



# R

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## Documentation of Project:

R is a programming language for statistical computing and graphics used to clean, analyze, and graph your data. In this tutorial, I plan to introduce the language, its IDE and usage in various forms. However, as it is not like other programming languages, rules and syntax will be covered. Concepts such as flow of control (loops and if statements) and packages built in R will be discussed. Data sets such as data frames, vectors, matrices and factors will be shown and will end with reading csv files and creating several different plots and graphs.

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## Script:

*15 minute video*

## Outline -

1. What is R? What is it used for? What IDE can I use (terminal vs. RStudio) ?
2. Important Functions in R
3. Graphs and plots
4. Data Frames, arrays, matrices, vectors
5. Reading csv file, manipulating and plotting

## Important Points -

- R can be used as a calculator
- library

- greater-than sign (>) is the prompt symbol which appears in the Console Pane
- : is used to mention a range of number → 10:20 would be [10-20] (both included), could also use 2: (3\*2+5) which would be [2-11]
- constants such as 'pi' → (1:3) + pi would give {414,5.14,6.14}
- power operator (^) 3^3 gives 27  
Modular arithmetic → 31%%7 gives 3 (remainder)  
integer part → 31%/7 gives 4
- Workspace known as the global environment that can be used to store the results of calculations, and other objects (variables)  
interest <- 10^20 (sign convention)  
could also use =, but <- is preferred as we are requesting an action rather than stating a relation
- we get around most of our work in R through functions.  
function to quit R is q()
- R is case-sensitive  
x <- 1:10  
MEAN(x) would not work unless you create a user-defined function MEAN() which uses the mean() function  
MEAN <- mean ; now the MEAN(x) function call would work
- A list of all objects in the current workspace can be printed to the screen using the objects() function; a synonym is ls() function
- Remember that if we quit our R session without saving the workspace image, then these objects will disappear. If we save the workspace image, then the workspace will be restored at our next R session.
- runif() ⇒ random number generator  
runif(4,min= , max= )
- Vectors in R →  
numeric vectors is a list of numbers.  
c() function is used to join values into a vector  
a <- c(10,2,5,8,3,4,7,2,7)  
ax <- c(a,x) (vectors can be internally joined too)

- extracting elements from vectors  
`ax[3]` or `a[c(2,5,1)]` or `ax[1:5]`  
negative indices can be used to avoid certain elements → `ax[-c(2,5)]`
- can easily perform vector arithmetic  
`x^3, y <- x + 6`
- patterned vectors :  
`seq(1,9, by = 2)` → 1 3 5 7 9 (if by is not specified, by default 1 will be used)  
`rep(3,5)` → 3 3 3 3 3  
`rep(seq(2,10,by=2),2)` → 2 4 6 8 10 2 4 6 8 10  
`rep(c(1,3), each = 4)` → 1 1 1 1 3 3 3 3  
`rep(c(1,2),c(4,9))` → 1 1 1 1 2 2 2 2 2 2 2 2 2 2  
`rep(1:5, rep(2,5))` → 1 1 2 2 3 3 4 4 5 5
- random pattern vectors :  
`sample(1:6, size = 5, replace = TRUE)`  
die is tossed 5 times to give random values between 1-6
- character vectors :  
all elements of a vector must be of the same type,  
`colors <- c("red","blue","green")`  
to take substrings, we use `substr(x, start, stop)` like `substr(colors,1,2)`  
`paste()` function to concatenate - `paste (colors, "flowers")`  
`collapse = ", "` used to make all into one string
- Factors :  
alternative way to store character data  
provide a level for the data  
`as.integer(grp)` - non repetitive
- Matrices and arrays  
`m <- matrix(1:6, nrow = 2, ncol = 3)`  
access element of a matrix using indices → `m[1,2]`  
access element of a array using indices → `m[4]`  
access row of a matrix using indices → `m[1,]`  
access column of a matrix using indices → `m[,2]`  
`a <- array(1:24, c(3,4,2))`
- use `?q` or `help(q)` to access the help facility for a function q

