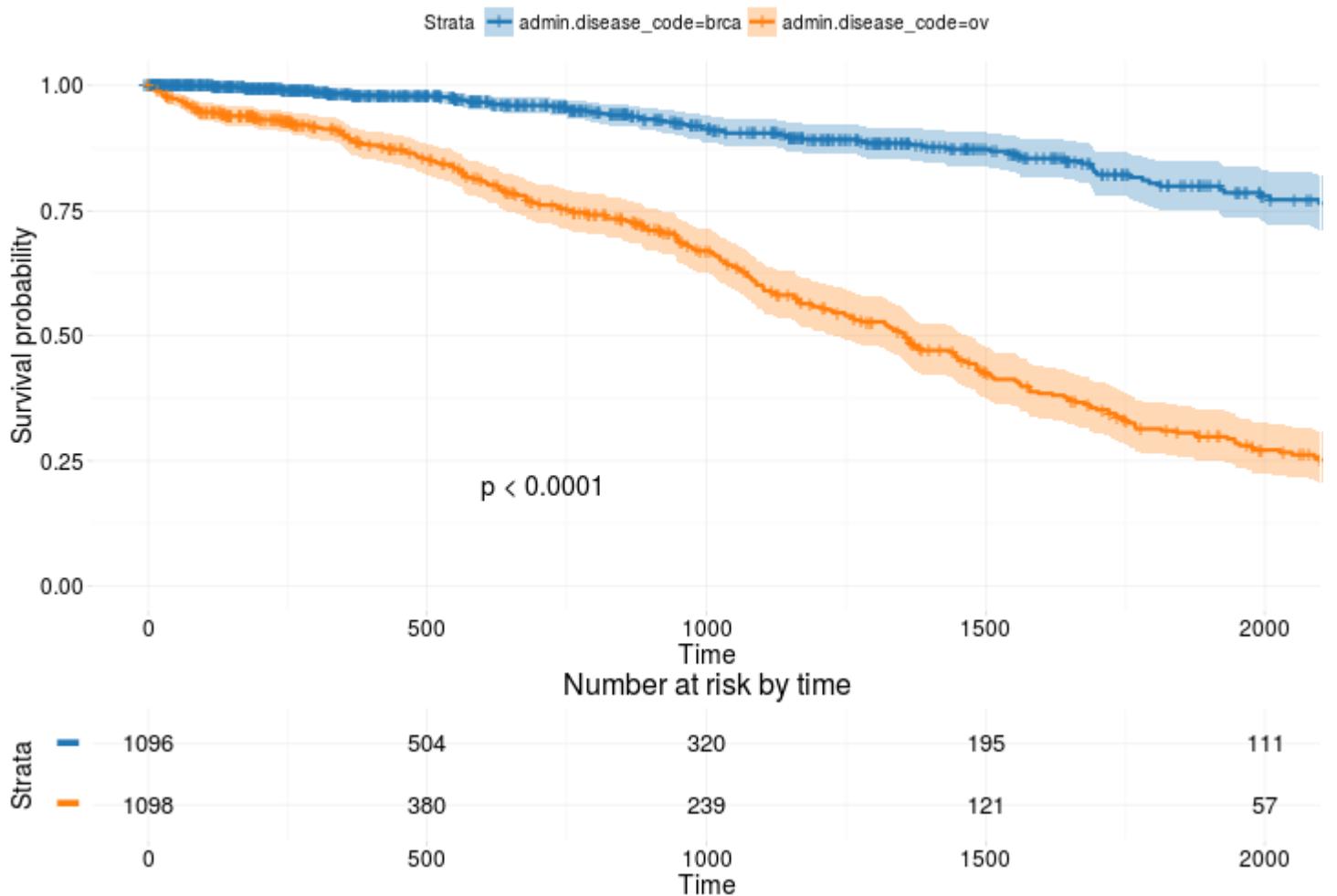
```



```

ggsurvplot(
 fit, # survfit object with calculated statistics.
 risk.table = TRUE, # show risk table.
 pval = TRUE, # show p-value of log-rank test.
 conf.int = TRUE, # show confidence intervals for
 # point estimates of survival curves.
 xlim = c(0,2000), # present narrower X axis, but not affect
 # survival estimates.
 break.time.by = 500, # break X axis in time intervals by 500.
 ggtheme = theme_RTCGA(), # customize plot and risk table with a theme.
 risk.table.y.text.col = T, # colour risk table text annotations.
 risk.table.y.text = FALSE # show bars instead of names in text annotations
 # in legend of risk table
)

```



<http://r-addict.com/2016/05/23/Informationative-Survival-Plots.html>

: [https://riptutorial.com/ko/r/topic/3788/-](https://riptutorial.com/ko/r/topic/3788/)

## 88: ()

- lm(, , , na.action, = "qr", = TRUE, x = FALSE, y = FALSE, qr = TRUE, singular.ok = TRUE, contrasts = NULL, offset, ...)

	Wilkinson-Rogers ; response ~ ... where ... data .
	: data .
	( )
na.action	( NA ) : see ?na.action
	. "qr" "model.frame" ( "model.frame" model=TRUE )
qr	QR
singular.ok	, ( NA )
	. ?model.matrix.default contrasts.arg . options() ( contrasts ) contrast (
	?contrasts options() .
	. . See ?model.offset
...	( lm.fit() lm.wfit() )

## Examples

### mtcars

```
mtcars , (), 32 (help(mtcars)).
```

```
(mpg) (wt) .
```

```
plot(mpg ~ wt, data = mtcars, col=2)
```

```
() !. , lm .
```

```
fit <- lm(mpg ~ wt, data = mtcars)
```

```
~ " " mpg ~ wt mpg ~ wt mpg . . .
```

```
summary(fit)
```

```
Call:
lm(formula = mpg ~ wt, data = mtcars)

Residuals:
 Min 1Q Median 3Q Max
-4.5432 -2.3647 -0.1252 1.4096 6.8727

Coefficients:
 Estimate Std. Error t value Pr(>|t|)
(Intercept) 37.2851 1.8776 19.858 < 2e-16 ***
wt -5.3445 0.5591 -9.559 1.29e-10 ***

Signif. codes: 0 '****' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Residual standard error: 3.046 on 30 degrees of freedom
Multiple R-squared: 0.7528, Adjusted R-squared: 0.7446
F-statistic: 91.38 on 1 and 30 DF, p-value: 1.294e-10
```

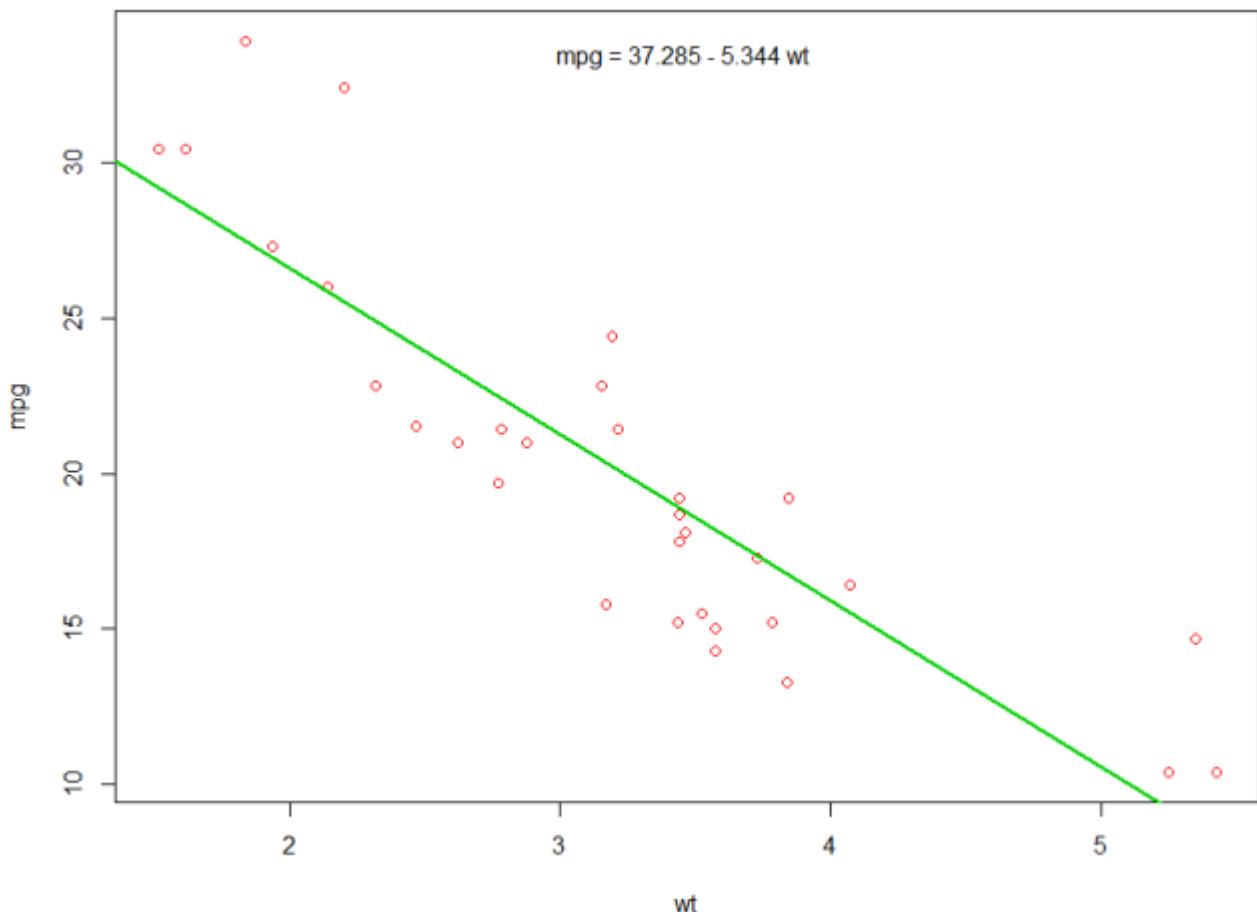
- mpg (wt y)  $37.2851 + (-5.3445) * \text{wt}$
- p-
- R^2 R^2 mpg

mpg

```
abline(fit,col=3,lwd=2)
```

```
. coef coef . paste0 +/- . mtext .
```

```
bs <- round(coef(fit), 3)
lmlab <- paste0("mpg = ", bs[1],
 ifelse(sign(bs[2]) == 1, " + ", " - "), abs(bs[2]), " wt ")
mtext(lmlab, 3, line=-2)
```



()

```
mtcars ,
```

```
fit <- lm(mpg ~ wt, data = mtcars)
```

```
plot(mtcars$wt, mtcars$mpg, pch=18, xlab = 'wt', ylab = 'mpg')
lines(c(min(mtcars$wt), max(mtcars$wt)),
as.numeric(predict(fit, data.frame(wt=c(min(mtcars$wt), max(mtcars$wt))))))
```

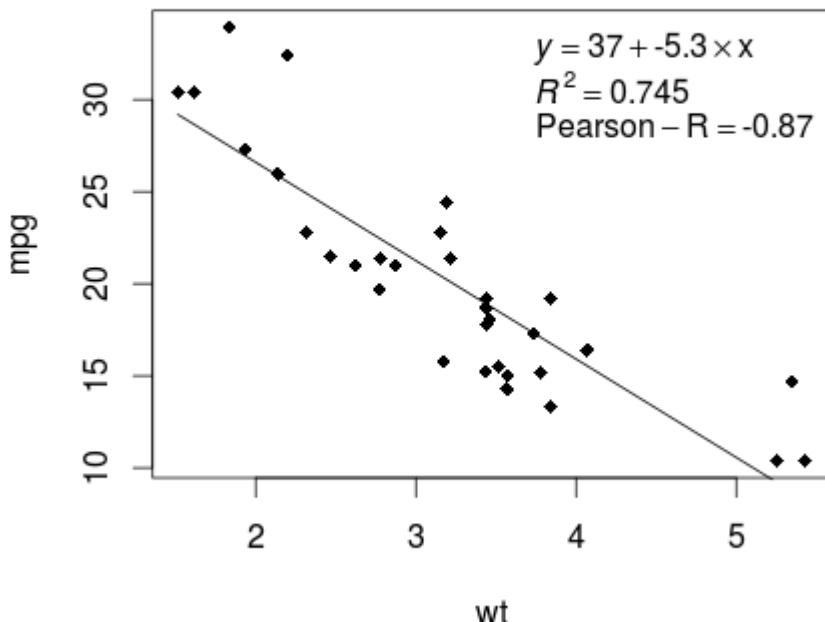
```
! , , rsquare . vector .
```

```
rp = vector('expression', 3)
rp[1] = substitute(expression(italic(y) == MYOTHERVALUE3 + MYOTHERVALUE4 %*% x),
list(MYOTHERVALUE3 = format(fit$coefficients[1], digits = 2),
MYOTHERVALUE4 = format(fit$coefficients[2], digits = 2))) [2]
rp[2] = substitute(expression(italic(R)^2 == MYVALUE),
list(MYVALUE = format(summary(fit)$adj.r.squared, dig=3))) [2]
rp[3] = substitute(expression(Pearson-R == MYOTHERVALUE2),
list(MYOTHERVALUE2 = format(cor(mtcars$wt, mtcars$mpg), digits = 2))) [2]
legend("topright", legend = rp, bty = 'n')
```

```
RMSE .10 . . .
```

```
rp = vector('expression',10)
```

```
r[1] ~ r[10] r[10]
```



- (Analytic Weights) : . , . . .
- (Inverse Probability Weights - IPW) : . . ( ) . .

```
lm() . survey svyglm() . survey . (: lm() svyglm() family gaussian()
```

```
data <- structure(list(lexptot = c(9.1595012302023, 9.86330744180814, 8.92372556833205, 8.58202430280175, 10.1133857229336), progvillm = c(1L, 1L, 1L, 1L, 0L), sexhead = c(1L, 1L, 0L, 1L, 1L), agehead = c(79L, 43L, 52L, 48L, 35L), weight = c(1.04273509979248, 1.01139605045319, 1.01139605045319, 1.01139605045319, 0.76305216550827)), .Names = c("lexptot", "progvillm", "sexhead", "agehead", "weight"), class = c("tbl_df", "tbl", "data.frame"), row.names = c(NA, -5L))
```

```
lm.analytic <- lm(lexptot ~ progvillm + sexhead + agehead, data = data, weight = weight)
summary(lm.analytic)
```

```
Call:
```

```

lm(formula = lexptot ~ progvillm + sexhead + agehead, data = data,
 weights = weight)

Weighted Residuals:
 1 2 3 4 5
9.249e-02 5.823e-01 0.000e+00 -6.762e-01 -1.527e-16

Coefficients:
 Estimate Std. Error t value Pr(>|t|)
(Intercept) 10.016054 1.744293 5.742 0.110
progvillm -0.781204 1.344974 -0.581 0.665
sexhead 0.306742 1.040625 0.295 0.818
agehead -0.005983 0.032024 -0.187 0.882

Residual standard error: 0.8971 on 1 degrees of freedom
Multiple R-squared: 0.467, Adjusted R-squared: -1.132
F-statistic: 0.2921 on 3 and 1 DF, p-value: 0.8386

```

## (IPW)

```

library(survey)
data$X <- 1:nrow(data) # Create unique id

Build survey design object with unique id, ipw, and data.frame
des1 <- svydesign(id = ~X, weights = ~weight, data = data)

Run glm with survey design object
prog.lm <- svyglm(lexptot ~ progvillm + sexhead + agehead, design=des1)

```

```

Call:
svyglm(formula = lexptot ~ progvillm + sexhead + agehead, design = des1)

Survey design:
svydesign(id = ~X, weights = ~weight, data = data)

Coefficients:
 Estimate Std. Error t value Pr(>|t|)
(Intercept) 10.016054 0.183942 54.452 0.0117 *
progvillm -0.781204 0.640372 -1.220 0.4371
sexhead 0.306742 0.397089 0.772 0.5813
agehead -0.005983 0.014747 -0.406 0.7546

Signif. codes: 0 '****' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

(Dispersion parameter for gaussian family taken to be 0.2078647)

Number of Fisher Scoring iterations: 2

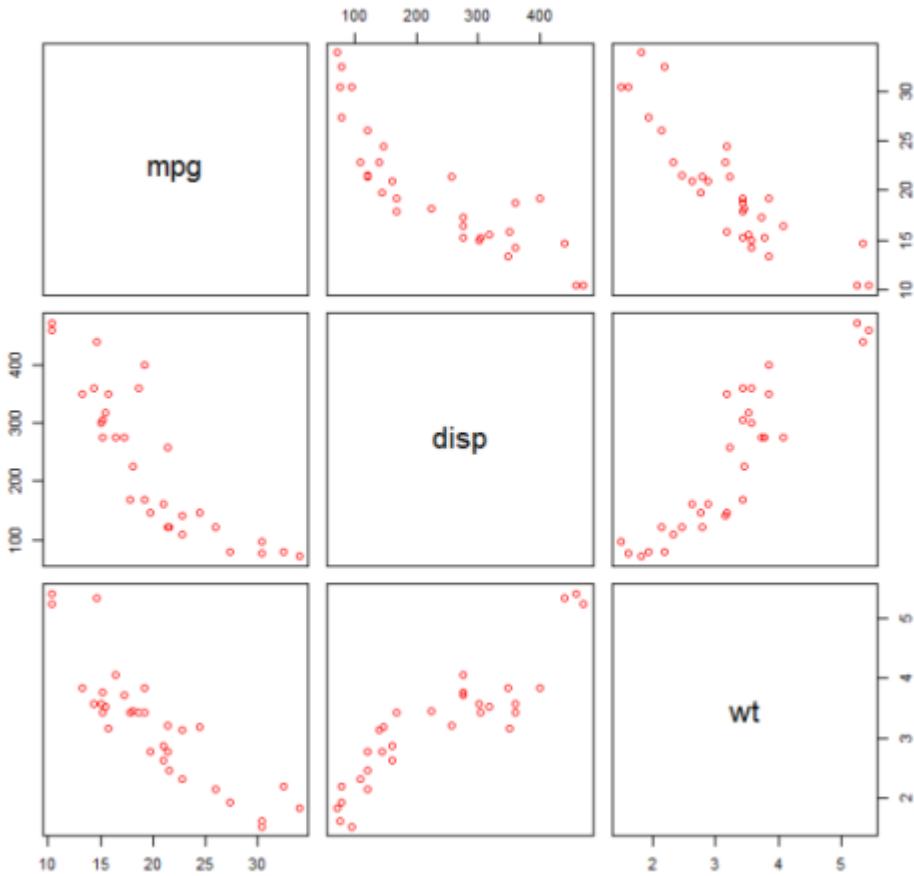
```

```

mtcars 2 mtcars . mtcars .
mpg (Miles / gallon), disp (Displacement (cu.in.)) wt (Weight (1000 lbs)) . mpg disp .

```

```
plot(mtcars[,c("mpg", "disp", "wt")])
```



disp .

```

fit0 = lm(mpg ~ wt+disp, mtcars)
summary(fit0)

Coefficients:
Estimate Std. Error t value Pr(>|t|)
#(Intercept) 34.96055 2.16454 16.151 4.91e-16 ***
#wt -3.35082 1.16413 -2.878 0.00743 **
#disp -0.01773 0.00919 -1.929 0.06362 .
#---
#Signif. codes: 0 '****' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
#Residual standard error: 2.917 on 29 degrees of freedom
#Multiple R-squared: 0.7809, Adjusted R-squared: 0.7658

```

2 I(disp^2) . R^2 .

```

fit1 = lm(mpg ~ wt+disp+I(disp^2), mtcars)
summary(fit1)

Coefficients:
Estimate Std. Error t value Pr(>|t|)
#(Intercept) 41.4019837 2.4266906 17.061 2.5e-16 ***
#wt -3.4179165 0.9545642 -3.581 0.001278 **
#disp -0.0823950 0.0182460 -4.516 0.000104 ***
#I(disp^2) 0.0001277 0.0000328 3.892 0.000561 ***
#---
#Signif. codes: 0 '****' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

```

```
#Residual standard error: 2.391 on 28 degrees of freedom
#Multiple R-squared: 0.8578, Adjusted R-squared: 0.8426
```

3 , .

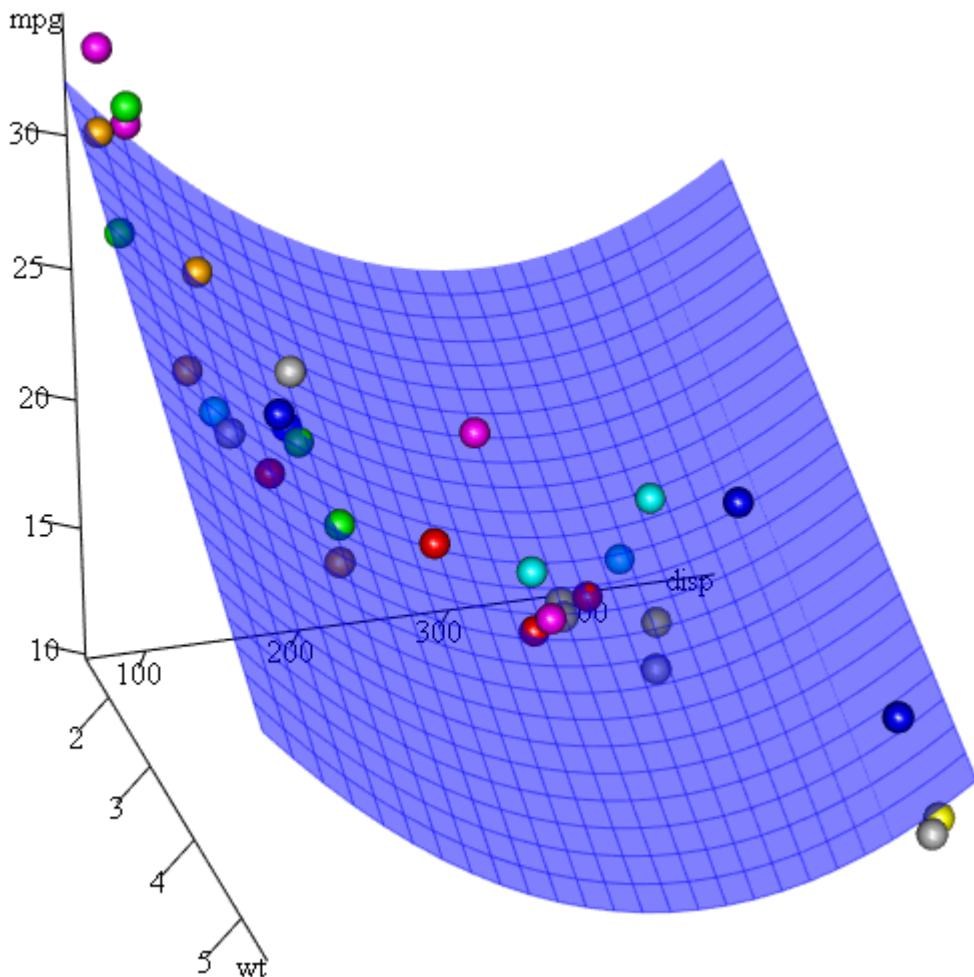
```
mpg = 41.4020 - 3.4179 * wt - 0.0824 * disp + 0.0001277 * disp^2
```

```
raw=TRUE poly . (help(poly)) . .
```

```
summary(lm(mpg ~ wt+poly(disp, 2, raw=TRUE),mtcars))
```

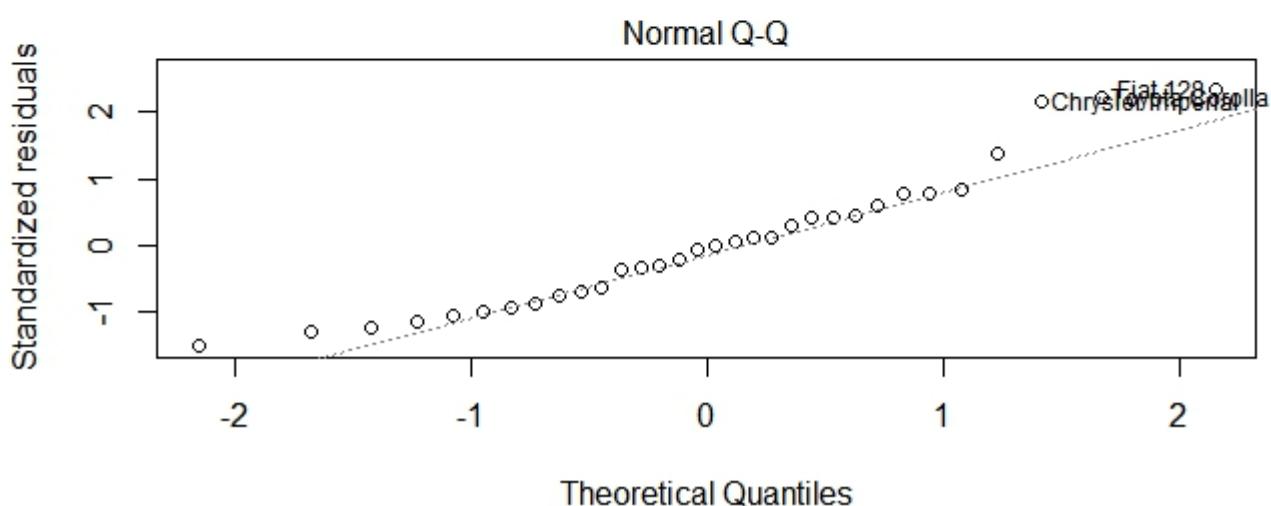
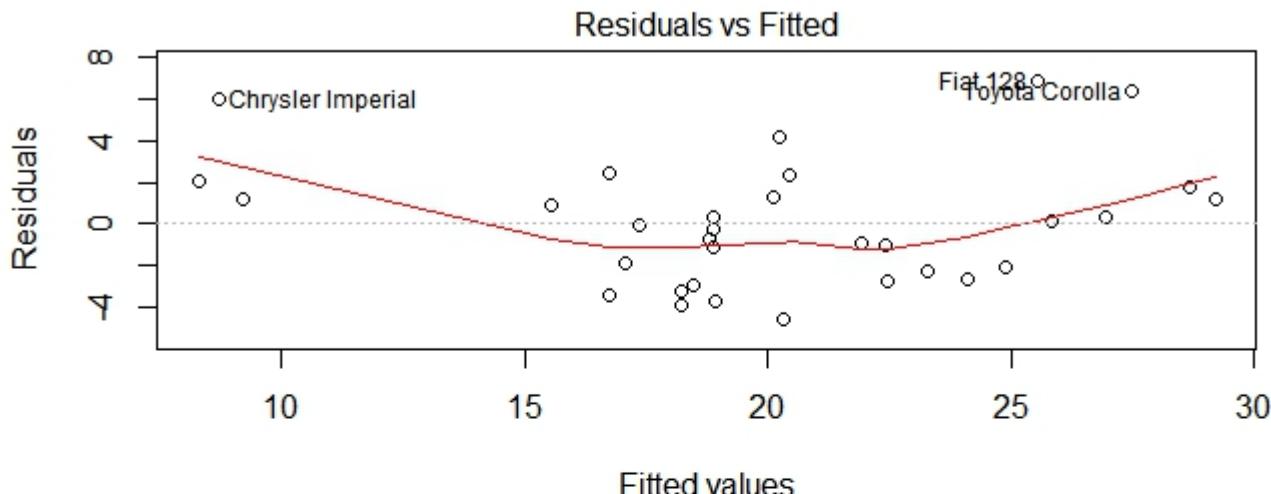
, ? , R 3D . Fit3d p3d .

```
library(p3d)
Init3d(family="serif", cex = 1)
Plot3d(mpg ~ disp+wt, mtcars)
Axes3d()
Fit3d(fit1)
```



```
fit the model
fit <- lm(mpg ~ wt, data = mtcars)
#
```

```
par(mfrow=c(2,1))
plot model object
plot(fit, which =1:2)
```



```
1. (mpg) (wt)
2. . . , QQ . .
```

"

```
predict . mtcars
```

```
my_mdl <- lm(mpg ~ disp, data=mtcars)
my_mdl
```

```
Call:
```

```
lm(formula = mpg ~ disp, data = mtcars)
```

```
Coefficients:
```

```
(Intercept) disp
 29.59985 -0.04122
```

```
set.seed(1234)
newdata <- sample(mtcars$disp, 5)
newdata
[1] 258.0 71.1 75.7 145.0 400.0

newdf <- data.frame(disp=newdata)
predict(my_mdl, newdf)
 1 2 3 4 5
18.96635 26.66946 26.47987 23.62366 13.11381
```

```
 disp
```

## 1. data.frame .

```
predict(my_mdl, newdata)
Error in eval(predvars, data, env) :
 numeric 'envir' arg not of length one
```

## 2. :

```
newdf2 <- data.frame(newdata)
predict(my_mdl, newdf2)
Error in eval(expr, envir, enclos) : object 'disp' not found
```

```
y . newdf 'mpg' 'disp' .
```

```
newdf <- data.frame(mpg=mtcars$mpg[1:10], disp=mtcars$disp[1:10])
mpg disp
1 21.0 160.0
2 21.0 160.0
3 22.8 108.0
4 21.4 258.0
5 18.7 360.0
6 18.1 225.0
7 14.3 360.0
8 24.4 146.7
9 22.8 140.8
10 19.2 167.6

p <- predict(my_mdl, newdf)

#root mean square error
sqrt(mean((p - newdf$mpg)^2, na.rm=TRUE))
[1] 2.325148
```

( ) : <https://riptutorial.com/ko/r/topic/801/>----

```
. class 0, . mode mode . class class<- function class<- . S3 S . S4 .
"" . class . mode . "" typeof "" " " mode .
```

## Examples

R . c() , . :

```
c(1, 2, 3)
[1] 1 2 3
c(TRUE, TRUE, FALSE)
[1] TRUE TRUE FALSE
c("a", "b", "c")
[1] "a" "b" "c"
```

c()

```
x <- c(1, 2, 5)
y <- c(3, 4, 6)
z <- c(x, y)
z
[1] 1 2 5 3 4 6
```

" "

R . class() str() . :

```
class(iris)
[1] "data.frame"

str(iris)
'data.frame': 150 obs. of 5 variables:
 $ Sepal.Length: num 5.1 4.9 4.7 4.6 5 5.4 4.6 5 4.4 4.9 ...
 $ Sepal.Width : num 3.5 3 3.2 3.1 3.6 3.9 3.4 3.4 2.9 3.1 ...
 $ Petal.Length: num 1.4 1.4 1.3 1.5 1.4 1.7 1.4 1.5 1.4 1.5 ...
 $ Petal.Width : num 0.2 0.2 0.2 0.2 0.2 0.4 0.3 0.2 0.2 0.1 ...
 $ Species : Factor w/ 3 levels "setosa","versicolor",...: 1 1 1 1 1 1 1 1 1 1 ...'

class(iris$Species)
[1] "factor"
```

data.frame str() . (Species) (Class Factor) . str() class() .

R . (:, ) . ?atomic : "logical", "integer", "numeric" (synonym "double"), "complex",
"character" "raw" . class .

```
x <- 1826
class(x) <- "Date"
x
```

```
[1] "1975-01-01"
x <- as.Date("1970-01-01")
class(x)
#[1] "Date"
is(x,"Date")
#[1] TRUE
is(x,"integer")
#[1] FALSE
is(x,"numeric")
#[1] FALSE
mode(x)
#[1] "numeric"
```

, , . R ::":

```
mylist <- list(A = c(5,6,7,8), B = letters[1:10], CC = list(5, "Z"))
```

```
f <- function(x) list(xplus = x + 10, xsq = x^2)

f(7)
$xplus
[1] 17
#
$xsq
[1] 49
```

```
L <- list(x = 1:2, y = c("A", "B"))
DF <- data.frame(L)
DF
x y
1 1 A
2 2 B
is.list(DF)
[1] TRUE
```

R ""-

: <https://riptutorial.com/ko/r/topic/3563/>

## Examples

C

```

teethVC = ToothGrowth[ToothGrowth$supp == 'VC',]
teethOJ = ToothGrowth[ToothGrowth$supp == 'OJ',]

permutationTest = function(vectorA, vectorB, testStat){
 N = 10^5
 fullSet = c(vectorA, vectorB)
 lengthA = length(vectorA)
 lengthB = length(vectorB)
 trials <- replicate(N,
 {index <- sample(lengthB + lengthA, size = lengthA, replace = FALSE)
 testStat((fullSet[index]), fullSet[-index]) })
 trials
}
vec1 =teethVC$len;
vec2 =teethOJ$len;
subtractMeans = function(a, b){ return (mean(a) - mean(b)) }
result = permutationTest(vec1, vec2, subtractMeans)
observedMeanDifference = subtractMeans(vec1, vec2)
result = c(result, observedMeanDifference)
hist(result)
abline(v=observedMeanDifference, col = "blue")
pValue = 2*mean(result <= (observedMeanDifference))
pValue

```

CSV

```

permutationTest = function(vectorA, vectorB, testStat){
 N = 10^5
 fullSet = c(vectorA, vectorB)
 lengthA = length(vectorA)
 lengthB = length(vectorB)
 trials <- replicate(N,
 {index <- sample(lengthB + lengthA, size = lengthA, replace = FALSE)
 testStat((fullSet[index]), fullSet[-index]) })
 trials
}

testStat . teststat trials .

N = 10^5 . N .

, trials , .

```

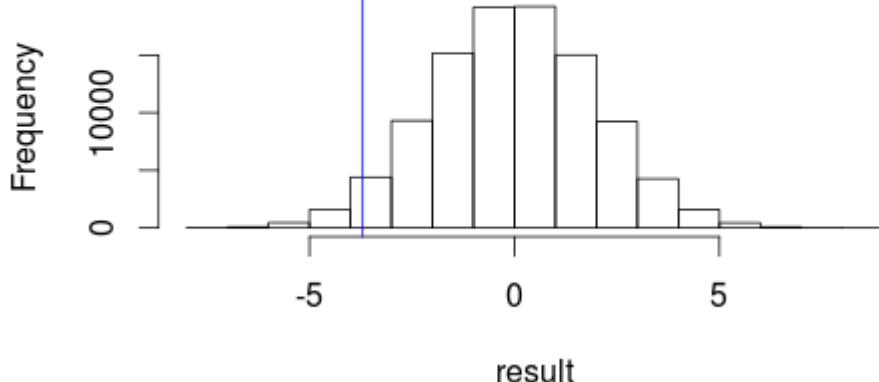
```
subtractMeans = function(a, b) { return (mean(a) - mean(b)) }
```

```
result = permutationTest(vec1, vec2, subtractMeans)
```

```
observedMeanDifference = subtractMeans(vec1, vec2)
```

```
hist(result)
abline(v=observedMeanDifference, col = "blue")
```

## Histogram of result



p- .

```
pValue = 2*mean(result >= (observedMeanDifference))
```

```
result >= (observedMeanDifference)
```

```
FALSE TRUE FALSE FALSE TRUE FALSE ...
```

TRUE result observedMean .

mean TRUE 1 , FALSE 0 1 ..

, 2 . " " .  
p- 0.06093939 . 0.06093939 . C .

: <https://riptutorial.com/ko/r/topic/3216/>--

## Examples

```
. is.numeric() . is.numeric() , as.numeric() .

x <- 12.3
y <- 12L

#confirm types
typeof(x)
[1] "double"
typeof(y)
[1] "integer"

confirm both numeric
is.numeric(x)
[1] TRUE
is.numeric(y)
[1] TRUE

logical to numeric
as.numeric(TRUE)
[1] 1

While TRUE == 1, it is a double and not an integer
is.integer(as.numeric(TRUE))
[1] FALSE
```

## Doubles R . 8 .R .

```
is.double(1)
TRUE
is.double(1.0)
TRUE
is.double(1L)
FALSE
```

## . L .L .

```
typeof(1)
[1] "double"
class(1)
[1] "numeric"
typeof(1L)
[1] "integer"
class(1L)
[1] "integer"
```

double double . 8 4 . .

