

Introduction to R, Rstudio & Project Management

Berry Boessenkool, uni-potsdam.de, May 2017

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swc-bb.github.io/2017-05-17-r-workshop

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ENCOURAGED

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Survey

knowledge survey to determine focus for this session

bit.ly/knowR

RStudio

The screenshot shows the RStudio environment with several annotations in red and orange text boxes:

- RUN CODE**: A red circle highlights the 'Run' button in the top toolbar.
- SCRIPTS**: A red circle highlights the 'Source' button in the top toolbar.
- OBJECTS IN WORKSPACE**: A red circle highlights the 'Environment' pane on the right, which lists objects like 'statlocsp', 'statnames', 'name', 'number', 'values', 'country', 'd', 'first', and 'man'.
- DOCUMENTATION**: A red circle highlights the 'Help' button in the bottom toolbar.
- PLOTS**: A red circle highlights the 'Plots' pane on the right, which displays a scatter plot of 'sort(as.Date(first))' vs 'Index'.
- CODE EXECUTION**: A red circle highlights the 'Console' pane at the bottom, which shows the execution of R code and the resulting output.

The R code in the editor includes:

```

country <- raster::shapefile("../RiverFiles/country", country.shp)
country <- sp::spTransform(country,
sp::plot(country, col=1:8)

# draw rivers depending on (strahler
riverlines <- function(w1=2.5, w2=2, w3=1)
{
  lines(river[river@data$rhine=1,], col="blue", lwd=w1)
  lines(      , col="blue", lwd=w2)
  lines(      , col="blue", lwd=w3)
}

library(OSMscale) # pointsMap, projectPoints
map <- pointsMap(yy.xx, statlocs@data, zoom=7, fx=0.3, type="maptoolkit-to
statlocsp <- projectPoints(yy.xx, statlocs@data, to=posm())
map2 <- pointsMap(yy.xx, statlocs@data, map=map, proj=pl1())

first <- sapply(d, function(x) as.character(x$date[1]))
plot(sort(as.Date(first)))
range(as.Date(first))
rm(first)
last <- sapply(d, function(x) as.character(x$date[1]))

```

The console output shows the execution of the code, including the download of a map and the plotting of the data.

RStudio configuration

keyboard shortcuts (ALT+SHIFT+K)

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Recommended settings for reproducible research under

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ON: Restore previously open source documents at startup

OFF: Restore .Rdata into workspace at startup

Save workspace to .RData on exit: **NEVER**

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Instead use `save(object, file="object.Rdata")` after long computations. You can load them later with `load("object.Rdata")`.

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Tools - Global Options - Code - Display

ON: Show margin (Margin column:80) *People hate horizontal scrolling!*

Tools - Global Options - Code - Saving

Line ending conversion: **Windows (CR/LF)**

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- ▶ comments: `# everything after a hashtag is not executed.`

Exercise 1: Basic R syntax

- ▶ Open Rstudio, start new script. Write comments about what you do, save the file in a useful place.
- ▶ Calculate $21+21$, $7*6$ and $\frac{0,3}{4} * \sqrt{313600}$
- ▶ Is $0.5 - 0.2$ equal to 0.3 ? Is $0.4 - 0.1$ equal to 0.3 ?
- ▶ With the `c` command, create a vector with body sizes of people around you. You can also use the values 1.75, 1.76, 1.83, 1.84, 1.77, 1.76, 1.77, 1.66, 1.86, 1.76
- ▶ What does `3:6` create? What does `YourObject[3:6]` do?
- ▶ What does `YourObject[-4]` do?
- ▶ BONUS (for fast people): Analyze the descriptive statistics: `mean(YourObject)`, `median`, `min`, `max`, `range`, `quantile`
- ▶ BONUS 2: Generate 150 random numbers from a normal distribution with $\mu = 170cm$ and $\sigma = 8cm$. Perform a Kolmogorov-Smirnov test for normality of that sample.

Solutions to Exercise 1: Basic R syntax

```
# simple introductory tasks
21+21 ; 7*6 ; 0.3/4*sqrt(313600)
0.5-0.2 == 0.3 # TRUE
0.4-0.1 == 0.3 # FALSE
print(0.4-0.1, digits=22) # Numerical accuracy limits
all.equal(0.4-0.1, 0.3) # TRUE

size <- c(1.75, 1.76, 1.83, 1.84, 1.77, 1.76, 1.77,
          1.66, 1.86, 1.76)
3:6 # A vector with consecutive integers
size[3:6] # Select the corresponding elements of a vector
size[-4] # Select all but the fourth value
mean(size); median(size); min(size); max(size)
range(size); quantile(size)
x <- rnorm(n=150, m=170, s=8)
ks.test(x, "pnorm", mean(x), sd(x) )
```

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- ▶ BONUS: What arguments for `read.table` seem useful?

