

Introduction to R for spatial modeling

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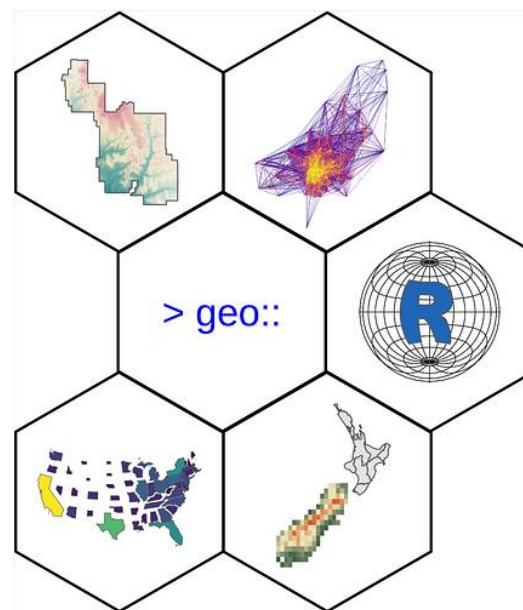
Content

□ Part 1

- R and RStudio
- Setting up an R session
- Data types
- Data structures
- Selecting subsets
- Functions
- Import/export of data
- Plotting
- Working with spatial data in R

□ Part 2

- Lets practice!!



What is R?

- R is a free programming language for statistics
- R is command line based
- R can be expanded through packages
- R is used to analyze and [visualize](#) (geographic) data
- R is comparatively easy to access



Why R?

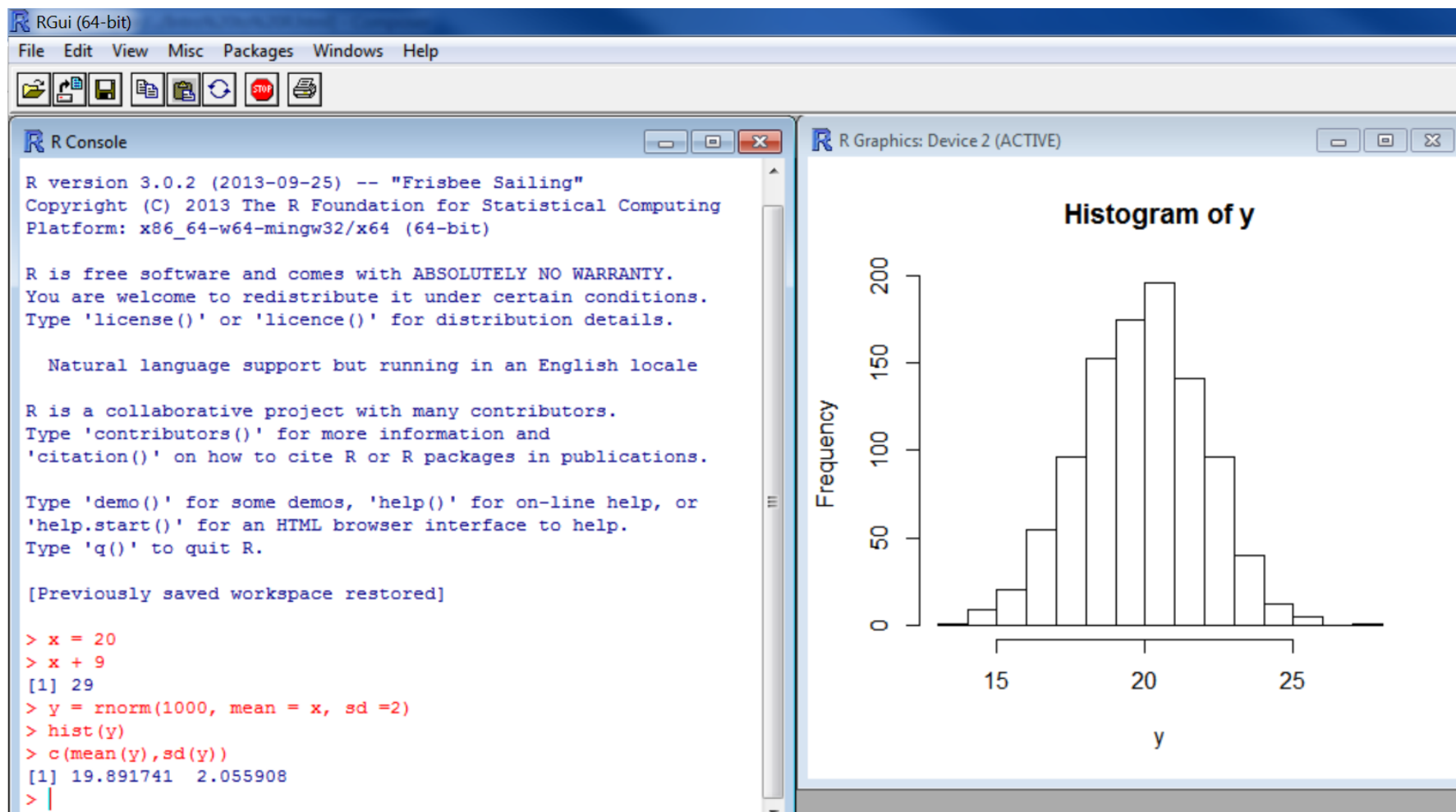
- It is free
- Available for the most popular operating systems
- Low system requirements
- Continuous [further development](#)
- Active user community with its own conference (useR!)
- Application in many natural sciences
- [PYPL](#) Popularity of Programming Language: 7th place



