Preamble

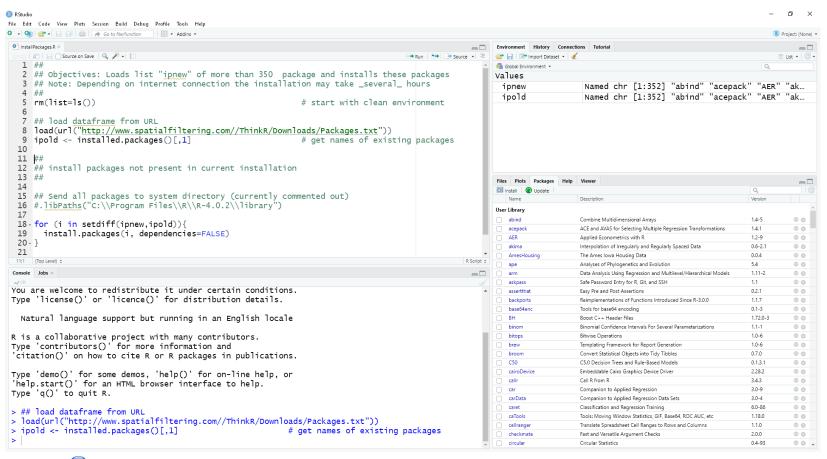
- An excellent website outlining data operations and basic data analysis with @ can be found at http://www.statmethods.net. It makes frequent references to the book by Kabacoff.
- Explore the \(\Pi \)-Project website: \(\frac{https://www.r-project.org/}{} \)
- Explore Microsoft's Open @ website: http://mran.microsoft.com/
- Explore RStudio's website: https://www.rstudio.com/

Reproducible Research

- Rather it mainly uses commands (i.e., functions), which can be pooled together into scripts.
- The disadvantage of this approach is a higher learning effort to know the commands for your specific task.
- A substantial advantage of this approach is that you or anyone else at any time can exactly reproduce and check what you were doing.
 - Thus is strongly embedded in the *academic paradigm of* REPRODUCIBLE RESEARCH.
- Furthermore, by analyzing your scripts instructors and teaching assistants can efficiently help you.

The RStudio Environment

• Discuss the Studio interface structure: EDITOR – CONSOLE – ENVIRONMENT/HISTORY/CONNECTIONS – FILES/PLOTS/PACKAGES/HELP/VIEWER



- Explore Studio's Tools and Help tab and 's help system.
- Concept of environment, history file and script. The associate file extensions are:
 - o *.RData or *.rda: Copy of the workspace with all its data objects and custom user functions.

- *.Rhistory contains all commands that have been issued during a session at the command prompt
 >.
- ★.R is a file that contains scripts, which can be a set of basic data analysis commands, individual functions, or an elaborate program
- Get and change the *working directory* were your scripts and workspace are stored and searched by default.

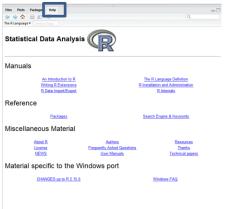
 Enter the command <code>getwd()</code> into the editor window and select "Run" from Your current working directory is displayed in the <code>CONSOLE</code> below

 Each command in an ends with parentheses, e.g., <code>getwd()</code>, which may include optional parameters. Even if no options are specified a function always ends with parentheses ()

 Comments:
 - The string "C:/Users/Michael/Documents" is the path to the working directory. In ♥ strings are always enclosed by quotation marks, i.e, "...".
 The WINDOWS convention is to separate sub-directories by a backward slash \.
 However, ♥ uses the forward slash / or alternatively a double backward slash \\.
 In ♥ the single backslash \ is reserved to start an escape character, for instance, \n becomes a line break and carriage return in text output, e.g., > cat("First line\n Second line") is shown in the CONSOLE on two lines.
 - The character > in the **CONSOLE** window is the command prompt which indicates that is ready to receive new commands. It shows up when completed executing a script.
 - o The **Esc** key or pressing

 in the **Console** window − available while a script is executing − can terminate the ongoing execution of a script.

- Receiving help at the command prompt > in the CONSOLE:
 - > help (FunctionName) or short > ?FunctionName
 This searches all active libraries (libraries currently linked to your session) for the help.
 - Or through the HTML-help menu system:



Fuzzy help can be obtained with the double question mark > ??PartName. This searches all installed libraries for the help on functions, data or vignettes containing the string "PartName".