```

test speed on lots of arithmetic
microbenchmark(
 for(i in 1:100000){
 2L * i
 10L + i
 },
 for(i in 1:100000){
 2.0 * i
 10.0 + i
 }
)
Unit: milliseconds
 expr min lq mean median uq
max neval
 for (i in 1:1e+05) { 2L * i 10L + i } 40.74775 42.34747 50.70543 42.99120 65.46864
94.11804 100
 for (i in 1:1e+05) { 2 * i 10 + i } 41.07807 42.38358 53.52588 44.26364 65.84971
83.00456 100

```

: [https://riptutorial.com/ko/r/topic/9018/----](https://riptutorial.com/ko/r/topic/9018/)

## 92: API (SparkR)

sparkR Spark . , , . SparkR SparkR

### Examples

#### R

Sparks R Spark Cluster .

```
library(SparkR)
sc <- sparkR.init() # connection to Spark context
sqlContext <- sparkRSQl.init(sc) # connection to SQL context
```

IDE Spark .

Apache Spark . , ( () () [ ], IBM ).

:

Spark . . . Spark RDD .

:

( ) . RDD .

:

( 3GB big csv ) .

```
library(SparkR)
next line is needed for direct csv import:
Sys.setenv('SPARKR_SUBMIT_ARGS'='"--packages" "com.databricks:spark-csv_2.10:1.4.0" "sparkr-
shell"')
sc <- sparkR.init()
sqlContext <- sparkRSQl.init(sc)

loading 3 GB big csv file:
train <- read.df(sqlContext, "/train.csv", source = "com.databricks.spark.csv", inferSchema =
"true")
cache(train)
system.time(head(train))
output: time elapsed: 125 s. This action invokes the caching at this point.
system.time(head(train))
output: time elapsed: 0.2 s (!!)
```

RDD( )

:

```
mtrdd <- createDataFrame(sqlContext, mtcars)
```

## CSV :

csv Spark CSV .

```
Sys.setenv('SPARKR_SUBMIT_ARGS'='"--packages" "com.databricks:spark-csv_2.10:1.4.0" "sparkr-shell"') # context for csv import read csv ->
sc <- sparkR.init()
sqlContext <- sparkRSQl.init(sc)
```

CSV .

```
train <- read.df(sqlContext, "/train.csv", header= "true", source =
"com.databricks.spark.csv", inferSchema = "true")
```

:

```
customSchema <- structType(
 structField("margin", "integer"),
 structField("gross", "integer"),
 structField("name", "string"))

train <- read.df(sqlContext, "/train.csv", header= "true", source =
"com.databricks.spark.csv", schema = customSchema)
```

API (SparkR) : <https://riptutorial.com/ko/r/topic/5349/-api--sparkr->

# 93:

```
stats glm() : glm() . CRAN .
```

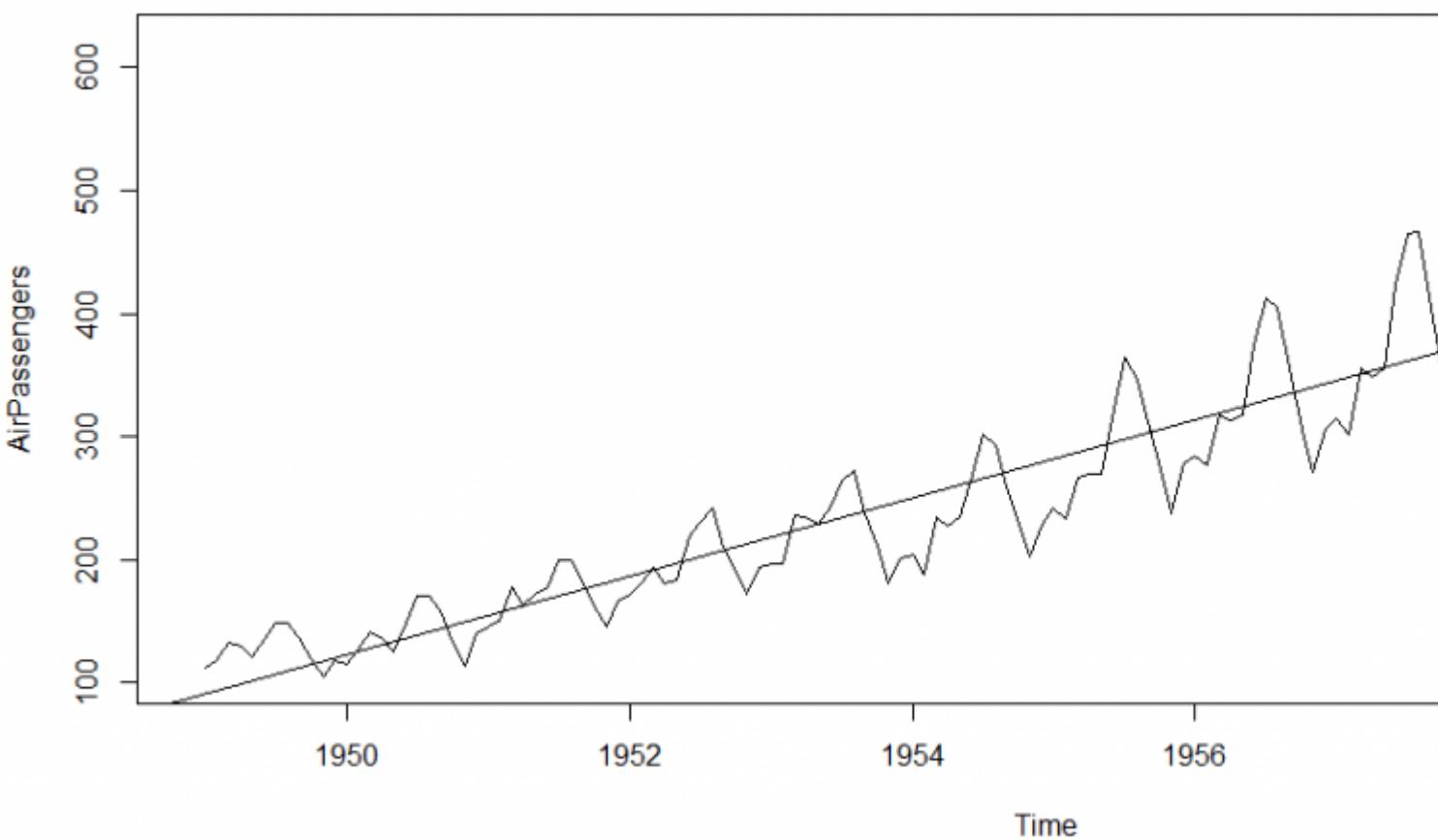
## Examples

```
data(AirPassengers)
class(AirPassengers)
```

```
1 "ts"
```

Exploratory Data Analysis (EDA) .

```
plot(AirPassengers) # plot the raw data
abline(reg=lm(AirPassengers~time(AirPassengers))) # fit a trend line
```



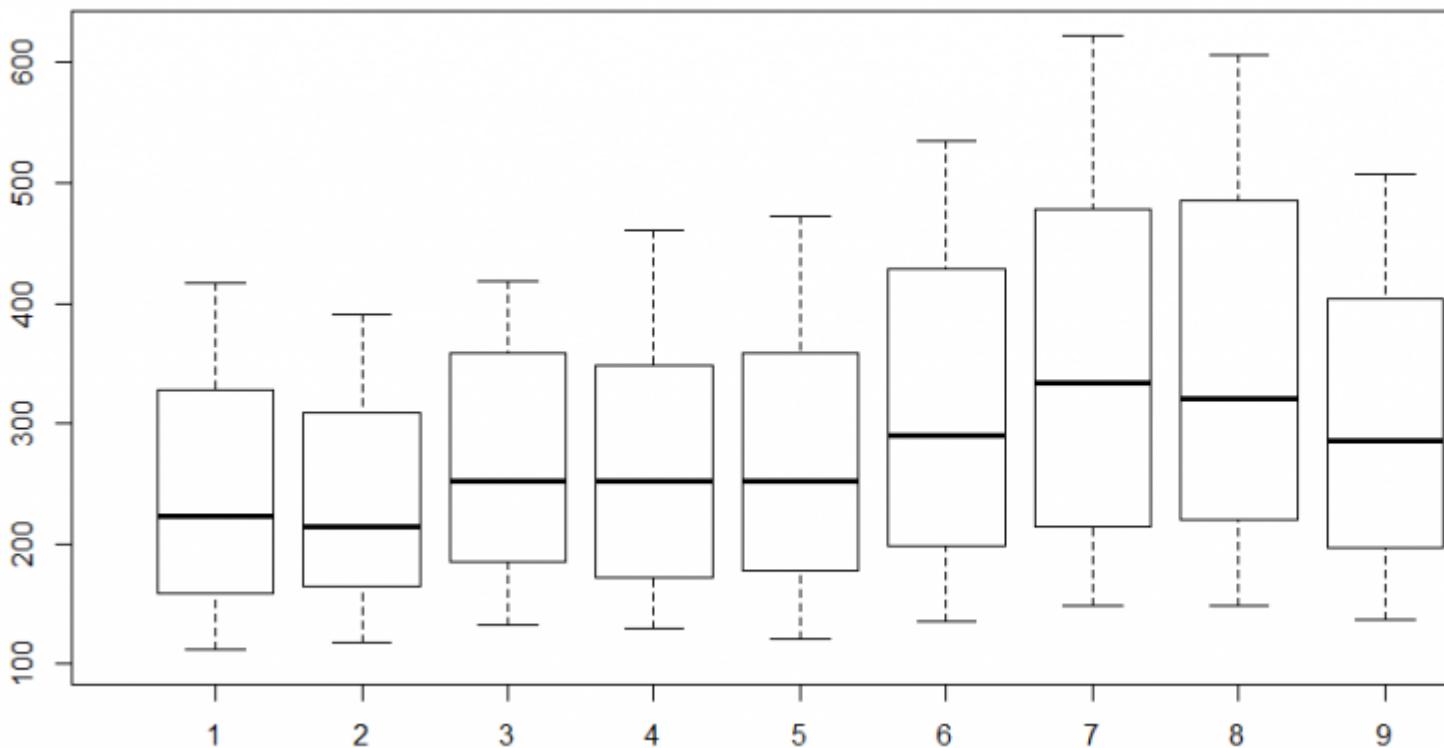
EDA : .

```
cycle(AirPassengers)
```

	Jan	Feb	Mar	Apr	May	Jun	Jul	Aug	Sep	Oct	Nov	Dec
1949	1	2	3	4	5	6	7	8	9	10	11	12
1950	1	2	3	4	5	6	7	8	9	10	11	12
1951	1	2	3	4	5	6	7	8	9	10	11	12

1952	1	2	3	4	5	6	7	8	9	10	11	12
1953	1	2	3	4	5	6	7	8	9	10	11	12
1954	1	2	3	4	5	6	7	8	9	10	11	12
1955	1	2	3	4	5	6	7	8	9	10	11	12
1956	1	2	3	4	5	6	7	8	9	10	11	12
1957	1	2	3	4	5	6	7	8	9	10	11	12
1958	1	2	3	4	5	6	7	8	9	10	11	12
1959	1	2	3	4	5	6	7	8	9	10	11	12
1960	1	2	3	4	5	6	7	8	9	10	11	12

```
boxplot(AirPassengers~cycle(AirPassengers)) #Box plot across months to explore seasonal effects
```



## TS

ts . ts ARIMA . window .

```
#Create a dummy dataset of 100 observations
x <- rnorm(100)

#Convert this vector to a ts object with 100 annual observations
x <- ts(x, start = c(1900), freq = 1)

#Convert this vector to a ts object with 100 monthly observations starting in July
x <- ts(x, start = c(1900, 7), freq = 12)

#Alternatively, the starting observation can be a number:
x <- ts(x, start = 1900.5, freq = 12)

#Convert this vector to a ts object with 100 daily observations and weekly frequency starting
#in the first week of 1900
x <- ts(x, start = c(1900, 1), freq = 7)
```

```
#The default plot for a ts object is a line plot
plot(x)

#The window function can call elements or sets of elements by date

#Call the first 4 weeks of 1900
window(x, start = c(1900, 1), end = (1900, 4))

#Call only the 10th week in 1900
window(x, start = c(1900, 10), end = (1900, 10))

#Call all weeks including and after the 10th week of 1900
window(x, start = c(1900, 10))
```

ts .

```
#Create a dummy matrix of 3 series with 100 observations each
x <- cbind(rnorm(100), rnorm(100), rnorm(100))

#Create a multi-series ts with annual observation starting in 1900
x <- ts(x, start = 1900, freq = 1)

#R will draw a plot for each series in the object
plot(x)
```

: <https://riptutorial.com/ko/r/topic/2701/>--

## 94:

```
Arima , R arima .
ARIMA () . . . ARIMA .
ARIMA .
```

## Examples

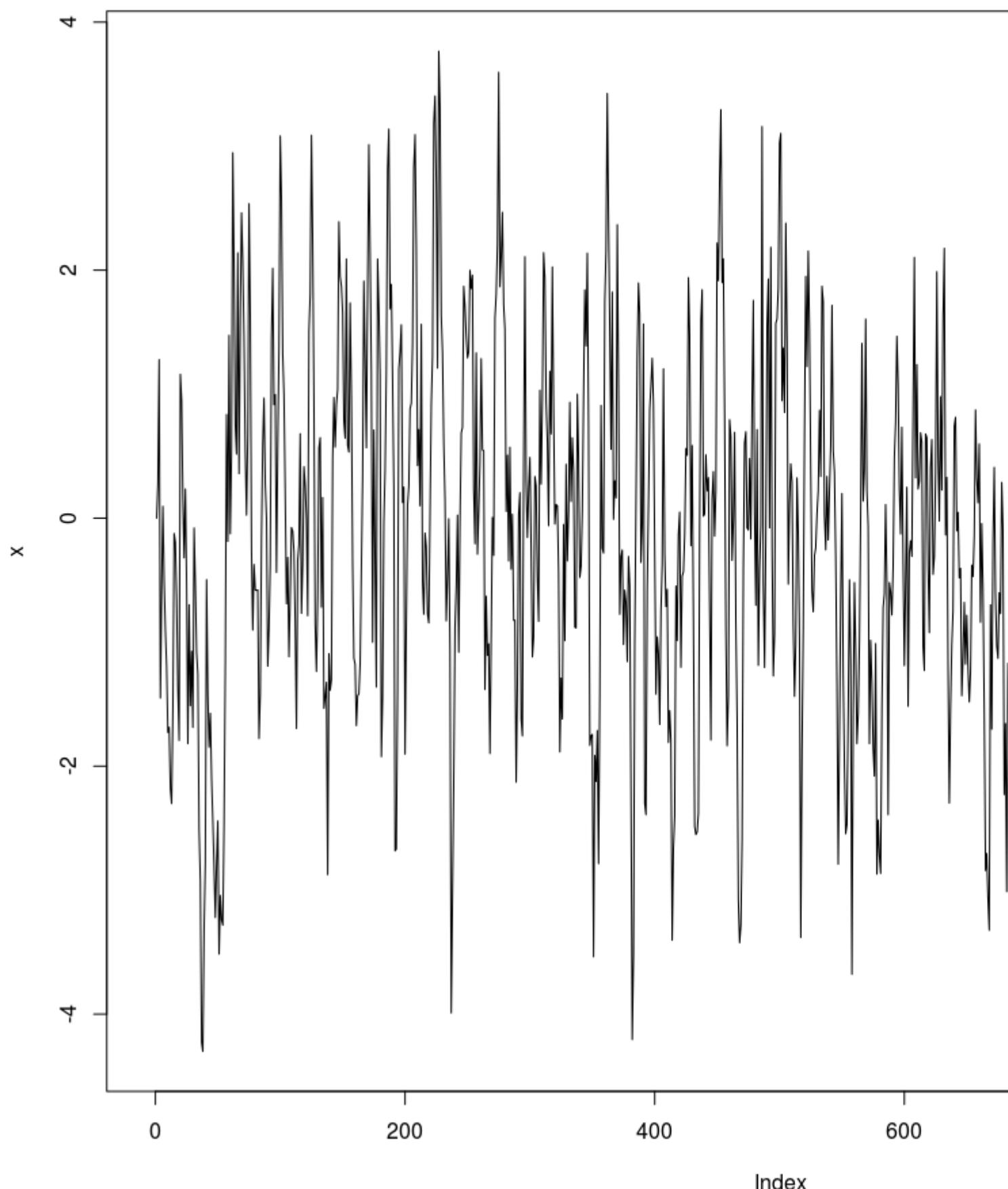
### Arima AR1

$$x_t = .7x_{t-1} + \epsilon \quad \epsilon \sim N(0, 1)$$

```
#Load the forecast package
library(forecast)

#Generate an AR1 process of length n (from Cowpertwait & Meltcalfe)
Set up variables
set.seed(1234)
n <- 1000
x <- matrix(0,1000,1)
w <- rnorm(n)

loop to create x
for (t in 2:n) x[t] <- 0.7 * x[t-1] + w[t]
plot(x,type='l')
```



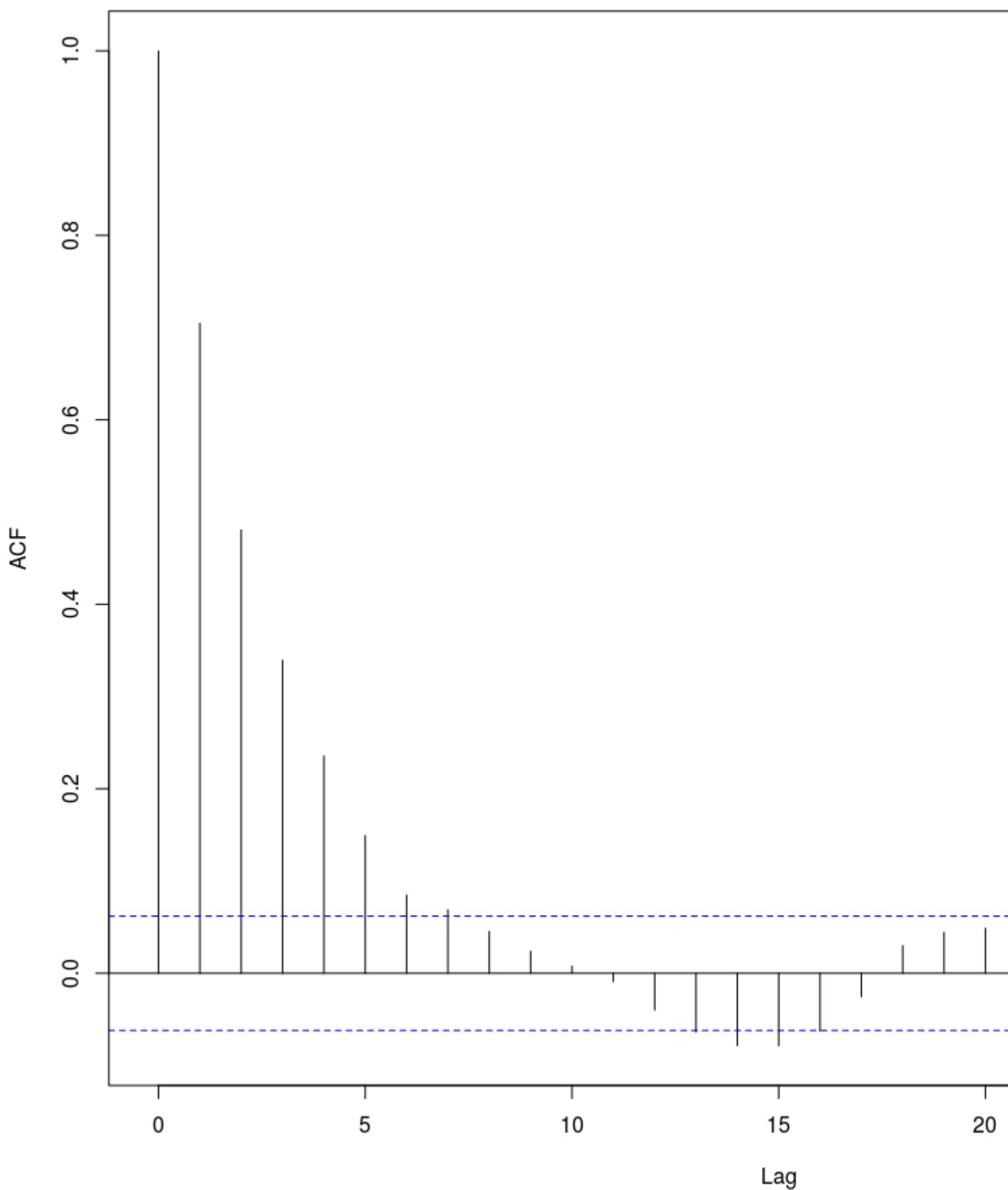
Arima 1, 0, MA 0 .

```
#Fit an AR1 model using Arima
fit <- Arima(x, order = c(1, 0, 0))
summary(fit)
Series: x
ARIMA(1,0,0) with non-zero mean
#
Coefficients:
ar1 intercept
0.7040 -0.0842
s.e. 0.0224 0.1062
#
sigma^2 estimated as 0.9923: log likelihood=-1415.39
AIC=2836.79 AICc=2836.81 BIC=2851.51
#
Training set error measures:
ME RMSE MAE MPE MAPE MASE ACF1
Training set -8.369365e-05 0.9961194 0.7835914 Inf Inf 0.91488 0.02263595
Verify that the model captured the true AR parameter
```

```
fit$coef[1]
ar1
0.7040085

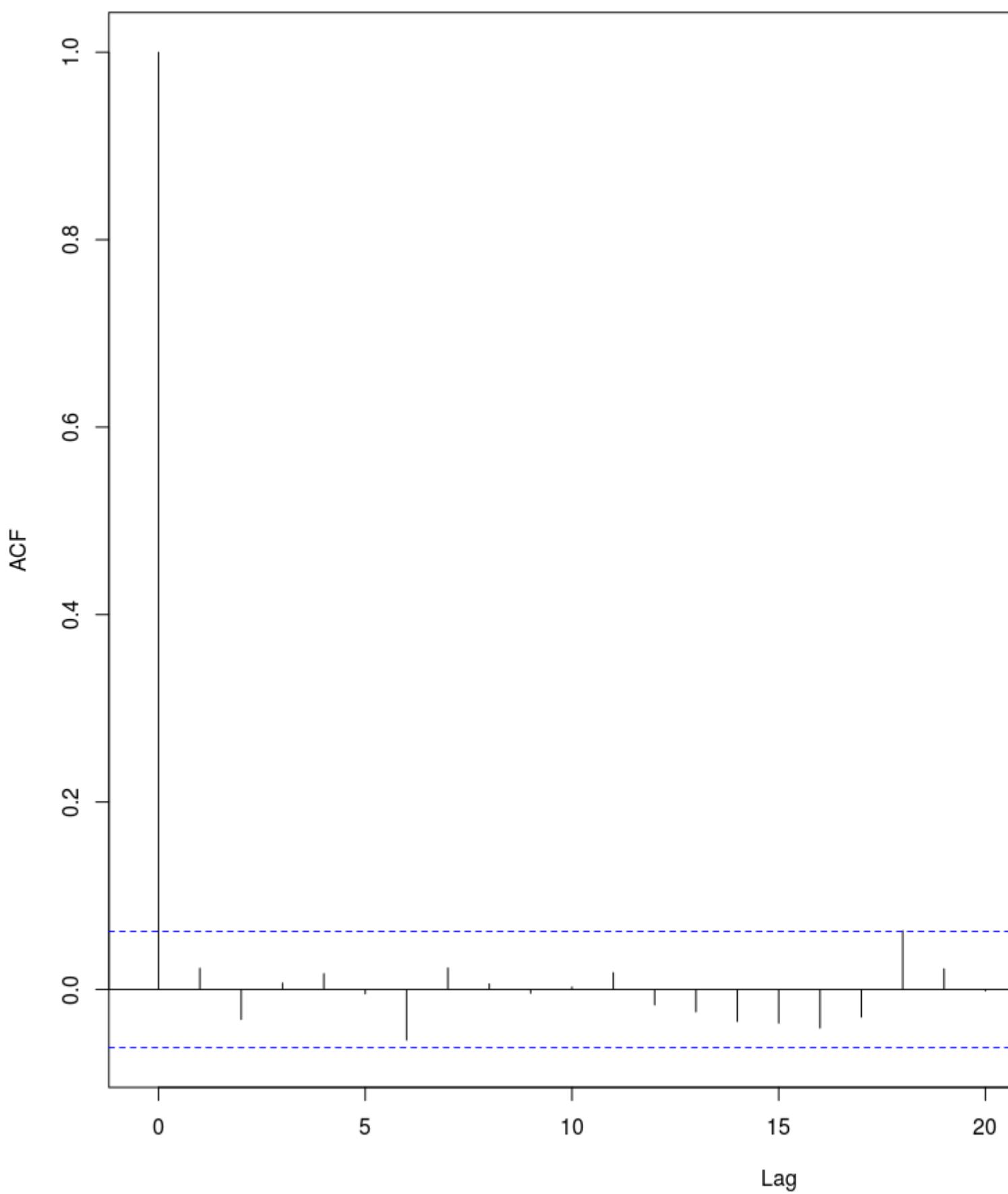
#Verify that the model eliminates the autocorrelation
acf(x)
```

## Series 1



```
acf(fit$resid)
```

**Series fit\$resid**

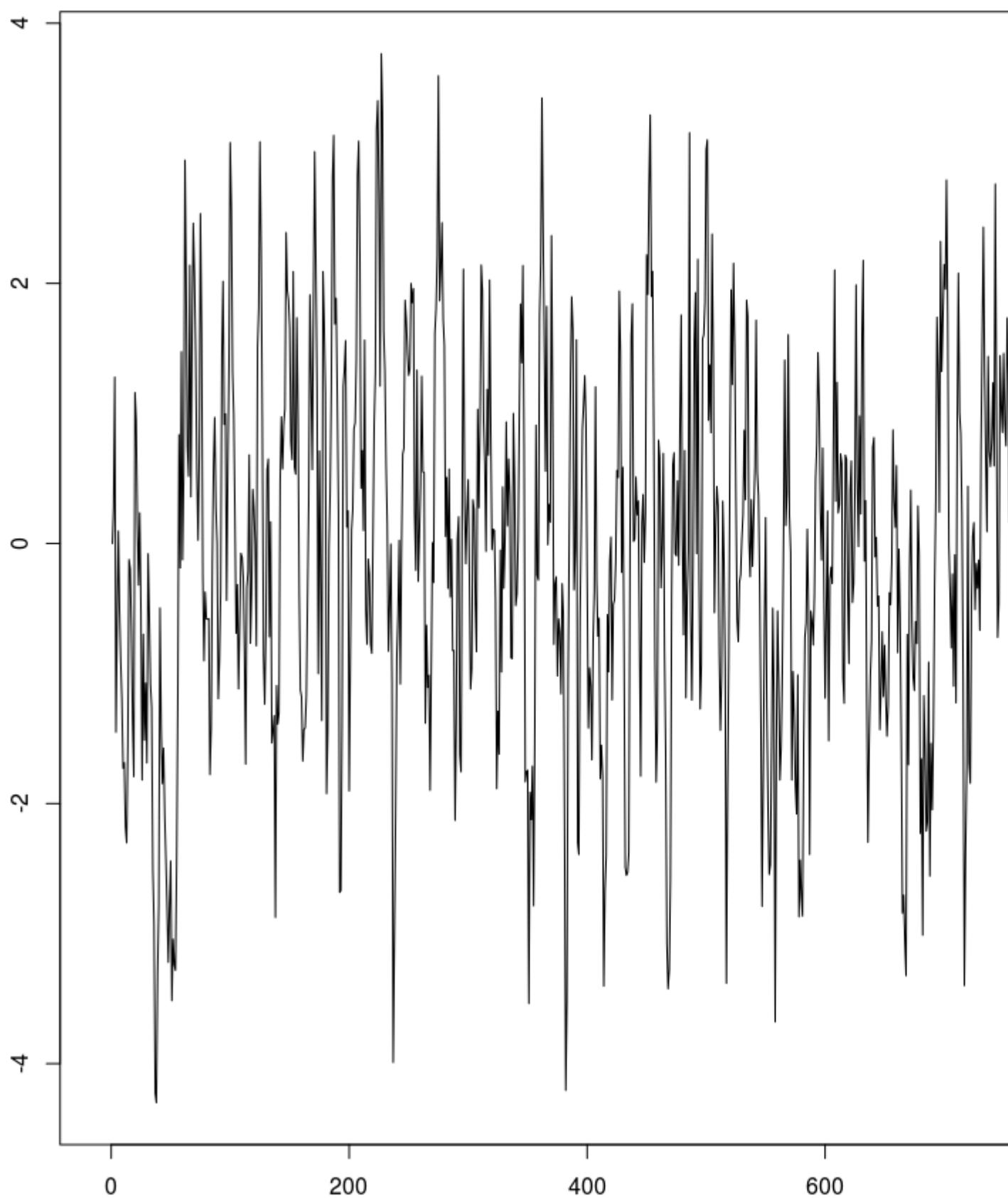


```

#Forecast 10 periods
fcst <- forecast(fit, h = 100)
fcst
 Point Forecast Lo 80 Hi 80 Lo 95 Hi 95
1001 0.282529070 -0.9940493 1.559107 -1.669829 2.234887
1002 0.173976408 -1.3872262 1.735179 -2.213677 2.561630
1003 0.097554408 -1.5869850 1.782094 -2.478726 2.673835
1004 0.043752667 -1.6986831 1.786188 -2.621073 2.708578
1005 0.005875783 -1.7645535 1.776305 -2.701762 2.713514
...
```
#Call the point predictions
fcst$mean
# Time Series:
# Start = 1001
# End = 1100
# Frequency = 1
[1] 0.282529070 0.173976408 0.097554408 0.043752667 0.005875783 -0.020789866 -
0.039562711 -0.052778954
[9] -0.062083302
...
```
#Plot the forecast
plot(fcst)

```

## Forecasts from ARIMA(1,0,0) with non-zero



: <https://riptutorial.com/ko/r/topic/1725/>

## Examples

### .zip

```
zip unzip unzip utils (base R) unzip .
```

```
unzip(zipfile = "bar.zip", exdir = "./foo")
```

```
"bar.zip" "foo" . . zip .
```

### .zip

```
zip utils (base R) unzip .
```

```
unzip(zipfile = "bar.zip", list = TRUE)
```

```
"bar.zip" "none" . . zip .
```

### .tar

```
tar utils (base R) untar .
```

```
untar(zipfile = "bar.tar", list = TRUE)
```

```
"bar.tar" "none" . . tar .
```

### .tar

```
tar utils (R) untar .
```

```
untar(tarfile = "bar.tar", exdir = "./foo")
```

```
"bar.tar" "foo" . . tar .
```

### .zip

```
for zip .
```

```
for (i in dir(pattern=".zip$"))
 unzip(i)
```

```
dir pattern . unzip zip i .
```

: <https://riptutorial.com/ko/r/topic/4323/>-----

## Examples

```
sum sum .

set.seed(20)
df1 <- data.frame(ID = rep(c("A", "B", "C"), each = 3), V1 = rnorm(9), V2 = rnorm(9))
m1 <- as.matrix(df1[-1])
```

. base R colSums

```
colSums(df1[-1], na.rm = TRUE)
```

na.rm = TRUE sum sum ( NAs )

matrix .

```
colSums(m1, na.rm = TRUE)
```

lapply/sapply/vapply .

```
lapply(df1[-1], sum, na.rm = TRUE)
```

list . vector

```
sapply(df1[-1], sum, na.rm = TRUE)
```

```
vapply(df1[-1], sum, na.rm = TRUE, numeric(1))
```

MARGIN = 1 apply apply

```
apply(m1, 2, FUN = sum, na.rm = TRUE)
```

dplyr data.table .

```
library(dplyr)
df1 %>%
 summarise_at(vars(matches("^V\\d+")), sum, na.rm = TRUE)
```

summarise\_at sum . v ( \\d+ ).

data.table .

```
library(data.table)
```

```
setDT(df1) [, lapply(.SD, sum, na.rm = TRUE), .SDcols = 2:ncol(df1)]
```

```
'data.table'('data.frame' setDT(df1)) .SDcols Data.table (.SD) sum.
```

```
/ .
```

```
df1 %>%
 group_by(ID) %>%
 summarise_at(vars(matches("^V\\d+")), sum, na.rm = TRUE)
```

```
sum summarise_each summarise_at
```

```
df1 %>%
 group_by(ID) %>%
 summarise_each(funs(sum(., na.rm = TRUE)))
```

```
data.table .
```

```
setDT(df1) [, lapply(.SD, sum, na.rm = TRUE), by = ID]
```

: <https://riptutorial.com/ko/r/topic/2212/>

R , .

## Examples

```
a <- c(1, 2, 3)
b <- c(4, 5, 6)
mean_ab <- (a + b) / 2

d <- c(1, 0, 1)
only_1_3 <- a[d == 1]
```

```
mat <- matrix(c(1,2,3,4), nrow = 2, ncol = 2)
dimnames(mat) <- list(c(), c("a", "b", "c"))
mat[,] == mat
```

```
df <- data.frame(qualifiers = c("Buy", "Sell", "Sell"),
 symbols = c("AAPL", "MSFT", "GOOGL"),
 values = c(326.0, 598.3, 201.5))
df$symbols == df[[2]]
df$symbols == df[["symbols"]]
df[[2, 1]] == "AAPL"
```

```
l <- list(a = 500, "aaa", 98.2)
length(l) == 3
class(l[1]) == "list"
class(l[[1]]) == "numeric"
class(l$a) == "numeric"
```

```
env <- new.env()
env[["foo"]] = "bar"
env2 <- env
env2[["foo"]] = "BAR"

env[["foo"]] == "BAR"
get("foo", envir = env) == "BAR"
rm("foo", envir = env)
env[["foo"]] == NULL
```

()

```

Creates a 1 row - 2 columns format
par(mfrow=c(1,2))

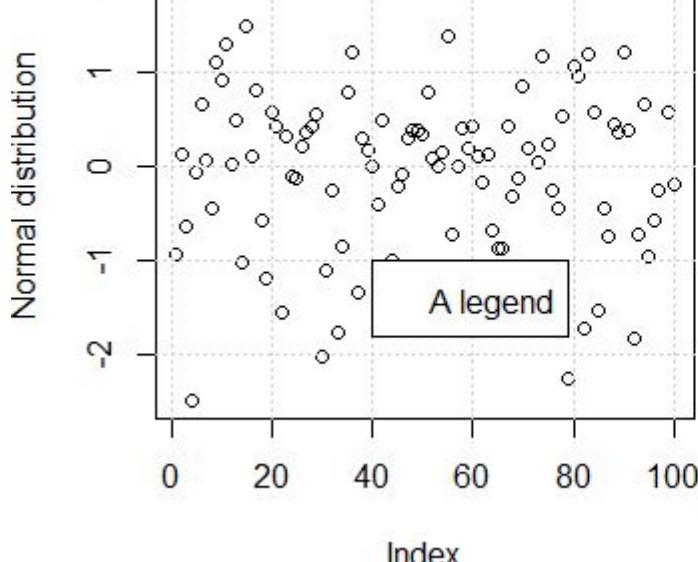
plot(rnorm(100), main = "Graph 1", ylab = "Normal distribution")
grid()
legend(x = 40, y = -1, legend = "A legend")

plot(rnorm(100), main = "Graph 2", type = "l")
abline(v = 50)

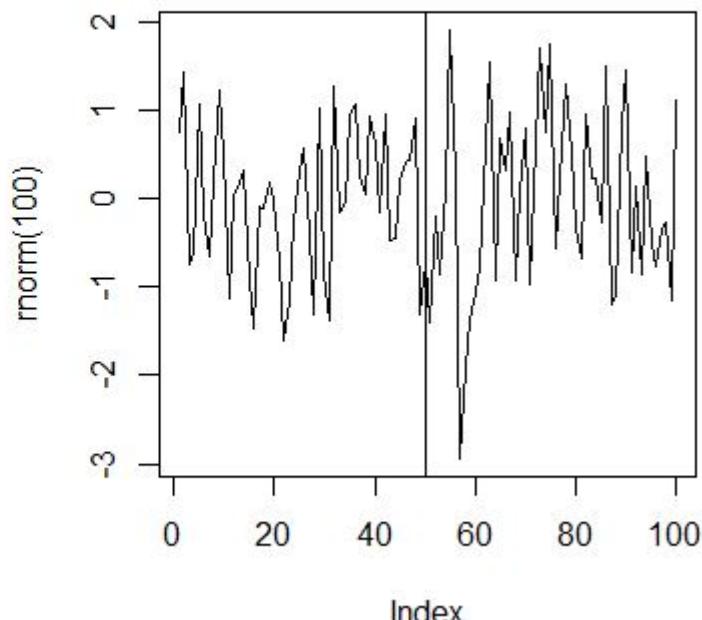
```

:

**Graph 1**



**Graph 2**



```

Create 100 standard normals in a vector
x <- rnorm(100, mean = 0, sd = 1)

Find the lenght of a vector
length(x)

Compute the mean
mean(x)

Compute the standard deviation
sd(x)

Compute the median value
median(x)

Compute the range (min, max)
range(x)

Sum an iterable
sum(x)

Cumulative sum (x[1], x[1]+x[2], ...)
cumsum(x)

```

```
Display the first 3 elements
head(3, x)

Display min, 1st quartile, median, mean, 3rd quartile, max
summary(x)

Compute successive difference between elements
diff(x)

Create a range from 1 to 10 step 1
1:10

Create a range from 1 to 10 step 0.1
seq(1, 10, 0.1)

Print a string
print("hello world")
```

: <https://riptutorial.com/ko/r/topic/10827/>--

# 98: (Excel, SAS, SPSS, Stata) I/O

## Examples

```
rio . import() . import() URL .
```

```
import("example.csv") # comma-separated values
import("example.tsv") # tab-separated values
import("example.dta") # Stata
import("example.sav") # SPSS
import("example.sas7bdat") # SAS
import("example.xlsx") # Excel
```

```
import() , URL (HTTP HTTPS) . rio github .
```

```
import()
```

```
import("example.csv", format = ",") #for csv file where comma is used as separator
import("example.csv", format = ";") #for csv file where semicolon is used as separator
```

## Excel

```
Excel R .
```

R	
xlsx	
XLconnect	
openxlsx	C ++
readxl	C ++
RODBC	ODBC
gdata	

```
Java ODBC R OS . R 64 xlsx XLconnect Java 64 .
```

```
Excel . package::function package::function . openxlsx openxlsx .
```

## xlsx

```
library(xlsx)
```

```
xlsx::read.xlsx("Book1.xlsx", sheetIndex=1)

xlsx::read.xlsx("Book1.xlsx", sheetName="Sheet1")
```

## XLconnect Excel

```
library(XLConnect)
wb <- XLConnect::loadWorkbook("Book1.xlsx")

Either, if Book1.xlsx has a sheet called "Sheet1":
sheet1 <- XLConnect::readWorksheet(wb, "Sheet1")
Or, more generally, just get the first sheet in Book1.xlsx:
sheet1 <- XLConnect::readWorksheet(wb, getSheets(wb)[1])
```

```
XLConnect Book1.xlsx Excel . Excel . Book1.xlsx myHeader , myBody myPcts . R ()
:
```

```
Headerstyle <- XLConnect::getCellStyle(wb, "myHeader")
Bodystyle <- XLConnect::getCellStyle(wb, "myBody")
Pctsstyle <- XLConnect::getCellStyle(wb, "myPcts")
```

```
R .
```

```
Headerrange <- expand.grid(row = 1, col = 1:8)
Bodyrange <- expand.grid(row = 2:6, col = c(1:5, 8))
Pctrange <- expand.grid(row = 2:6, col = c(6, 7))

XLConnect::setCellStyle(wb, sheet = "sheet1", row = Headerrange$row,
 col = Headerrange$col, cellstyle = Headerstyle)
XLConnect::setCellStyle(wb, sheet = "sheet1", row = Bodyrange$row,
 col = Bodyrange$col, cellstyle = Bodystyle)
XLConnect::setCellStyle(wb, sheet = "sheet1", row = Pctrange$row,
 col = Pctrange$col, cellstyle = Pctsstyle)
```

```
XLConnect openxlsx .
```

## openxlsx .