First steps

- <https://www.r-project.org/>
 - R website, where you will find the program itself, FAQs, manuals and other useful information about and around R:
- CRAN (Comprehensive R Archive Network; cran.r-project.org)
- Manuals (cran.r-project.org/manuals.html)
- Mailing lists
- Books list

R software



First steps

- Other user interfaces

- R Commander (<https://www.rcommander.com/>)

- RStudio(<https://rstudio.com/>)

- (<https://rstudio.com/products/rstudio/download/>)

- And some others (e.g., Tinn-R)

- Open source and commercial user interfaces

- Desktop and server variants

RStudio



The screenshot shows the RStudio interface with four numbered callouts highlighting key components:

- 1- Code Editor**: The top-left pane showing R code for loading ggplot2, viewing the diamonds dataset, and creating a faceted plot.
- 2- R Console**: The bottom-left pane showing the output of the R code, including summary statistics for the diamonds dataset.
- 3- Workspace and History**: The top-right pane showing the current workspace with the 'diamonds' dataset and its summary.
- 4 - Plots and files**: The bottom-right pane showing a faceted scatter plot titled 'Diamond Pricing' with 'Price' on the y-axis and 'Carat' on the x-axis, faceted by clarity.

```
1 library(ggplot2)
2
3 view(diamonds)
4 summary(diamonds)
5
6 summary(diamonds$price)
7 aveSize <- round(mean(diamonds$carat), 4)
8 clarity <- levels(diamonds$clarity)
9
10 p <- ggplot(diamonds, aes(carat, price, color=clarity))
11
12 # Create a faceted plot
13 p <- facet_wrap(p, ~ clarity,
14               main="Diamond Pricing")
```

Console output:

	x	y	z
Min.	0.000	0.000	0.000
1st Qu.	4.710	4.720	2.910
Median	5.700	5.710	3.530
Mean	5.700	5.710	3.539
3rd Q	6.700	6.710	4.040
Max.	18.820	18.820	1.800

Workspace summary:

diamonds	53940 obs. of 10 variables	
aveSize	0.7979	

Plot title: Diamond Pricing

Plot axes: Price (y-axis), Carat (x-axis)