Basic Mechanics

Make your Working Directory

• My suggestion is that you setup a specific directory on your hard- or jump-drive for this course, e.g.,

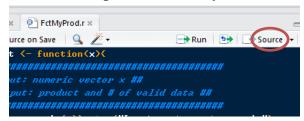


You store your scripts, data etc. in this directory or its subdirectories.

- Then set your working directory to this location with the command
 - > setwd("E:\\Lectures2020\\WorkingWithR").
- Notice the double backward slash \\ here as substitute for the single forward slash / and the use of the quotation "..." to enclose the directory string.

Interacting with the R-Console

- Collections of commands (or programs) can be stored in external *.R script-files (see FILE menu)
- Single commands in the script editor can be run from the command prompt or with **RUN** for a highlighted line in a script.
- To run a script (all lines in your editor) use the **SOURCE** button:



- In the **CONSOLE** window the arrow **UP** and **DOWN** keys scrolls through the history of previously issued © commands.
- Previously issued commands can be edited by moving with the arrow keys the edit prompt to a particular location of your command line.
- While being in the **CONSOLE** window, commands with their list of parameters can be broken over several lines. Then the continuing prompt "+" will be displayed *automatically* at the beginning of a new line until the command is completed.

Variables

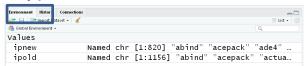
• The assignment operator to a variable is the backward arrow "<-", i.e., it requires two key-strokes, > my.name <- "Michael" (The variable my.name will now show up in the ENVIRONMENT) with the content "Michael")

Note: The backward arrow <- is preferred over the equal sign = , which is reserved for parameter assignments in function calls.

• Variable names:

- o Names must begin with a letter and can consist of an alpha/numeric combination of letters including the period "." and/or the underscore " "
- O Note: Specific characters and keywords such "\$" "@" "&" and "%" or "+" "-" "*" /" and "^" cannot be used because they have special meanings. For a full list see > ?Reserved
- Warning: Variable and function names in are case sensitive, e.g., my. Var and my. var are different objects (\Rightarrow this can lead to typos while writing a script. These are extremely difficult to spot)
- Tip: Name variables properly so an external reader or you, after a few weeks have passed, can understand what you were doing
- O Use the dot to structure associated variable names, e.g., sales.plano and sales.dallas, or the camelback convention salesPlano and salesDallas
- o Just for experts: The document GOOGLE-R-STYLE.PDF. suggests professional naming and typesetting conventions of your @ code.

• Any *function* or *data structure* that is defined during a @-session becomes an object in the **ENVIRONMENT**



- These can be removed from the **Environment** with the remove function
 - > rm(ipold)
- <u>Warning:</u> if you happen to name a variable or function identically to an existing @ object, which already exists in the *search path* of your session, therefore, that object will be masked and is no longer directly accessible:

```
> pi  # gives the system constant 3.141593
> pi <- 2.71  # this masks the system constant pi
> base::pi  # pi still be found in its library base. Note the "::"
> rm(pi)  # removes user's pi and makes the constant pi available
```

- Some hard-wired values in @ are:
 - o *Logical* values **T** and **F** (alternatively **TRUE** and **FALSE** can be used)
 - o An object without content has the value **NULL**
 - o *Impossible* operations, such as log(-1), lead to a *not-a-number* NaN.
 - o *Missing* numbers have the value **NA** which stands for not available.
 - o Some *predefined* numbers are infinity Inf and pi (that is, $\pi = 3.141593$)

Data Representation in @

Data-Sets

For statistical analyses data-sets are usually arranged in rectangular data-frames and imported from external files (such as SPSS, STATA, EXCEL or DBASE) or embedded in workspaces (with the extension *.RData) of libraries.

For instance, to read an SPSS file use:

- > library(foreign) # makes import functions available during a session
- > setwd("E:\\Lectures2020\\WorkingWithR")
- > MyPower <- read.spss("DallasTempPower.sav", to.data.frame=T)</pre>
- Some data file types can also be imported using FILE ► IMPORT DATASET or with Import Dataset ▼ in the ENVIRONMENT window.
- Notice that the object MyPower is added as data-frame to your Environment.
- Check the class of an object use
 - > class(MyPower)
 - [1] "data.frame"
- To preview the data-frame double click on it in the **Environment**.