```
Excel openxlsx .
```

```
library(openxlsx)

openxlsx::read.xlsx("spreadsheet1.xlsx", colNames=TRUE, rowNames=TRUE)

#colNames: If TRUE, the first row of data will be used as column names.
#rowNames: If TRUE, first column of data will be used as row names.
```

```
R sheet sheet .
```

```
openxlsx::read.xlsx("spreadsheet1.xlsx", sheet = 1)

::: openxlsx:::read.xlsx("spreadsheet1.xlsx", sheet = "Sheet1")

openxlsx . detectDates TRUE TRUE .

openxlsx::read.xlsx("spreadsheet1.xlsx", sheet = "Sheet1", detectDates= TRUE)
```

# **readxl** **read**

## Excel readxl R

```
library(readxl)

.xls .xlsx .

readxl::read_excel("spreadsheet1.xls")
readxl::read_excel("spreadsheet2.xlsx")
```

```
readxl::read_excel("spreadsheet.xls", sheet = 1)
readxl::read_excel("spreadsheet.xls", sheet = "summary")
```

```
readxl::read_excel("spreadsheet.xls", sheet = 1, col_names = TRUE)
```

```
readxl::read_excel("spreadsheet.xls", sheet = 1, col_names = TRUE,
 col_types = c("text", "date", "numeric", "numeric"))
```

# Reading RODBC

Excel Windows Access Database Engine (ACE) ( JET) ODBC Excel . RODBC R  
. : JET / ACE .dll Windows / PC .

```
df <- sqlQuery(xlconn, "SELECT * FROM [SheetName$]")
close(xlconn)
```

**SQL JOIN UNION    Excel . JET / ACE SQL . : DML , SELECT**

```
joindf <- sqlQuery(xlconn, "SELECT t1.*, t2.* FROM [Sheet1$] t1
 INNER JOIN [Sheet2$] t2
 ON t1.[ID] = t2.[ID]")

uniondf <- sqlQuery(xlconn, "SELECT * FROM [Sheet1$]
 UNION
 SELECT * FROM [Sheet2$]")
```

**ODBC .**

```
otherwkbkdf <- sqlQuery(xlconn, "SELECT * FROM
 [Excel 12.0 Xml;HDR=Yes;
 Database=C:\\Path\\To\\Other\\Workbook.xlsx].[Sheet1$];")
```

## Reading gdata Excel .

**Stata, SPSS SAS**

foreign haven Stata, SPSS SAS . . . read .

```
loading the packages
library(foreign)
library(haven)
library(readstata13)
library(Hmisc)
```

```
reading Stata files with `foreign`
read.dta("path\to\your\data")
reading Stata files with `haven`
read_dta("path\to\your\data")
```

foreign Stata 7-12 stata (.dta) . . read.dta 13 . . Stata readstata13 haven . readstata13

```
reading recent Stata (13+) files with `readstata13`
read.dta13("path\to\your\data")
```

**SPSS SAS**

```
reading SPSS files with `foreign`
read.spss("path\to\your\data.sav", to.data.frame = TRUE)
reading SPSS files with `haven`
```

```

read_spss("path\to\your\data.sav")
read_sav("path\to\your\data.sav")
read_por("path\to\your\data.por")

reading SAS files with `foreign`
read.ssd("path\to\your\data")
reading SAS files with `haven`
read_sas("path\to\your\data")
reading native SAS files with `Hmisc`
sas.get("path\to\your\data") #requires access to saslib
Reading SA XPORT format (*.XPT) files
sasxport.get("path\to\your\data.xpt") # does not require access to SAS executable

```

SAS SET read.fwf . . . <https://github.com/ajdamico/SASci>.

```

write.foreign() . . .

writing to Stata, SPSS or SAS files with `foreign`
write.foreign(dataframe, datafile, codefile,
 package = c("SPSS", "Stata", "SAS"), ...)
write.foreign(dataframe, "path\to\data\file", "path\to\instruction\file", package = "Stata")

writing to Stata files with `foreign`
write.dta(dataframe, "file", version = 7L,
 convert.dates = TRUE, tz = "GMT",
 convert.factors = c("labels", "string", "numeric", "codes"))

writing to Stata files with `haven`
write_dta(dataframe, "path\to\your\data")

writing to Stata files with `readstata13`
save.dta13(dataframe, file, data.label = NULL, time.stamp = TRUE,
 convert.factors = TRUE, convert.dates = TRUE, tz = "GMT",
 add.rownames = FALSE, compress = FALSE, version = 117,
 convert.underscore = FALSE)

writing to SPSS files with `haven`
write_sav(dataframe, "path\to\your\data")

```

SPSS read.spss .

```

foreign::read.spss('data.sav', to.data.frame=TRUE, use.value.labels=FALSE,
 use.missings=TRUE, reencode='UTF-8')
to.data.frame if TRUE: return a data frame
use.value.labels if TRUE: convert variables with value labels into R factors with those
levels
use.missings if TRUE: information on user-defined missing values will be used to set the
corresponding values to NA.
reencode character strings will be re-encoded to the current locale. The default, NA, means
to do so in a UTF-8 locale, only.

```

**Feather** (:) **Python** **R** **Apache Arrow** . **data.frame** **tibble** .

```

library(feather)

path <- "filename.feather"
df <- mtcars

```

```

write_feather(df, path)

df2 <- read_feather(path)

head(df2)
A tibble: 6 x 11
mpg cyl disp hp drat wt qsec vs am gear carb
<dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl>
1 21.0 6 160 110 3.90 2.620 16.46 0 1 4 4
2 21.0 6 160 110 3.90 2.875 17.02 0 1 4 4
3 22.8 4 108 93 3.85 2.320 18.61 1 1 4 1
4 21.4 6 258 110 3.08 3.215 19.44 1 0 3 1
5 18.7 8 360 175 3.15 3.440 17.02 0 0 3 2
6 18.1 6 225 105 2.76 3.460 20.22 1 0 3 1

head(df)
A tibble: 6 x 11
mpg cyl disp hp drat wt qsec vs am gear carb
<dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl>
1 Mazda RX4 21.0 6 160 110 3.90 2.620 16.46 0 1 4 4
2 Mazda RX4 Wag 21.0 6 160 110 3.90 2.875 17.02 0 1 4 4
3 Datsun 710 22.8 4 108 93 3.85 2.320 18.61 1 1 4 1
4 Hornet 4 Drive 21.4 6 258 110 3.08 3.215 19.44 1 0 3 1
5 Hornet Sportabout 18.7 8 360 175 3.15 3.440 17.02 0 0 3 2
6 Valiant 18.1 6 225 105 2.76 3.460 20.22 1 0 3 1

```

.....

(Excel, SAS, SPSS, Stata) I / O : <https://riptutorial.com/ko/r/topic/5536---excel--sas--spss--stata--i---o>

# 99:

1. (x = character (), , = , = NA, = is.ordered (x), nmax = NA)  
2. ?factor .

factor .

1. integer .  
2. levels .  
3. factor .

, 1,000 .

```
set.seed(1)
Color <- sample(x = c("Red", "Blue", "Green", "Yellow"),
 size = 1000,
 replace = TRUE)
Color <- factor(Color)
```

Color .

```
#* 1. It is stored internally as an `integer` vector
typeof(Color)
```

```
[1] "integer"
```

```
#* 2. It maintains a `levels` attribute that shows the character representation of the values.
#* 3. Its class is stored as `factor`
attributes(Color)
```

```
$levels
[1] "Blue" "Green" "Red" "Yellow"
$class
[1] "factor"
```

( stringsAsFactors : Unauthorized Biography ). Color ., Color factor 1.7 .

```
#* Amount of memory required to store Color as a factor.
object.size(Color)
```

```
4624 bytes
```

```
#* Amount of memory required to store Color as a character
object.size(as.character(Color))
```

```
8232 bytes
```

```
head(Color)
```

```
[1] Blue Blue Green Yellow Red Yellow
Levels: Blue Green Red Yellow
```

```
head(as.numeric(Color))
```

```
[1] 1 1 2 4 3 4
```

R

```
head(levels(Color) [as.numeric(Color)])
```

```
[1] "Blue" "Blue" "Green" "Yellow" "Red" "Yellow"
```

```
head(Color)
```

```
[1] Blue Blue Green Yellow Red Yellow
Levels: Blue Green Red Yellow
```

2007 R (:stringsAsFactors ). 1.7 , R , 2007 .

R . R .

1. /

2.

. (: "", "" "") . , ( ) .

## Examples

R . . . R .

```
charvar <- rep(c("n", "c"), each = 3)
f <- factor(charvar)
f
levels(f)
```

```
> f
[1] n n n c c c
Levels: c n
> levels(f)
[1] "c" "n"
```

```
levels(factor(charvar, levels = c("n", "c")))

> levels(factor(charvar, levels = c("n", "c")))
[1] "n" "c"
```

```
> f <- factor(charvar, levels=c("n", "c"), labels=c("Newt", "Capybara"))
> f
[1] Newt Newt Newt Capybara Capybara Capybara
Levels: Newt Capybara
```

```
> Weekdays <- factor(c("Monday", "Wednesday", "Thursday", "Tuesday", "Friday", "Sunday",
"Saturday"))
> Weekdays
[1] Monday Wednesday Thursday Tuesday Friday Sunday Saturday
Levels: Friday Monday Saturday Sunday Thursday Tuesday Wednesday
> Weekdays <- factor(Weekdays, levels=c("Monday", "Tuesday", "Wednesday", "Thursday",
"Friday", "Saturday", "Sunday"), ordered=TRUE)
> Weekdays
[1] Monday Wednesday Thursday Tuesday Friday Sunday Saturday
Levels: Monday < Tuesday < Wednesday < Thursday < Friday < Saturday < Sunday
```

, droplevels()

```
> Weekend <- subset(Weekdays, Weekdays == "Saturday" | Weekdays == "Sunday")
> Weekend
[1] Sunday Saturday
Levels: Monday < Tuesday < Wednesday < Thursday < Friday < Saturday < Sunday
> Weekend <- droplevels(Weekend)
> Weekend
[1] Sunday Saturday
Levels: Saturday < Sunday
```

```
set.seed(1)
colorful <- sample(c("red", "Red", "RED", "blue", "Blue", "BLUE", "green", "gren"),
 size = 20,
 replace = TRUE)
colorful <- factor(colorful)
```

R

```
table(colorful)
```

```
colorful
blue Blue BLUE green gren red Red RED
 3 1 4 2 4 1 3 2
```

## factor ( factor\_approach )

```
factor(as.character(colorful),
 levels = c("blue", "Blue", "BLUE", "green", "gren", "red", "Red", "RED"),
 labels = c("Blue", "Blue", "Blue", "Green", "Green", "Red", "Red", "Red"))
```

```
[1] Green Blue Red Red Blue Red Red Red Blue Red Green Green Green
Blue Red Green
[17] Red Green Green Red
Levels: Blue Blue Blue Green Green Red Red Red
Warning message:
In `levels<-`(`*tmp*`, value = if (nl == nL) as.character(labels) else
paste0(labels, :
 duplicated levels in factors are deprecated
```

## ifelse ( ifelse\_approach )

```
factor(ifelse(colorful %in% c("blue", "Blue", "BLUE"),
 "Blue",
 ifelse(colorful %in% c("green", "gren"),
 "Green",
 "Red")))
```

```
[1] Green Blue Red Red Blue Red Red Red Blue Red Green Green Green
Blue Red Green
[17] Red Green Green Red
Levels: Blue Green Red
```

```
ifelse . ifelse
```

## ( list\_approach )

```
 factor levels
```

```
levels(colorful) <-
 list("Blue" = c("blue", "Blue", "BLUE"),
 "Green" = c("green", "gren"),
 "Red" = c("red", "Red", "RED"))
```

```
[1] Green Blue Red Red Blue Red Red Red Blue Red Green Green Green
Blue Red Green
[17] Red Green Green Red
Levels: Blue Green Red
```

```
Unit: microseconds
expr min lq mean median uq max neval cld
factor 78.725 83.256 93.26023 87.5030 97.131 218.899 100 b
ifelse 104.494 107.609 123.53793 113.4145 128.281 254.580 100 c
list_approach 49.557 52.955 60.50756 54.9370 65.132 138.193 100 a
```

ifelse . . .

**Factor R** as.character() x factor() as.factor()

```
, ,

standard
factor(c(1,1,2,2,3,3))
[1] 1 1 2 2 3 3
Levels: 1 2 3
```

factor()

```
factor(c(1,1,2,2,3,3),
 levels = c(1,2,3,4,5))
[1] 1 1 2 2 3 3
Levels: 1 2 3 4 5
```

```
factor(c(1,1,2,2,3,3),
 levels = c(1,2,3,4,5),
 labels = c("Fox", "Dog", "Cow", "Brick", "Dolphin"))
[1] Fox Fox Dog Dog Cow Cow
Levels: Fox Dog Cow Brick Dolphin
```

== != . . .

```
factor(c(1,1,2,2,3,3), levels = c(1,2,3)) == factor(c(1,1,2,2,3,3), levels = c(1,2,3,4,5))
Error in Ops.factor(factor(c(1, 1, 2, 2, 3, 3), levels = c(1, 2, 3)), :
 level sets of factors are different
```

**RHS R** . . .

< , <= , > = . . . ordered = TRUE factor ordered . . .

```
x <- factor(1:3, labels = c('low', 'medium', 'high'), ordered = TRUE)
print(x)
[1] low medium high
```

```

Levels: low < medium < high

y <- ordered(3:1, labels = c('low', 'medium', 'high'))
print(y)
[1] high medium low
Levels: low < medium < high

x < y
[1] TRUE FALSE FALSE

```

## Factor .

```

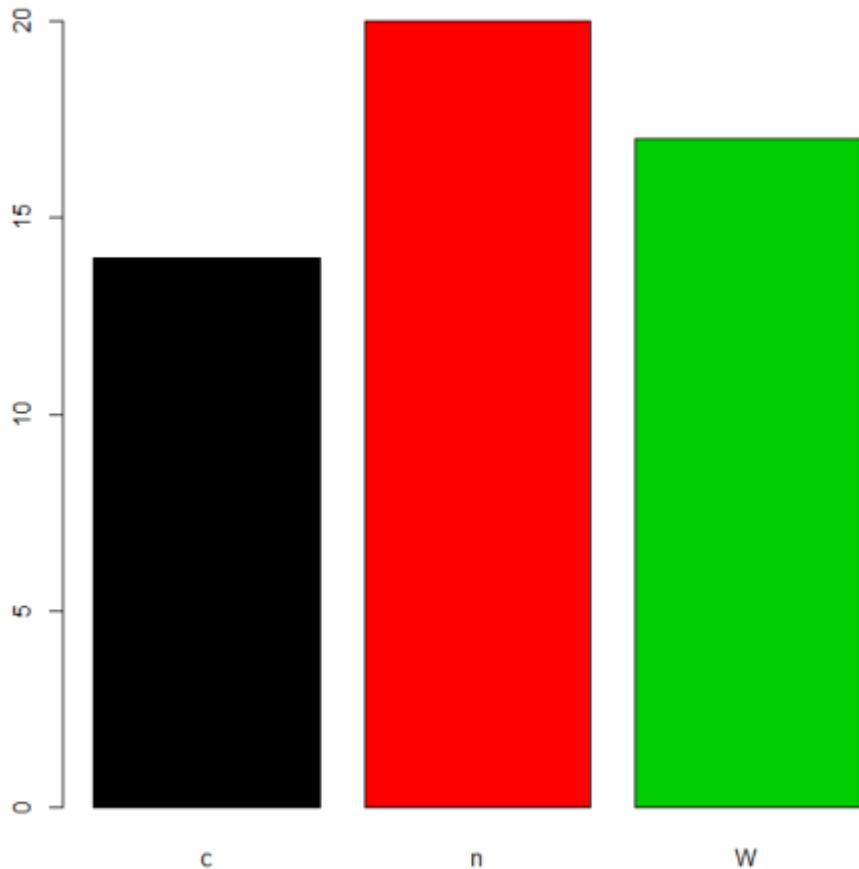
levels as.character .

charvar <- rep(c("W", "n", "c"), times=c(17,20,14))
f <- factor(charvar)
levels(f)
[1] "c" "n" "W"

levels (/) . , plot , :

plot(f,col=1:length(levels(f)))

```



```

levels levels labels ("" .). .

```

## 1.

```
levels .

ff <- factor(charvar, levels = c("n", "W", "c"))
levels(ff)
[1] "n" "W" "c"

gg <- factor(charvar, levels = c("W", "c", "n"))
levels(gg)
[1] "W" "c" "n"

, labels levels "" "" labels:

fm <- factor(as.numeric(f), levels = c(2,3,1),
 labels = c("nn", "WW", "cc"))
levels(fm)
[1] "nn" "WW" "cc"

fm <- factor(LETTERS[1:6], levels = LETTERS[1:4], # only 'A'-'D' as input
 labels = letters[1:4]) # but assigned to 'a'-'d'
fm
#[1] a b c d <NA> <NA>
#Levels: a b c d
```

## 2. relevel

```
level relevel ., base .

g<-relevel(f, "n") # moves n to be the first level
levels(g)
[1] "n" "c" "W"

f g .

all.equal(f, g)
[1] "Attributes: < Component \"levels\": 2 string mismatches >"
all.equal(f, g, check.attributes = F)
[1] TRUE
```

## 3.

```
, , levels reorder .levels .

table(g)
g
n c W
20 14 17

reorder help(reordered), :x, factor; x x . FUN x x levels . . .

g.ord <- reorder(g, rep(1, length(g)), FUN=sum) #increasing
levels(g.ord)
```

```
[1] "c" "W" "n"
```

( -1 )

```
g.ord.d <- reorder(g, rep(-1,length(g)), FUN=sum)
levels(g.ord.d)
[1] "n" "W" "c"
```

```
data.frame(f,g,g.ord,g.ord.d)[seq(1,length(g),by=5),] #just same lines
f g g.ord g.ord.d
1 W W W W
6 W W W W
11 W W W W
16 W W W W
21 n n n n
26 n n n n
31 n n n n
36 n n n n
41 c c c c
46 c c c c
51 c c c c
```

levels levels . iris ( help("iris") help("iris") Species Sepal.Width Species .

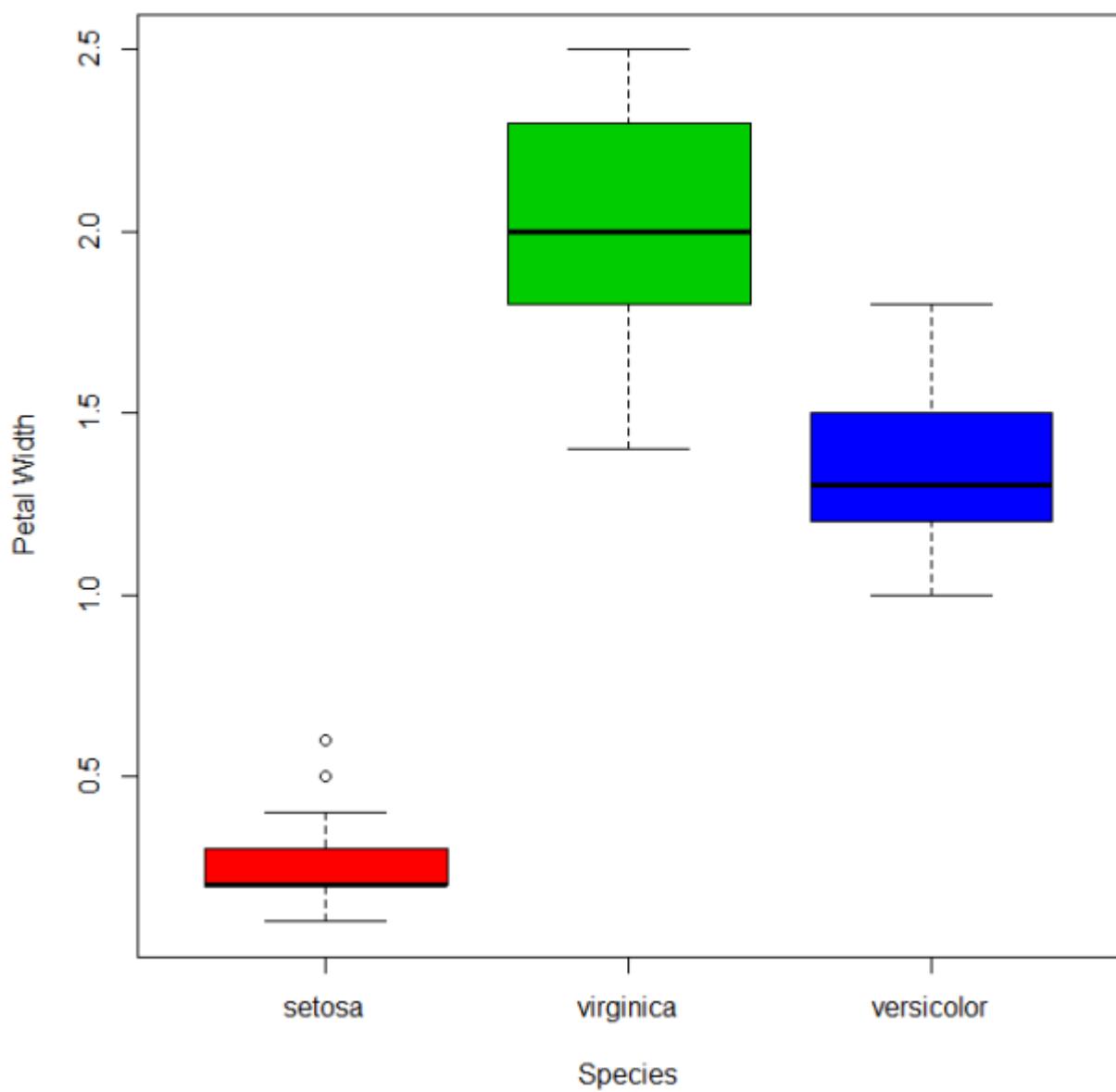
```
miris <- iris #help("iris") # copy the data
with(miris, tapply(Sepal.Width,Species,mean))
setosa versicolor virginica
3.428 2.770 2.974

miris$Species.o<-with(miris,reorder(Species,-Sepal.Width))
levels(miris$Species.o)
[1] "setosa" "virginica" "versicolor"
```

boxplot ( with(miris, boxplot(Petal.Width~Species) ) : **setosa , versicolor , virginica** .
Sepal.Width :

```
boxplot(Petal.Width~Species.o, data = miris,
 xlab = "Species", ylab = "Petal Width",
 main = "Iris Data, ordered by mean sepal width", varwidth = TRUE,
 col = 2:4)
```

### Iris Data, ordered by mean sepal width



```
levels levels levels . levels .
```

```
f1<-f
levels(f1)
[1] "c" "n" "W"
levels(f1) <- c("upper", "upper", "CAP") #rename and grouping
levels(f1)
[1] "upper" "CAP"

f2<-f1
levels(f2) <- c("upper", "CAP", "Number") #add Number level, which is empty
levels(f2)
[1] "upper" "CAP" "Number"
f2[length(f2):(length(f2)+5)]<-"Number" # add cases for the new level
table(f2)
f2
upper CAP Number
33 17 6
```

```
f3<-f1
levels(f3) <- list(G1 = "upper", G2 = "CAP", G3 = "Number") # The same using list
levels(f3)
[1] "G1" "G2" "G3"
f3[length(f3):(length(f3)+6)]<-"G3" ## add cases for the new level
table(f3)
f3
G1 G2 G3
33 17 7
```

-  
ordered . factors . levels , labels .

```
ordvar<-rep(c("Low", "Medium", "High"), times=c(7,2,4))

of<-ordered(ordvar,levels=c("Low", "Medium", "High"))
levels(of)
[1] "Low" "Medium" "High"

of1<-of
levels(of1)<- c("LOW", "MEDIUM", "HIGH")
levels(of1)
[1] "LOW" "MEDIUM" "HIGH"
is.ordered(of1)
[1] TRUE
of1
[1] LOW LOW LOW LOW LOW LOW MEDIUM MEDIUM HIGH HIGH HIGH HIGH HIGH

Levels: LOW < MEDIUM < HIGH
```

0

R . ( , , ) . 5 .

, , ,

. , 1 = , 2 = , 3 = , 4 = 5 : :

**4 , 4 , 3 , 1 , 2**

(:) . R ?

20 . .

```
set.seed(18)
ii <- sample(1:4, 20, replace=T)
ii
```

[1] 4 3 4 1 1 3 2 3 2 1 3 4 1 2 4 1 3 1 4 1

1 4 .

```
fii <- factor(ii, levels=1:4) # it is necessary to indicate the numeric levels
fii
```

```
[1] 4 3 4 1 1 3 2 3 2 1 3 4 1 2 4 1 3 1 4 1
: 1 2 3 4
```

```
levels(fii) <- c("empty", "low", "normal", "full")
fii
```

```
[1]
[11]
:
```

: <https://riptutorial.com/ko/r/topic/1104/>

# 100:

R

R (RSelenium, httr, curl, RCurl), (XML, xml2) (rvest).

API.

## Examples

### rvest

rvest Hadley Wickham Python BeautifulSoup . Hadley xml2 HTML libxml2 .

tidyverse , rvest .

- xml2::read\_html HTML xml2::read\_html ,
- CSS XPath html\_node html\_nodes .
- html\_text html\_table R .

### R Wikipedia

```
library(rvest)

url <- 'https://en.wikipedia.org/wiki/R_(programming_language)'

scrape HTML from website
url %>% read_html() %>%
 # select HTML tag with class="wikitable"
 html_node(css = '.wikitable') %>%
 # parse table into data.frame
 html_table() %>%
 # trim for printing
 dplyr::mutate(Description = substr(Description, 1, 70))

Release Date Description
1 0.16 This is the last alpha version developed primarily by Ihaka
2 0.49 1997-04-23 This is the oldest source release which is currently availab
3 0.60 1997-12-05 R becomes an official part of the GNU Project. The code is h
4 0.65.1 1999-10-07 First versions of update.packages and install.packages funct
5 1.0 2000-02-29 Considered by its developers stable enough for production us
6 1.4 2001-12-19 S4 methods are introduced and the first version for Mac OS X
7 2.0 2004-10-04 Introduced lazy loading, which enables fast loading of data
8 2.1 2005-04-18 Support for UTF-8 encoding, and the beginnings of internatio
9 2.11 2010-04-22 Support for Windows 64 bit systems.
10 2.13 2011-04-14 Adding a new compiler function that allows speeding up funct
11 2.14 2011-10-31 Added mandatory namespaces for packages. Added a new paralle
12 2.15 2012-03-30 New load balancing functions. Improved serialization speed f
13 3.0 2013-04-03 Support for numeric index values 231 and larger on 64 bit sy
```

```
data.frame . , NA
 . jQuery , read_html , RSelenium .
```

## rvest

```
library(rvest)

#Address of the login webpage
login<-
"https://stackoverflow.com/users/login?ssrc=head&returnurl=http%3a%2f%2fstackoverflow.com%2f"

#create a web session with the desired login address
pgsession<-html_session(login)
pgform<-html_form(pgsession)[[2]] #in this case the submit is the 2nd form
filled_form<-set_values(pgform, email="*****", password="*****")
submit_form(pgsession, filled_form)

#pre allocate the final results dataframe.
results<-data.frame()

#loop through all of the pages with the desired info
for (i in 1:5)
{
 #base address of the pages to extract information from
 url<-"http://stackoverflow.com/users/**********?tab=answers&sort=activity&page="
 url<-paste0(url, i)
 page<-jump_to(pgsession, url)

 #collect info on the question votes and question title
 summary<-html_nodes(page, "div .answer-summary")
 question<-matrix(html_text(html_nodes(summary, "div")), trim=TRUE), ncol=2, byrow = TRUE)

 #find date answered, hyperlink and whether it was accepted
 dateans<-html_node(summary, "span") %>% html_attr("title")
 hyperlink<-html_node(summary, "div a") %>% html_attr("href")
 accepted<-html_node(summary, "div") %>% html_attr("class")

 #create temp results then bind to final results
 rtemp<-cbind(question, dateans, accepted, hyperlink)
 results<-rbind(results, rtemp)
}

#Dataframe Clean-up
names(results)<-c("Votes", "Answer", "Date", "Accepted", "HyperLink")
results$Votes<-as.integer(as.character(results$Votes))
results$Accepted<-ifelse(results$Accepted=="answer-votes default", 0, 1)
```

5 . \*\*\*\*\* , .

: <https://riptutorial.com/ko/r/topic/2890--->

# 101:

- `ymd_hms` (..., `quiet = FALSE`, `tz = "UTC"`, `locale = Sys.getlocale ("LC_TIME")`)
- `(tzone = "")`
- `(, , tzone = attr (, "tzone"))`
- `(num = NULL, units = "seconds", ...)`
- `(num = NULL, units = "second", ...)`

CRAN .

```
install.packages("lubridate")
```

Github

```
library(devtools)
dev mode allows testing of development packages in a sandbox, without interfering
with the other packages you have installed.
dev_mode(on=T)
install_github("hadley/lubridate")
dev_mode(on=F)
```

**lubridate :**

```
vignette("lubridate")
```

**foo :**

```
help(foo) # help about function foo
?foo # same thing

Example
help("is.period")
?is.period
```

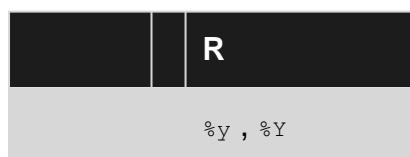
**foo :**

```
example("foo")

Example
example("interval")
```

## Examples

`lubridate` `datetime` `lubridate` .



R	
m (y d)	%m , %b , %h , %B
	%d , %e
h	%H , %I%p
m (h s)	%M
	%S

```
, , , , , "2016-07-22" ymd_hms() ymd() , : "2016-07-22 13:04:47" .
(: / , -) . .
```

---

**date** Date .

```
library(lubridate)

mdy(c(' 07/02/2016 ', '7 / 03 / 2016', ' 7 / 4 / 16 '))
[1] "2016-07-02" "2016-07-03" "2016-07-04"

ymd(c("20160724","2016/07/23","2016-07-25")) # inconsistent separators
[1] "2016-07-24" "2016-07-23" "2016-07-25"
```

---

## Datetimes

```
ymd_hms ymd_hm ymd_h . datetime as.POSIXct strptime tz "UTC" .
```

**datetime** POSIXct .

```
x <- c("20160724 130102","2016/07/23 14:02:01","2016-07-25 15:03:00")
ymd_hms(x, tz="EST")
[1] "2016-07-24 13:01:02 EST" "2016-07-23 14:02:01 EST"
[3] "2016-07-25 15:03:00 EST"

ymd_hms(x)
[1] "2016-07-24 13:01:02 UTC" "2016-07-23 14:02:01 UTC"
[3] "2016-07-25 15:03:00 UTC"
```

lubridate as.POSIXct strptime **datetimes** strptime .

parse_date_time	POSIXct	. % lubridate <b>datetime</b> ( :"ymd_hms" strptime strptime .
parse_date_time2	POSIXct; lt = TRUE ,	. strptime ( % ) .

## POSIXlt

```
fast.strptime POSIXlt; lt = FALSE . (- , / , :) % strptime .
POSIXct
```

```
x <- c('2016-07-22 13:04:47', '07/22/2016 1:04:47 pm')

parse_date_time(x, orders = c('mdy Imsp', 'ymd hms'))
[1] "2016-07-22 13:04:47 UTC" "2016-07-22 13:04:47 UTC"

x <- c('2016-07-22 13:04:47', '2016-07-22 14:47:58')

parse_date_time2(x, orders = 'Ymd HMS')
[1] "2016-07-22 13:04:47 UTC" "2016-07-22 14:47:58 UTC"

fast.strptime(x, format = '%Y-%m-%d %H:%M:%S')
[1] "2016-07-22 13:04:47 UTC" "2016-07-22 14:47:58 UTC"
```

parse\_date\_time2 fast.strptime C .

?parse\_date\_time .

## lubridate

Lubridate ymd() . y, m d - , .

```
mdy("07-21-2016") # Returns Date

[1] "2016-07-21"

mdy("07-21-2016", tz = "UTC") # Returns a vector of class POSIXt

"2016-07-21 UTC"

dmy("21-07-2016") # Returns Date

[1] "2016-07-21"

dmy(c("21.07.2016", "22.07.2016")) # Returns vector of class Date

[1] "2016-07-21" "2016-07-22"
```

## lubridate

```
date <- now()
date
"2016-07-22 03:42:35 IST"

year(date)
2016

minute(date)
42
```

```
wday(date, label = T, abbr = T)
[1] Fri
Levels: Sun < Mon < Tues < Wed < Thurs < Fri < Sat

day(date) <- 31
"2016-07-31 03:42:35 IST"

If an element is set to a larger value than it supports, the difference
will roll over into the next higher element
day(date) <- 32
"2016-08-01 03:42:35 IST"
```

. - . is.instant is.instant .