Plot legend: VS2, VS1, VVS2, VVS1, IF

First steps-Help

- R-help(Mailing-Listen)

Most [important list](#) for user questions
(Pay attention to [posting](#))

- R Community, most problems are not new and solutions can be found online

[Stackoverflow](#)

[R-bloggers](#),

[Quick-R](#),

- Books and eBooks

[R Cookbook](#)

[R for Data Science](#), [Spatial Data Science](#),

[Geocomputationwith R](#), [The R Inferno](#),
[Rstudio Cheatsheets](#)

Essentials of R

- Basic arithmetic functions:

+ (addition), **-** (subtraction), ***** (multiplication), **/** (division),
^ (exponentiation)

7 + 4 # => 11

7 - 4 # => 3

7 / 2 # => 3.5

7 * 2 # => 14

7 ^ 2 # => 49

Essentials of R

- Basic arithmetic operations:

Logarithms and exponentials: `log2(x)`, `log10(x)`, `exp(x)`

Trigonometric functions: `cos(x)`, `sin(x)`, `tan(x)`, `acos(x)`, `asin(x)`, `atan(x)`

Other mathematical functions: `abs(x)`: absolute value; `sqrt(x)`: square root.

```
log2 (4) # => 2
```

```
abs (-4) # => 4
```

```
sqrt (4) # => 2
```

Essentials of R

▪Assigning values to variables:

```
s <- 5
```

```
5
```

```
s <- seq(from=20, to=0, by=-2)
```

```
20 18 16 14 12 10 8 6 4 2 0
```

```
s <- rep(1:4, 2)
```

```
1 2 3 4 1 2 3 4
```

```
s <- rnorm(10, mean = 10, sd = 1)
```

```
9.3 10.5 10.4 13.2 10.1 10.9 11.5 10.4 9.1 9.6
```

Essentials of R

- Assigning values to variables:

```
Clay <- 40
```

```
Sand <- 40
```

```
Silt <- 20
```

```
Sum <- Clay + Sand + Silt
```

```
print(Sum) # => 100
```

Note: R is case sensitive!

Data types

▪ Basic data types:

▪ Numeric

```
> x <- 10.1
> x
[1] 10.1
> class(x)
[1] "numeric"
```

▪ Integer

```
> y <- as.integer(x)
> y
[1] 10
> class(y)
[1] "integer"
```

▪ Character

```
> z <- "Bas"
> z
[1] "Bas"
> class(z)
[1] "character"
```

▪ Logical

```
> x <- 1; y <- -2
> z <- x > y
> z
[1] FALSE
> class(z)
[1] "logical"
>
```

class(x)

Data structures

▪ Vectors:

- Sequence of elements of the same basic type (character, logical, integer or numeric).

```
x <- c(1, 2, 3)
```

```
x <- 1:3
```

```
y <- c(1, 1, 1)
```

```
y <- rep(2, 10)
```

```
z <- as.character(1:3)
```

```
z <- c("a", "b", "c")
```

```
length(z)
```

```
names(x) <- z
```

```
x[2:3]
```

```
x["b"]
```

```
as.vector(x)
```

```
is.vector(x)
```


Data structures

- **Factors:**

- **A vector with categorical values (classes)**

```
x <- c(1, 2, 3)
```

```
y <- as.factor(x)
```

```
> y
```

```
[1] 1 2 3
```

```
Levels: 1 2 3
```

Data structures

▪Matrix:

- Matrices are similar to vectors, but have two dimensions

```
x <- 1:20
x <- matrix(x, 5, 4)           # matrix(x, nrow = 5, ncol = 4)
x[2, 3]
x[c(1, 5), 2:4]
x[, 2:4]

dim(x); nrow(x); ncol(x)
length(x)
as.matrix(x)
is.matrix(x)
x[, "b"]
x[, c("a", "b")]
```

Data structures

▪ Arrays:

