### Epidemiologic data analysis using R

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#### Contents

- ► Basic properties of R
- Script files
- Data structures and objects
- ▶ Data input and output
- Functions
- ► Tabulation functions

#### What is R

- Statistical software or "package" and a lot more
- R is a languageand environment for statistical computing and graphics (www.r-project.org/)
- Developed by volunteers, coordinated by the R Development Core Team.
- Available for Windows, Linux, Mac, Unix, . . . .
- Is expanding rapidly: new version every 6 months.
- ▶ No licence fee(!) & source code open.

For further information and download: {http://www.r-project.org/}

### Properties of R

- ► Large repertory of basic and advanced methods.
- Versatile graphics of high quality.
- R Reads datasets from Stata, SAS, SPSS, Epi-Info even Excel
- Deals simultaneously with different objects or data structures newline – not just a single data matrix.
- Results of analysis saved as **objects**, readily available for further processing.
- Parsimonious output listing!
- For advanced users! Easy to expand and tailor to specific needs using the **object-oriented**programming tools.

#### To learn more about R

Hills, M., Plummer, M., Carstensen, B. A Short Introduction to R for Epidemiology, 2011. http://bendixcarstensen.com/Epi/R-intro.pdf

- ▶ Dalgaard, P. Introductory Statistics with R, 2nd Ed. Springer, New York, 2008.
- Statistical Practice in Epidemiology Using R. An international course, IARC, Tarto, 2020. http://bendixcarstensen.com/SPE/
- R blog
- Masses of books, articles, websites, etc . . .

## What does R offer for epidemiologists?

- Descriptive tools
  - Versatile tabulation
  - ► High-quality graphics
- Analytic methods
  - Basic epidemiologic statistics
  - Generalized linear models and their extensions
  - Survival analysis methods
  - Other . . .

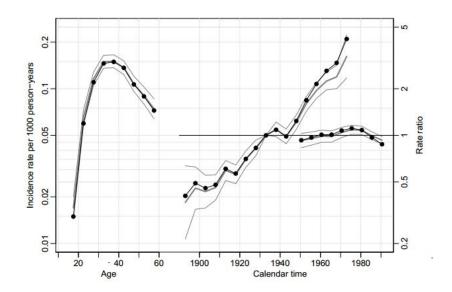
These are provided by SAS and Stata, too, so why R ...?

Many features of R are more appealing in the long run.

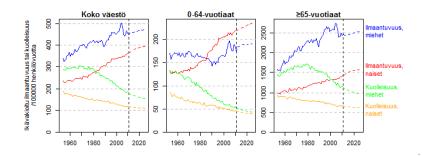
### Graphics in R

- Versatile, flexible, high quality, . . .
- Easy to add items (points, lines, text, legends . . . ) to an existing graph.
- ► Fine tuning of symbols, lines, axes, colours, etc. by graphical parameters (> 67 of them!)
- ▶ Interactive tools using the mouse
  - Put new things on a graph
  - Identify points
- ► Modern lattice or *Trellis* graphics.
- Saving formats: Metafile, .pdf, .png, .bmp, .jpg, . . .

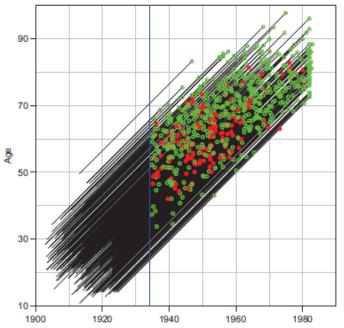
# Age-period-cohort incidence in DK



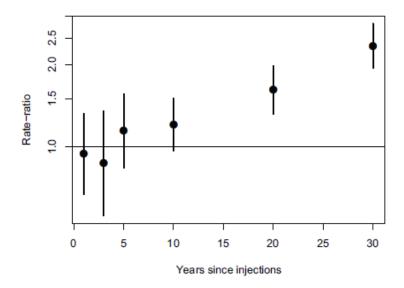
#### Cancer Predictions in Finland



# Follow-up of Welsh nickel cohort in Lexis diagram



# RRs & CIs by exposure in a cohort study



# Package or library

- ► Collection of functions pertaining to some specialized application area, e.g. survival, boot
- Contributed by users of R.
- Available after loading:library(survival)
- Alternatively load from the menu bar: Packages - Load package... - Select one
- New versions easily updated from Internet. (https://www.rdocumentation.org/trends)

### R script - R Studio - commands in a file

**R** script file is an ASCII file containing a sequence of R commands to be executed.

The **script editor** – use R-Studio

▶ In R-Studio open the script editor window: New file - R script, or when editing an existing script file: File - Recent Files,

- Save the script file: Save e.g. or Save As \*.R
- Excecute a line Ctrl-Enter

# R script (cont'd)}

▶ Paint the lines to be excecuted and Ctrl -Enter will execute lines.