```
library(lubridate)

today_start <- dmy_hms("22.07.2016 12:00:00", tz = "IST") # default tz="UTC"
today_start
[1] "2016-07-22 12:00:00 IST"
is.instant(today_start)
[1] TRUE

now_dt <- ymd_hms(now(), tz="IST")
now_dt
[1] "2016-07-22 13:53:09 IST"
is.instant(now_dt)
[1] TRUE

is.instant("helloworld")
[1] FALSE
is.instant(60)
[1] FALSE
```

,

**lubridate** . .

```
create interval by subtracting two instants
today_start <- ymd_hms("2016-07-22 12:00:00", tz="IST")
today_start
[1] "2016-07-22 12:00:00 IST"
today_end <- ymd_hms("2016-07-22 23:59:59", tz="IST")
today_end
[1] "2016-07-22 23:59:59 IST"
span <- today_end - today_start
span
Time difference of 11.99972 hours
as.interval(span, today_start)
[1] 2016-07-22 12:00:00 IST--2016-07-22 23:59:59 IST

create interval using interval() function
span <- interval(today_start, today_end)
[1] 2016-07-22 12:00:00 IST--2016-07-22 23:59:59 IST
```

```
duration(60, "seconds")
[1] "60s"

duration(2, "minutes")
[1] "120s (~2 minutes)"
```

```
dseconds, dminutes . .
?quick_durations .
```

```
dseconds(60)
[1] "60s"

dhours(2)
[1] "7200s (~2 hours)"

dyears(1)
[1] "31536000s (~365 days)"
```

```
today_start + dhours(5)
[1] "2016-07-22 17:00:00 IST"

today_start + dhours(5) + dminutes(30) + dseconds(15)
[1] "2016-07-22 17:30:15 IST"
```

```
as.duration(span)
[1] "43199s (~12 hours)"
```

```
period seconds, hours . .
?quick_periods .
```

```
period(1, "hour")
[1] "1H 0M 0S"

hours(1)
[1] "1H 0M 0S"

period(6, "months")
[1] "6m 0d 0H 0M 0S"

months(6)
[1] "6m 0d 0H 0M 0S"

years(1)
[1] "1y 0m 0d 0H 0M 0S"
```

```
is.period . .
```

```
is.period(years(1))
[1] TRUE
```

```
is.period(dyears(1))
[1] FALSE
```

```
now_dt <- ymd_hms(now(), tz="IST")
now_dt
[1] "2016-07-22 13:53:09 IST"
```

```
round_date() - .
```

```
round_date(now_dt, "minute")
[1] "2016-07-22 13:53:00 IST"

round_date(now_dt, "hour")
[1] "2016-07-22 14:00:00 IST"

round_date(now_dt, "year")
[1] "2017-01-01 IST"
```

```
floor_date() - .
```

```
floor_date(now_dt, "minute")
[1] "2016-07-22 13:53:00 IST"

floor_date(now_dt, "hour")
[1] "2016-07-22 13:00:00 IST"

floor_date(now_dt, "year")
[1] "2016-01-01 IST"
```

```
ceiling_date() - .
```

```
ceiling_date(now_dt, "minute")
[1] "2016-07-22 13:54:00 IST"

ceiling_date(now_dt, "hour")
[1] "2016-07-22 14:00:00 IST"

ceiling_date(now_dt, "year")
[1] "2017-01-01 IST"
```

, DST .

```
start_2012 <- ymd_hms("2012-01-01 12:00:00")
[1] "2012-01-01 12:00:00 UTC"

period() considers leap year calculations.
start_2012 + period(1, "years")
[1] "2013-01-01 12:00:00 UTC"

Here duration() doesn't consider leap year calculations.
start_2012 + duration(1)
[1] "2012-12-31 12:00:00 UTC"
```

```
with_tz - .
```

```
nyc_time <- now("America/New_York")
nyc_time
[1] "2016-07-22 05:49:08 EDT"

corresponding Europe/Moscow time
with_tz(nyc_time, tzzone = "Europe/Moscow")
[1] "2016-07-22 12:49:08 MSK"
```

```
force_tz X .
```

```
nyc_time <- now("America/New_York")
nyc_time
[1] "2016-07-22 05:49:08 EDT"

force_tz(nyc_time, tzzone = "Europe/Moscow") # only timezone changes
[1] "2016-07-22 05:49:08 MSK"
```

: <https://riptutorial.com/ko/r/topic/2496/>

# 102:

## Examples

(      ).

, *logit log-odds* .      :

$$\sigma(t) = \frac{e^t}{e^t + 1} = \frac{1}{1 + e^{-t}}$$

-Inf; + Inf[ 0 1 . . .

```
family = binomial (family = binomial(link="logit") : logit family = binomial(link="logit")
glm .
```

RMS .

:

```
url <- "http://biostat.mc.vanderbilt.edu/wiki/pub/Main/DataSets/titanic.txt"
titanic <- read.csv(file = url, stringsAsFactors = FALSE)
```

.

.

```
titanic$age[is.na(titanic$age)] <- mean(titanic$age, na.rm = TRUE)
```

:

```
titanic.train <- glm(survived ~ pclass + sex + age,
 family = binomial, data = titanic)
```

:

```
summary(titanic.train)
```

:

```
Call:
glm(formula = survived ~ pclass + sex + age, family = binomial, data = titanic)
```

```
Deviance Residuals:
Min 1Q Median 3Q Max
-2.6452 -0.6641 -0.3679 0.6123 2.5615
```

```
Coefficients:
Estimate Std. Error z value Pr(>|z|)
```

```

(Intercept) 3.552261 0.342188 10.381 < 2e-16 ***
pclass2nd -1.170777 0.211559 -5.534 3.13e-08 ***
pclass3rd -2.430672 0.195157 -12.455 < 2e-16 ***
sexmale -2.463377 0.154587 -15.935 < 2e-16 ***
age -0.042235 0.007415 -5.696 1.23e-08 ***

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

(Dispersion parameter for binomial family taken to be 1)

Null deviance: 1686.8 on 1312 degrees of freedom
Residual deviance: 1165.7 on 1308 degrees of freedom
AIC: 1175.7

Number of Fisher Scoring iterations: 5

```

- . .
- . .
- , , z- ( Wald z- ) p- .
  - "dummified". . I .
  - 4 0.1 % .
  - .
  - ( ) .
  - (CI) confint confint .
- null Akaike Information Criterion (AIC) . .
  - AIC .
  - . (, ) .
- :

```

exp(coef(titanic.train)[3])

pclass3rd
0.08797765

```

1 3 1/10 .

```

confint(titanic.train)

Waiting for profiling to be done...
 2.5 % 97.5 %
(Intercept) 2.89486872 4.23734280
pclass2nd -1.58986065 -0.75987230
pclass3rd -2.81987935 -2.05419500
sexmale -2.77180962 -2.16528316
age -0.05695894 -0.02786211

```

(, ) .

```
with(titanic.train, pchisq(null.deviance - deviance, df.null - df.residual
, lower.tail = FALSE))
[1] 1.892539e-111
```

p- 0 .

: <https://riptutorial.com/ko/r/topic/2892-->

# 103:

```
file.path file.path .
```

```
dir .
```

## Examples

```
R .
```

```
. .
```

```
. getwd() ?setwd .
```

```
set.seed(1)
for (i in 1:3)
 write.table(
 data.frame(id = 1:2, v = sample(letters, 2)),
 file = sprintf("file201%s.csv", i)
)
```

```
CSV .
```

```
. .
```

```
3 . list list .
```

```
file_names = c("file2011.csv", "file2012.csv", "file2013.csv")
file_contents = lapply(setNames(file_names, file_names), read.table)

$file2011.csv
id v
1 1 g
2 2 j
#
$file2012.csv
id v
1 1 o
2 2 w
#
$file2013.csv
id v
1 1 f
2 2 w
```

```
str(file_contents) ?rbind ?lapply .
```

```
. .
```

```
?read.table ?write.table .:
```

- R 2 ( )
- ○ CSV
- ○ TSV
- ○ ○
- ○ SAS
- ○ SPSS
- ○ ○
- ○ MySQL
- ○ SQLite
- ○ PostgreSQL

: <https://riptutorial.com/ko/r/topic/5543/>

# 104:

(NLP)

## Examples

( ) . . . : ( ) . . .

. R tm ( ) .

```
require(tm)
doc1 <- "drugs hospitals doctors"
doc2 <- "smog pollution environment"
doc3 <- "doctors hospitals healthcare"
doc4 <- "pollution environment water"
corpus <- c(doc1, doc2, doc3, doc4)
tm_corpus <- Corpus(VectorSource(corpus))
```

Corpus VectorSource tm Corpus Corpus . VectorSource . tm\_corpus ( ) .

```
str(tm_corpus)
List of 4
$ 1:List of 2
..$ content: chr "drugs hospitals doctors"
..$ meta :List of 7
...$. author : chr(0)
...$. timestamp: POSIXlt[1:1], format: "2017-06-03 00:31:34"
...$. description : chr(0)
...$. heading : chr(0)
...$. id : chr "1"
...$. language : chr "en"
...$. origin : chr(0)
...- attr(*, "class")= chr "TextDocumentMeta"
..- attr(*, "class")= chr [1:2] "PlainTextDocument" "TextDocument"
[truncated]
```

Corpus , Corpus ( ) . tm tm\_map . tm\_map apply .

```
tm_corpus <- tm_map(tm_corpus, tolower)
tm_corpus <- tm_map(tm_corpus, removeWords, stopwords("english"))
tm_corpus <- tm_map(tm_corpus, removeNumbers)
tm_corpus <- tm_map(tm_corpus, PlainTextDocument)
tm_corpus <- tm_map(tm_corpus, stemDocument, language="english")
tm_corpus <- tm_map(tm_corpus, stripWhitespace)
tm_corpus <- tm_map(tm_corpus, PlainTextDocument)
```

```
tdm <- TermDocumentMatrix(tm_corpus)
```

```
<<TermDocumentMatrix (terms: 8, documents: 4)>>
```

```
Non-/sparse entries: 12/20
Sparsity : 62%
Maximal term length: 9
Weighting : term frequency (tf)
```

```
as.matrix(tdm)
```

	Docs				
Terms	character(0)	character(0)	character(0)	character(0)	character(0)
doctor	1	0	1	0	
drug	1	0	0	0	
environ	0	1	0	1	
healthcar	0	0	1	0	
hospit	1	0	1	0	
pollut	0	1	0	1	
smog	0	1	0	0	
water	0	0	0	1	

```
. (4 , 4) (:environment - environ).
```

```
/ (,).
```

: <https://riptutorial.com/ko/r/topic/10119/>--

# **105: % <> % : ?**

NAMESPACE .

## **Examples**

```
." " zzz.R utils.R . , :
```

```
#' Pipe operator
#'
#' @name %>%
#' @rdname pipe
#' @keywords internal
#' @export
#' @importFrom magrittr %>%
#' @usage lhs \%>\% rhs
NULL
```

% <> % : ? : <https://riptutorial.com/ko/r/topic/10547/-----lt--gt----->

# 106: R

" ( ) . .  
. , () . .

- Peng, RD (2011). . Science, 334 (6060), 1226-1227] .  
<http://doi.org/10.1126/science.1213847>
- Peng, Roger D. R. Leanpub, 2015. . <https://leanpub.com/reportwriting> .

## Examples

```
dput() dget()
() dput() .R
: .getwd() ?setwd .
dput(mtcars, file = 'df.txt')
```

```
dget() R GlobalEnvironment .
```

```
df <- dget('df.txt')
```

R . .

R . . R . . checkpoint .

2014-09-17 CRAN Microsoft R Archived Network . R . .

1. (R) .
2. checkpoint::checkpoint('YYYY-MM-DD') .

```
checkpoint R_home (~/) .checkpoint . . , checkpoint .R library() require() CRAN
```

PRO .

CONTRA .

R : <https://riptutorial.com/ko/r/topic/4087/--r>

# 107:

R ?

```
c(1,2,3) + c(1,2,3,4,5,6)
[1] 2 4 6 5 7 9
```

```
c(1,2,3,1,2,3) + c(1,2,3,4,5,6)
```

```
c(1,2,3) + c(1,2,3,4,5,6,7)
[1] 2 4 6 5 7 9 8
Warning message:
In c(1, 2, 3) + c(1, 2, 3, 4, 5, 6, 7) :
 longer object length is not a multiple of shorter object length
```

```
matrix(nrow =5, ncol = 2, 1:5)
 [,1] [,2]
 [1,] 1 1
 [2,] 2 2
 [3,] 3 3
 [4,] 4 4
 [5,] 5 5
```

## Examples

```
my_vec <- c(1,2,3,4,5,6,7,8,9,10)
my_vec[c(TRUE, FALSE)]
[1] 1 3 5 7 9
```

```
my_vec <- c("foo", "bar", "soap", "mix")
```

```
my_vec == "bar"
[1] FALSE TRUE FALSE FALSE
```

"" .

: <https://riptutorial.com/ko/r/topic/5649/>

# 108: ()

( "regex" "regexp" ) . R ?regex [Regex Docs](#) . SO / " R-regex pattern .

- "[AB]" A B .
- "[[:alpha:]]"
- "[[:lower:]]" . "[az]" . :ú .
- "[[:upper:]]" . "[AZ]" (:ú .
- "[[:digit:]]" 0, 1, 2, ... 9 "[0-9]" .

+ , \* ? . - + , \* 0 ? 0 1 .

- "^..." .
- "...\$" .

R

- R - ( "\\" R ). \s R \\s .
- R UTF-8 U . [\U{1F600}] [\U{1F600}] match , Ruby u .

[reg101](#) R .

[R Programming wikibook](#) .

## Examples

```
string <- ' some text on line one;
and then some text on line two '
```

" " . gsub .

R 3.2.0

```
gsub(pattern = "(^ | +$)",
 replacement = "",
 x = string)

[1] "some text on line one; \nand then some text on line two"
```

R 3.2.0

```
trimws(x = string)
[1] "some text on line one; \nand then some text on line two"
```

R 3.2.0

```
sub(pattern = "^\n",
 replacement = "",
 x = string)

[1] "some text on line one; \nand then some text on line two"
```

R3.2.0

```
trimws(x = string,
 which = "left")

[1] "some text on line one; \nand then some text on line two"
```

R 3.2.0

```
sub(pattern = " +$",
 replacement = "",
 x = string)

[1] " some text on line one; \nand then some text on line two"
```

R 3.2.0

```
trimws(x = string,
 which = "right")

[1] " some text on line one; \nand then some text on line two"
```

```
gsub(pattern = "\\s",
 replacement = "",
 x = string)
```

( \t ), ( \r \n ) .

"YYYYMMDD"

YYYYMMDD (: 20170101 results.csv .

\d{4} (0[1-9] | 1[012]) (0[1-9] | [12][0-9] | 3[01])

: 0000-9999 , 01-12 01-31 01-31 .

```
> grepl("\d{4}(0[1-9]|1[012])(0[1-9]|12)[0-9]|3[01])", "20170101")
[1] TRUE
> grepl("\d{4}(0[1-9]|1[012])(0[1-9]|12)[0-9]|3[01])", "20171206")
[1] TRUE
> grepl("\d{4}(0[1-9]|1[012])(0[1-9]|12)[0-9]|3[01])", "29991231")
[1] TRUE
```

: (: 20170229 (2017 )).

```
> grep1("\d{4}(0[1-9]|1[012])(0[1-9]|12)[0-9]|3[01])", "20170229")
[1] TRUE
```

```
is.Date <- function(x) {return(!is.na(as.Date(as.character(x), format = '%Y%m%d')))}
```

```
> is.Date(c("20170229", "20170101", 20170101))
[1] FALSE TRUE TRUE
```

regex 50 Commonwealth / Territory ([www.50states.com](http://www.50states.com)).

```
regex <-
```

"(A[LKSZR]) | (C[AOT]) | (D[EC]) | (F[ML]) | (G[AU]) | (HI) | (I[DLNA]) | (K[SY]) | (LA) | (M[EHDAINSOT]) | (N[EVHJMYCD]) | (N[EVHJMYCD])

```
> test <- c("AL", "AZ", "AR", "AJ", "AS", "DC", "FM", "GU", "PW", "FL", "AJ", "AP")
> grepl(us.states.pattern, test)
[1] TRUE TRUE TRUE FALSE TRUE TRUE TRUE TRUE TRUE FALSE FALSE
>
```

50 state.abb R- state.abb from state (:

```
> data(state)
> test %in% state.abb
[1] TRUE TRUE TRUE FALSE FALSE FALSE FALSE FALSE FALSE FALSE TRUE FALSE FALSE
```

AL, AZ, AR, FL 50 TRUE .

```
us.phones.regex <- "^\s*(\+\s*\1(-?\s+))*[0-9]\{3\}\s*-?\s*[0-9]\{3\}\s*-?\s*[0-9]\{4\}$"
```

```
us.phones.regex <- "^\s*(\s*\s*(-?|\s+))*[0-9]\{3\}\s*-?\s*[0-9]\{3\}\s*-?\s*[0-9]\{4\$"

phones.OK <- c("305-123-4567", "305 123 4567", "+1-786-123-4567",
 "+1 786 123 4567", "7861234567", "786 - 123 4567", "+ 1 786 - 123 4567")

phones.NOK <- c("124-456-78901", "124-456-789", "124-456-78 90",
 "124-45 6-7890", "12 4-456-7890")
```

```
> grepl(us.phones.regex, phones.OK)
[1] TRUE TRUE TRUE TRUE TRUE TRUE TRUE TRUE
>
```

```
> grep(us.phones.regex, phones.NOK)
[1] FALSE FALSE FALSE FALSE FALSE FALSE
>
```

• \s , .

R

R " " grep , sub , gsub . 3 4 . .

```

x <- c("a\nb", "c\td", "e f")
x # how it's stored
[1] "a\nb" "c\td" "e f"
cat(x) # how it will be seen with cat
#a
#b c d e f

gsub(patt="\n|\t", repl=" ", x)
#[1] "a b" "c d" "e f"

```

```
() . . . 4
```

Perl POSIX

R      POSIX R      perl = TRUE

1

```
perl = TRUE look-ahead look-behind .
● " (?<=A)B" B , A "ABACADABRA" , "abacadabra" "aBacadabra" .
```

( ) : <https://riptutorial.com/ko/r/topic/5748/--->

# 109:

for . for .

```
for ([index] in [domain]) {
 [body]
}
```

1. [index] [domain] .
2. [domain] .
3. [body] .

, for .

```
x <- 1:4
cumulative_sum <- 0
for (i in x){
 cumulative_sum <- cumulative_sum + x[i]
}
cumulative_sum
```

# For

For . , apply . for . , for .

' 'for . . .

```
[output] <- [vector_of_length]
for ([index] in [length_safe_domain]) {
 [output][index] <- [body]
}
```

( . " x\_squared <- x^2 ).

```
x <- 1:100
x_squared <- vector("numeric", length = length(x))
for (i in seq_along(x)){
 x_squared[i] <- x[i]^2
}
```

, x\_squared x "numeric" x\_squared . seq\_along " " . seq\_along for . for (i in  
1:length(x)) , x 0 1:0 (0 R ). ).

apply for apply . for .

For . .

```
mtcars for (colMeans).
```

```
column_mean_loop <- vector("numeric", length(mtcars))
for (k in seq_along(mtcars)){
 column_mean_loop[k] <- mean(mtcars[[k]])
}
```

for

```
col_mean_fn <- function(x) mean(x)
column_mean_apply <- vapply(mtcars, col_mean_fn, numeric(1))
```

apply

## Examples

( mtcars .

A :

```
squared_deviance <- vector("list", length(mtcars))
for (i in seq_along(mtcars)){
 squared_deviance[[i]] <- (mtcars[[i]] - mean(mtcars[[i]]))^2
}
```

squared deviance 11 .

```
class(squared_deviance)
length(squared deviance)
```

B:

```
squared_deviance <- vector("list", length(mtcars))
Squared_deviance <- setNames(squared_deviance, names(mtcars))
for (k in names(mtcars)){
 squared_deviance[[k]] <- (mtcars[[k]] - mean(mtcars[[k]]))^2
}
```

```
data.frame ? . . . data.frame for .
```

```

squared_deviance <- mtcars #copy the original
squared_deviance[TRUE] <- NA #replace with NA or do squared_deviance[,] <- NA
for (i in seq_along(mtcars)){
 squared_deviance[[i]] <- (mtcars[[i]] - mean(mtcars[[i]]))^2
}

```

```
dim(squared_deviance)
[1] 32 11
```

(B) .

For

- 1. for for
- 2. for
- 3. \*apply
- 4. colMeans

```
column_mean_poor <- NULL
for (i in 1:length(mtcars)){
 column_mean_poor[i] <- mean(mtcars[[i]])
}
```

```
column_mean_optimal <- vector("numeric", length(mtcars))
for (i in seq_along(mtcars)){
 column_mean_optimal <- mean(mtcars[[i]])
}
```

vapply

```
column_mean_vapply <- vapply(mtcars, mean, numeric(1))
```

colMeans

```
column_mean_colMeans <- colMeans(mtcars)
```

( )

```
Unit: microseconds
 expr min lq mean median uq max neval cld
 poor 240.986 262.0820 287.1125 275.8160 307.2485 442.609 100 d
optimal 220.313 237.4455 258.8426 247.0735 280.9130 362.469 100 c
 vapply 107.042 109.7320 124.4715 113.4130 132.6695 202.473 100 a
colMeans 155.183 161.6955 180.2067 175.0045 194.2605 259.958 100 b
```

for for . for .

```
for .
```

```
vapply . ().
```

```
colMeans vapply . colMeans as.matrix(mtcars A data.frame) vapply .
```

## : while repeat

```
R while repeat .
```

---

```
while
```

```
while .
```

```
while (condition) {
 ## do something
 ## in loop body
}
```

```
condition . condition TRUE condition FALSE (break). for while . . .
```

```
for (i in 0:4) {
 cat(i, "\n")
}
0
1
2
3
4

i <- 0
while (i < 5) {
 cat(i, "\n")
 i <- i + 1
}
0
1
2
3
4
```

```
while i <- i + 1 .
```

---

```
break while .
```

```
iter <- 0
while (TRUE) {
 if (runif(1) < 0.25) {
 break
 } else {
 iter <- iter + 1
 }
```

```
}
```

```
iter
```

```
[1] 4
```

```
condition TRUE break . iter PRNG () .
```

---

### repeat

```
repeat while (TRUE) { ## something } .
```

```
repeat ({
 ## do something
 ## in loop body
})
```

```
{ } () . repeat ,
```

```
iter <- 0
repeat ({
 if (runif(1) < 0.25) {
 break
 } else {
 iter <- iter + 1
 }
})
iter
#[1] 2
```

---

### break break

```
break . , .
```

```
while (TRUE) {
 while (TRUE) {
 cat("inner loop\n")
 break
 }
 cat("outer loop\n")
}
```

```
. , .
```

```
while (TRUE) {
 cat("outer loop body\n")
 while (TRUE) {
 cat("inner loop body\n")
 x <- runif(1)
 if (x < .3) {
 break
 } else {
 cat(sprintf("x is %.5f\n", x))
```

```
 }
 }
}

break return . return break return() .
```

```
function() {
 while (TRUE) {
 cat("outer loop body\n")
 while (TRUE) {
 cat("inner loop body\n")
 x <- runif(1)
 if (x < .3) {
 return()
 } else {
 cat(sprintf("x is %.5f\n", x))
 }
 }
 }
})()
```

```
(exit) <<- .
```

```
exit <- FALSE
while (TRUE) {
 cat("outer loop body\n")
 while (TRUE) {
 cat("inner loop body\n")
 x <- runif(1)
 if (x < .3) {
 exit <<- TRUE
 break
 } else {
 cat(sprintf("x is %.5f\n", x))
 }
 }
 if (exit) break
}
```

: <https://riptutorial.com/ko/r/topic/2201/>--

# 110:

```
. , R . R . set , setdiff , intersect , union , setequal %in% . v %in% S S v .
```

```
R , Rcpp .
```

## Examples

```
R .
```

```
v = "A"
w = c("A", "A")
```

```
. R .
```

```
setequal(v, w)
TRUE
```

```
:
```

```
x = c(1, 2, 3)
y = c(2, 4)

union(x, y)
1 2 3 4

intersect(x, y)
2

setdiff(x, y)
1 3
```

```
, ?union .
```

```
%in% .
```

```
v = "A"
w = c("A", "A")

w %in% v
TRUE TRUE

v %in% w
TRUE
```

```
().
```

```
%in% TRUE FALSE FALSE .
```

```
c(1, NA) %in% c(1, 2, 3, 4)
TRUE FALSE
```

```
?`%in%` .
```

```
"""
```

```
(x, y) x X Y Y expand.grid expand.grid expand.grid:
```

```
X = c(1, 1, 2)
Y = c(4, 5)

expand.grid(X, Y)

Var1 Var2
1 1 4
2 1 4
3 2 4
4 1 5
5 1 5
6 2 5
```

```
data.frame. "" . unique lapply do.call .
```

```
m = do.call(expand.grid, lapply(list(X, Y), unique))

Var1 Var2
1 1 4
2 2 4
3 1 5
4 2 5
```

```

```

```
f(x, y) .
```

```
m$p = with(m, Var1*Var2)
Var1 Var2 p
1 1 4 4
2 2 4 8
3 1 5 5
4 2 5 10
```

```
, 2 . outer .
```

```
uX = unique(X)
uY = unique(Y)

outer(setNames(uX, uX), setNames(uY, uY), `*`)

4 5
1 4 5
```

```
2 8 10
```

/ /

unique ( ).

```
x = c(2, 1, 1, 2, 1)
unique(x)
2 1
```

duplicated duplicated :

```
duplicated(x)
FALSE FALSE TRUE TRUE TRUE
```

anyDuplicated(x) > 0L

/

```
xtab_set <- function(A, B){
 both <- union(A, B)
 inA <- both %in% A
 inB <- both %in% B
 return(table(inA, inB))
}

A = 1:20
B = 10:30

xtab_set(A, B)

inB
inA FALSE TRUE
FALSE 0 10
TRUE 9 11
```

Venn

: <https://riptutorial.com/ko/r/topic/1383/>

# 111: I/O()

## Examples

```
rgdal rgdal rgdal . readOGR readOGR . ArcGIS dsn shapefile . layer shapefile (map
map.shp).
```

```
library(rgdal)
readOGR(dsn = "path\to\the\folder\containing\the\shapefile", layer = "map")
```

**shapefile** writeOGR . R. dsn layer . **obligatory 4.** ogrDrivers() . **shapefile** ArcGis QGis  
driver = "ESRI Shapefile" .

```
writeOGR(Rmap, dsn = "path\to\the\folder\containing\the\shapefile", layer = "map",
 driver = "ESRI Shapefile")
```

```
tmap rgdal::readOGR() read_shape() . read_shape() read_shape() . tmap .
```

I/O() : <https://riptutorial.com/ko/r/topic/5538/---i---o---->

# 112:

I/O

## Examples

**map ()**

```
maps map() R
```

:

```
require(maps)
map()
```



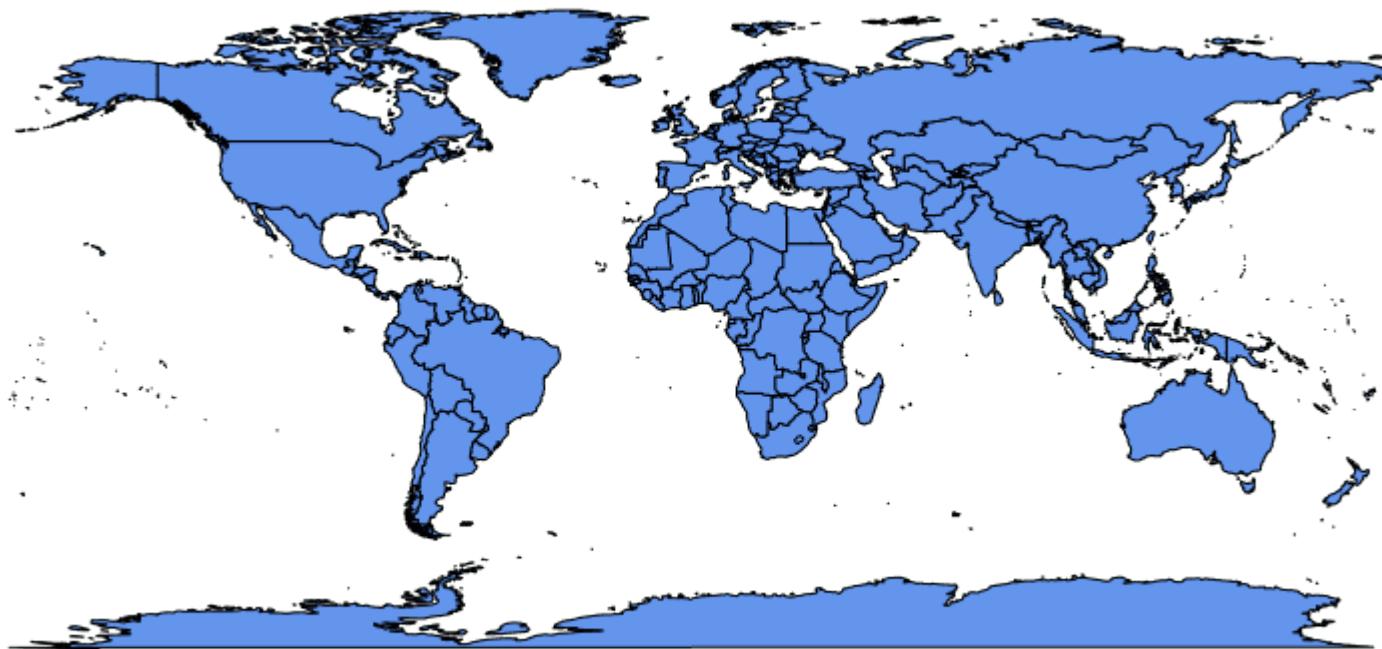
col 16

```
require(maps)
map(col = "cornflowerblue")
```



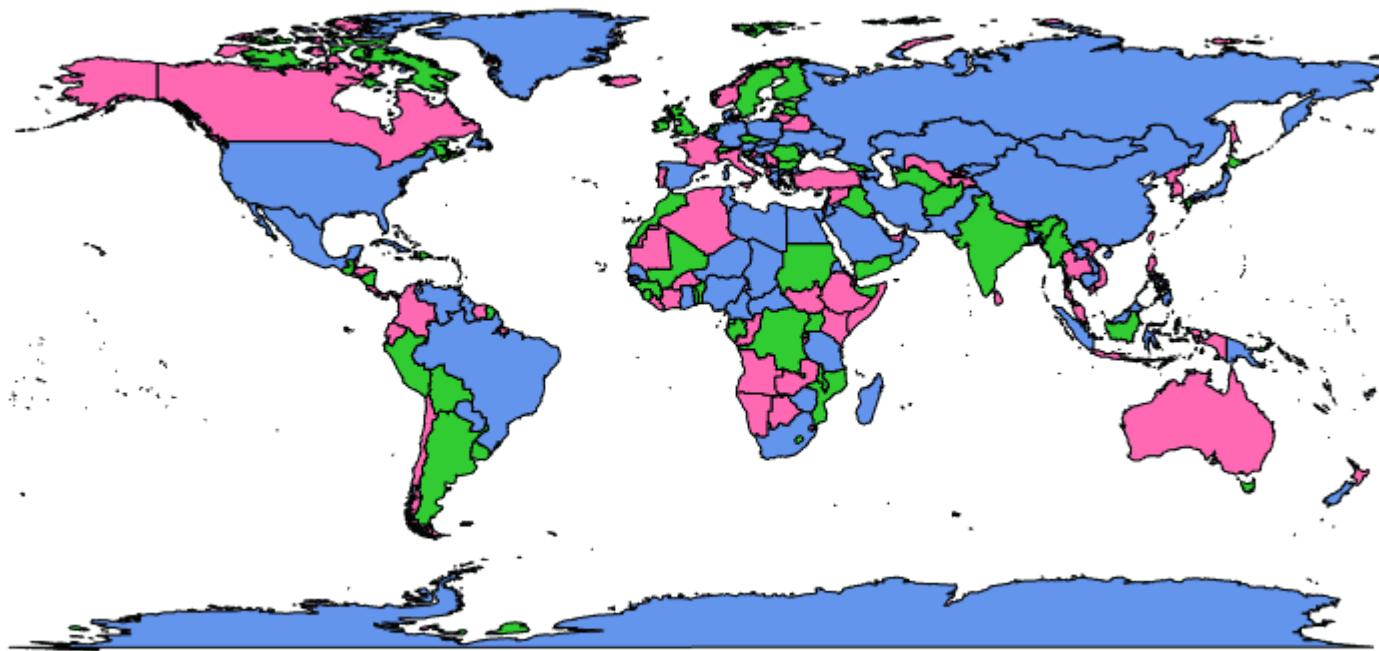
```
col fill = TRUE
```

```
require(maps)
map(fill = TRUE, col = c("cornflowerblue"))
```



```
fill = TRUE , col .
```

```
require(maps)
map(fill = TRUE, col = c("cornflowerblue", "limegreen", "hotpink"))
```



```

col
.
"choropleth"
.

choropleth database "county" "state" map() unemp county.fips "state" .