- Storage of numbers : in the form of binary  $\Rightarrow 2^0, 2^1, 2^2, \dots$
- `values <- NULL` (keeps the variable empty)  
`xyz[seq(2, 20, 2)] <- seq(2, 20, 2)`  
`## [1] NA 2 NA 4 NA 6 NA 8 NA 10 NA 12 NA 14 NA 16 NA 18 NA 20`  
`is.na(xyz)`  $\Rightarrow$  detects if there are null values
- elementary built in functions:  
`var(x)`, `summary(x)`, `length(x)`, `min(x)`, `max(x)`, `pmin(x)` {pairwise minima}, `pmax(x)` {pairwise maxima}, `range(x)`, `IQR(x)`
- Logical operations in R:  
`a <- c(TRUE, FALSE, FALSE, TRUE)`  
`b <- c(13, 6, 3, 2)`  
`b[a]` gives 13 and 2  
`sum(a)` gives 2  $\rightarrow$  number of true values  
`!a` gives FALSE TRUE TRUE FALSE  
`&&` and `||` are similar to `&` and `|`; they are not vectorized: only one calculation is done
- Relational operators:  
`v <- c(4, 6, 2, 9, 7)`  
`v > 5` gives you FALSE TRUE FALSE TRUE TRUE
- Data frames and lists:  
data sets are stored in R as data frames, and we get some with R, such as `women`.  
`summary(women)`, `nrow(women)`, `ncol(women)`, `dim(women)`, `str(women)`  
extract from data frames similar to matrices  
columns can be accessed using `$` operator  
`women$height[women$weight > 150]`  
`with()` function allows us to access columns of data frame without using `$`,  
`with(women, weight/height)`  
constructing data frames: `xy <- data.frame(x, y)`
- `barplot(WorldPhones51, cex.names = 0.75, cex.axis = 0.75,`  
`main = "Numbers of Telephones in 1951")`
- `dotchart(WorldPhones51, xlab = "Numbers of Phones ('000s)")`
- `groupsizes <- c(18, 30, 32, 10, 10)`  
`labels <- c("A", "B", "C", "D", "F")`

```
pie(groupsizes, labels,col = c("grey40", "white", "grey", "black", "grey90"))
```

- `boxplot(Sepal.Length ~Species, data = iris, ylab = "Sepal length (cm)", main = "Iris measurements", boxwex = 0.5)`
- Flow control:  
for() loop - for (name in vector) { commands }  
factorial function -  
n <- 10  
res <- 1  
for (i in 1:n)  
    res <- res \* i  
res
- if() statement -  
if (condition) {commands when TRUE}  
if (condition) {commands when TRUE} else {commands when FALSE}  
x <- 3 if (x > 2) y <- 2 \* x else y <- 3 \* x
- while() loop-  
Fib1 <- 1  
Fib2 <- 1  
Fibonacci <- c(Fib1, Fib2)  
while (Fib2 < 300) {  
    Fibonacci <- c(Fibonacci, Fib2)  
    oldFib2 <- Fib2  
    Fib2 <- Fib1 + Fib2  
    Fib1 <- oldFib2  
}
- replicate() function:  
evaluate the statements n times - `replicate(n, {statements})`

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## Personal Notes -

|          |  |          |
|----------|--|----------|
| <b>2</b> | <b>Introduction to the R language</b>    | <b>7</b> |
| 2.1      | First steps                              | 7        |
| 2.2      | Basic features of R                      | 11       |
| 2.3      | Vectors in R                             | 13       |
| 2.4      | Data storage in R                        | 22       |
| 2.5      | Packages, libraries, and repositories    | 27       |
| 2.6      | Getting help                             | 28       |
| 2.7      | Logical vectors and relational operators | 34       |
| 2.8      | Data frames and lists                    | 37       |
| 2.9      | Data input and output                    | 43       |

|          |   |           |
|----------|---|-----------|
| <b>3</b> | <b>Programming statistical graphics</b> | <b>49</b> |
| 3.1      | High level plots                        | 50        |
| 3.2      | Choosing a high level graphic           | 62        |
| 3.3      | Low level graphics functions            | 63        |
| 3.4      | Other graphics systems                  | 70        |

|          |                                       |           |
|----------|---------------------------------------|-----------|
| <b>4</b> | <b>Programming with R</b>             | <b>76</b> |
| 4.1      | Flow control                          | 76        |
| 4.2      | Managing complexity through functions | 91        |
| 4.3      | The <code>replicate()</code> function | 97        |
| 4.4      | Miscellaneous programming tips        | 97        |
| 4.5      | Some general programming guidelines   | 100       |
| 4.6      | Debugging and maintenance             | 107       |
| 4.7      | Efficient programming                 | 113       |