- Each column in a data-frame has an associated elementary data type.
- Some 🖫 libraries also include their own data-frame. These can be opened by the command
 - > data("CPS1985", package="AER")

Elementary Data Types

- Within a data-frame the variables can be of an elementary @ data types:
 - o logical: FALSE or TRUE (also binary 0 or 1), e.g., am.I.logical <- TRUE
 - o character strings: always enclosed by single or double quotation marks,
 - $e.g., {\tt my.name} < {\tt -} "Michael".$
 - o <u>factors:</u> Internally each value is stored with a specific integer numbers. Each integer has a descriptive label assigned to it. When listed, factor values are not enclosed within quotation marks. For instance:
 - > MyPower\$Month
 - [1] JAN FEB MAR APR MAY JUN JUL AUG SEP OCT NOV DEC
 - [13] JAN FEB MAR APR MAY JUN JUL AUG SEP OCT NOV DEC
 - [25] JAN FEB MAR APR MAY JUN JUL AUG

Levels: JAN FEB MAR APR MAY JUN JUL AUG SEP OCT NOV DEC

- o <u>numeric</u>: either integer, double precision or complex numbers, which are not used in this course.
 - True integer values can be enforced with the added symbol "L"
 - > two <- 2L.
 - > typeof(two)
 - [1] "integer"

 Integers, under specific circumstance, are treated as real numbers or may be *coerced* to real numbers

```
> two.sq <- two * 2.0
> typeof(two.sq)
[1] "double"
```

Very small remainders after operations are interpreted as zeros, such as

[1] 4.440892e-16.

This discrepancy is due to *rounding errors* associated with floating point computations, i.e. taking the square root and squaring the remaining results. This problem applies to all software environments and computer chips dealing with rational numbers.

Rational numbers with and infinite number of digits can only be *digitally approximated* due to the limited bit-depth implemented in operating systems.

E.g., while the constant $\pi = 3.14$... has an infinite number of digits, only the first 16 can be numerically represented:

```
> options(digits=20)
> pi
[1] 3.1415926535897931
```

o <u>Dates:</u> Dates come in different formats. This course will *not* cover dates.

Basic Data Objects in @:

o scalar: an individual datum. All data objects are composed of a collection of scalars.

- o vector: *atomic data structure* that collects more than one value of *identical type*. Example:
 - > score <- c(23,53,45,30,53,60) or
 - > catName <- c("Austin", "Gretchen", "Charlie")</pre>

where the function c () concatenates – combines –several scalars into a vector.

> length (score) gives the number of elements in the vector.

Note: if a printed vector stretches over several rows in the console the position of the first value in each row will be numbered by [elementNumber]

Mixture of scalars with different data types will be *coerced* into characters

- > mixture <- c(1,2,3,T,pi,"A")
- > mixture

```
[1] "1" "2" "3" "TRUE" "3.14159265358979" "A"
```

- o <u>matrix</u>: two-dimensional arrangement of a set of vectors that are all of the *same data type and length*.
- o data frames: collection of vectors of the same length but different data types per column are allowed.
- o <u>list:</u> collection of vectors of any type and length.
- To obtain information about a data object and to look at its structure use the function str():
 - > str(MyPower)

```
'data.frame': 32 obs. of 8 variables:
$ SeqID : num 1 2 3 4 5 6 7 8 9 10 ...
$ Year : num 2009 2009 2009 2009 2009 ...
$ Month : Factor w/ 12 levels "JAN", "FEB", "MAR", ...: 1 2 3 4 5 6 7...
$ DaysBill: num 34 29 30 32 29 30 32 29 30 31 ...
- attr(*, "codepage") = int 1252
```

Object Oriented Philosophy

- All data objects have an *object class* assigned to them. This may be fairly rudimentary, such as being a numeric vector, or highly advanced, such as the output from a regression model.
- All © commands are *implemented as functions* with parentheses at the end of the function name. A function may not use [a] specific input arguments, e.g., the function <code>getwd()</code>, or [b] a selection of arguments, [c] some of these arguments may have default values and only need to be issued when they are overwritten, and [d] some arguments are positional (without a leading keyword) or are introduced by a specific keyword.

The general syntax of a function is:

```
ResultObject<sub>Class</sub> <- function(List of InputArguments<sub>Class</sub>)
```

• Depending on the class of the input arguments, an object-specific function will be used.