require(maps)
if(require(mapproj)) { # mapproj is used for projection="polyconic"
color US county map by 2009 unemployment rate
match counties to map using FIPS county codes
Based on J's solution to the "Choropleth Challenge"
Code improvements by Hack-R (hack-r.github.io)

load data
unemp includes data for some counties not on the "lower 48 states" county
map, such as those in Alaska, Hawaii, Puerto Rico, and some tiny Virginia
cities
data(unemp)
data(county.fips)

define color buckets
colors = c("paleturquoise", "skyblue", "cornflowerblue", "blueviolet", "hotpink",
"darkgrey")
unemp$colorBuckets <- as.numeric(cut(unemp$unemp, c(0, 2, 4, 6, 8, 10, 100)))
leg.txt <- c("<2%", "2-4%", "4-6%", "6-8%", "8-10%", ">10%")

align data with map definitions by (partial) matching state, county
names, which include multiple polygons for some counties

```

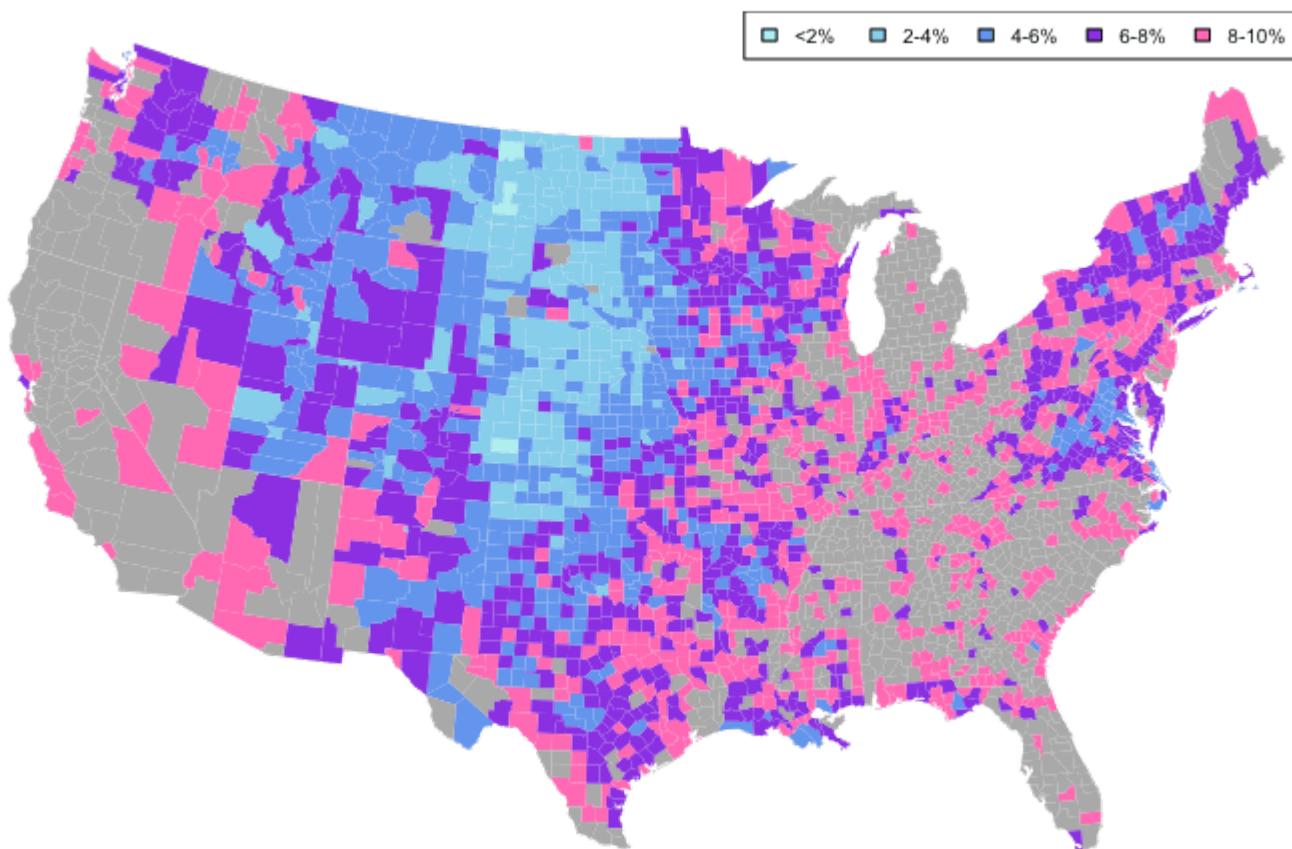
```

cnty.fips <- county.fips$fips[match(map("county", plot=FALSE)$names,
 county.fips$polyname)]
colorsmatched <- unemp$colorBuckets[match(cnty.fips, unemp$fips)]

draw map
par(mar=c(1, 1, 2, 1) + 0.1)
map("county", col = colors[colorsmatched], fill = TRUE, resolution = 0,
 lty = 0, projection = "polyconic")
map("state", col = "white", fill = FALSE, add = TRUE, lty = 1, lwd = 0.1,
 projection="polyconic")
title("unemployment by county, 2009")
legend("topright", leg.txt, horiz = TRUE, fill = colors, cex=0.6)
}

```

## unemployment by county, 2009



## Google Viz 50

50 ( ) choropleth () .

50 Google . Google API googleVis , ggmap RgoogleMaps .

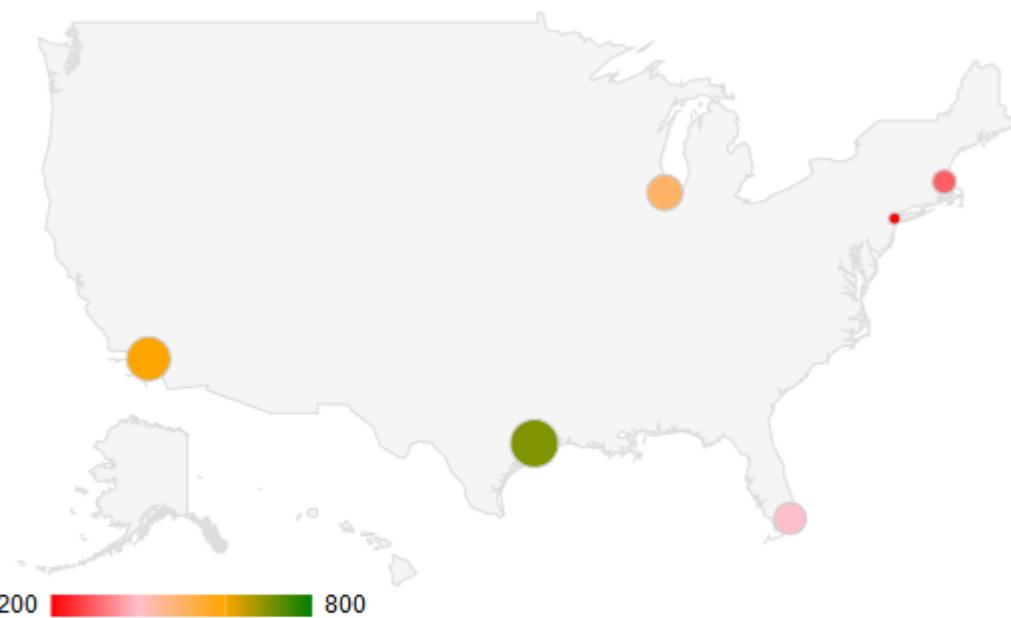
```

require(googleVis)

G4 <- gvisGeoChart(CityPopularity, locationvar='City', colorvar='Popularity',
 options=list(region='US', height=350,
 displayMode='markers',
 colorAxis="{values:[200,400,600,800],
 colors:[\red, \pink, \orange, \green]}"))

```

```
)
plot(G4)
```

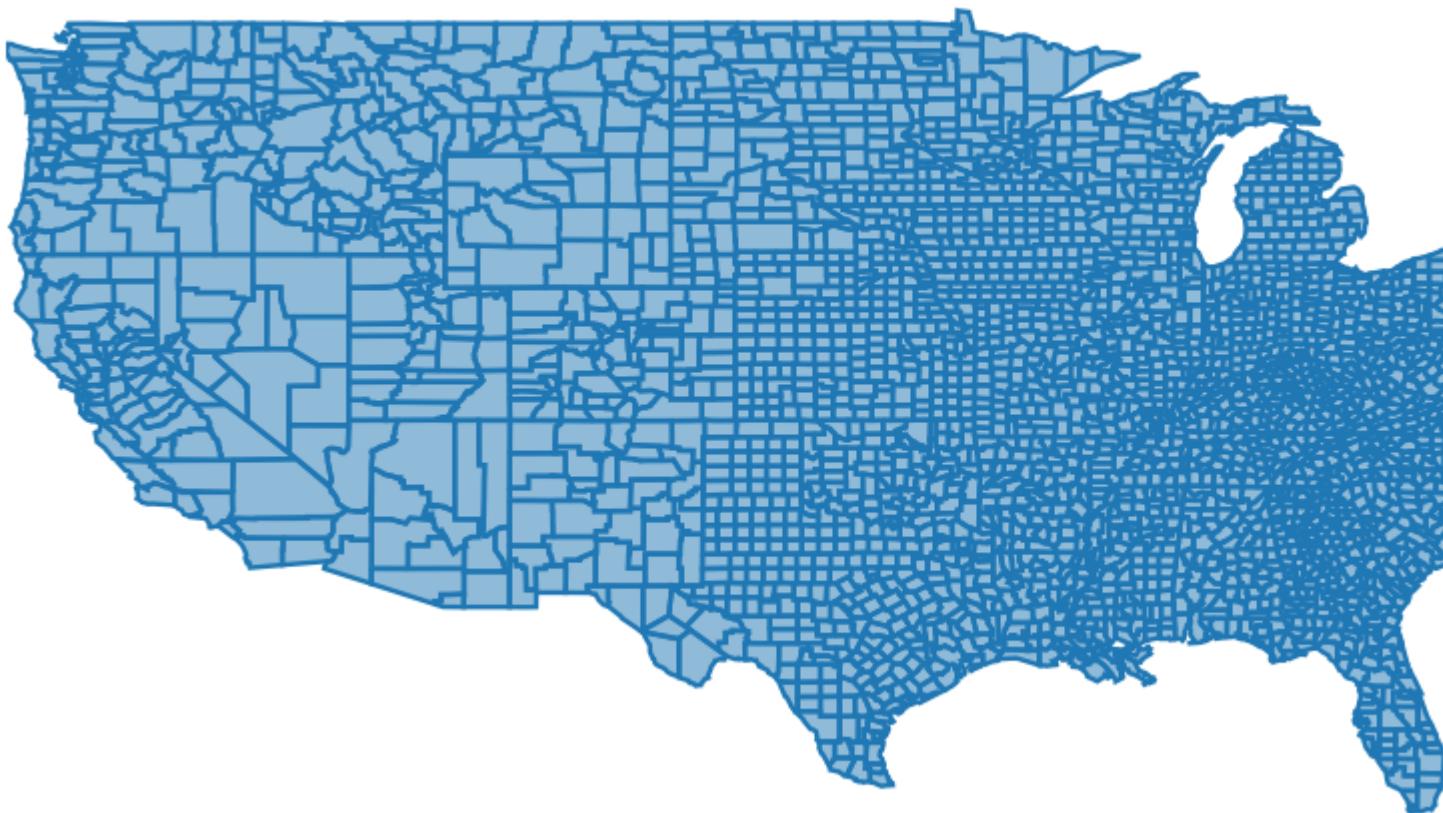


Data: CityPopularity • Chart ID: GeoChartID28504adb439a • googleVis-0.5.2  
R version 3.1.0 (2014-04-10) • Google Terms of Use • Documentation and Data Policy

```
gvisGeoChart() maps map() choropleth maps . colorvar locationvar colorvar . options
(height), (markers) (colors colorAxis colors) .
```

```
plotly , . plotly . (plot_ly() ggplotly()), plotly "" (plot_geo() plot_mapbox()), . .
```

```
library(plotly)
map_data("county") %>%
 group_by(group) %>%
 plot_ly(x = ~long, y = ~lat) %>%
 add_polygons() %>%
 layout(
 xaxis = list(title = "", showgrid = FALSE, showticklabels = FALSE),
 yaxis = list(title = "", showgrid = FALSE, showticklabels = FALSE)
)
```



```
plot_geo() plot_mapbox() plot_ly() plot_ly() . . .
```

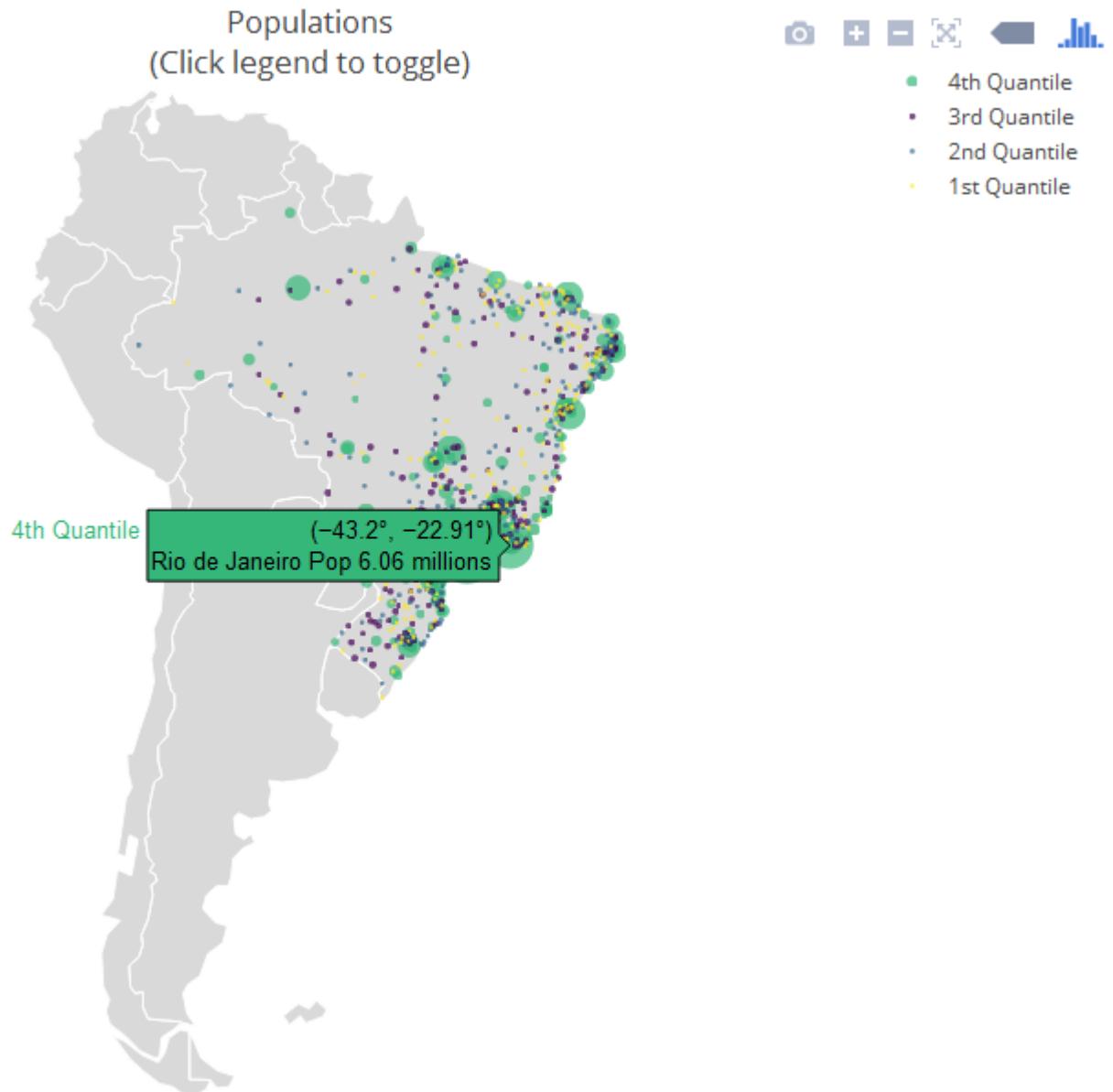
```
layout.geo / "" . maps world.cities "" .
```

```
:poph (poph .q quantile .ge . . .
```

```
library(maps)
dfb <- world.cities[world.cities$country.etc=="Brazil",]
library(plotly)
dfb$poph <- paste(dfb$name, "Pop", round(dfb$pop/1e6,2), " millions")
dfb$q <- with(dfb, cut(pop, quantile(pop), include.lowest = T))
levels(dfb$q) <- paste(c("1st", "2nd", "3rd", "4th"), "Quantile")
dfb$q <- as.ordered(dfb$q)

ge <- list(
 scope = 'south america',
 showland = TRUE,
 landcolor = toRGB("gray85"),
 subunitwidth = 1,
 countrywidth = 1,
 subunitcolor = toRGB("white"),
 countrycolor = toRGB("white")
)

plot_geo(dfb, lon = ~long, lat = ~lat, text = ~poph,
 marker = ~list(size = sqrt(pop/10000) + 1, line = list(width = 0)),
 color = ~q, locationmode = 'country names') %>%
 layout(geo = ge, title = 'Populations
(Click legend to toggle)')
```



## HTML

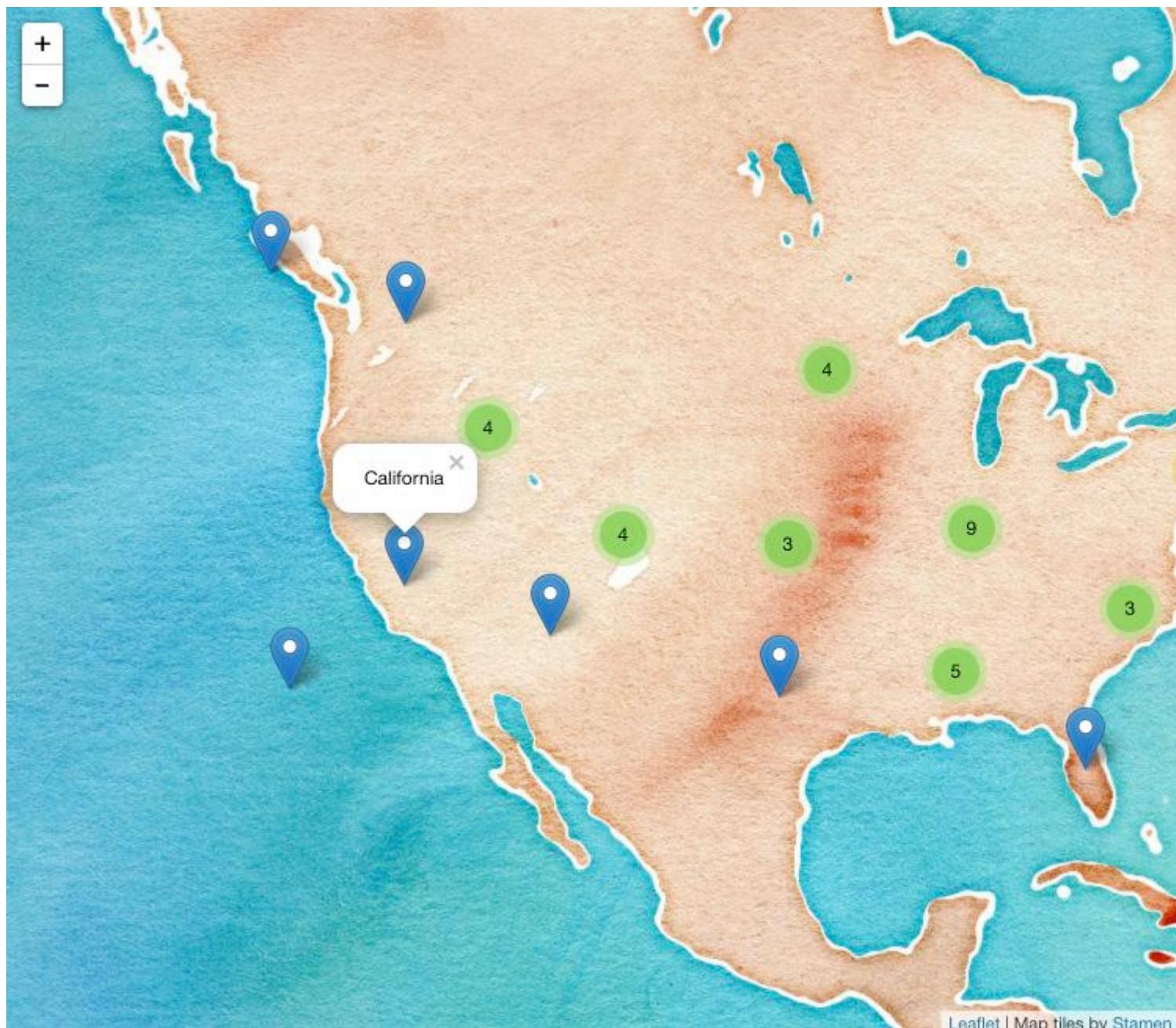
```
. RStudio htmlwidgets leaflet htmlwidgets RMarkdown Shiny .

leaflet() () . . (choroplet) . leaflet() data.frame function-style ~ quotation .

state.name state.center :

library(leaflet)

data.frame(state.name, state.center) %>%
 leaflet() %>%
 addProviderTiles('Stamen.Watercolor') %>%
 addMarkers(lng = ~x, lat = ~y,
 popup = ~state.name,
 clusterOptions = markerClusterOptions())
```



Leaflet | Map tiles by Stamen

(, .)

## Dynamic Leaflet

Shiny .

```
ui leafletOutput() renderLeaflet() .

library(shiny)
library(leaflet)

ui <- fluidPage(
 leafletOutput("my_leaf")
)

server <- function(input, output, session){

 output$my_leaf <- renderLeaflet({
```

```

 leaflet() %>%
 addProviderTiles('Hydda.Full') %>%
 setView(lat = -37.8, lng = 144.8, zoom = 10)

 })

}

shinyApp(ui, server)

renderLeaflet

leafletProxy()

leaflet leafletProxy

library(shiny)
library(leaflet)

ui <- fluidPage(
 sliderInput(inputId = "slider",
 label = "values",
 min = 0,
 max = 100,
 value = 0,
 step = 1),
 leafletOutput("my_leaf")
)

server <- function(input, output, session){
 set.seed(123456)
 df <- data.frame(latitude = sample(seq(-38.5, -37.5, by = 0.01), 100),
 longitude = sample(seq(144.0, 145.0, by = 0.01), 100),
 value = seq(1,100))

 ## create static element
 output$my_leaf <- renderLeaflet({

 leaflet() %>%
 addProviderTiles('Hydda.Full') %>%
 setView(lat = -37.8, lng = 144.8, zoom = 8)

 })

 ## filter data
 df_filtered <- reactive({
 df[df$value >= input$slider,]
 })

 ## respond to the filtered data
 observe({

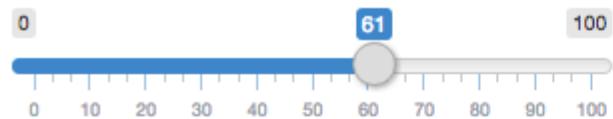
 leafletProxy(mapId = "my_leaf", data = df_filtered()) %>%
 clearMarkers() %>% ## clear previous markers
 addMarkers()

 })
}

```

```
shinyApp(ui, server)
```

### values



: <https://riptutorial.com/ko/r/topic/1372/>

## 113: 2

	.
	-
	.
importFrom	( )

## Examples

roxygen2

## roxygen2

roxygen2 Hadley Wickham

```
#' R . . .
```

```
#' @author The Author
```

, :

```
mean<-function(x) sum(x)/length(x)
```

( ).

```
#' Mean
#'
#' A function to compute the mean of a vector
#' @param x A numeric vector
#' @keyword mean
#' @importFrom base sum
#' @export
#' @examples
#' mean(1:3)
#' \dontrun{ mean(1:1e99) }
mean<-function(x) sum(x)/length(x)
```

- #' Mean , .
- @param . @export @export .
- @keyword .
- @importFrom . @import .

- @example .
    - .
    - \dontrun - - -.
- 

devtools::document() . devtools::check()

2 : <https://riptutorial.com/ko/r/topic/5171/-2>

# 114:

R , .

R . R .

## Examples

```
"table" (data.frame , table ,
```

```

(plain-text) () :
```

```
: . getwd() ?setwd .
```

```
..w = options()$width
options(width = 500) # reduce text wrapping
sink(file = "mytab.txt")
 summary(mtcars)
sink()
options(width = ..w)
rm(..w)
```

```
CSV() .
```

```
: . getwd() ?setwd .
```

```
write.csv(mtcars, file="mytab.csv")
```

- ```
_____  
• knitr::kable  
•  
• tables::tabular  
•  
• xtable
```

```
utils Sweave LaTeX , .
```

- ```

• Knitr RMarkdown
```

: <https://riptutorial.com/ko/r/topic/9039/>

# 115:

## Examples

R CPU (:

```
system.time(print("hello world"))

[1] "hello world"
user system elapsed
0 0 0
```

```
system.time({
 library(numbers)
 Primes(1,10^5)
})
```

```
fibb <- function (n) {
 if (n < 3) {
 return(c(0,1)[n])
 } else {
 return(fibb(n - 2) + fibb(n -1))
 }
}

system.time(fibb(30))
```

proc.time ()

proc.time()      CPU ( proc.time() . . .

```
proc.time()

user system elapsed
284.507 120.397 515029.305
```

```
t1 <- proc.time()
fibb <- function (n) {
 if (n < 3) {
 return(c(0,1)[n])
 } else {
 return(fibb(n - 2) + fibb(n -1))
 }
}
print("Time one")
```

```
print(proc.time() - t1)

t2 <- proc.time()
fibb(30)

print("Time two")
print(proc.time() - t2)
```

```
source('~/active-rstudio-document')
```

```
[1] "Time one"
user system elapsed
0 0 0

[1] "Time two"
user system elapsed
1.534 0.012 1.572
```

```
system.time() / proc.time() .
```

```
print(t1 <- system.time(replicate(1000, 12^2)))
user system elapsed
0.000 0.000 0.002
```

```
proc.time .
```

```
str(t1)
Class 'proc_time' Named num [1:5] 0 0 0.002 0 0
..- attr(*, "names")= chr [1:5] "user.self" "sys.self" "elapsed" "user.child" ...
```

Hadley Wickham [lineprof](#). auto.arima .

```
library(lineprof)
library(forecast)

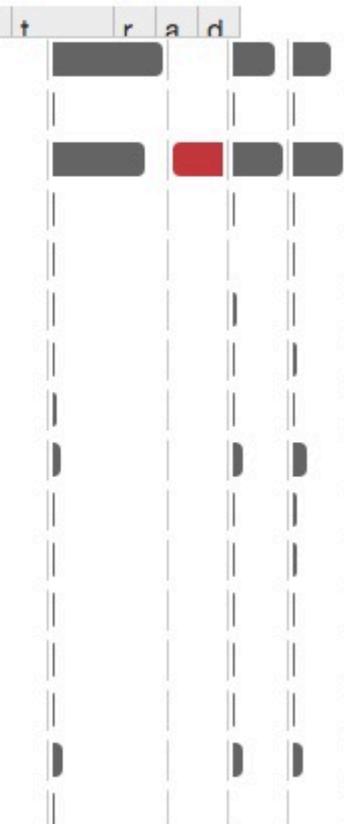
l <- lineprof(auto.arima(AirPassengers))
shine(l)
```

```
. R . . .
```

# Line profiling

Back

```
| Source code
1 | nsdiffs/OCSBtest
2 | nsdiffs/diff
3 | nsdiffs/OCSBtest
4 | diff/diff.ts
5 | ndiffs/suppressWarnings
6 | ndiffs/diff
7 | diff/diff.ts
8 | try/tryCatch
9 | myarima/suppressWarnings
10 |
11 | myarima/suppressWarnings
12 | myarima
13 | myarima/suppressWarnings
14 |
15 | myarima/suppressWarnings
16 | data.frame
```



## Microbenchmark

```
system.time(print("hello world"))

[1] "hello world"
user system elapsed
0 0 0
```

```
system.time proc.time proc.time . "hello world" 1 1 . . .
```

```
library(microbenchmark)
microbenchmark(print("hello world"))

Unit: microseconds
expr min lq mean median uq max neval
print("hello world") 26.336 29.984 44.11637 44.6835 45.415 158.824 100
```

```
print("hello world") 100 44 . "hello world" 100 .
```

```
print("hello world") cat("hello world\n") .
```

```
microbenchmark(cat("hello world\n"))

Unit: microseconds
```

```
expr min lq mean median uq max neval
cat("hello world\\n") 14.093 17.6975 23.73829 19.319 20.996 119.382 100
```

```
cat() print()
```

```
microbenchmark
```

```
microbenchmark(print("hello world"), cat("hello world\\n"))
Unit: microseconds
expr min lq mean median uq max neval
print("hello world") 29.122 31.654 39.64255 34.5275 38.852 192.779 100
cat("hello world\\n") 9.381 12.356 13.83820 12.9930 13.715 52.564 100
```

```
microbenchmark " "
```

```
6 data.table .
```

```
:
```

```
id , time status data.table . id . . . 7 false .
```

```
library(microbenchmark)
library(data.table)

set.seed(20160723)
dt <- data.table(id = c(rep(seq(1:10000), each = 10)),
 time = c(rep(seq(1:10000), 10)),
 status = c(sample(c(TRUE, FALSE), 10000*10, replace = TRUE)))
setkey(dt, id, time) ## create copies of the data so the 'updates-by-reference' don't affect
other expressions
dt1 <- copy(dt)
dt2 <- copy(dt)
dt3 <- copy(dt)
dt4 <- copy(dt)
dt5 <- copy(dt)
dt6 <- copy(dt)

microbenchmark(

 expression_1 = {
 dt1[dt1[order(time), .I[.N], by = id]$V1, status := status * time < 7]
 },

 expression_2 = {
 dt2[,status := c(.SD[-.N, status], .SD[.N, status * time > 7]), by = id]
 },

 expression_3 = {
 dt3[dt3[, .N, by = id][,cumsum(N)], status := status * time > 7]
 },

 expression_4 = {
 y <- dt4[, .SD[.N], by=id]
 dt4[y, status := status & time > 7]
 },

 expression_5 = {
```

```

y <- dt5[, .SD[.N, .(time, status)], by = id][time > 7 & status]
dt5[y, status := FALSE]
},
expression_6 = {
 dt6[dt6[, .I == .I[which.max(time)], by = id]$V1 & time > 7, status := FALSE]
},
times = 10L ## specify the number of times each expression is evaluated
)
Unit: milliseconds
expr min lq mean median uq max neval
expression_1 11.646149 13.201670 16.808399 15.643384 18.78640 26.321346 10
expression_2 8051.898126 8777.016935 9238.323459 8979.553856 9281.93377 12610.869058 10
expression_3 3.208773 3.385841 4.207903 4.089515 4.70146 5.654702 10
expression_4 15.758441 16.247833 20.677038 19.028982 21.04170 36.373153 10
expression_5 7552.970295 8051.080753 8702.064620 8861.608629 9308.62842 9722.234921 10
expression_6 18.403105 18.812785 22.427984 21.966764 24.66930 28.607064 10

```

expression\_3 .

**data.table -**

**data.table - data.table**

: <https://riptutorial.com/ko/r/topic/2149/>--

# 116:

## Examples

combn .

```
combn(LETTERS, 3)

Showing only first 10.
[,1] [,2] [,3] [,4] [,5] [,6] [,7] [,8] [,9] [,10]
[1,] "A" "A" "A" "A" "A" "A" "A" "A" "A" "A"
[2,] "B" "B" "B" "B" "B" "B" "B" "B" "B" "B"
[3,] "C" "D" "E" "F" "G" "H" "I" "J" "K" "L"
```

expand.grid .

```
expand.grid(LETTERS, LETTERS, LETTERS)
or
do.call(expand.grid, rep(list(LETTERS), 3))

Showing only first 10.
Var1 Var2 Var3
1 A A A
2 B A A
3 C A A
4 D A A
5 E A A
6 F A A
7 G A A
8 H A A
9 I A A
10 J A A
```

, outer .

```
FUN here is used as a function executed on each resulting pair.
in this case it's string concatenation.
outer(LETTERS, LETTERS, FUN=paste0)

Showing only first 10 rows and columns
[,1] [,2] [,3] [,4] [,5] [,6] [,7] [,8] [,9] [,10]
[1,] "AA" "AB" "AC" "AD" "AE" "AF" "AG" "AH" "AI" "AJ"
[2,] "BA" "BB" "BC" "BD" "BE" "BF" "BG" "BH" "BI" "BJ"
[3,] "CA" "CB" "CC" "CD" "CE" "CF" "CG" "CH" "CI" "CJ"
[4,] "DA" "DB" "DC" "DD" "DE" "DF" "DG" "DH" "DI" "DJ"
[5,] "EA" "EB" "EC" "ED" "EE" "EF" "EG" "EH" "EI" "EJ"
[6,] "FA" "FB" "FC" "FD" "FE" "FF" "FG" "FH" "FI" "FJ"
[7,] "GA" "GB" "GC" "GD" "GE" "GF" "GG" "GH" "GI" "GJ"
[8,] "HA" "HB" "HC" "HD" "HE" "HF" "HG" "HH" "HI" "HJ"
[9,] "IA" "IB" "IC" "ID" "IE" "IF" "IG" "IH" "II" "IJ"
[10,] "JA" "JB" "JC" "JD" "JE" "JF" "JG" "JH" "JI" "JJ"
```

```
choose(length(LETTERS), 5)
[1] 65780
```

```
length(letters)^5
[1] 11881376
```

: <https://riptutorial.com/ko/r/topic/5836/>

# 117:

```
caret R . . . , , , .
```

```
caret .
```

## Examples

```
preProcess() . . . x preProcess()
```

```
preProcess() method . .
```

- 1.
- 2.
3. Box-Cox / Yeo-Johnson /
- 4.
- 5.
- 6.
- 7.
8. PCA
9. ICA
- 10.

```
mtcars , .
```

```
auto_index <- createDataPartition(mtcars$mpg, p = .8,
 list = FALSE,
 times = 1)

mt_train <- mtcars[auto_index,]
mt_test <- mtcars[-auto_index,]

process_mtcars <- preProcess(mt_train, method = c("center", "scale", "spatialSign"))

mtcars_train_transf <- predict(process_mtcars, mt_train)
mtcars_test_tranf <- predict(process_mtcars, mt_test)
```

: <https://riptutorial.com/ko/r/topic/4271/>

# 118: -

## Examples

### Rcpp

```
(first) x_i cos(x_{i-1} + 1) len .
```

```
repeatedCosPlusOne <- function(first, len) {
 x <- numeric(len)
 x[1] <- first
 for (i in 2:len) {
 x[i] <- cos(x[i-1] + 1)
 }
 return(x)
}
```

```
(cos(x[i-1]+1)) for , . R "x+1" R .
```

### Rcpp C++ .