- To run a whole script file, write in console window:
  - > source("c:/.../mycmds.R", echo=TRUE)

► The script can also be written and edited by any external editor programs (like Notepad).

### Data objects of different kinds

- vector: ordered set of similar elementse.g. real numbers or character sequences,
- factor: categorical variable with levels
  e.g. gender, levels: c(1,2) or c('male', 'female');
- matrix, array: 2- and k-dimensional tables,
- data.frame: "data matrix" (more of this soon!),
- ts: time series object,
- list: sequence of different types of objects.

### Attributes of data objects

Functions that extract some key properties of objects:

- length(): number of elements,
- mode(): basic type of elements,
- dim(): dimensions of arrays, matrices and data frames,
- str(): overall structure,
- class(): property that determines how certain generic functions (e.g. summary(); plot()) work when the object is given as argument.

#### Data frame – data matrix

- ▶ [Data frame] = a **list** of column vectors
- Rows correspond to observational units, and columns (same length) refer to variables.
- Column vectors can be numeric, character or logical
- Columns are subobjects of the data frame. Their names are not directly accessible. Two possibilities:
  - Use "surname\$firstname", e.g. mydata\$var1 ,
  - Place the data frame in the search path at position 2: attach(mydata); then use just "firstname": var1

## Data frame import from external files

- Common ASCII files, for example: read.table("C:/owndir/rfiles/mydata.txt", ...); read.table("http://cc.oulu.fi/~tilel/esan.txt",...)
- ► Files with fixed-width format: read.fwf();
- Files created in SPSS, SAS, Stata etc.: functions read.spss(), read.ssd(), read.dta(), etc. in package foreign,
- Excel-files: either {read.table("clipboard", ...)}, or
  (1) save the Excel-file in .csv or .txt format,
  (2) in R: read.csv2() or read.table()
- Relational DBMSs: several R packages available.

## Data frame import from external files with R

#### Choose Import Datasets

- ▶ from text (base) for **text** files
- ► from text (readr) for csv files
- from excel for excel files
- from SPSS for spss files
- from SAS for sas files
- ► from STATA for stata files

### Dealing with output

- ► The console contents, *i.e.* the flow of input commands and output results from the console window, can be
  - \$ \$ printed on paper: File Print...
  - \$ \$ saved to an ASCII file: File Save to file...
- Selected parts can be copied from the console and pasted to an external file.
- ► Function sink("results.txt") diverts all subsequent output to an external text file. Back to console: sink().
- ► Choose *New File R Markdown* output to MS-Word
- Graphs saved in desired format: File Save. . .

### R is a functional language

Most computations in R involve the {invocation} or {call} of functions. They are called by name with a set of arguments separated by commas, e.g. fun(x, y, z); Function

- = sequence of rules on how to produce desired output:
- ▶ value of the function, from given input, i.e.
- **arguments** of the function.

Example: Function sqrt() computes square roots:

argument vector defined

$$x \leftarrow c(0,1,2,3,4)$$

call with argument x

```
sqrt(x)
```

[1] 0.000000 1.000000 1.414214 1.732051 2.000000

# Defining a new function (1)

*Example.* Function CIapp to calculate an approximate confidence interval from point estimate (estim) and std error (SE) by formula estim  $\pm z_{\gamma/2} \times$  SE.

Defining code (without prompts):

```
Clapp <- function(estim, SE, level = 0.95)
  {z <- qnorm(1- (1-level)/2 ) # setting the quantile
lower <- estim - z*SE
upper <- estim + z*SE
Clapp <- c(lower, upper)
Clapp }</pre>
```

▶ Formal arguments, here estim, SE, level

# Calling the new function (1)

▶ Actual arguments, used in function call:

```
CIapp(3, 1, 0.9) #90\% limits: $3 \pm 1.645\times 1$
```

[1] 1.355146 4.644854

NB! Positional matching: order of actual arguments.