- **Arrays are similar to matrices, but can have more than two dimensions**

```
x <- 1:60
```

```
x <- array(x, c(5, 4, 3))
```

```
x[2, 3, 1]
```

```
x[1, 2:4, 3]
```

```
x[, , 1]
```

```
dim(x); dimnames(x)
```

```
nrow(x); ncol(x)
```

```
length(x)
```

```
as.array(x)
```

```
is.array(x)
```

Data structures

▪List:

- Ordered sequence of objects of the same or dissimilar type

```
x <- list(Eins = 11:15, Zwei = c("a", "b", "c"), Drei = (1:4) > 0)
```

```
y <- list(x = x, Vier = 1:3)
```

```
x$Eins
```

```
y$x$Eins
```

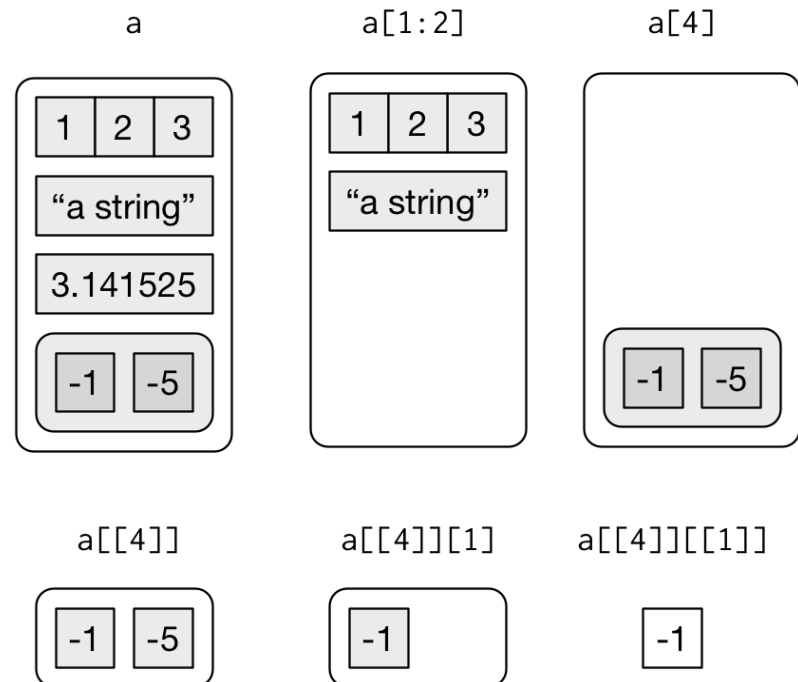
```
y$Vier
```

```
y[[2]]
```

```
length(x)
```

```
length(y)
```

```
y$Fuenf <- names(x)
```



Data structures

▪Data frame:

- Data frames are similar to matrices but with the same or different types

```
x <- data.frame(N = 11:14, C = c("a", "b", "c", "d"), L = (1:4) > 0)
```

```
dim(x)
```

```
nrow(x)
```

```
ncol(x)
```

```
length(x)
```

```
names(x)
```

```
as.data.frame(x)
```

```
is.data.frame(x)
```

```
## Select row 1 in column 2
```

```
df[1,2]
```

	ID	items	store	price
1	10	book	TRUE	2.5
2	20	pen	FALSE	8.0
3	30	textbook	TRUE	10.0
4	40	pencil_case	FALSE	7.0

```
## Select Rows 1 to 3 and columns 3 to 4
```

```
df[1:3, 3:4]
```

```
## Select Rows 1 to 2
```

```
df[1:2,]
```