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- ▶ `str(YourObject)` must yield the column data types: num, num, factor.
- ▶ BONUS: What arguments for `read.table` seem useful?
- ▶ BONUS 2: What commands are useful to read csv files, excel sheets or dataset with fix column widths?

Solution to Exercise 2: Reading files

```
treesize <- read.table(file="treesize.txt", header=TRUE)
```

header = TRUE

read first line as column names

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treesize <- read.table(file="treesize.txt", header=TRUE)
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dec = ","

comma as decimal mark

Solution to Exercise 2: Reading files

```
treesize <- read.table(file="treesize.txt", header=TRUE)
```

header = TRUE

dec = ","

sep = "_"

read first line as column names

comma as decimal mark

underscore as column separator ("\\t" for tabstop)

Solution to Exercise 2: Reading files

```
treesize <- read.table(file="treesize.txt", header=TRUE)
```

header = TRUE

dec = ","

sep = "_"

fill = T

read first line as column names

comma as decimal mark

underscore as column separator ("\\t" for tabstop)

fill incomplete rows with NAs at the end

Solution to Exercise 2: Reading files

```
treesize <- read.table(file="treesize.txt", header=TRUE)
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dec = ","

sep = "_"

fill = T

skip = 12

read first line as column names

comma as decimal mark

underscore as column separator ("\\t" for tabstop)

fill incomplete rows with NAs at the end

ignore the first 12 lines (eg with meta data)

Solution to Exercise 2: Reading files

```
treesize <- read.table(file="treesize.txt", header=TRUE)
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header = TRUE

dec = ","

sep = "_"

fill = T

skip = 12

comment.char = "%"

read first line as column names

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identify NA entries (missing values)

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Alternatives to read.table:

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Alternatives to read.table:

scan At the core of read.table - for complicated things

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`read.csv` comma separated values (different defaults than read.table)

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header = TRUE	read first line as column names
dec = ","	comma as decimal mark
sep = "\t"	underscore as column separator ("\\t" for tabstop)
fill = T	fill incomplete rows with NAs at the end
skip = 12	ignore the first 12 lines (eg with meta data)
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read_excel Excel files (install package, see github.com/hadley/readxl)

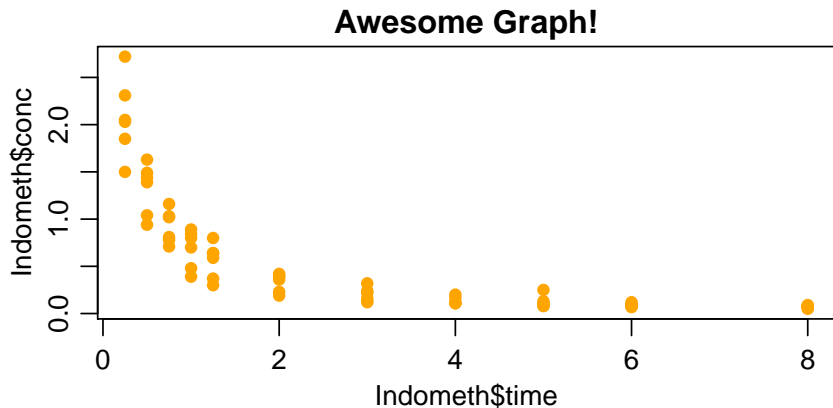
Plotting I

General code for scatterplots: `plot(x, y, ...)`

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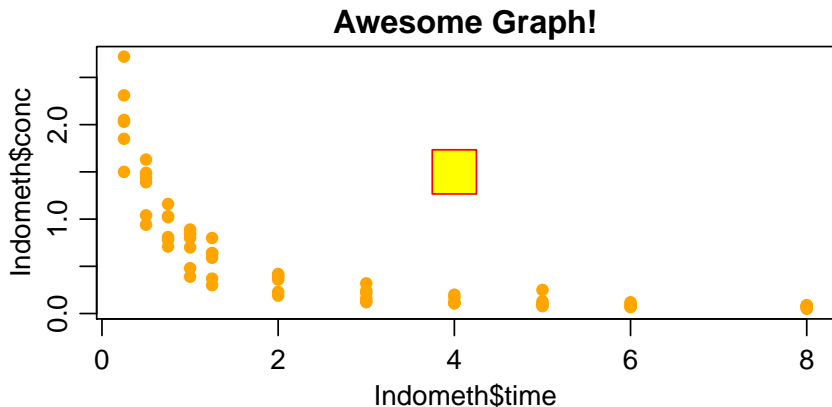
```
plot(x=Indometh$time, y=Indometh$conc,  
     col="orange", pch=16, main="Awesome Graph!")
```