```
library(Rcpp)
cppFunction("NumericVector repeatedCosPlusOneRcpp(double first, int len) {
 NumericVector x(len);
 x[0] = first;
 for (int i=1; i < len; ++i) {
 x[i] = cos(x[i-1]+1);
 }
 return x;
}")
```

```
all.equal(repeatedCosPlusOne(1, 1e6), repeatedCosPlusOneRcpp(1, 1e6))
[1] TRUE
system.time(repeatedCosPlusOne(1, 1e6))
user system elapsed
1.274 0.015 1.310
system.time(repeatedCosPlusOneRcpp(1, 1e6))
user system elapsed
0.028 0.001 0.030
```

```
, Rcpp R 1.31 0.03 1 .
```

```
Rcpp , len (first) x_i cos(x_{i-1} + 1) :
```

```
repeatedCosPlusOne <- function(first, len) {
 x <- numeric(len)
 x[1] <- first
 for (i in 2:len) {
 x[i] <- cos(x[i-1] + 1)
```

```

 }
 return(x)
}

```

R

```

library(compiler)
repeatedCosPlusOneCompiled <- cmpfun(repeatedCosPlusOne)

```

```

all.equal(repeatedCosPlusOne(1, 1e6), repeatedCosPlusOneCompiled(1, 1e6))
[1] TRUE
system.time(repeatedCosPlusOne(1, 1e6))
user system elapsed
1.175 0.014 1.201
system.time(repeatedCosPlusOneCompiled(1, 1e6))
user system elapsed
0.339 0.002 0.341

```

1.20 0.34 1

repeatedCosPlusOne , Reduce :

```

iterFunc <- function(init, n, func) {
 funcs <- replicate(n, func)
 Reduce(function(.., f) f(..), funcs, init = init, accumulate = TRUE)
}
repeatedCosPlusOne_vec <- function(first, len) {
 iterFunc(first, len - 1, function(.) cos(. + 1))
}

```

repeatedCosPlusOne\_vec "" repeatedCosPlusOne . 2

```

library(microbenchmark)
microbenchmark(
 repeatedCosPlusOne(1, 1e4),
 repeatedCosPlusOne_vec(1, 1e4)
)
#> Unit: milliseconds
#>
#> expr min lq mean median uq max
neval cld
#> repeatedCosPlusOne(1, 10000) 8.349261 9.216724 10.22715 10.23095 11.10817 14.33763
100 a
#> repeatedCosPlusOne_vec(1, 10000) 14.406291 16.236153 17.55571 17.22295 18.59085 24.37059
100 b

```

- : [https://riptutorial.com/ko/r/topic/1203/-----](https://riptutorial.com/ko/r/topic/1203/)

## Examples

N

```
tm

require(RWeka)
require(tau)
require(tm)
require(tm.plugin.webmining)
require(wordcloud)

Scrape Google Finance -----
googlefinance <- WebCorpus(GoogleFinanceSource("NASDAQ:LFVN"))

Scrape Google News -----
lv.googlenews <- WebCorpus(GoogleNewsSource("LifeVantage"))
p.googlenews <- WebCorpus(GoogleNewsSource("Protandim"))
ts.googlenews <- WebCorpus(GoogleNewsSource("TrueScience"))

Scrape NYTimes -----
lv.nytimes <- WebCorpus(NYTimesSource(query = "LifeVantage", appid = nytimes_appid))
p.nytimes <- WebCorpus(NYTimesSource("Protandim", appid = nytimes_appid))
ts.nytimes <- WebCorpus(NYTimesSource("TrueScience", appid = nytimes_appid))

Scrape Reuters -----
lv.reutersnews <- WebCorpus(ReutersNewsSource("LifeVantage"))
p.reutersnews <- WebCorpus(ReutersNewsSource("Protandim"))
ts.reutersnews <- WebCorpus(ReutersNewsSource("TrueScience"))

Scrape Yahoo! Finance -----
lv.yahoofinance <- WebCorpus(YahooFinanceSource("LFVN"))

Scrape Yahoo! News -----
lv.yahoonews <- WebCorpus(YahooNewsSource("LifeVantage"))
p.yahoonews <- WebCorpus(YahooNewsSource("Protandim"))
ts.yahoonews <- WebCorpus(YahooNewsSource("TrueScience"))

Scrape Yahoo! Inplay -----
lv.yahooinplay <- WebCorpus(YahooInplaySource("LifeVantage"))

Text Mining the Results -----
corpus <- c(googlefinance, lv.googlenews, p.googlenews, ts.googlenews, lv.yahoofinance,
lv.yahoonews, p.yahoonews,
ts.yahoonews, lv.yahooinplay) #lv.nytimes, p.nytimes, ts.nytimes, lv.reutersnews,
p.reutersnews, ts.reutersnews,

inspect(corpus)
wordlist <- c("lfvn", "lifevantage", "protandim", "truescience", "company", "fiscal",
"nasdaq")

ds0.1g <- tm_map(corpus, content_transformer(tolower))
ds1.1g <- tm_map(ds0.1g, content_transformer(removeWords), wordlist)
ds1.1g <- tm_map(ds1.1g, content_transformer(removeWords), stopwords("english"))
```

```

ds2.1g <- tm_map(ds1.1g, stripWhitespace)
ds3.1g <- tm_map(ds2.1g, removePunctuation)
ds4.1g <- tm_map(ds3.1g, stemDocument)

tdm.1g <- TermDocumentMatrix(ds4.1g)
dtm.1g <- DocumentTermMatrix(ds4.1g)

findFreqTerms(tdm.1g, 40)
findFreqTerms(tdm.1g, 60)
findFreqTerms(tdm.1g, 80)
findFreqTerms(tdm.1g, 100)

findAssocs(dtm.1g, "skin", .75)
findAssocs(dtm.1g, "scienc", .5)
findAssocs(dtm.1g, "product", .75)

tdm89.1g <- removeSparseTerms(tdm.1g, 0.89)
tdm9.1g <- removeSparseTerms(tdm.1g, 0.9)
tdm91.1g <- removeSparseTerms(tdm.1g, 0.91)
tdm92.1g <- removeSparseTerms(tdm.1g, 0.92)

tdm2.1g <- tdm92.1g

Creates a Boolean matrix (counts # docs w/terms, not raw # terms)
tdm3.1g <- inspect(tdm2.1g)
tdm3.1g[tdm3.1g>=1] <- 1

Transform into a term-term adjacency matrix
termMatrix.1gram <- tdm3.1g %*% t(tdm3.1g)

inspect terms numbered 5 to 10
termMatrix.1gram[5:10,5:10]
termMatrix.1gram[1:10,1:10]

Create a WordCloud to Visualize the Text Data -----
notsparse <- tdm2.1g
m = as.matrix(notsparse)
v = sort(rowSums(m), decreasing=TRUE)
d = data.frame(word = names(v), freq=v)

Create the word cloud
pal = brewer.pal(9,"BuPu")
wordcloud(words = d$word,
 freq = d$freq,
 scale = c(3,.8),
 random.order = F,
 colors = pal)

```



```
RColorBrewer random.order
```

```
1-gram .
```

```
n-gram . TDM n-gram .
```

```
R n-gram tm bi-gram n-gram . , ()
```

```
, tm . tau textcnt() tokenizer .
```

```
tokenize_ngrams <- function(x, n=3)
return(rownames(as.data.frame(unclass(textcnt(x, method="string", n=n)))))
```

```
tm RWeka :
```

```
BigramTokenizer
BigramTokenizer <- function(x) NGramTokenizer(x, Weka_control(min = 2, max = 2))
```

```
1-gram .
```

```
Create an n-gram Word Cloud -----
tdm.ng <- TermDocumentMatrix(ds5.1g, control = list(tokenize = BigramTokenizer))
dtm.ng <- DocumentTermMatrix(ds5.1g, control = list(tokenize = BigramTokenizer))

Try removing sparse terms at a few different levels
tdm89.ng <- removeSparseTerms(tdm.ng, 0.89)
tdm9.ng <- removeSparseTerms(tdm.ng, 0.9)
tdm91.ng <- removeSparseTerms(tdm.ng, 0.91)
tdm92.ng <- removeSparseTerms(tdm.ng, 0.92)

notsparse <- tdm91.ng
m = as.matrix(notsparse)
v = sort(rowSums(m), decreasing=TRUE)
```

```

d = data.frame(word = names(v), freq=v)

Create the word cloud
pal = brewer.pal(9, "BuPu")
wordcloud(words = d$word,
 freq = d$freq,
 scale = c(3,.8),
 random.order = F,
 colors = pal)

```



## Hack-R

: [https://riptutorial.com/ko/r/topic/3579/-](https://riptutorial.com/ko/r/topic/3579/)

# 120: R .

( ) . . .

## Examples

. . . (2017 5 ).

2017 5 .

1

2

## R

R .

```
library("devtools")
library("twitteR")
library("ROAuth")
```

```
api_key <- XXXXXXXXXXXXXXXXXXXXXXXX
api_secret <- XXXXXXXXXXXXXXXXXXXXXXXX
access_token <- XXXXXXXXXXXXXXXXXXXXXXXX
access_token_secret <- XXXXXXXXXXXXXXXXXXXXXXXX

setup_twitter_oauth(api_key, api_secret)
```

XXXXXXXXXXXXXXXXXXXXXX ( ).

```
search.string <- "#coffee"
no.of.tweets <- 1000

c_tweets <- searchTwitter(search.string, n=no.of.tweets, lang="en")
```

"" 1000 .

. ( sapply ).

```
coffee_tweets = sapply(c_tweets, function(t) t$getText())

coffee_tweets <- sapply(coffee_tweets, function(row) iconv(row, "latin1", "ASCII", sub=""))
```

```
head .
```

```
head(coffee_tweets)
```

R . : <https://riptutorial.com/ko/r/topic/10086/-r-->

# 121: (%>% )

magrittr , dplyr R R

- lhs %>% rhs # rhs(lhs)
- rhs(lhs, a = 1) lhs %>% rhs (a = 1) #
- lhs %>% rhs (a = 1, b =.) # rhs(a = 1, b = lhs) rhs(a = 1, b = lhs)
- lhs %<>% rhs # lhs <- rhs(lhs) lhs <- rhs(lhs)
- lhs %\$% rhs (a) # with(lhs, rhs(lhs\$a))
- lhs %T>% rhs # { rhs(lhs); lhs } { rhs(lhs); lhs }

lhs	rhs
magrittr . magrittr	

%>%

magrittr dplyr ( dplyr magrittr ) . API " " tidyverse .  
magrittr %<>% , %\$% T %T>% . ( + , [ , [ , etc. ] )  
( + , \* , ^ , & , %in% ) ?' %>% ' help(' %>% ') pkg: magrittr ).

RStudio Ctrl+Shift+M ( Windows Linux ), Cmd+Shift+M ( Mac ) Cmd+Shift+M .

- ()
- ( object %>% rm() object )

## Examples

%>% . magrittr . (LHS) (RHS) . :

```
library(magrittr)

1:10 %>% mean
[1] 5.5

is equivalent to
mean(1:10)
```

```
[1] 5.5

. years . .

years <- factor(2008:2012)

nesting
as.numeric(as.character(years))

piping
years %>% as.character %>% as.numeric
```

## RHS (Right Hand Side) LHS () . . .

```
example with grepl
its syntax:
grepl(pattern, x, ignore.case = FALSE, perl = FALSE, fixed = FALSE, useBytes = FALSE)

note that the `substring` result is the *2nd* argument of grepl
grepl("Wo", substring("Hello World", 7, 11))

piping while naming other arguments
"Hello World" %>% substring(7, 11) %>% grepl(pattern = "Wo")

piping with .
"Hello World" %>% substring(7, 11) %>% grepl("Wo", .)

piping with . and curly braces
"Hello World" %>% substring(7, 11) %>% { c(paste('Hi', .)) }
#[1] "Hi World"

#using LHS multiple times in argument with curly braces and .
"Hello World" %>% substring(7, 11) %>% { c(paste(. , 'Hi', .)) }
#[1] "World Hi World"
```

. %>% RHS

```
, . . .
```

```
library(magrittr) # needed to include the pipe operators
library(lubridate)
read_year <- . %>% as.character %>% as.Date %>% year

Creating a dataset
df <- data.frame(now = "2015-11-11", before = "2012-01-01")
now before
1 2015-11-11 2012-01-01

Example 1: applying `read_year` to a single character-vector
df$now %>% read_year
[1] 2015

Example 2: applying `read_year` to all columns of `df`
df %>% lapply(read_year) %>% as.data.frame # implicit `lapply(df, read_year)
```

```

now before
1 2015 2012

Example 3: same as above using `mutate_all`
library(dplyr)
df %>% mutate_all(funs(read_year))
if an older version of dplyr use `mutate_each`
now before
1 2015 2012

```

functions functions .

```

read_year
Functional sequence with the following components:
#
1. as.character(..)
2. as.Date(..)
3. year(..)
#
Use 'functions' to extract the individual functions.

```

```

read_year[[2]]
function (.)
as.Date(.)


```

## % <> %

magrittr    rhs                 **infix-operator** %<>% .                 ( <- ). %<>% .

```

library(magrittr)
library(dplyr)

df <- mtcars

```

```

df <- df %>% select(1:3) %>% filter(mpg > 20, cyl == 6)

```

```

df %>% select(1:3) %>% filter(mpg > 20, cyl == 6) -> df

```

df .

```

df %<>% select(1:3) %>% filter(mpg > 20, cyl == 6)

```

## % \$ %

%%%    R . data ( , lm )    data.frame    ( dplyr ).

%%%            . cor.test    cor.test            cor.test .

```

library(magrittr)
library(dplyr)
mtcars %>%
 filter(wt > 2) %$%
 cor.test(hp, mpg)

#>
#> Pearson's product-moment correlation
#>
#> data: hp and mpg
#> t = -5.9546, df = 26, p-value = 2.768e-06
#> alternative hypothesis: true correlation is not equal to 0
#> 95 percent confidence interval:
#> -0.8825498 -0.5393217
#> sample estimates:
#> cor
#> -0.7595673

```

%>% **data.frame** filter() , %\$% cor.test() .

**base R** with()

## dplyr ggplot2

%>% **dplyr ggplot** . . . (EDA) . . ggplot . . .

```

library(dplyr)
library(ggplot)

diamonds %>%
 filter(depth > 60) %>%
 group_by(cut) %>%
 summarize(mean_price = mean(price)) %>%
 ggplot(aes(x = cut, y = mean_price)) +
 geom_bar(stat = "identity")

```

**% T> %**

**R** ( : , , ) . . .

%T>% (**tee** ) lhs . . , tee %>% . , rhs / lhs .

: , , . %T>% %>% all\_letters NULL NULL .

```

all_letters <- c(letters, LETTERS) %>%
 sort %T>%
 write.csv(file = "all_letters.csv")

read.csv("all_letters.csv") %>% head()
x
1 a
2 A
3 b
4 B

```

```
5 c
6 C
```

```
: save() . load() load() . ().
```

```
all_letters <- c(letters, LETTERS) %>%
 sort %T>%
 save(file = "all_letters.RData")

load("all_letters.RData", e <- new.env())

get("all_letters", envir = e)
Error in get("all_letters", envir = e) : object 'all_letters' not found

get(".", envir = e)
[1] "a" "A" "b" "B" "c" "C" "d" "D" "e" "E" "f" "F" "g" "G" "h" "H" "i" "I" "j" "J"
[21] "k" "K" "l" "L" "m" "M" "n" "N" "o" "O" "p" "P" "q" "Q" "r" "R" "s" "S" "t" "T"
[41] "u" "U" "v" "V" "w" "W" "x" "X" "y" "Y" "z" "Z"

Work-around
save2 <- function(. = ., name, file = stop("'file' must be specified")) {
 assign(name, .)
 call_save <- call("save", ... = name, file = file)
 eval(call_save)
}

all_letters <- c(letters, LETTERS) %>%
 sort %T>%
 save2("all_letters", "all_letters.RData")
```

(%> % ) : <https://riptutorial.com/ko/r/topic/652/----gt----->

# 122:

CRAN (Comprehensive R Archive Network)

## Examples

dplyr

```
help(package = "dplyr")
```

dplyr

```
data(package = "dplyr")
```

dplyr

```
library(dplyr)
ls("package:dplyr")
```

```
Checking package version which was installed at past or
installed currently but not loaded in the current session
```

```
packageVersion("seqinr")
[1] '3.3.3'
packageVersion("RWeka")
[1] '0.4.29'
```

```
search()
```

```
(.packages())
```

: <https://riptutorial.com/ko/r/topic/7408/>

# 123:

- `install.packages` (`pkgs`, `lib`, `repos`, `method`, `destdir`, `dependencies`, ...)

```
pkgs . repos = NULL .
lib .
repos , URL, NULL
destdir
/install / import / suggest (). repos = NULL .
... 'download.file' OS X Windows .
```

## Examples

`R` , . () `R` . `R` CRAN .

## CRAN

CRAN .

```
install.packages("dplyr")
```

"dplyr" .

`c()` .

```
install.packages(c("dplyr", "tidyverse", "ggplot2"))
```

`install.packages` `getOption("repos")` CRAN . CRAN repos .

```
install.packages("dplyr", repos = "https://cloud.r-project.org/")
```

repos . ?`install.packages` ?`install.packages` .

(`:package` `data.table`) . ( ) `dependencies` TRUE TRUE .

```
install.packages("data.table", dependencies = TRUE)
```

## Bioconductor

Bioconductor Bioinformatics . biocLite .

```
Try http:// if https:// URLs are not supported
source("https://bioconductor.org/biocLite.R")
biocLite()
```

. Bioconductor RImmPort RImmPort .

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

```
install.packages(path_to_source, repos = NULL, type="source")

install.packages("~/Downloads/dplyr-master.zip", repos=NULL, type="source")
```

path\_to\_source .

zip tar.gz .

```
install.packages(file.choose(), repos=NULL)
```

---

GUI RStudio .

1 : .

2 : .

3 : (.zip; .tar.gz)

4 : Browse () ( : crayon\_1.3.1.zip) ( )

---

R devtools install\_local() .

```
library(devtools)
install_local("~/Downloads/dplyr-master.zip")
```

GitHub

GitHub devtools .

```
library(devtools)
install_github("authorName/repositoryName")

github ggplot2 ,

devtools::install_github("tidyverse/ggplot2")

master ggplot2 . ref . , googleway dev_general .

devtools::install_github("SymbolixAU/googleway", ref = "dev_general")

ghit ghit . github .

install.packages("ghit")
ghit::install_github("google/CausalImpact")
```

Github <http://www.github.com/settings/tokens/> ( ? install\_github ). :

1. install.packages(c("curl", "httr"))
2. config = httr::config(ssl\_verifypeer = FALSE)
3. install.packages("RCurl")
options(RCurlOptions = c(getOption("RCurlOptions"), ssl.verifypeer = FALSE,
ssl.verifyhost = FALSE) )
4. getOption("RCurlOptions")

.

```
ssl.verifypeer ssl.verifyhost
FALSE FALSE
```

5. library(httr)
set\_config(config(ssl\_verifypeer = 0L))

. " CA "

6. .

```
install_github("username/package_name", auth_token="abc")
```

GITHUB\_PAT .

```
Sys.setenv(GITHUB_PAT = "access_token")
devtools::install_github("organisation/package_name")
```

```
Github PAT , .Rprofile .
```

## CLI -

```
pacman R .
```

```
pacman p_load () . pacman pacman .
```

```
p_load(data.table, dplyr, ggplot2)
```

```
library, require install.packages pacman .
```

```
library(pacman)
p_load(data.table, dplyr, ggplot2)
```

```
pacman::p_load(data.table, dplyr, ggplot2)
```

```
pacman
```

```
pacman () pacman :
```

```
if(!require(pacman)) install.packages("pacman")
pacman::p_load(data.table, dplyr, ggplot2)
```

```
R . ()
```

```
R CMD build my_package
```

```
R . R .
```

```
unloadNamespace("my_package")
library(my_package)
```

```
devtools . R
```

```
devtools::install()
```

: <https://riptutorial.com/ko/r/topic/1719/>

# 124:

- grep ( "", "", optional\_args)
- grepl ( "query", "subject", optional\_args)
- gsub ( "(group1) (group2)", "\\ group #", "subject")

```
(: \1) pattern sub gsub replacement \\1 .
(grep, sub, regexp) PCRE (Perl Compatible Regular Expression) . . perl=TRUE . R
```

- 
- stringr

## Examples

```
example data
test_sentences <- c("The quick brown fox quickly", "jumps over the lazy dog")
```

```
sub("brown", "red", test_sentences)
#[1] "The quick red fox quickly" "jumps over the lazy dog"
```

```
, "fast" "fastly" . :
```

```
sub("quick", "fast", test_sentences)
#[1] "The fast red fox quickly" "jumps over the lazy dog"
```

```
sub gsub .
```

```
gsub("quick", "fast", test_sentences)
#[1] "The fast red fox fastly" "jumps over the lazy dog"
```

```
example data
test_sentences <- c("The quick brown fox", "jumps over the lazy dog")
```

?

```
grepl() . TRUE / FALSE ("Boolean") .
```

```
""
```

```
grepl("fox", test_sentences)
#[1] TRUE FALSE
```

```
grep . . "fox" .
```

```
grep("fox", test_sentences)
#[1] 1
```

```
,
```

```
each of the following lines does the job:
test_sentences[grep("fox", test_sentences)]
test_sentences[grepl("fox", test_sentences)]
grep("fox", test_sentences, value = TRUE)
[1] "The quick brown fox"
```

```
"fox" fixed = TRUE (grep grepl).
```

```
grep("fox", test_sentences, fixed = TRUE)
#[1] 1
```

```
grep with invert = TRUE . -grep(...) !grepl(...) -grep(...)
```

```
grepl(pattern, x) grep(pattern, x) x pattern . pattern[1] x[1], pattern[2] x[2], .
```

```
grepl TRUE FALSE . . summary :
```

```
example data
test_sentences <- c("The quick brown fox", "jumps over the lazy dog")

find matches
matches <- grepl("fox", test_sentences)

overview
summary(matches)
```

```
PCRE g .
```

```
R : :
```

- `sub(pattern,replacement,text)`    `sub(pattern,replacement,text)`    **pattern** .
- `gsub(pattern,replacement,text)`    **sub** .
- `regexp(pattern,text)` .
- `gregexpr(pattern,text)` .

```
set.seed(123)
teststring <- paste0(sample(letters,20),collapse="")

teststring
#[1] "htjuwakqzxpgrsbncvyo"
```

```
sub("[aeiouy]," ** HERE WAS A VOWEL** ",teststring)
#[1] "htj ** HERE WAS A VOWEL** wakqzxpgrsbncvyo"

gsub("[aeiouy]," ** HERE WAS A VOWEL** ",teststring)
#[1] "htj ** HERE WAS A VOWEL** w ** HERE WAS A VOWEL** kqzxpgrsbncv ** HERE WAS A VOWEL** **
HERE WAS A VOWEL** "
```

```
regexp("[^aeiou][aeiou]+",teststring)
#[1] 3
#attr(),"match.length")
#[1] 2
#attr(),"useBytes")
#[1] TRUE
```

2 3 . : ju

```
gregexpr("[^aeiou][aeiou]+",teststring)
#[[1]]
#[1] 3 5 19
#attr(),"match.length")
#[1] 2 2 2
#attr(),"useBytes")
#[1] TRUE
```

. regmatches **regexp**

```
matches <- gregexpr("[^aeiou][aeiou]+",teststring)
regmatches(teststring,matches)
#[[1]]
```

```
[1] "ju" "wa" "yo"
```

```
, (.) :
```

```
teststring2 <- "this is another string to match against"
regmatches(teststring2, matches)
#[[1]]
#[1] "is" " i" "ri"
```

```
: Perl
```

```
perl=TRUE .
```

```
, grep("fox", test_sentences) . . .
```

```
perl = TRUE . fixed = TRUE . .
```

```
example data
test_sentences <- c("The quick brown fox", "jumps over the lazy dog")

grep("fox", test_sentences, perl = TRUE)
#[1] TRUE FALSE
```

```
. grep . .
```

```
searchCorpus <- function(corpus, pattern) {
 return(tm_index(corpus, FUN = function(x) {
 grep(pattern, x, ignore.case = TRUE, perl = TRUE)
 }))
}
```

```
: https://riptutorial.com/ko/r/topic/1123/--
```

# 125: (CSV, TSV )

- (, = TRUE, sep = ",", quote = "" ", dec =". ", = , comment.char =" ", ...)
- read.csv2 (, = TRUE, sep = ";", quote = "" ", dec =", ", = TRUE, comment.char =" ", ...)
- read\_csv (file, col\_names = TRUE, col\_types = NULL, locale = default\_locale (), na = c ( "", "NA"), comment = "", trim\_ws = TRUE, skip = 0, n\_max = -1 , progress = interactive ())
- stringAsFactors = FALSE, verbose =getOption ( "datatable", "auto", nrows = -1L, header = "auto", na.strings = "NA" ( : "integer") dec = if (sep! = "0"), = 1L, = 0L, select = NULL, drop = NULL, colClasses = NULL, integer64 =getOption ( "datatable.integer64" ( "datatable.showProgress"), # default : TRUE . ". ". ". ", col.names, check.names = FALSE, = " ", strip.white = TRUE, showProgress =getOption table =getOption ( "datatable.fread.datatable") #default : TRUE)

CSV	
	logical : .csv ?
	:
	:
	:
	logical : TRUE .
comment.char	character : csv . . .
...	read.table

.RData Feather .  
/ Input Output .

## Examples

.CSV

R

(CSV) read.csv sep = "," read.table .

```

get the file path of a CSV included in R's utils package
csv_path <- system.file("misc", "exDIF.csv", package = "utils")

path will vary based on installation location
csv_path
[1] "/Library/Frameworks/R.framework/Resources/library/utils/misc/exDIF.csv"

df <- read.csv(csv_path)

df
Var1 Var2
1 2.70 A
2 3.14 B
3 10.00 A
4 -7.00 A

```

file.choose .

```
df <- read.csv(file.choose())
```

- `read.table , read.csv header = TRUE .`
  - `as.is = TRUE stringsAsFactors = FALSE factor stringsAsFactors = FALSE .`
  - `read.csv2 sep = ";" dec = ","`
- 

**readr** `read_csv , stringsAsFactors = FALSE` `read.csv stringsAsFactors = FALSE .`

```

library(readr)

df <- read_csv(csv_path)

df
A tibble: 4 x 2
Var1 Var2
<dbl> <chr>
1 2.70 A
2 3.14 B
3 10.00 A
4 -7.00 A

```

## data.table

`data.table fread . read.table fread .`

```

get the file path of a CSV included in R's utils package
csv_path <- system.file("misc", "exDIF.csv", package = "utils")

path will vary based on R installation location
csv_path
[1] "/Library/Frameworks/R.framework/Resources/library/utils/misc/exDIF.csv"

dt <- fread(csv_path)

```

```
dt
Var1 Var2
1: 2.70 A
2: 3.14 B
3: 10.00 A
4: -7.00 A
```

```
input .
```

- ( :"filename.csv"),
- ( :"grep 'word' filename")
- ( :"input1, input2 \n A, B \n C, D").

```
fread data.table data.frame data.table [] . data.table data.table FALSE FALSE .
```

```
df <- fread(csv_path, data.table = FALSE)

class(df)
[1] "data.frame"

df
Var1 Var2
1 2.70 A
2 3.14 B
3 10.00 A
4 -7.00 A
```

- fread read.table read.table . na.comment . #
- fread " quote .
- fread .

## .tsv ( R )

```
file.path . Windows, Mac Linux paste .
```

```
FilePath <- file.path(AVariableWithFullPath, "SomeSubfolder", "SomeFileName.txt.gz")

Data <- as.matrix(read.table(FilePath, header=FALSE, sep ="\t"))
```

```
. FilePath 0 79 % 8970 8970 .
```

```
system.time(expr=Data<-as.matrix(read.table(file=FilePath,header=FALSE,sep=" ")))
```

```
system.time 267 .
```

user	system	elapsed
265.563	1.949	267.563

```
FilePath <- "SomeFile"
connection<- gzfile(FilePath,open="r")
TableList <- list()
Counter <- 1
system.time(expr= while (length(Vector<-as.matrix(scan(file=connection, sep=" ", nlines=1,
quiet=TRUE))) > 0) {
 TableList[[Counter]]<-Vector
 Counter<-Counter+1
})
 user system elapsed
165.976 0.060 165.941
close(connection)
system.time(expr=(Data <- do.call(rbind,TableList)))
 user system elapsed
 0.477 0.088 0.565
```

read.matrix futile.matrix . 1 .

## .CSV



write.csv() CSV .

```
write.csv(mtcars, "mtcars.csv")
```

row.names = FALSE na = "" .

---

readr::write\_csv write.csv .

```
library(readr)

write_csv(mtcars, "mtcars.csv")
```

## CSV

```
files = list.files(pattern="*.csv")
data_list = lapply(files, read.table, header = TRUE)
```

. data.frame data.frame .

```
df <- do.call(rbind, data_list)
```

; , . . ; , ( ) . .

```

Column1 Column2 Column3 Column4Column5
1647 pi 'important' 3.141596.28318
1731 euler 'quite important' 2.718285.43656
1979 answer 'The Answer.' 42 42

```

constants.txt .

---

## R

```

df <- read.fwf('constants.txt', widths = c(8,10,18,7,8), header = FALSE, skip = 1)

df
#> V1 V2 V3 V4 V5
#> 1 1647 pi 'important' 3.14159 6.28318
#> 2 1731 euler 'quite important' 2.71828 5.43656
#> 3 1979 answer 'The Answer.' 42 42.0000

```

:

- (Column4Column5).
  - widths .
  - read.fwf() .
- 

## reader

```

library(readr)

df <- read_fwf('constants.txt',
 fwf_cols(Year = 8, Name = 10, Importance = 18, Value = 7, Doubled = 8),
 skip = 1)
df
#> # A tibble: 3 x 5
#> Year Name Importance Value Doubled
#> <int> <chr> <chr> <dbl> <dbl>
#> 1 1647 pi 'important' 3.14159 6.28318
#> 2 1731 euler 'quite important' 2.71828 5.43656
#> 3 1979 answer 'The Answer.' 42.00000 42.00000

```

:

- **readr** `fwf_*` ( `fwf_empty` ) .
- R .
- .

(CSV, TSV) : <https://riptutorial.com/ko/r/topic/481/-----csv--tsv-->

## 126: : +

parse .

eval .

### Examples

```
the string
str <- "1+1"

A string is not an expression.
is.expression(str)
[1] FALSE

eval(str)
[1] "1+1"

parse convert string into expressions
parsed.str <- parse(text="1+1")

is.expression(parsed.str)
[1] TRUE

eval(parsed.str)
[1] 2
```

: + : <https://riptutorial.com/ko/r/topic/5746/----plus->

# 127:

( ) ( ) . . . .  
. . . , , . , : (Fourier synthesis)  
( ) . . . , . , .  
. .  
. (:) (Joseph Fourier)  
(Liemann) ,  
, Dirac . . . , 3 3 ( 4 - ) .  
. Fourier .  $\mathbb{R}$   $\mathbb{R}^n$  ( ) (DTFT,  $= \mathbb{Z}$ ), DFT, group =  $\mathbb{Z}$  mod N ( = S1, ).  
(FFT) DFT .

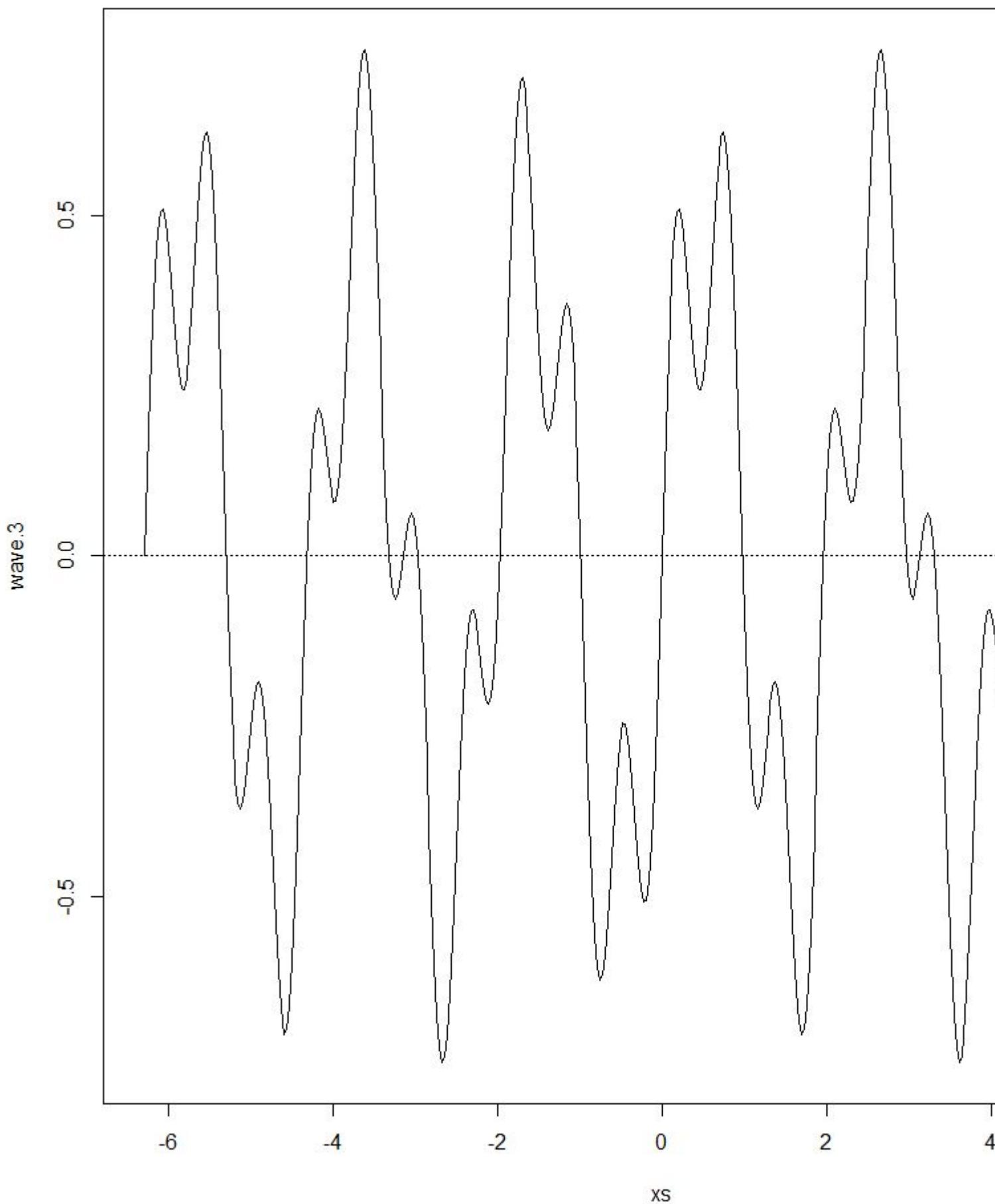
## Examples

(Joseph Fourier)

```
Sine waves
xs <- seq(-2*pi, 2*pi, pi/100)
wave.1 <- sin(3*xs)
wave.2 <- sin(10*xs)
par(mfrow = c(1, 2))
plot(xs, wave.1, type="l", ylim=c(-1, 1)); abline(h=0, lty=3)
plot(xs, wave.2, type="l", ylim=c(-1, 1)); abline(h=0, lty=3)

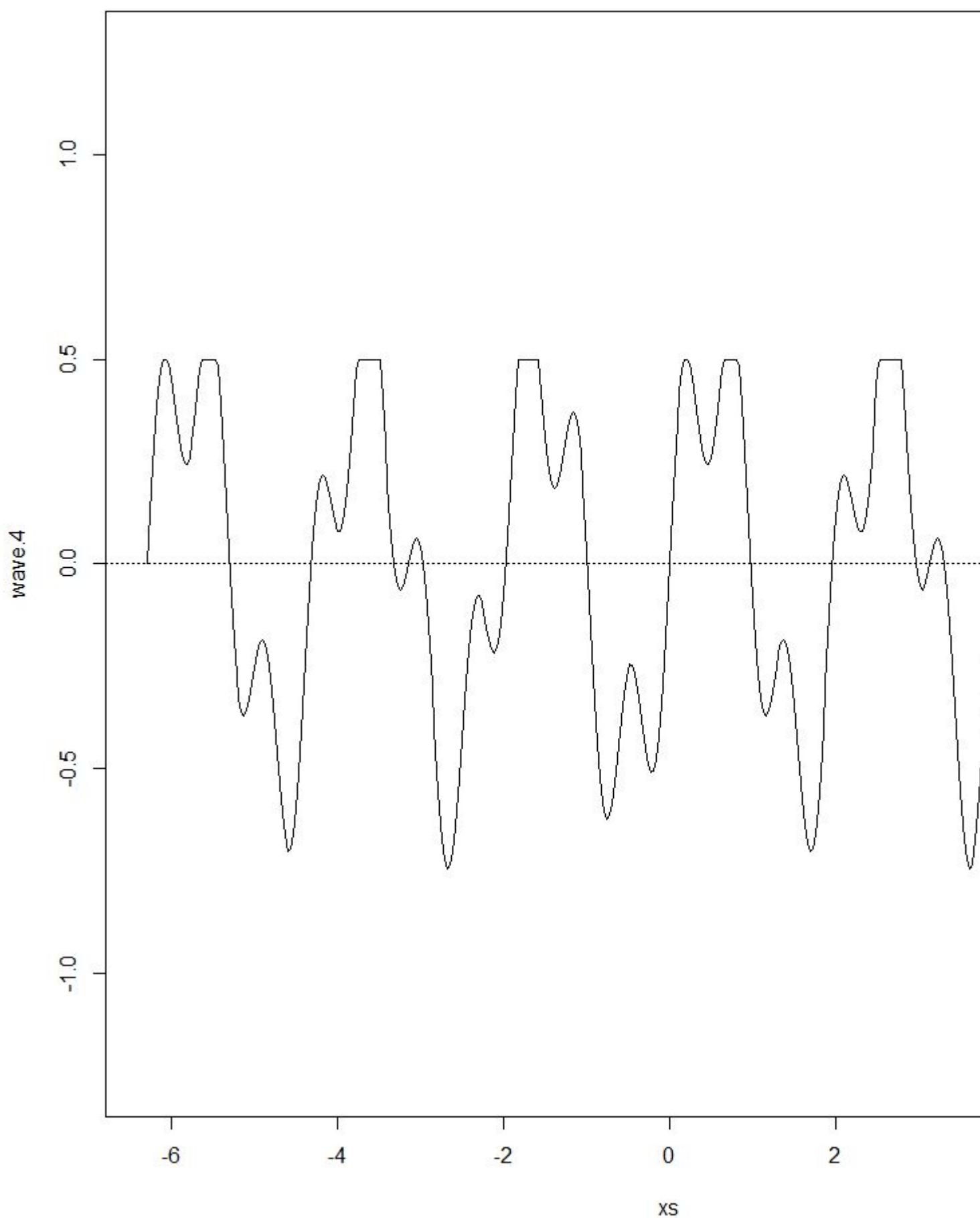
Complex Wave
wave.3 <- 0.5 * wave.1 + 0.25 * wave.2
plot(xs, wave.3, type="l"); title("Eg complex wave"); abline(h=0, lty=3)
```

### Eg complex wave



```
wave.4 <- wave.3
wave.4[wave.3>0.5] <- 0.5
plot(xs,wave.4,type="l",ylim=c(-1.25,1.25))
title("overflowed, non-linear complex wave")
abline(h=0,lty=3)
```

## overflowed, non-linear complex wave



$\dots, (\dots, \dots) \quad f \lambda(\dots, \lambda = v/f_0, v) \dots$

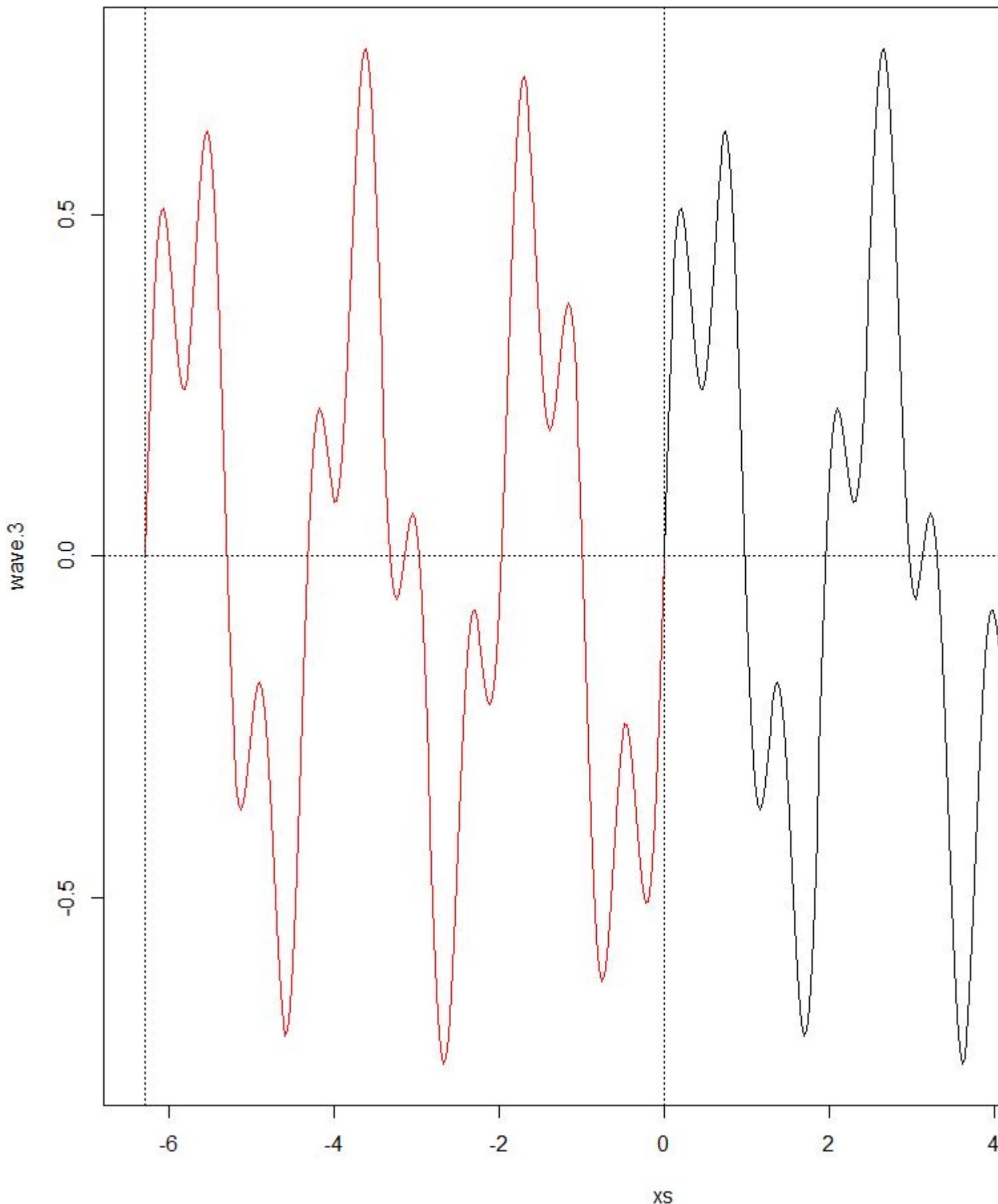
$\vdots$

- $T$ ,
- (sr)  $(\dots)$ .  
si,  
 $N$ .  
 $si = TN$
- $f_0 1T.$   
 $\dots, 12\pi.$   
 $\dots, f_0 1, 2 2^* f_0, 3 3^* f_0.$

```
repeat.xs <- seq(-2*pi,0,pi/100)
wave.3.repeat <- 0.5*sin(3*repeat.xs) + 0.25*sin(10*repeat.xs)
plot(xs, wave.3, type="l")

title("Repeating pattern")
points(repeat.xs, wave.3.repeat, type="l", col="red");
abline(h=0, v=c(-2*pi, 0), lty=3)
```

## Repeating pattern



R .