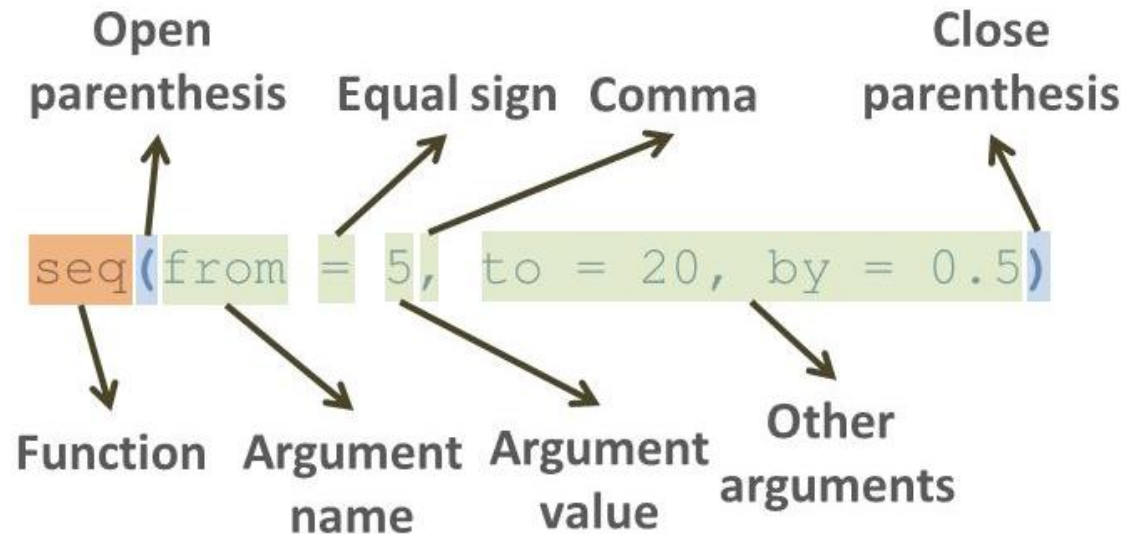
```
## Select Column 1
```

```
df[,1]
```

Functions, Arguments, and Packages

▪Functions:

Data analyses and modelling is done through functions.



hist(x)
plot(x)

args(plot)

Functions, Arguments, and Packages

■ Packages:

install.packages("package name")

install.packages("ggplot2")

```
library(ggplot2)
```

```
detach("ggplot2 ")
```

installed.packages()

remove.packages()

update.packages()



Useful commands

Command	Purpose
<code>help()</code>	Obtain documentation for a given R command
<code>example()</code>	View some examples on the use of a command
<code>c()</code> , <code>scan()</code>	Enter data manually to a vector in R
<code>seq()</code>	Make arithmetic progression vector
<code>rep()</code>	Make vector of repeated values
<code>data()</code>	Load (often into a <code>data.frame</code>) built-in dataset
<code>View()</code>	View dataset in a spreadsheet-type format
<code>str()</code>	Display internal structure of an R object
<code>read.csv()</code> , <code>read.table()</code>	Load into a <code>data.frame</code> an existing data file
<code>library()</code> , <code>require()</code>	Make available an R add-on package
<code>dim()</code>	See dimensions (# of rows/cols) of <code>data.frame</code>
<code>length()</code>	Give length of a vector
<code>ls()</code>	Lists memory contents
<code>rm()</code>	Removes an item from memory
<code>names()</code>	Lists names of variables in a <code>data.frame</code>
<code>hist()</code>	Command for producing a histogram
<code>histogram()</code>	Lattice command for producing a histogram
<code>stem()</code>	Make a stem plot
<code>table()</code>	List all values of a variable with frequencies
<code>xtabs()</code>	Cross-tabulation tables using formulas
<code>mosaicplot()</code>	Make a mosaic plot
<code>cut()</code>	Groups values of a variable into larger bins
<code>mean()</code> , <code>median()</code>	Identify “center” of distribution
<code>by()</code>	apply function to a column split by factors
<code>summary()</code>	Display 5-number summary and mean
<code>var()</code> , <code>sd()</code>	Find variance, sd of values in vector
<code>sum()</code>	Add up all values in a vector
<code>quantile()</code>	Find the position of a quantile in a dataset
<code>barplot()</code>	Produces a bar graph
<code>barchart()</code>	Lattice command for producing bar graphs
<code>boxplot()</code>	Produces a boxplot
<code>bwplot()</code>	Lattice command for producing boxplots