Plotting II

General code for scatterplots: `plot(x, y, ...)`

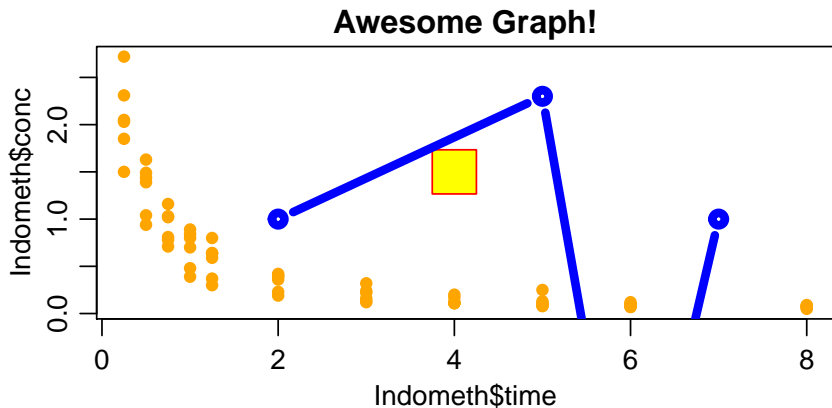
```
points(4, 1.5, pch=22, bg="yellow", cex=4, col="red")  
# PointCharacter, BackGround, Character EXpansion
```



Plotting III

General code for scatterplots: `plot(x, y, ...)`

```
lines(x=c(2,5,6,7), y=c(1,2.3,-3,1),  
      col=4, type="b", lwd=5)
```



Exercise 3: Plotting the treesize dataset

```
treesize <- read.table(file="treesize.txt", header=TRUE)
```

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```
treesize <- read.table(file="treesize.txt", header=TRUE)
```

```
plot(x=xvalues, y=y_values, xlab="nice axis label",  
     main="graph title", las=1)
```

Exercise 3: Plotting the treesize dataset

```
treesize <- read.table(file="treesize.txt", header=TRUE)
```

```
plot(x=xvalues, y=y_values, xlab="nice axis label",  
     main="graph title", las=1)
```

- ▶ Plot tree height over age.
- ▶ Add labels to the plot.
- ▶ Change the point character (`pch`) and color (`col`).
- ▶ BONUS 1: Use a vector for colors, e.g. subset by tree measurement
- ▶ BONUS 2: Compare the histogram (`hist`) of the heights with the `boxplot` and `quantile(x, probs=c(0.1, 0.8))`.

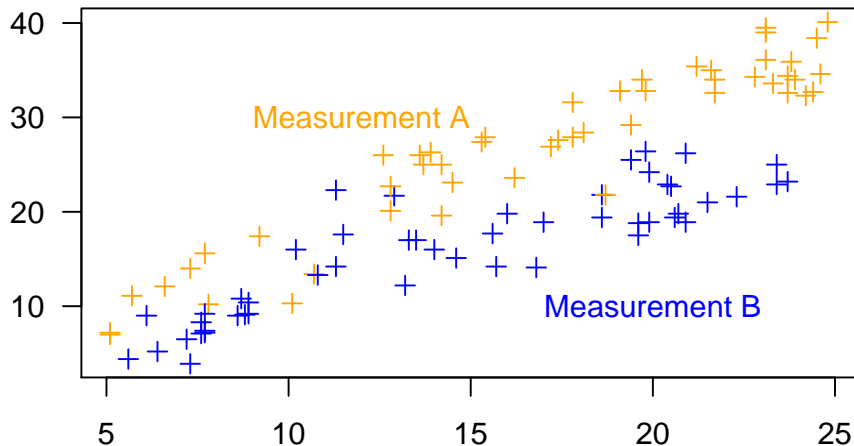
Solution to Exercise 3: Plotting the treesize dataset I

```
treesize <- read.table(file="treesize.txt", header=TRUE)

plot(treesize$age, treesize$height)
cols <- c("orange", "blue")
plot(treesize$age, treesize$height, las=1, ylab="Tree height [m]",
      xlab="Tree age [years]", col=cols[treesize$measurement],
      main="Older trees are larger", pch=3)
text(x=c(12,20), y=c(30,10),
      labels=paste("Measurement", levels(treesize$measurement)), col=cols)
quantile(treesize$height, probs=c(0.1, 0.8))

##    10%    80%
##  8.93 32.36
```

Solution to Exercise 3: Plotting the treesize dataset II

Older trees are larger

Solution to Exercise 3: Plotting the treesize dataset III

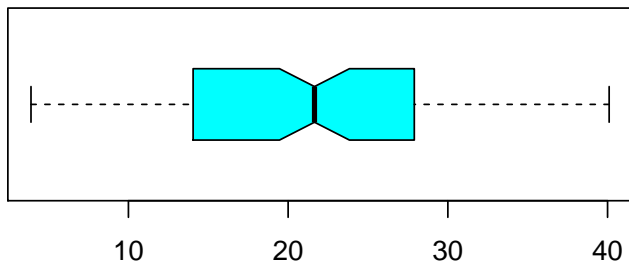
```
hist(treesize$height, col=6, breaks=20, las=1)
```

Histogram of treesize\$height



Solution to Exercise 3: Plotting the treesize dataset IV