- **Keyword matching**: the order of arguments in the call is irrelevant if the names of formal arguments are given

```
CIapp(SE=1.0, level=0.90, estim=3)
```

- [1] 1.355146 4.644854
  - ► If a **default value** for an argument is given in the definition and is OK, it can be omitted in calling

```
Clapp(3, 1) # 95\% limits: $3 \pm 1.96 \times 1$
```

### Function call & value object

- Simple call: Evaluates the value of the function with given arguments and prints value items (according to the print method specific to the class of the value object).
- ► Call of function and assignment of its value to an object.

To extract information & items from the value object, e.g.

- {str()}: overall structure,
- {names()}: names of the components,
- fprint()): selective printing of value items,
- \$\ \summary()\}: selective print (not available for all functions). % - \{plot()\}: certain graphical display, -"-

## Example, function {range()}}

Returns the minimum and maximum values of a data vector.

```
{> v <- c(15.3, 10.8, 8.1, 19.5, 5.3) #} data vector
{> range(y) #} simple call with argument {y}
\{[1] 5.3 19.5\}
{> ra <- range(y) #} call with assignment of value
{> ra #} or {print(ra)}, equivalent to simple
call
\{[1] 5.3 19.5\}
{> str(ra) #} structure of the value object
{num [1:2] 5.3 19.5}
{> ra[1] #} extracting an item from the value
object
{[1] 5.3}
```

#### Different kinds of functions

- Mathematical, e.g. {sqrt(x); log(x); exp(x)}. Arguments and values typically numeric vectors.
- Data handling, e.g.
  {dafr <- data.frame(x, y);
   adata <- read.table("a.dat", header=T, ...);
   % redc1 <- subset(redc, group == "24 h");}
  Main argument(s): data object(s). Value: data object.</pre>
- ▶ Graphical, e.g.
  {plot(y ~ x); stripchart(y, xlim=c(0,3))}
  Main argument(s): data object(s). Value:
  graph.
  Ancillary arguments: e.g. graphical parameters.

#### Value of the function

- numeric object (e.g. vector, matrix) for many mathematical and statistical functions,
- data object (e.g. vector, data frame) for data handling functions,
- graph for graphical functions,
- table for tabulating functions,
- \*\*list} = a sequence of objects of different kinds, for many statistical functions.

#### Statistical functions

- ► {Main} argument(s): Typically data object(s).

  Often a {model formula} like {y ~ x} with

  y representing the response variable and
  expression {x} = explanatory variable(s) or
  factor(s).
- Ancillary arguments or parameters: additional specifications. Some default values usually offered for these.
- ► {Value}: Usually a {list} object consisting of several components of different types. %— Note however graphics functions.
- % Example (ks. vanha luentomoniste, t.test)

### Function values as list objects

- List = object consisting of an ordered collection of component objects, maybe of different types.
- Provides a convenient way to return the results of statistical computation.
- A list with named components formed from existing objects:
   \${ } \$ {Lista <- list(name=obj1,title=obj2,addr=obj3)}
  A single component identified:
   {Lista{\$}name};</pre>
- Concatenation of several lists into one: {longlist <- c(list1, list2, ...)}.</pre>

# Ex: Function *t.test()*

```
Description of syntax in the {help()} page Default S3 method:
t.test(x, y = NULL, alternative = c("two.sided", "less",
"greater"), mu = 0, paired = FALSE, var.equal = FALSE,
conf.level = 0.95, ...)
t.test(formula, data, subset, na.action, ...)
\begin{itemize} - Main argument(s): data vector(s)
{x} (and {y}) or formula - Ancillary arguments, like
{var.equal, conf.level}:
Default values given. - **NB.} Dots '{...}'
```

### Example. Red cell folate levels

The data describe red cell folate levels (variable folate,  $\mu g/I$ ) in two groups

of cardiac bypass surgery patients given two different nitrous oxide ventilation ( $50\% \text{ NO} + 50\% \text{ O}_2$ ) treatments (variable group):

- ▶ group 1  $(n_1 = 8)$  continuously for 24 h (label "24 h"),
- ▶ group 2  $(n_2 = 9)$  only during the operation ("oper").