Command	Purpose
<code>plot()</code>	Produces a scatterplot
<code>xyplot()</code>	Lattice command for producing a scatterplot
<code>lm()</code>	Determine the least-squares regression line
<code>anova()</code>	Analysis of variance (can use on results of <code>lm()</code>)
<code>predict()</code>	Obtain predicted values from linear model
<code>nls()</code>	estimate parameters of a nonlinear model
<code>residuals()</code>	gives (observed - predicted) for a model fit to data
<code>sample()</code>	take a sample from a vector of data
<code>replicate()</code>	repeat some process a set number of times
<code>cumsum()</code>	produce running total of values for input vector
<code>ecdf()</code>	builds empirical cumulative distribution function
<code>dbinom()</code> , etc.	tools for binomial distributions
<code>dpois()</code> , etc.	tools for Poisson distributions
<code>pnorm()</code> , etc.	tools for normal distributions
<code>qt()</code> , etc.	tools for student <i>t</i> distributions
<code>pchisq()</code> , etc.	tools for chi-square distributions
<code>binom.test()</code>	hypothesis test and confidence interval for 1 proportion
<code>prop.test()</code>	inference for 1 proportion using normal approx.
<code>chisq.test()</code>	carries out a chi-square test
<code>fisher.test()</code>	Fisher test for contingency table
<code>t.test()</code>	student <i>t</i> test for inference on population mean
<code>qqnorm()</code> , <code>qqline()</code>	tools for checking normality
<code>addmargins()</code>	adds marginal sums to an existing table
<code>prop.table()</code>	compute proportions from a contingency table
<code>par()</code>	query and edit graphical settings
<code>power.t.test()</code>	power calculations for 1- and 2-sample <i>t</i>
<code>anova()</code>	compute analysis of variance table for fitted model

Base R

Cheat Sheet

Getting Help

Accessing the help files

?mean

Get help of a particular function.

help.search('weighted mean')

Search the help files for a word or phrase.

help(package = 'dplyr')

Find help for a package.

More about an object

str(iris)

Get a summary of an object's structure.

class(iris)

Find the class an object belongs to.

Using Libraries

install.packages('dplyr')

Download and install a package from CRAN.

library(dplyr)

Load the package into the session, making all its functions available to use.

dplyr::select

Use a particular function from a package.

data(iris)

Load a built-in dataset into the environment.

Working Directory

getwd()

Find the current working directory (where inputs are found and outputs are sent).

setwd('C://file/path')

Change the current working directory.

Use projects in RStudio to set the working directory to the folder you are working in.

Vectors

Creating Vectors

c(2, 4, 6)	2 4 6	Join elements into a vector
2:6	2 3 4 5 6	An integer sequence
seq(2, 3, by=0.5)	2.0 2.5 3.0	A complex sequence
rep(1:2, times=3)	1 2 1 2 1 2	Repeat a vector
rep(1:2, each=3)	1 1 1 2 2 2	Repeat elements of a vector

Vector Functions

sort(x) Return x sorted.	rev(x) Return x reversed.
table(x) See counts of values.	unique(x) See unique values.

Selecting Vector Elements

By Position

x[4]	The fourth element.
x[-4]	All but the fourth.
x[2:4]	Elements two to four.
x[-(2:4)]	All elements except two to four.
x[c(1, 5)]	Elements one and five.

By Value

x[x == 10]	Elements which are equal to 10.
x[x < 0]	All elements less than zero.
x[x %in% c(1, 2, 5)]	Elements in the set 1, 2, 5.