```
boxplot(treesize$height, col=5, horizontal=TRUE, notch=TRUE)
```



Commonly needed plot arguments

```
plot(x, y, # point coordinates
col="lightblue", # point color
pch=0, # point character (symbol)
xlab="My label [km]", ylab="", # axis labels
main="Graph title", # title
cex=1.8, # character expansion (symbol size)
type="l", # draw lines instead of points
lwd=3, # line width (thickness of lines)
las=1, # label axis type (axis numbers upright)
xaxt="n" # axis type (none to suppress axis)
)
```

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- ▶ Or just the Rstudio button
- ▶ To make sure your script is reproducible (you may rename objects, for example, and miss one occurrence):
restart R (**CTRL** + **SHIFT** + **F10**) every once in a while (Make sure Rstudio settings are reproducible as shown on slide 4).

Overview: data types

In order of coercion (if mixed, TRUE is converted to 1, 3.14 to "3.14" etc)

Description	example	typeof	class
empty set	NULL	NULL	NULL
not available	NA	logical	logical
logical	c(T, F, FALSE, TRUE)	logical	logical
category	factor("left")	integer	factor
integer number	4:6	integer	integer
decimal	8.7	double	numeric
complex	5+3i	complex	complex
character string	"homer rocks"	character	character
time	Sys.time()	double	POSIXct
date	as.Date("2017-05-02")	double	Date
function	ncol	closure	function

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adv-r.had.co.nz/Data-structures.

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function	<code>ncol</code>	closure	function

adv-r.had.co.nz/Data-structures. `as.character(3.14)` converts a data type;

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character string	"homer rocks"	character	character
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In order of coercion (if mixed, TRUE is converted to 1, 3.14 to "3.14" etc)

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empty set	NULL	NULL	NULL
not available	NA	logical	logical
logical	c(T, F, FALSE, TRUE)	logical	logical
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Object	example	<code>typeof</code>	<code>class</code>
vector	<i>see data types</i>
matrix	<code>matrix(9:15, ncol=2)</code>	...	matrix
array	<code>array(letters[1:24], dim=c(2,6,4))</code>	...	array
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`as.matrix(Object)` converts the class of an object by force.

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- ▶ Rarely needed: `remove.packages("packagename")`

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- ▶ Feed the data into `lm`, assign the output to an object (useful name!).
- ▶ Briefly explain the `summary` of the linear model.

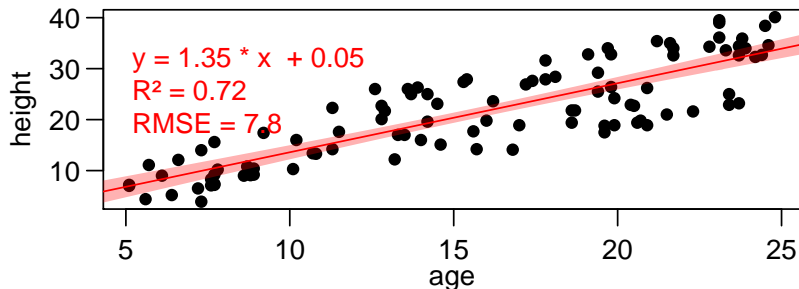
Solution to Exercise 4: Linear regression

```
library("berryFunctions")  
linReg(height~age, data=treesize)  
linReg # nicer: berryFunctions::funSource(linReg)  
browseURL("https://github.com/brry/berryFunctions") # R/linReg.R -> lm  
linear_model <- lm(height~age, data=treesize)  
summary(linear_model)
```

blog.yhathq.com/posts/r-lm-summary.html

stats.stackexchange.com/questions/5135/interpretation-of-rs-lm-output

linear regression of treesize



More things

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- ▶ Data.frames

Objects: data.frames

- ▶ For tables with different data types (numbers, characters, categories, integers), R has the object type `data.frame`:

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data.frame(count=c(2,6,5), type=c("a","k","k"))
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- ▶ `read.table` also returns a `data.frame`
- ▶ If we have the object `df`, we can subset with `df[rows,columns]`
- ▶ `df[1,2:4]; df[2,]; df[, "name"]; df$name`
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From the dataset `treecsize` from the previous exercise, obtain:

- ▶ The first 5 values in column 2
- ▶ The maximum "Height" (the maximum of the values in that column)
- ▶ For each entry: is the measurement equal to (`==`) A?
- ▶ BONUS 1: The height entries for trees older than 23.5 years
- ▶ BONUS 2: All rows, excluding rows 3, 7,8,9,...,20