Observed folate levels in the two groups: > folate[group=="24 h"] [1] 243 251 275 292 347 354 380 392 > folate[group=="oper"] [1] 206 210 226 249 255 273 285 295 309

# Ex: Call of {t.test()} by formula argument

t.test(folate ~ group, var.equal=TRUE, conf.level=0.9)

#### Output:

Two Sample t-test

data: folate by group t = 2.5653, df = 15, p-value = 0.02153

alternative hypothesis: true difference in means is not equal to  $\boldsymbol{0}$ 

90 percent confidence interval: 19.09502 101.51610

sample estimates: mean in group 24 h mean in group oper 316.7500 256.4444

## Ex: Value returned by {t.test()} is a list}

Function value assigned to an object and examined: > tfol <- t.test(folate  $\sim$  group, var.equal=TRUE, + conf.level=0.9) > str(tfol) # The structure of the object

```
{List of 9 $ statistic : Named num 2.57 ..- attr(, "names")= chr "t" $ parameter : Named num 15 ..- attr(, "names")= chr "df" $ p.value : num 0.0215 $ conf.int : atomic [1:2] 19.1 101.5 ..- attr(, "conf.level")= num 0.9 $ estimate : Named num [1:2] 317 256 ..- attr(, "names")= chr [1:2] "mean in group 24 h" "mean in group oper" $ null.value : Named num 0 ..- attr(, "names")= chr "difference in means" $ alternative: chr "two.sided" $ method : chr " Two Sample t-test" $ data.name : chr "folate by group" - attr(, "class")= chr "htest"
```

names(tfol) [1] "statistic" "parameter" "p.value" "conf.int" "estimate" [6] "null.value" "alternative" "method" "data.name"

# Ex: Value of {t.test()} utilized}

► Extracting items for further processing: > tfol\$estimate # contents of the 'estimate' component mean in group 24 h mean in group oper 316.7500 256.4444

▶ Utilizing the component value in further calculations: mean.diff <- tfolestimate[1] - tfolestimate[2] - Item names in the parent object "inherited". Can be renamed: names(mean.diff) <- c("Mean difference"); mean.diff Mean difference 60 30556

# Defining a new function (2)}

We now create a new function {T.estimCI()}. It will return only the mean difference between the groups (which is not reported by {t.test()}!) and its confidence interval.

The function is defined as follows:

```
T.estimCI <- function(x, ... ) { tt <- t.test(x,
...) mean.diff <- ttestimate[1] - ttestimate[2]
names(mean.diff) <- c("Mean difference") T.estimCI <-
list(Meandiff = mean.diff, Conflimits = tt$conf.int)
T.estimCI }</pre>
```

# Calling the new function (2)

```
When {t.estimCI()} is called, a list with 2 named components is returned and printed: > T.estimCI(folate ~ group, var.equal=T, conf.level=0.9)

$Meandiff Mean difference 60.30556 $Conflimits [1]
19.09502 101.51610 attr(,"conf.level") [1] 0.90
```

## Dealing with functions

- Defining code can (mostly) be viewed by typing the function name without parentheses and arguments.
- ► Functions can be saved into a separate script or source file, {e.g.} {myfuns.R}, which may contain several functions.
- Source file accessible in an R run after
  {> source("C:/.../myfuns.R") }
- ▶ Alternatively from menu bar: {File Source R code ...}
- Loading from Internet:
  {> source("http://.../myfuns.R")}

#### Tabulation functions

- ▶ {table(c1, c2)}: simple contingency tables
- ▶ {xtabs()}: more elaborate tabulation features
- fftable(c1, c2, c3)}: "flat" contingency tables
  % {apply()} for e.g. calculating margins in a
  cont. table % {sweep()} for e.g. calculating
  percentages % in table cells
- {tapply(var,fac,fun)} tabulates values of function {fun()} (for example {mean()}) applied to values of variable {var} in categories of factor {fac},

in package {Epi} for more informative tabulation.

- package plyr and ddply-funtion ...
- ▶ package data.table for BIG data ...
- missing variables...
- ▶ other ...

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