Named Vectors

x['apple']	Element with name 'apple'.
-------------------	----------------------------

Programming

For Loop

```
for (variable in sequence){  
  Do something  
}
```

Example

```
for (i in 1:4){  
  j <- i + 10  
  print(j)  
}
```

If Statements

```
if (condition){  
  Do something  
} else {  
  Do something different  
}
```

Example

```
if (i > 3){  
  print('Yes')  
} else {  
  print('No')  
}
```

While Loop

```
while (condition){  
  Do something  
}
```

Example

```
while (i < 5){  
  print(i)  
  i <- i + 1  
}
```

Functions

```
function_name <- function(var){  
  Do something  
  return(new_variable)  
}
```

Example

```
square <- function(x){  
  squared <- x*x  
  return(squared)  
}
```

Reading and Writing Data

Input	Output	Description
df <- read.table('file.txt')	write.table(df, 'file.txt')	Read and write a delimited text file.
df <- read.csv('file.csv')	write.csv(df, 'file.csv')	Read and write a comma separated value file. This is a special case of read.table/write.table.
load('file.Rdata')	save(df, file = 'file.Rdata')	Read and write an R data file, a file type special for R.

Conditions

a == b	Are equal	a > b	Greater than	a >= b	Greater than or equal to	is.na(a)	Is missing
a != b	Not equal	a < b	Less than	a <= b	Less than or equal to	is.null(a)	Is null

Types

Converting between common data types in R. Can always go from a higher value in the table to a lower value.

<code>as.logical</code>	TRUE, FALSE, TRUE	Boolean values (TRUE or FALSE).
<code>as.numeric</code>	1, 0, 1	Integers or floating point numbers.
<code>as.character</code>	'1', '0', '1'	Character strings. Generally preferred to factors.
<code>as.factor</code>	'1', '0', '1', levels: '1', '0'	Character strings with preset levels. Needed for some statistical models.

Maths Functions

<code>log(x)</code>	Natural log.	<code>sum(x)</code>	Sum.
<code>exp(x)</code>	Exponential.	<code>mean(x)</code>	Mean.
<code>max(x)</code>	Largest element.	<code>median(x)</code>	Median.
<code>min(x)</code>	Smallest element.	<code>quantile(x)</code>	Percentage quantiles.
<code>round(x, n)</code>	Round to n decimal places.	<code>rank(x)</code>	Rank of elements.
<code>signif(x, n)</code>	Round to n significant figures.	<code>var(x)</code>	The variance.
<code>cor(x, y)</code>	Correlation.	<code>sd(x)</code>	The standard deviation.

Variable Assignment

```
> a <- 'apple'
> a
[1] 'apple'
```




The Environment

<code>ls()</code>	List all variables in the environment.
<code>rm(x)</code>	Remove x from the environment.
<code>rm(list = ls())</code>	Remove all variables from the environment.

You can use the environment panel in RStudio to browse variables in your environment.

Matrixes

```
m <- matrix(x, nrow = 3, ncol = 3)
Create a matrix from x.
```

 <code>m[2,]</code>	- Select a row	<code>t(m)</code>	Transpose
 <code>m[, 1]</code>	- Select a column	<code>m %*% n</code>	Matrix Multiplication
 <code>m[2, 3]</code>	- Select an element	<code>solve(m, n)</code>	Find x in: $m \cdot x = n$

Lists

```
l <- list(x = 1:5, y = c('a', 'b'))
A list is collection of elements which can be of different types.
```

<code>l[[2]]</code>	<code>l[1]</code>	<code>l\$x</code>	<code>l['y']</code>
Second element of l.	New list with only the first element.	Element named x.	New list with only element named y.



Also see the **dplyr** library.