Example notice amorph behavior of the **summary** () function:

```
> summary(dataframe)  # Get summary statistics for all variables
> regObject <- lm(...)  # run a regression model and store in regObject
> summary(regObject)  # view key results of the regression analysis
```

• If just the object name is entered at the command prompt its default print method will be used. E.g.,

```
> regObject  # print content of regObject on console
```

Saving Analysis Results, Plots and Data

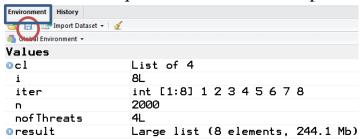
• Analysis Results and Plots: The easiest way of saving [a] **CONSOLE** output and [b] generated plots in the graphics window is to *copy and past* them into a graphically enhanced text editor such as **WORD**.

Important: Any text output needs to be typeset in a <u>fix pitch font</u> such as Courier New.

Otherwise the formatting of the output will be lost, e.g., columns of a matrix will not line up properly.

Perhaps reduce the font size and line-spacing of imported output as well as switch to a landscape layout.

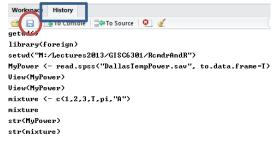
• <u>Data:</u> The collection of all variables, data-frames and functions, which were created during a session, can be saved for subsequent sessions in a workspace



Before saving the workspace non-desired data-objects, variables and functions should be dropped with the remove command:

> rm(objectName).

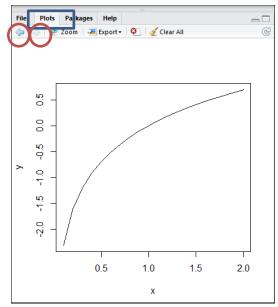
- Data-frames within the workspace can also be *exported* into different file formats (see the package **foreign** or connections to **SQL** servers).
- Command History: All commands, which were issued during session, can be saved into a history file



Working with some functions

• Exercise:

• Select in the Plots window scroll through your list of plots with the arrow icons:



• Exercise (cont.):

```
> z <- rnorm(length(x)) # vector of standard normal random numbers
> mat <- cbind(x,y,z) # merge vectors of same length into a matrix
> dim(mat) # see the dimensions of the matrix
> summary(mat) # get statistics of each matrix column
> class(mat) # evaluate object type
[1] "matrix"
> df <- data.frame(x=x, xlog=y, rand=z) # build dataframe
> class(df)
[1] "data.frame"
```

Working with Data

Data-Frames

- Data-frames can pool several vectors of *same length* but potentially of *different data-types* together.
- Almost all statistical analysis functions are defined on data-frames.

- The variables catName and catAge are stored now in the data frame my.data under new names Name and Age
- To access individual variables in the data-frame several commands can be used

```
> my.data$Name
> my.data["Name"]
> my.data[1]
> with(my.data, Name)
> my.data[ , "Name"]
> my.data[ , 1]
```

List Objects

- Note: © functions can only return one data object. However, different data object can be bundled into a list and returned by the function.
- List objects allow to link data objects of *different types* and *different length* together into a container:

```
> A <- matrix(c(1,2,3,4,5,6),2,3)  # 2 x 3 matrix
> my.list <- list(name = catName, age = catAge, mat = A)
> my.list
$name
[1] "Austin" "Gretchen" "Charlie"
$age
[1] 9 10 20
```

\$mat

```
[,1] [,2] [,3]
[1,] 1 3 5
[2,] 2 4 6
```

• Individual objects of the list can be addressed either by

```
> my.list$name
[1] "Austin" "Gretchen" "Charlie"
> my.list[[1]]
[1] "Austin" "Gretchen" "Charlie"
```

• Individual elements of an object in a list can be addressed by

```
> my.list$name[1]
[1] "Austin"
```

• To delete an object in a list assign the **NULL** value:

```
> my.list$mat <- NULL
```

• To get information about an object use the attributes function:

```
> attributes(my.list)
$names
[1] "name" "age"
```

• Remove the objects my.list and the matrix A from the workspace

```
> rm(my.list,A)
```

Matrix Objects

• Matrices can store vectors of *same length* and *same data-type* in an rectangular arrangement

- To generate a matrix:
 - O Vector of 12 elements:

```
> b < -c(10,20,30,40,15,25,35,45,1,2,3,4)
```

• Place elements into 4x3 matrix:

```
> mat <- matrix(b, nrow=4, ncol=3)</pre>
```

• One element at location row and col mat[row, col]

A sequence of values mat[1:2,] (here the first and second row)

Exclusion of elements mat[-1,] (here the first row)

- One row mat[1,] or one column mat[,2]
- Also logical operations are permitted (here the first and second column):

```
> select <- c(T,T,F)</pre>
```

> mat[,select]