```
plot.fourier <- function(fourier.series, f.0, ts) {
 w <- 2*pi*f.0 trajectory <- sapply(ts, function(t)
fourier.series(t,w))
 plot(ts, trajectory, type="l", xlab="time", ylab="f(t)");
 abline(h=0,lty=3)}
```

: <https://riptutorial.com/ko/r/topic/4139/>--

# 128:

```
R , . subsetting .
:
NA [NA] (NA $)
NA .
NA "" "(typeof(NA)), "" . x[NA] x[rep_len(as.logical(NA), length(x))] x[as.logical(NA)]
(NA) x x .:

x <- 1:3
x[NA]
[1] NA NA NA

""/ "" NA NA (NA).

x[as.integer(NA)]
[1] NA

x[c(NA, 1, NA, NA)]
[1] NA 1 NA NA

:
[> length(x) NA , NULL . [[when [(, length(dim(x)) > 2 bounds) .

(1:3)[10]
[1] NA
(1:3)[[10]]
Error in (1:3)[[10]] : subscript out of bounds
as.matrix(1:3)[10]
[1] NA
as.matrix(1:3)[, 10]
Error in as.matrix(1:3)[, 10] : subscript out of bounds
list(1, 2, 3)[10]
[[1]]
NULL
list(1, 2, 3)[[10]]
Error in list(1, 2, 3)[[10]] : subscript out of bounds

"names" "" .

c(a = 1, b = 2) ["c"]
<NA>
NA
list(a = 1, b = 2) ["c"]
<NA>
NULL
```

```
?Extract .
```

## Examples

```
() [:
```

```
create an example vector
v1 <- c("a", "b", "c", "d")

select the third element
v1[3]
[1] "c"
```

```
[. . .]
```

```
v1 <- c("a", "b", "c", "d")

v1[c(1, 3)]
[1] "a" "c"
```

```
(-) . , v1 v1[-1] . . . , v1[-c(1, 3)] .
```

```
> v1[-1]
[1] "b" "c" "d"
> v1[-c(1, 3)]
[1] "b" "d"
```

```
> v1=="c"
[1] FALSE FALSE TRUE FALSE
> which(v1=="c")
[1] 3
```

```
(names) : .
```

```
v <- 1:3
names(v) <- c("one", "two", "three")

v
one two three
1 2 3

v["two"]
two
2
```

```
[[1 . . .]]
```

```
v[[c(1, 2)]]
```

```

Error in v[[c(1, 2)]] :
attempt to select more than one element in vectorIndex

v[["two"]]
[1] 2

. , . . y x x[y] , if length(y) < length(x) y length(x) .

v[c(TRUE, FALSE, TRUE)]
one three
1 3

v[c(FALSE, TRUE)] # recycled to 'c(FALSE, TRUE, FALSE)'
two
2

v[TRUE] # recycled to 'c(TRUE, TRUE, TRUE)'
one two three
1 2 3

v(FALSE) # handy to discard elements but save the vector's type and basic structure
named integer(0)

```

[ :

```

l1 <- list(c(1, 2, 3), 'two' = c("a", "b", "c"), list(10, 20))
l1
[[1]]
[1] 1 2 3
##
$two
[1] "a" "b" "c"
##
[[3]]
[[3]][[1]]
[1] 10
##
[[3]][[2]]
[1] 20

l1[1]
[[1]]
[1] 1 2 3

l1['two']
$two
[1] "a" "b" "c"

l1[[2]]
[1] "a" "b" "c"

l1[['two']]
[1] "a" "b" "c"

```

[ 11[2] . [[ . .
() . [ . . length > 1 [ [[ ( ) ""

```
l1[c(3, 1)]
[1]
[[1]][[1]]
[1] 10

[[1]][[2]]
[1] 20

[[2]]
[1] 1 2 3
```

```
l1[[c(3, 1)]]
[1] 10
```

```
l1[[3]][[1]]
[1] 10
```

```
$ [[, . $:
```

```
l1$two
[1] "a" "b" "c"
```

```
$
```

```
l1$t
[1] "a" "b" "c"
```

```
[[
```

```
l1[["t"]]
NULL
l1[["t", exact = FALSE]]
[1] "a" "b" "c"
```

```
options(warnPartialMatchDollar = TRUE) $ "" .
```

```
l1$t
[1] "a" "b" "c"
Warning message:
In l1$t : partial match of 't' to 'two'
```

```
[. . . [i, j] [i, j] i j . 1 .
```

```
a sample matrix
mat <- matrix(1:6, nrow = 2, dimnames = list(c("row1", "row2"), c("col1", "col2", "col3")))
mat
```

```
col1 col2 col3
row1 1 3 5
row2 2 4 6
```

```
mat[i,j] i , j mat . i 2 j 1 . i j .
```

```
mat[, 3]
row1 row2
5 6

mat[1,]
col1 col2 col3
1 3 5
```

```
() .
```

```
mat[, 'col1']
row1 row2
1 2
```

```
. 2 1 . drop = FALSE [::
```

```
This selects the first row as a vector
class(mat[1,])
[1] "integer"

Whereas this selects the first row as a 1x3 matrix:
class(mat[1, , drop = F])
[1] "matrix"
```

---

```
mat[1:2, 2:3] ## A 2x2 matrix
col2 col3
row1 3 5
row2 4 6
```

Nx2 N ( ). (1st row, 1st column), (1st row, 3rd column), (2nd row, 3rd column), (2nd row, 1st column) :

```
mat
col1 col2 col3
row1 1 3 5
row2 2 4 6

ind = rbind(c(1, 1), c(1, 3), c(2, 3), c(2, 1))
ind
[,1] [,2]
[1,] 1 1
[2,] 1 3
[3,] 2 3
[4,] 2 1
```

```
mat[ind]
[1] 1 5 6 2
```

1 ind mat ind mat .

```
> df3 <- data.frame(x = 1:3, y = c("a", "b", "c"), stringsAsFactors = FALSE)

> df3
x y
1 1 a
2 2 b
3 3 c

> df3[1] # Subset a variable by number
x
1 1
2 2
3 3

> df3["x"] # Subset a variable by name
x
1 1
2 2
3 3

> is.data.frame(df3[1])
TRUE

> is.list(df3[1])
TRUE
```

[[[]]] \$ .

```
> df3[[2]] # Subset a variable by number using [[]]
[1] "a" "b" "c"

> df3[["y"]] # Subset a variable by name using [[]]
[1] "a" "b" "c"

> df3$x # Subset a variable by name using $
[1] 1 2 3

> typeof(df3$x)
"integer"

> is.vector(df3$x)
TRUE
```

2 i j .

```
> df3[1, 2] # Subset row and column by number
[1] "a"

> df3[1, "y"] # Subset row by number and column by name
[1] "a"
```

```

> df3[2,] # Subset entire row by number
x y
2 2 b

> df3[, 1] # Subset all first variables
[1] 1 2 3

> df3[, 1, drop = FALSE]
x
1 1
2 2
3 3

```

```
: j() i data.frame . drop FALSE .
```

```

> is.vector(df3[, 2])
TRUE

> is.data.frame(df3[2,])
TRUE

> is.data.frame(df3[, 2, drop = FALSE])
TRUE

```

```
[[[. , R (isTRUE(is.object(x)) "class") . / [[.
, [.data.frame (is.object(iris)) [.data.frame [[.data.frame "matrix" "" . "data.frame"
[.data.frame [.data.frame [.
```

```
iris[invalidArgument,]
Error in ` [.data.frame` (iris, invalidArgument,) :
object 'invalidArgument' not found
```

```
[:
```

```
x = structure(1:5, class = "myClass")
x[c(3, 2, 4)]
[1] 3 2 4
'[.myClass' = function(x, i) cat(sprintf("We'd expect '%s[%s]' to be returned but this a
custom `[' method and should have a `? [.myClass` help page for its behaviour\n",
deparse(substitute(x)), deparse(substitute(i))])

x[c(3, 2, 4)]
We'd expect 'x[c(3, 2, 4)]' to be returned but this a custom `[' method and should have a
`? [.myClass` help page for its behaviour
NULL
```

```
.subset ([[] .subset2 [. "" ("class" () "" unclass(x)) .
.subset(x, c(3, 2, 4))
[1] 3 2 4
```

```
> x <- 11:20
> x
[1] 11 12 13 14 15 16 17 18 19 20
```

R 1-indexed,  $x[1]$  11 . . .

```
> x[c(2,4,6)]
[1] 12 14 16
```

R . .

```
> x[c(-1,-3)]
[1] 12 14 15 16 17 18 19 20
```

. TRUE .

```
> x[c(rep(TRUE,5),rep(FALSE,5))]
[1] 11 12 13 14 15 16
```

```
> x[c(TRUE,FALSE)]
[1] 11 13 15 17 19
> x[c(TRUE,FALSE,FALSE)]
[1] 11 14 17 20
```

A B . + , - , / , \* , ^ . .

	B	
+	$A + B$	$A B$
-	$A - B$	$A B .$
/	$A / B$	$A B .$
*	$A * B$	$A B .$
$\wedge$	$A^{\wedge}(-1)$	, A .

(Linear Algebra) " (true)" %\*% . , A B A %\*% B . A ncol() nrow() B nrow() .

nrow ()	(A)	A .
ncol ()	ncol (A)	A .
rownames ()	rownames (A)	A .

colnames ()	colnames (A)	A .
rowMeans ()	rowMeans (A)	A .
colMeans ()	colMeans (A)	A .
upper.tri ()	upper.tri (A)	.
		A
lower.tri ()	lower.tri (A)	.
		A
det ()	det (A)	A .
()	(A)	- A .
diag ()	(A)	0
		A
()	)	A .
()	(A)	A .
is.matrix ()	is.matrix (A)	A TRUE FALSE .
as.matrix ()	as.matrix (x)	x .

: <https://riptutorial.com/ko/r/topic/1686/>

# 129:

## Examples

```
: , .
R new.env environment () . . hash TRUE .
```

```
H <- new.env(hash = TRUE)
H <- new.env()
```

```
, 29 size . R environment size . size .
```

```
object.size(new.env())
56 bytes

object.size(new.env(size = 10e4))
56 bytes
```

---

```
[[<- $<- environment, "" ([<-):
```

```
H <- new.env()

H[["key"]] <- rnorm(1)

key2 <- "xyz"
H[[key2]] <- data.frame(x = 1:3, y = letters[1:3])

H$another_key <- matrix(rbinom(9, 1, 0.5) > 0, nrow = 3)

H["error"] <- 42
#Error in H["error"] <- 42 :
object of type 'environment' is not subsettable
```

```
R , (object[[key]] <- value) (object$key <- value) (: key2).
```

```
, environment . - .
```

```
H[["key3"]] <- "original value"

H[["key3"]] <- "new value"

H[["key3"]]
#[1] "new value"
```

---

```
[[$, [:
```

```
H[["key"]]
#[1] 1.630631
```

```

H[[key2]] ## assuming key2 <- "xyz"
x y
1 1 a
2 2 b
3 3 c

H$another_key
[,1] [,2] [,3]
[1,] TRUE TRUE TRUE
[2,] FALSE FALSE FALSE
[3,] TRUE TRUE TRUE

H[1]
#Error in H[1] : object of type 'environment' is not subsettable

```

---

environment .

```

names(H)
#[1] "another_key" "xyz" "key" "key3"

ls(H)
#[1] "another_key" "key" "key3" "xyz"

str(H)
#<environment: 0x7828228>

ls.str(H)
another_key : logi [1:3, 1:3] TRUE FALSE TRUE TRUE FALSE TRUE ...
key : num 1.63
key3 : chr "new value"
xyz : 'data.frame': 3 obs. of 2 variables:
$ x: int 1 2 3
$ y: chr "a" "b" "c"

```

rm .

```

rm(list = c("key", "key3"), envir = H)

ls.str(H)
another_key : logi [1:3, 1:3] TRUE FALSE TRUE TRUE FALSE TRUE ...
xyz : 'data.frame': 3 obs. of 2 variables:
$ x: int 1 2 3
$ y: chr "a" "b" "c"

```

---

environment , environment .

```

H2 <- new.env()

H2[["a"]] <- LETTERS
H2[["b"]] <- as.list(x = 1:5, y = matrix(rnorm(10), 2))
H2[["c"]] <- head(mtcars, 3)
H2[["d"]] <- Sys.Date()
H2[["e"]] <- Sys.time()
H2[["f"]] <- (function() {

```

```

H3 <- new.env()
for (i in seq_along(names(H2))) {
 H3[[names(H2)[i]]] <- H2[[names(H2)[i]]]
}
H3
})()

ls.str(H2)
a : chr [1:26] "A" "B" "C" "D" "E" "F" "G" "H" "I" "J" "K" ...
b : List of 5
$: int 1
$: int 2
$: int 3
$: int 4
$: int 5
c : 'data.frame': 3 obs. of 11 variables:
$ mpg : num 21 21 22.8
$ cyl : num 6 6 4
$ disp: num 160 160 108
$ hp : num 110 110 93
$ drat: num 3.9 3.9 3.85
$ wt : num 2.62 2.88 2.32
$ qsec: num 16.5 17 18.6
$ vs : num 0 0 1
$ am : num 1 1 1
$ gear: num 4 4 4
$ carb: num 4 4 1
d : Date[1:1], format: "2016-08-03"
e : POSIXct[1:1], format: "2016-08-03 19:25:14"
f : <environment: 0x91a7cb8>

ls.str(H2$f)
a : chr [1:26] "A" "B" "C" "D" "E" "F" "G" "H" "I" "J" "K" ...
b : List of 5
$: int 1
$: int 2
$: int 3
$: int 4
$: int 5
c : 'data.frame': 3 obs. of 11 variables:
$ mpg : num 21 21 22.8
$ cyl : num 6 6 4
$ disp: num 160 160 108
$ hp : num 110 110 93
$ drat: num 3.9 3.9 3.85
$ wt : num 2.62 2.88 2.32
$ qsec: num 16.5 17 18.6
$ vs : num 0 0 1
$ am : num 1 1 1
$ gear: num 4 4 4
$ carb: num 4 4 1
d : Date[1:1], format: "2016-08-03"
e : POSIXct[1:1], format: "2016-08-03 19:25:14"

```

environment      R    / .

```

names(H2)
#[1] "a" "b" "c" "d" "e" "f"

```

```

H2[[c("a", "b")]]
#Error in H2[[c("a", "b")]] :
wrong arguments for subsetting an environment

Keys <- c("a", "b")
H2[[Keys]]
#Error in H2[[Keys]] : wrong arguments for subsetting an environment

```

vapply list2env .

```

E1 <- new.env()
invisible({
 vapply(letters, function(x) {
 E1[[x]] <- rnorm(1)
 logical(0)
 }, FUN.VALUE = logical(0))
})

all.equal(sort(names(E1)), letters)
#[1] TRUE

Keys <- letters
E2 <- list2env(
 setNames(
 as.list(rnorm(26)),
 nm = Keys),
 envir = NULL,
 hash = TRUE
)

all.equal(sort(names(E2)), letters)
#[1] TRUE

```

- for .

:

R . . . R . .

:

```

Generic unique string generator
unique_strings <- function(n){
 string_i <- 1
 string_len <- 1
 ans <- character(n)
 chars <- c(letters, LETTERS)
 new_strings <- function(len,pfx) {
 for(i in 1:length(chars)){
 if (len == 1){
 ans[string_i] <- paste(pfx,chars[i],sep='')
 string_i <- string_i + 1
 } else {
 new_strings(len-1,pfx=paste(pfx,chars[i],sep=''))
 }
 if (string_i > n) return ()
 }
 }
}
```

```

 }
 while(string_i <= n){
 new_strings(string_len,'')
 string_len <- string_len + 1
 }
 sample(ans)
 }

Generate timings using an environment
timingsEnv <- plyr::adply(2^(10:15),.mar=1,.fun=function(i){
 strings <- unique_strings(i)
 ht1 <- new.env(hash=TRUE)
 lapply(strings, function(s){ ht1[[s]] <- 0L})
 data.frame(
 size=c(i,i),
 seconds=c(
 system.time(for (j in 1:i) ht1[[strings[j]]]==0L)[3]),
 type = c('1_hashedEnv')
)
})
}

timingsHash <- plyr::adply(2^(10:15),.mar=1,.fun=function(i){
 strings <- unique_strings(i)
 ht <- hash::hash()
 lapply(strings, function(s) ht[[s]] <- 0L)
 data.frame(
 size=c(i,i),
 seconds=c(
 system.time(for (j in 1:i) ht[[strings[j]]]==0L)[3]),
 type = c('3_stringHash')
)
})
}

```

## : listenv

```

package:listenv , . . package:future package:future package:future .
package : hash .

```

```

timingsListEnv <- plyr::adply(2^(10:15),.mar=1,.fun=function(i){
 strings <- unique_strings(i)
 le <- listenv::listenv()
 lapply(strings, function(s) le[[s]] <- 0L)
 data.frame(
 size=c(i,i),
 seconds=c(
 system.time(for (k in 1:i) le[[k]]==0L)[3]),
 type = c('2_numericListEnv')
)
})

```

: <https://riptutorial.com/ko/r/topic/5179/>

# 130:

## Examples

```
., . matrix matrix .

matrix(data = 1:6, nrow = 2, ncol = 3)
[,1] [,2] [,3]
[1,] 1 3 5
[2,] 2 4 6
```

```
, 1 6 2 3 . data , nrow , ncol . . . byrow .
```

```
matrix(data = 1:6, nrow = 2, ncol = 3, byrow = TRUE)
[,1] [,2] [,3]
[1,] 1 2 3
[2,] 4 5 6
```

```
. . :

matrix(data = c(TRUE, TRUE, TRUE, FALSE, FALSE, FALSE), nrow = 3, ncol = 2)
[,1] [,2]
[1,] TRUE FALSE
[2,] TRUE FALSE
[3,] TRUE FALSE
matrix(data = c("a", "b", "c", "d", "e", "f"), nrow = 3, ncol = 2)
[,1] [,2]
[1,] "a" "d"
[2,] "b" "e"
[3,] "c" "f"
```

```
. . rownames colnames . . . NULL NULL . .
```

```
mat1 <- matrix(data = 1:6, nrow = 2, ncol = 3, byrow = TRUE)
rownames(mat1)
NULL
colnames(mat1)
NULL
rownames(mat1) <- c("Row 1", "Row 2")
colnames(mat1) <- c("Col 1", "Col 2", "Col 3")
mat1
Col 1 Col 2 Col 3
Row 1 1 2 3
Row 2 4 5 6
```

```
. .
class, is as 1 .
```

```
class(mat1)
[1] "matrix"
is.matrix(mat1)
```

```
[1] TRUE
as.vector(mat1)
[1] 1 4 2 5 3 6
```

: <https://riptutorial.com/ko/r/topic/9019/>

# 131: .2

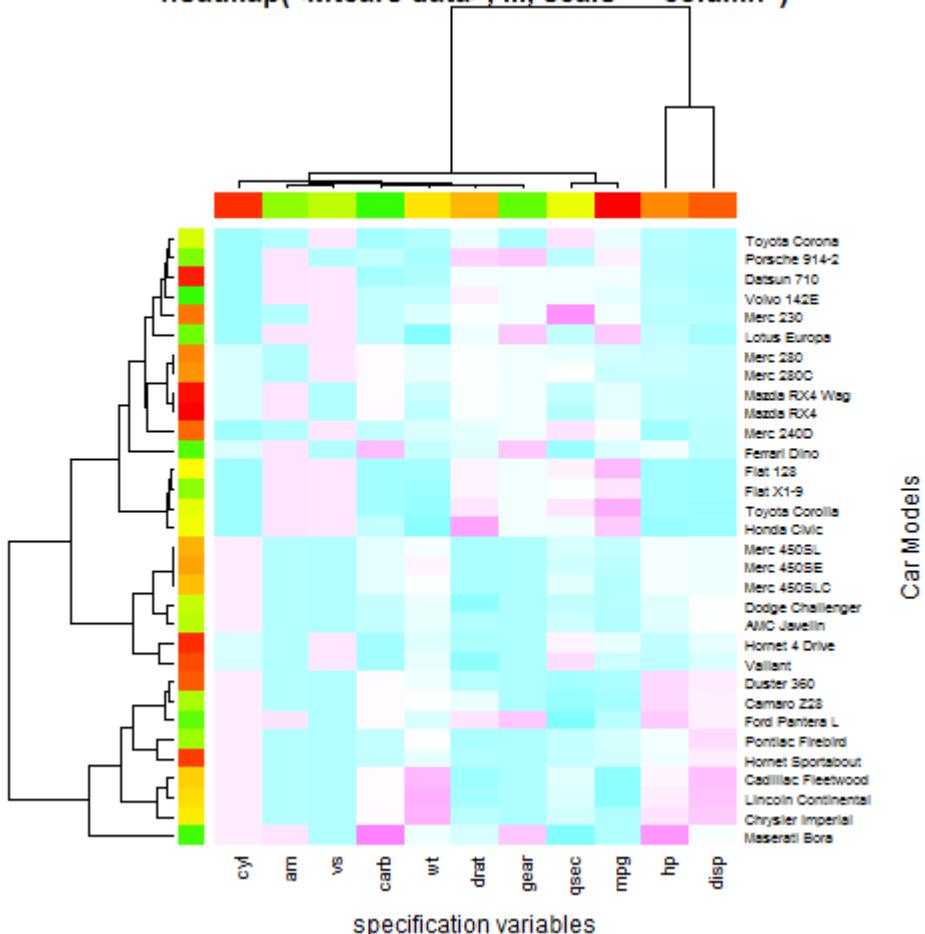
## Examples

### :: heatmap

1 ()

```
require(graphics); require(grDevices)
x <- as.matrix(mtcars)
rc <- rainbow(nrow(x), start = 0, end = .3)
cc <- rainbow(ncol(x), start = 0, end = .3)
hv <- heatmap(x, col = cm.colors(256), scale = "column",
 RowSideColors = rc, ColSideColors = cc, margins = c(5,10),
 xlab = "specification variables", ylab = "Car Models",
 main = "heatmap(<Mtcars data>, ..., scale = \"column\")")
```

heatmap(<Mtcars data>, ..., scale = "column")

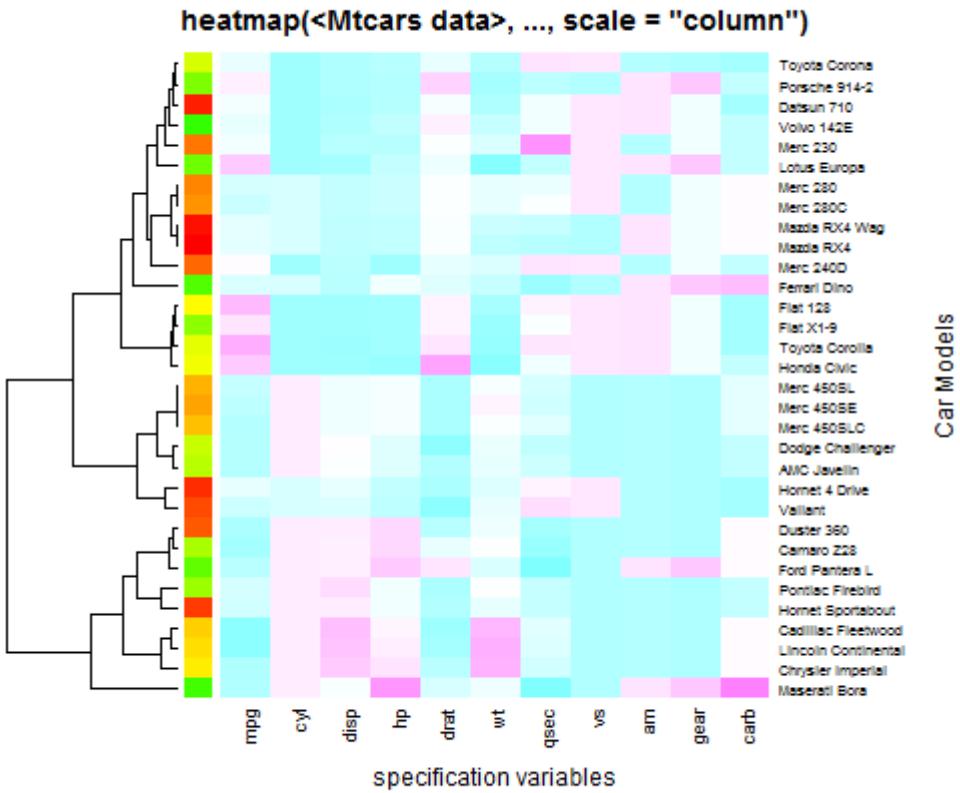


```
utils::str(hv) # the two re-ordering index vectors
List of 4
$ rowInd: int [1:32] 31 17 16 15 5 25 29 24 7 6 ...
$ colInd: int [1:11] 2 9 8 11 6 5 10 7 1 4 ...
$ Rowv : NULL
```

```
$ Colv : NULL
```

## 2 ( ( ) )

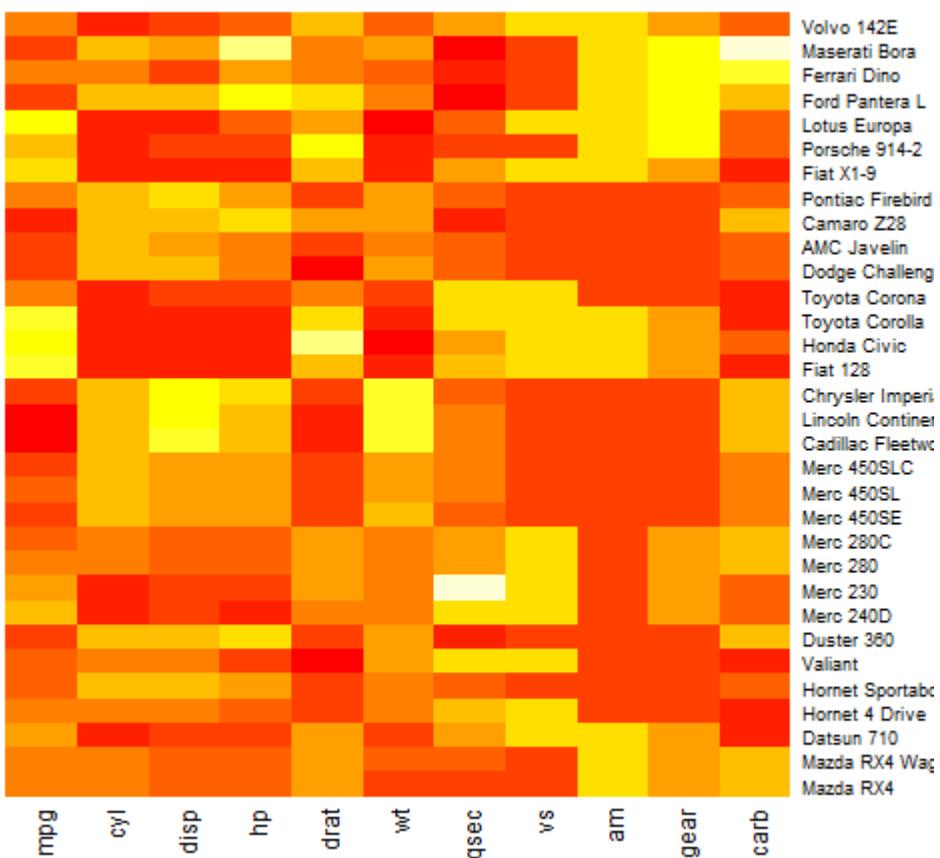
```
heatmap(x, Colv = NA, col = cm.colors(256), scale = "column",
 RowSideColors = rc, margins = c(5,10),
 xlab = "specification variables", ylab = "Car Models",
 main = "heatmap(<Mtcars data>, ..., scale = \"column\")")
```



## 3 ( " " )

```
heatmap(x, Rowv = NA, Colv = NA, scale = "column",
 main = "heatmap(*, NA, NA) ~ image(t(x))")
```

**heatmap(\*, NA, NA) ≈ image(t(x))**

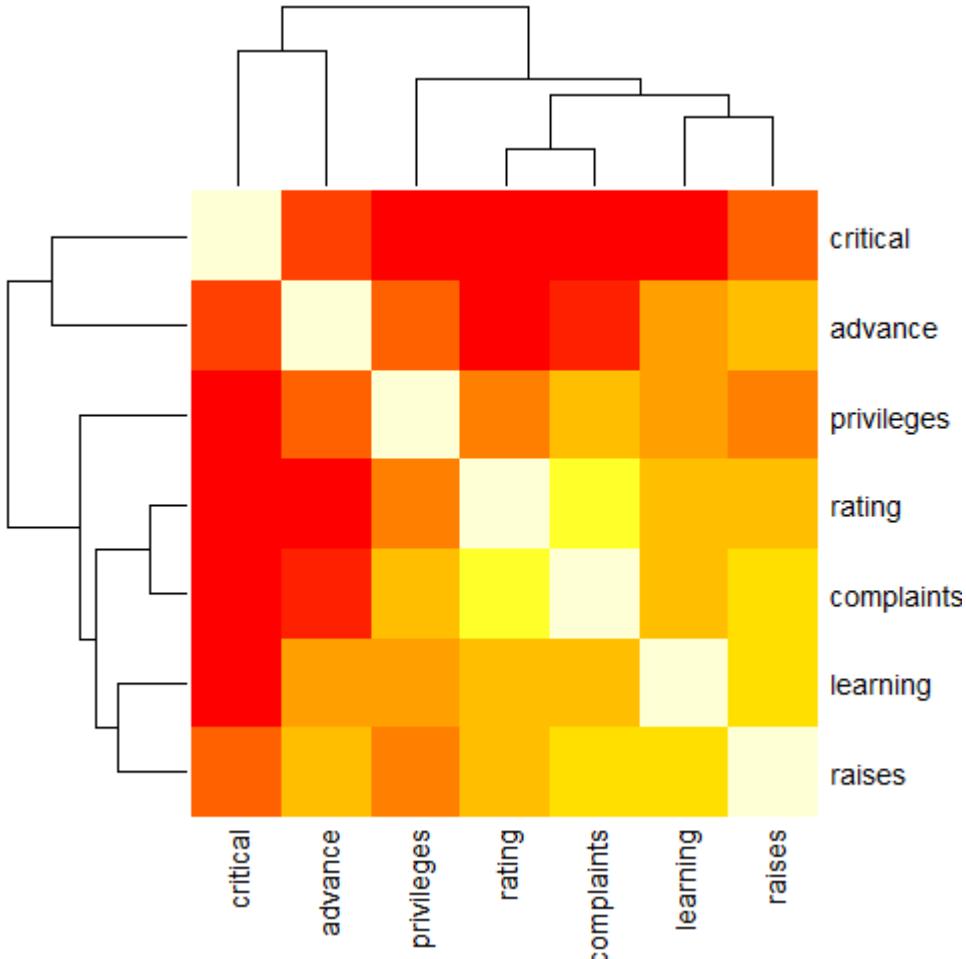


## 4 (reorder ()) )