Data Frames



```
df <- data.frame(x = 1:3, y = c('a', 'b', 'c'))
A special case of a list where all elements are the same length.
```

x	y
1	a
2	b
3	c

Matrix subsetting

<code>df[, 2]</code>	
<code>df[2,]</code>	
<code>df[2, 2]</code>	

List subsetting

<code>df\$x</code>		<code>df[[2]]</code>	
<i>Understanding a data frame</i>			
<code>View(df)</code>	See the full data frame.		
<code>head(df)</code>	See the first 6 rows.		

`nrow(df)`
Number of rows.

`ncol(df)`
Number of columns.

`dim(df)`
Number of columns and rows.

`cbind` - Bind columns.



`rbind` - Bind rows.



Strings

Also see the **stringr** library.

<code>paste(x, y, sep = ' ')</code>	Join multiple vectors together.
<code>paste(x, collapse = ' ')</code>	Join elements of a vector together.
<code>grep(pattern, x)</code>	Find regular expression matches in x.
<code>gsub(pattern, replace, x)</code>	Replace matches in x with a string.
<code>toupper(x)</code>	Convert to uppercase.
<code>tolower(x)</code>	Convert to lowercase.
<code>nchar(x)</code>	Number of characters in a string.

Factors

<code>factor(x)</code>	Turn a vector into a factor. Can set the levels of the factor and the order.
<code>cut(x, breaks = 4)</code>	Turn a numeric vector into a factor but 'cutting' into sections.

Statistics

<code>lm(x ~ y, data=df)</code> Linear model.	<code>t.test(x, y)</code> Perform a t-test for difference between means.	<code>prop.test</code> Test for a difference between proportions.
<code>glm(x ~ y, data=df)</code> Generalised linear model.	<code>pairwise.t.test</code> Perform a t-test for paired data.	<code>aov</code> Analysis of variance.
<code>summary</code> Get more detailed information out a model.		

Distributions

	Random Variates	Density Function	Cumulative Distribution	Quantile
Normal	<code>rnorm</code>	<code>dnorm</code>	<code>pnorm</code>	<code>qnorm</code>
Poisson	<code>rpois</code>	<code>dpois</code>	<code>ppois</code>	<code>qpois</code>
Binomial	<code>rbinom</code>	<code>dbinom</code>	<code>pbinom</code>	<code>qbinom</code>
Uniform	<code>runif</code>	<code>dunif</code>	<code>punif</code>	<code>qunif</code>

Plotting

Also see the **ggplot2** library.

 <code>plot(x)</code> Values of x in order.	 <code>plot(x, y)</code> Values of x against y.	 <code>hist(x)</code> Histogram of x.
---	---	---

Dates

See the **lubridate** library.

Working with spatial data in R

- **Spatial Data in R:**

- **R offers a wide variety of packages and tools that can handle**

- **Relevant packages:**

- **sp** : handling spatial data
 - **raster** : reading/manipulating/writing spatial raster data
 - **rgdal** : reading/writing spatial data

Working with spatial data in R

▪ Spatial data class summary:

–**sp**:

- SpatialPointsDataFrame
- SpatialPixelDataFrame
- SpatialGridDataFrame
- SpatialPolygonDataFrame
- SpatialLinesDataFrame

• Format: **shapefile**

• Format: **GeoTiff**

–**raster**:

- RasterLayer(single layer)
- RasterStack/ RasterBrick(multiple layers)

gridded

fullgridded

Working with spatial data in R

■ Importing and exporting spatial data:

- rgdal: **readOGR**(vector), **readGDAL**(raster)
- raster: **raster**

- rgdal: **writeOGR**(vector), **writeGDAL**(raster)
- raster: **writeRaster**

Working with spatial data in R

▪Projections:

- Once you have loaded your spatial data in R, you might need to tell R its geographic projection.
- Check the current projection: **proj4string** function (sppackage).
- Setting a projection: **CRS**function (sppackage).
- Reprojectingto another coordinate system:
spTransformfunction (sppackage) or **projectRaster**(raster package).

Please install [R](#) and [RStudio](#)

Please install [SAGA](#) and [Google Earth](#)