```

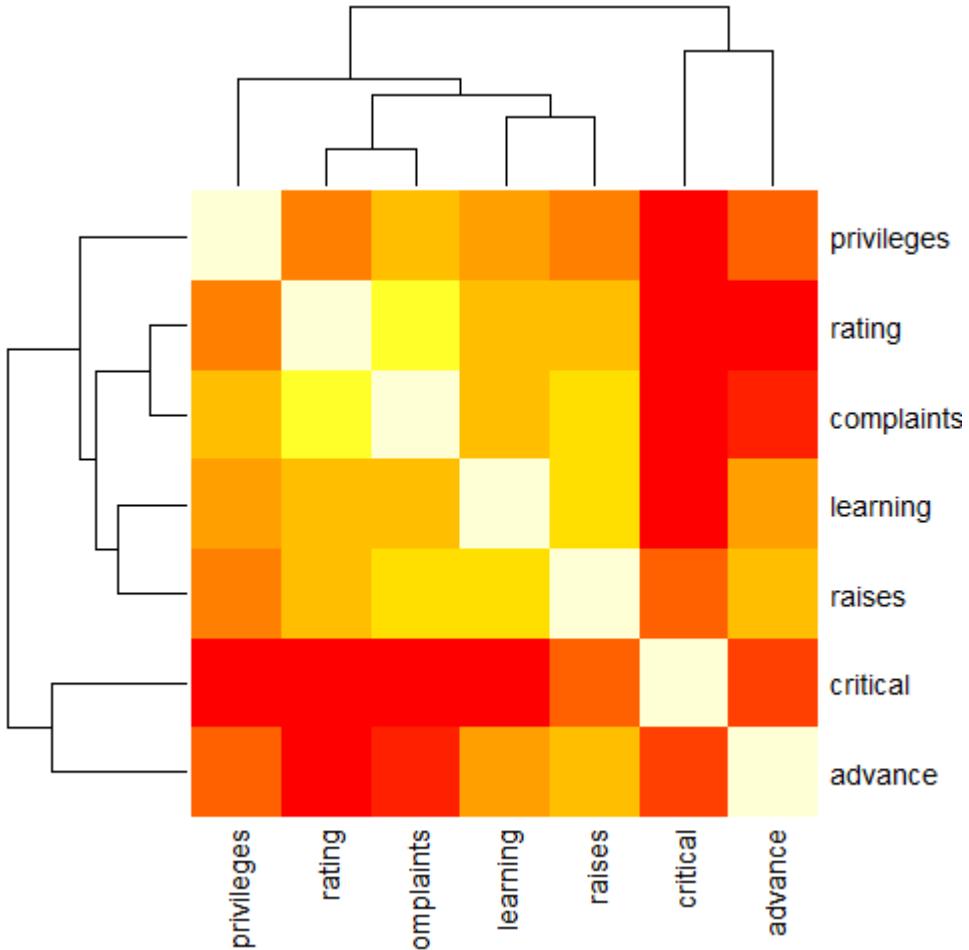
round(Ca <- cor(attitude), 2)
rating complaints privileges learning raises critical advance
rating 1.00 0.83 0.43 0.62 0.59 0.16 0.16
complaints 0.83 1.00 0.56 0.60 0.67 0.19 0.22
privileges 0.43 0.56 1.00 0.49 0.45 0.15 0.34
learning 0.62 0.60 0.49 1.00 0.64 0.12 0.53
raises 0.59 0.67 0.45 0.64 1.00 0.38 0.57
critical 0.16 0.19 0.15 0.12 0.38 1.00 0.28
advance 0.16 0.22 0.34 0.53 0.57 0.28 1.00
symnum(Ca) # simple graphic
rt cm p l rs cr a
rating 1
complaints + 1
privileges . . 1
learning , . . 1
raises . , . , 1
critical . 1
advance . . . 1
attr(),"legend")
[1] 0 ` ' 0.3 ` .' 0.6 ` , 0.8 ` +' 0.9 ` '*' 0.95 `B` 1
heatmap(Ca, symm = TRUE, margins = c(6,6))

```



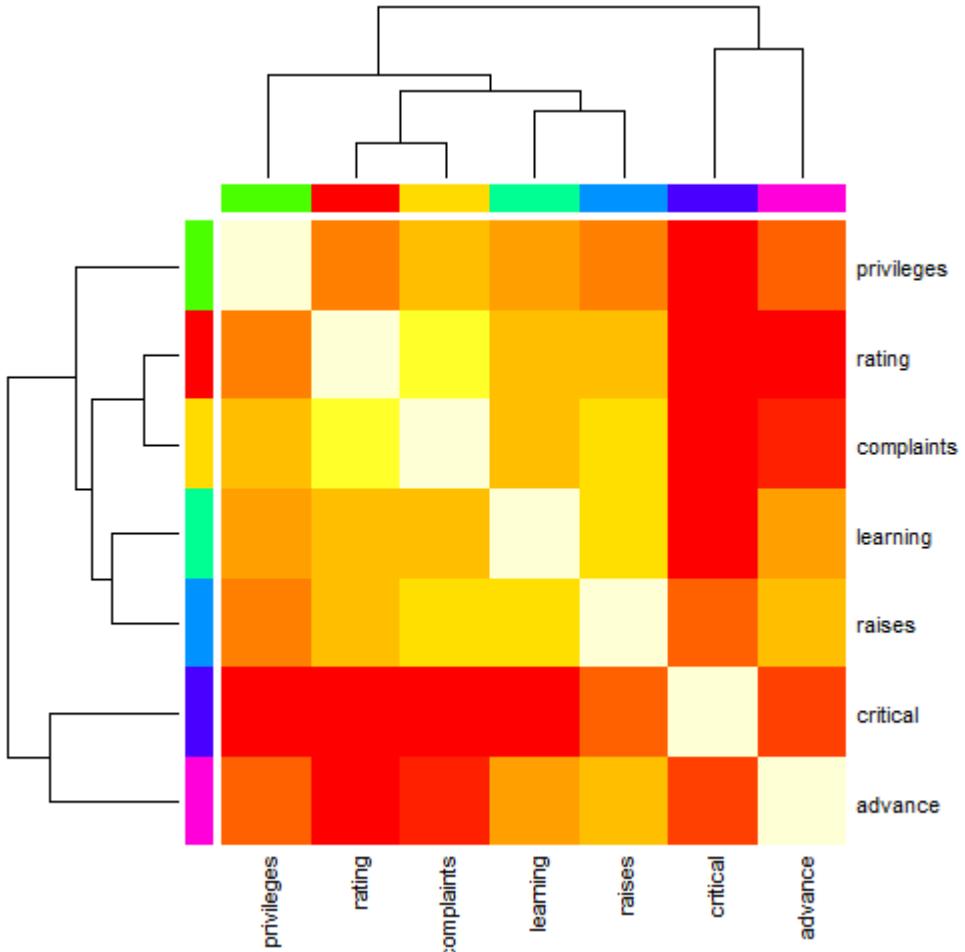
## 5 ( NO reorder ())

```
heatmap(Ca, Rowv = FALSE, symm = TRUE, margins = c(6, 6))
```



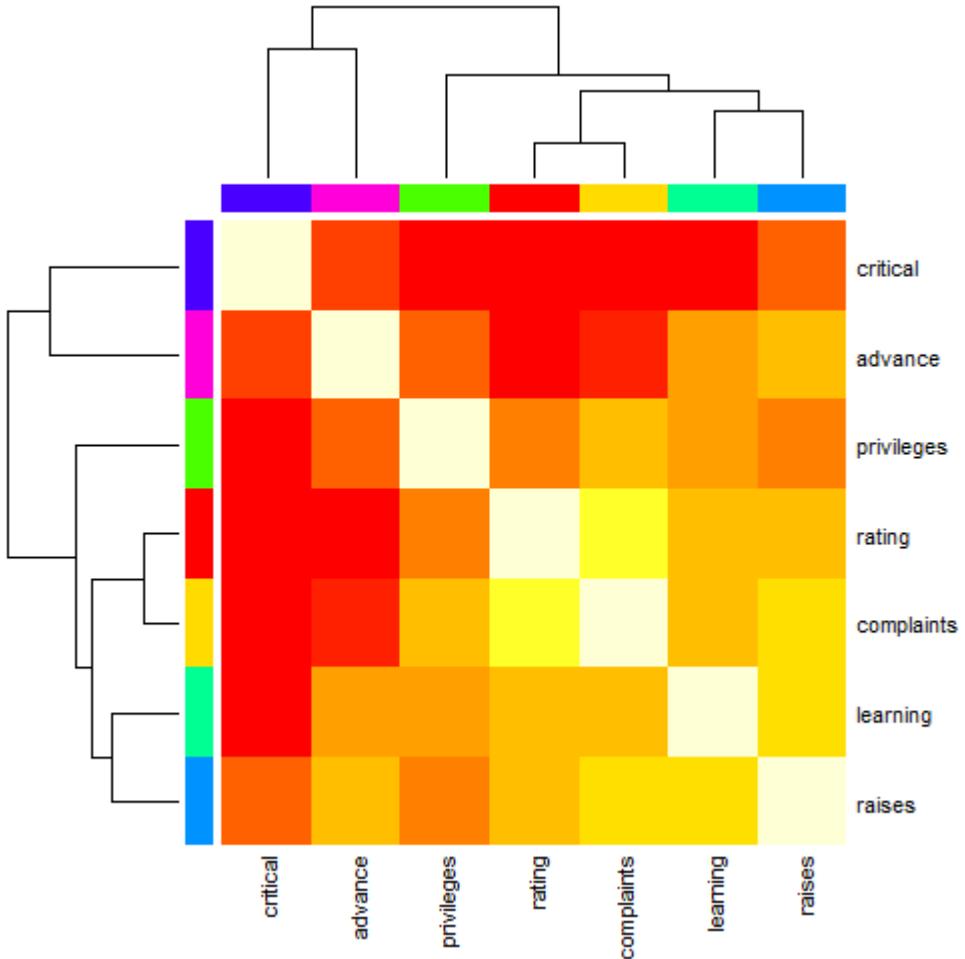
6 ( , )

```
cc <- rainbow(nrow(Ca))
heatmap(Ca, Rowv = FALSE, symm = TRUE, RowSideColors = cc, ColSideColors = cc,
 margins = c(6,6))
```



7 ( )

```
heatmap(Ca, symm = TRUE, RowSideColors = cc, ColSideColors = cc,
 margins = c(6,6))
```



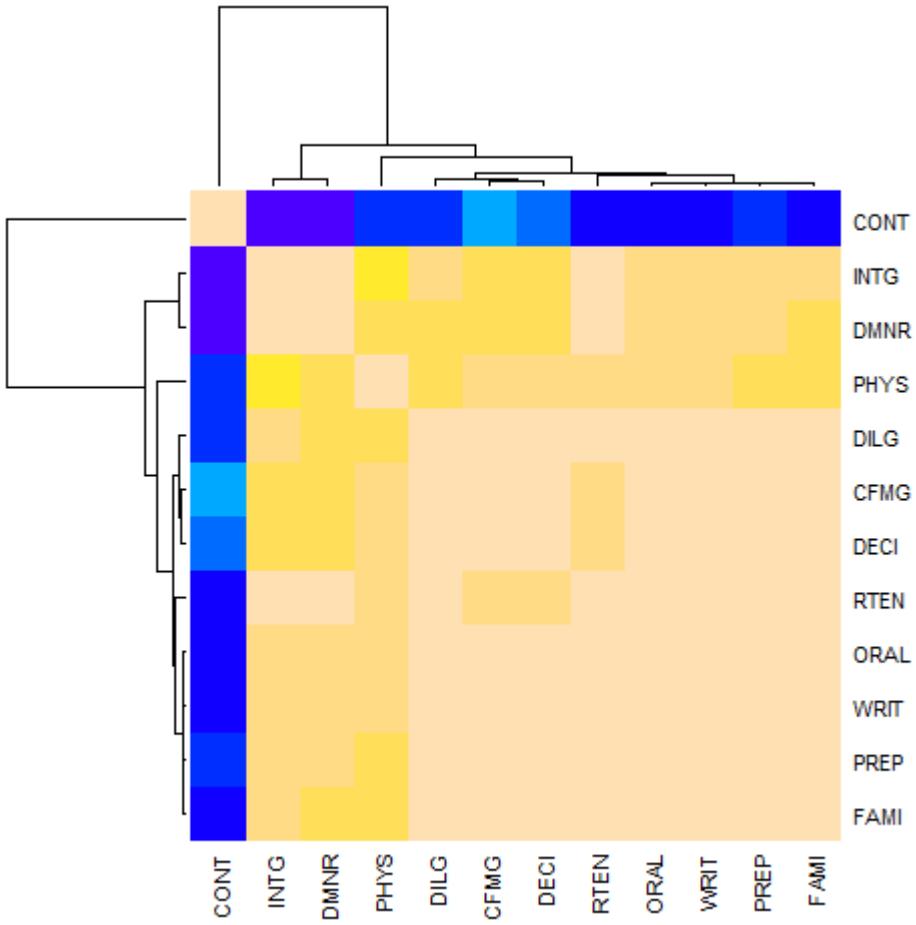
## 8 ( cor() )

```

symnum(cU <- cor(USJudgeRatings))
CO I DM DI CF DE PR F O W PH R
CONT 1
INTG 1
DMNR B 1
DILG + + 1
CFMG + + B 1
DECI + + B B 1
PREP + + B B B 1
FAMI + + B * * B 1
ORAL * * B B * B B 1
WRIT * + B * * B B B 1
PHYS , , + + + + + + + 1
RTEN * * * * B * B B * 1
attr(),"legend")
[1] 0 ' 0.3 ' 0.6 ' 0.8 ' 0.9 ' 0.95 'B' 1

hU <- heatmap(cU, Rowv = FALSE, symm = TRUE, col = topo.colors(16),
 distfun = function(c) as.dist(1 - c), keep.dendro = TRUE)

```



```
The Correlation matrix with same reordering:
round(100 * cU[hU[[1]], hU[[2]]])
CONT INTG DMNR PHYS DILG CFMG DECI RTEN ORAL WRIT PREP FAMI
CONT 100 -13 -15 5 1 14 9 -3 -1 -4 1 -3
INTG -13 100 96 74 87 81 80 94 91 91 88 87
DMNR -15 96 100 79 84 81 80 94 91 89 86 84
PHYS 5 74 79 100 81 88 87 91 89 86 85 84
DILG 1 87 84 81 100 96 96 93 95 96 98 96
CFMG 14 81 81 88 96 100 98 93 95 94 96 94
DECI 9 80 80 87 96 98 100 92 95 95 96 94
RTEN -3 94 94 91 93 93 92 100 98 97 95 94
ORAL -1 91 91 89 95 95 95 98 100 99 98 98
WRIT -4 91 89 86 96 94 95 97 99 100 99 99
PREP 1 88 86 85 98 96 96 95 98 99 100 99
FAMI -3 87 84 84 96 94 94 94 98 99 99 100
```

```
The column dendrogram:
utils::str(hU$Colv)
--[dendrogram w/ 2 branches and 12 members at h = 1.15]
|--leaf "CONT"
`--[dendrogram w/ 2 branches and 11 members at h = 0.258]
|--[dendrogram w/ 2 branches and 2 members at h = 0.0354]
| |--leaf "INTG"
| `--leaf "DMNR"
`--[dendrogram w/ 2 branches and 9 members at h = 0.187]
|--leaf "PHYS"
`--[dendrogram w/ 2 branches and 8 members at h = 0.075]
|--[dendrogram w/ 2 branches and 3 members at h = 0.0438]
| |--leaf "DILG"
```

```

| `--[dendrogram w/ 2 branches and 2 members at h = 0.0189]
| |--leaf "CFMG"
| `--leaf "DECI"
`--[dendrogram w/ 2 branches and 5 members at h = 0.0584]
|--leaf "RTEN"
`--[dendrogram w/ 2 branches and 4 members at h = 0.0187]
|--leaf "ORAL"
`--leaf "WRIT"
`--[dendrogram w/ 2 branches and 2 members at h = 0.0101]
|--leaf "PREP"
`--leaf "FAMI"

```

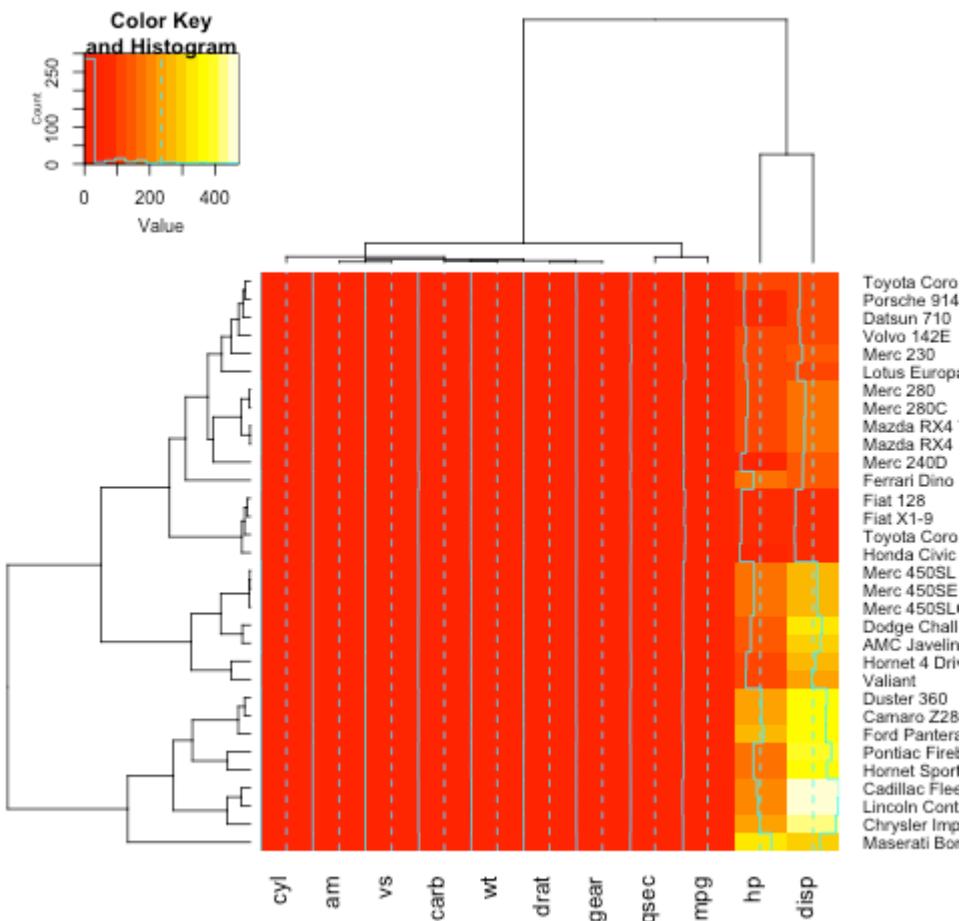
2

:

```
x <- as.matrix(mtcars)
```

```
heatmap.2 - heatmap ,
```

```
require(gplots)
heatmap.2(x)
```

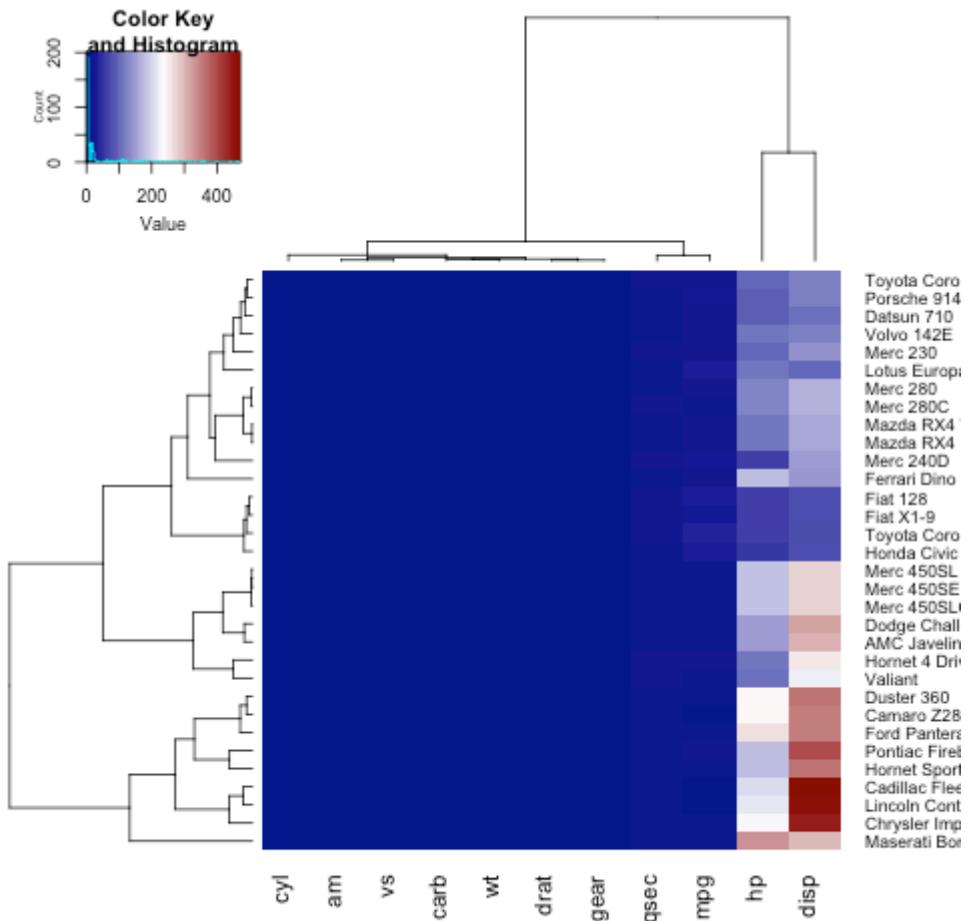


, x y main , xlab ylab .

```
heatmap.2(x, main = "My main title: Overview of car features", xlab="Car features", ylab =
"Car brands")
```

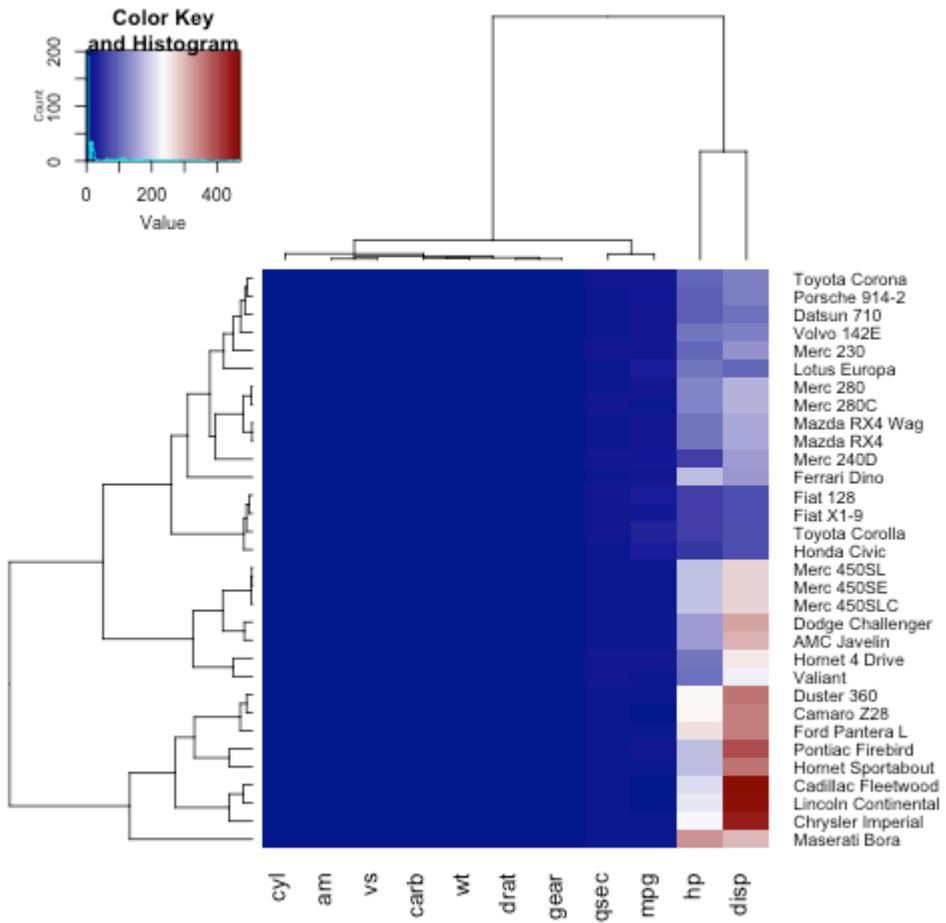
```
colorRampPalette col .
```

```
heatmap.2(x, trace="none", key=TRUE, Colv=FALSE,dendrogram = "row",col =
colorRampPalette(c("darkblue","white","darkred"))(100))
```

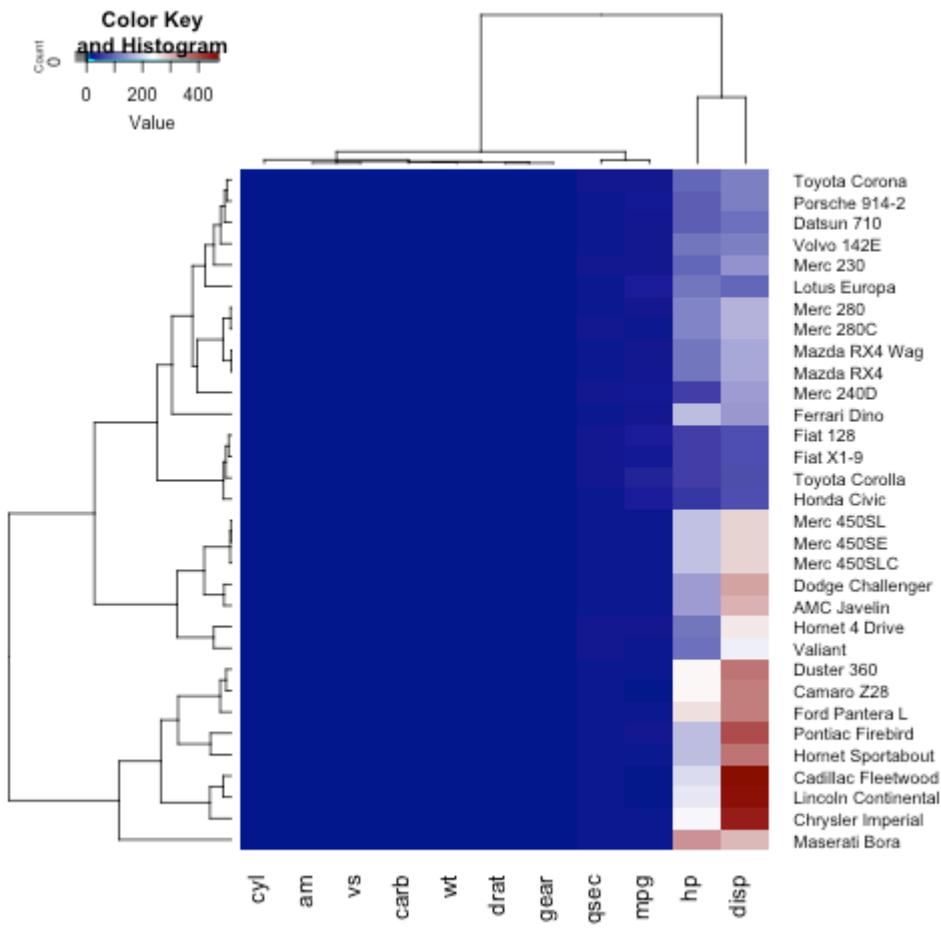


```
y () . margins .
```

```
heatmap.2(x, trace="none", key=TRUE,col =
colorRampPalette(c("darkblue","white","darkred"))(100), margins=c(5,8))
```

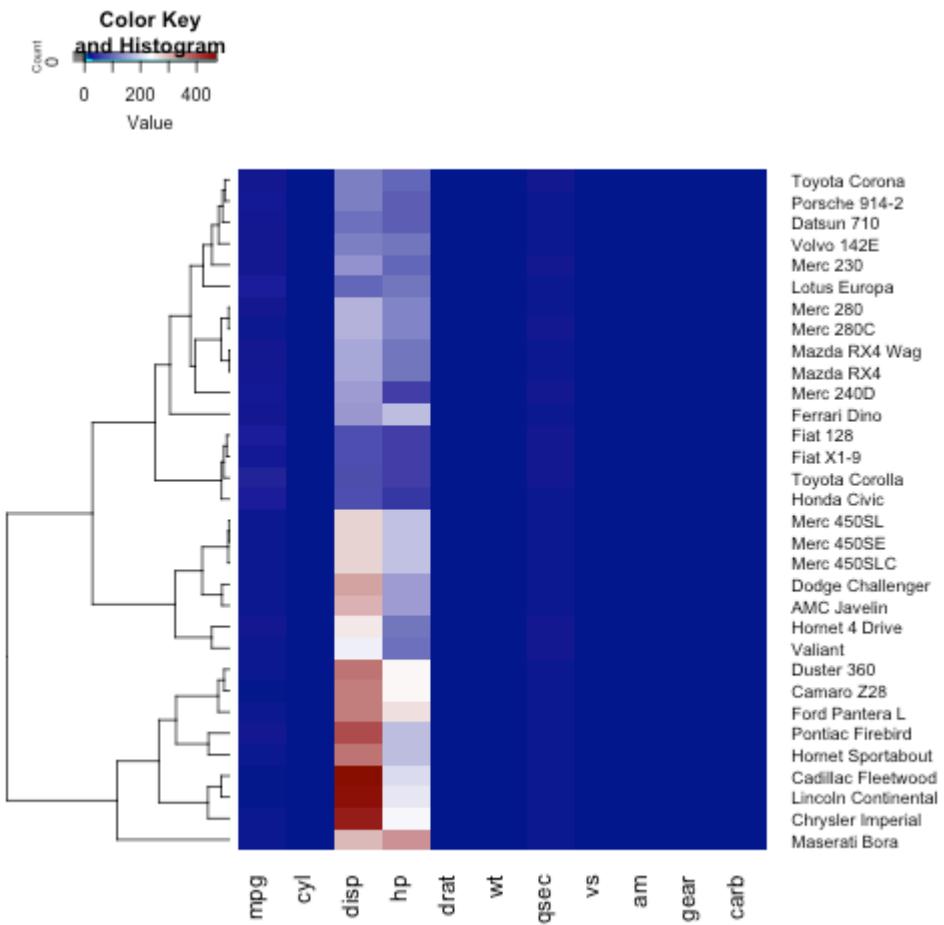


```
lhei lwid (,) .
```



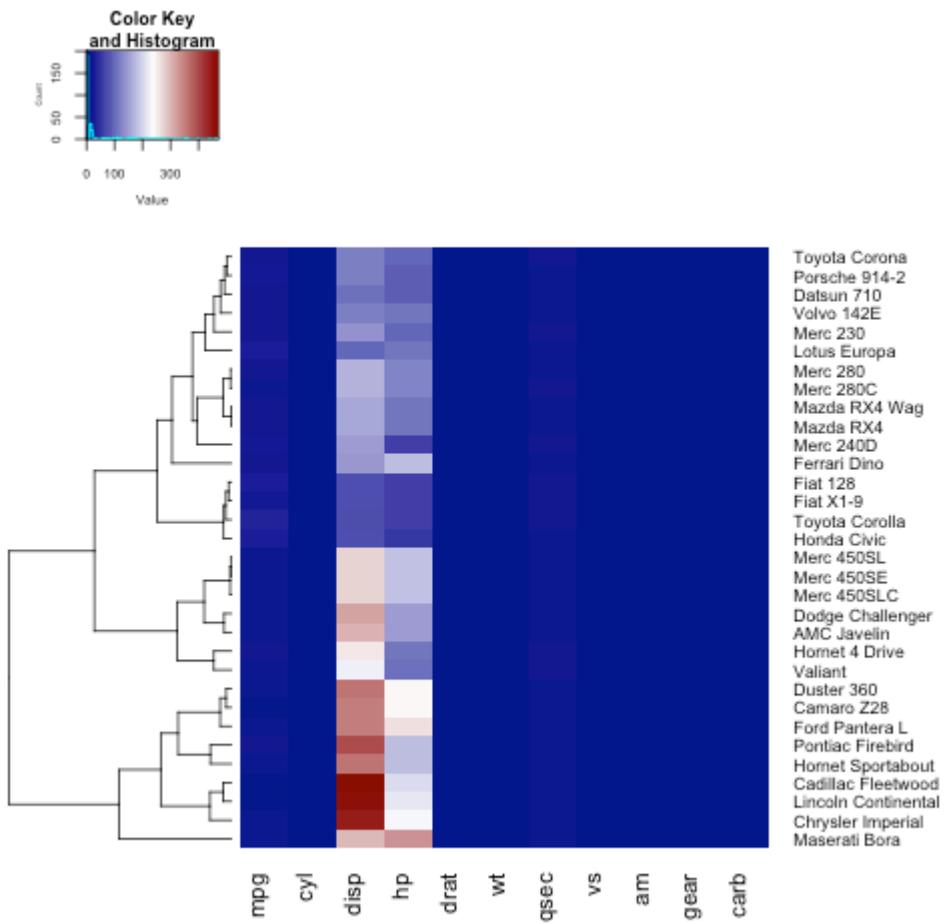
```
() Colv=FALSE (Rowv=FALSE) dendrogram .
```

```
heatmap.2(x, trace="none", key=TRUE, Colv=FALSE, dendrogram = "row", col =
colorRampPalette(c("darkblue","white","darkred"))(100), margins=c(5,8), lwid = c(5,15), lhei =
c(3,15))
```



```
, par(cex.main, cex.lab, cex.axis).
```

```
par(cex.main=1, cex.lab=0.7, cex.axis=0.7)
heatmap.2(x, trace="none", key=TRUE, Colv=FALSE, dendrogram = "row", col =
colorRampPalette(c("darkblue","white","darkred"))(100), margins=c(5,8), lwid = c(5,15), lhei = c(5,15))
```



.2 : <https://riptutorial.com/ko/r/topic/4814/-----2>

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		<a href="#">Padmanabhan</a> , <a href="#">seasmith</a> , <a href="#">SymbolixAU</a> , <a href="#">theArun</a> , <a href="#">user890739</a> , <a href="#">xamgore</a> , <a href="#">zx8754</a>
124		<a href="#">Abdou</a> , <a href="#">Alex</a> , <a href="#">Artem Klevtsov</a> , <a href="#">David Arenburg</a> , <a href="#">David Leal</a> , <a href="#">Frank</a> , <a href="#">Gavin Simpson</a> , <a href="#">Jaap</a> , <a href="#">NWaters</a> , <a href="#">R. Schifini</a> , <a href="#">SommerEngineering</a> , <a href="#">Steve_Corrin</a> , <a href="#">Tensibai</a> , <a href="#">thelatemail</a> , <a href="#">user2100721</a>
125	(CSV, TSV )	<a href="#">a.powell</a> , <a href="#">Aaghaz Hussain</a> , <a href="#">abhileor</a> , <a href="#">Alex</a> , <a href="#">alistaire</a> , <a href="#">Andrea</a> , <a href="#">Cirillo</a> , <a href="#">bartektartanus</a> , <a href="#">Carl Witthoft</a> , <a href="#">Carlos Cinelli</a> , <a href="#">catastrophicfailure</a> , <a href="#">cdrini</a> , <a href="#">Charmgoggles</a> , <a href="#">Crops</a> , <a href="#">DaveRGP</a> , <a href="#">David</a> , <a href="#">Arenburg</a> , <a href="#">Dawny33</a> , <a href="#">Derwin McGeary</a> , <a href="#">EDi</a> , <a href="#">Eric Lecoutre</a> , <a href="#">FoldedChromatin</a> , <a href="#">Frank</a> , <a href="#">Gavin Simpson</a> , <a href="#">gitblame</a> , <a href="#">Hairizuan</a> , <a href="#">Noorazman</a> , <a href="#">herbaman</a> , <a href="#">ikashnitsky</a> , <a href="#">Jaap</a> , <a href="#">Jeromy Anglim</a> , <a href="#">JHowlX</a> , <a href="#">joeyreid</a> , <a href="#">Jordan Kassof</a> , <a href="#">K.Daisey</a> , <a href="#">kitman0804</a> , <a href="#">kneijenhuijs</a> , <a href="#">Imo</a> , <a href="#">Ioki</a> , <a href="#">Miha</a> , <a href="#">PAC</a> , <a href="#">polka</a> , <a href="#">russellpierce</a> , <a href="#">Sam</a> , <a href="#">Firke</a> , <a href="#">stats-hb</a> , <a href="#">Thomas</a> , <a href="#">Uwe</a> , <a href="#">zacdav</a> , <a href="#">zelite</a> , <a href="#">zx8754</a>
126	: +	<a href="#">YCR</a>
127		<a href="#">Hack-R</a>
128		<a href="#">42-</a> , <a href="#">Agriculturist</a> , <a href="#">alexis_laz</a> , <a href="#">alistaire</a> , <a href="#">dayne</a> , <a href="#">Frank</a> , <a href="#">Gavin</a> , <a href="#">Simpson</a> , <a href="#">Gregor</a> , <a href="#">L.V.Rao</a> , <a href="#">Mario</a> , <a href="#">mrip</a> , <a href="#">RamenChef</a> , <a href="#">smci</a> , <a href="#">user2100721</a> , <a href="#">zx8754</a>
129		<a href="#">nrussell</a> , <a href="#">russellpierce</a>
130		<a href="#">dayne</a> , <a href="#">Frank</a>
131	.2	<a href="#">AndreyAkinshin</a> , <a href="#">Nanami</a>