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Basic R Lookup Script

```
# ----- header -----
## Introduction to R programming
# Author: Johannes Schielein
# Date: 2016-04-13
## Contents of the script:
# (1) Starting a new session
# (2) Installing and Activating New Packages
# (3) Getting Help
# (4) Object Types and Simple Operations
# (5) Indexing Specific Parts of Data
# (6) Importing, Exporting
# (7) Functions
# (8) Control Structures (loops, if-else)
# (9) Plotting
# (10) Closing a Session
# Additional Remarks and Data Sources

#----- (1) Starting a new session -----

## create a new working directory for R and your R projects (do this only once!)
# Note: Generally avoid using whitespace and special characters, use only lowercase letters!
# You can do this step as well outside R

dir.create("/home/jschielein/R/projects/intro-classes")
dir.create("/home/jschielein/R/projects/intro-classes/input")
dir.create("/home/jschielein/R/projects/intro-classes/output")
dir.create("/home/jschielein/R/projects/intro-classes/scripts")
dir.create("/home/jschielein/R/projects/intro-classes/rdata")

## set your workspace to the new directory
setwd("/home/jschielein/R/projects/intro-classes/")
# check if it worked
getwd()

## clean up your environment and memory
rm(list=ls()) # remove eventuall remaining objects from the global environment
gc(reset=TRUE) # free the memory from eventuall load

## load your workspace (Only if previously saved at the end of the session)
load("rdata/lookup-script-2016.Rdata")

#----- (2) Installing and Activating New Packages -----
## installing a new package (only once). You can also do this in R-Studio under the tab "Packages".
install.packages("ggplot2")
install.packages("xlsx")
install.packages("highlight")

## installing packages from source (if they can't be installed due to compatability issues)
# First locate and download the package source on http://cran.r-project.org/web/packages/
# For windows use the Windows binaries and locate the download link under "r-release" (ZIP file)
install.packages("/home/jschielein/R/packages-source/raster_2.3-24.zip", repos=NULL, type="source")

# activate the package and its functions for the current session (doe this every session that you n
library("ggplot2")
library("xlsx") # known java issues on windows 7 can be solved by installing and activating xlsx,
library("highlight")

#----- (3) Getting Help -----
# show help for any function
help(boxplot)
# alternativly
?boxplot

# note: for further tips on how to solve problems refer to the handout of the class!
```

Basic R Lookup Script

```
#----- (4) Object Types and Simple Operations -----  
## (4.1) Use of Simple Mathematical Operators and object creation  
3*(3+1)-6/3  
# create a new object from this operation  
result.example<-3*(3+1)-6/3  
  
# show the result in the command prompt  
result.example  
  
# we can further operate on our object  
result.example + 4  
  
### (4.2) Vectors  
# are a combination of observations that can be concatenated e.g. like this  
vector.example<-c(1,2,3,4,5)  
  
# show it  
vector.example  
  
# R operates on vectors  
vector.example*5  
  
# matrices are a combination of several vectors (as columns) -> details see below  
  
# (4.3) Different data types  
# (4.3.1) numerical data  
# integer (full rounded numbers)  
integer.vector.example<-1:5  
  
# double precision numbers  
double.vector.example<-c(1.32,2.54,3.72,4.89,5.01)  
  
# How numeric numbers are stored influences processing speed.  
  
## (4.3.2) Character Strings and Factor (=Categorical) data  
# simple character string  
string.example<-"Fischer Fritz fischt frische Fische" # thats a german tongue twister by the way!  
  
# several strings in a vector  
string.vector.example<-c("R is great! I love it",  
                          "R seems like chinese to mee!",  
                          "R? Isn't that the letter before S?",  
                          "I want an ice-cream!!",  
                          "I want an ice-cream!!",  
                          "I use R every day. I even write my emails in R!")  
  
# show it in the console  
string.vector.example  
  
# convert the string data to factor data/categorical data  
factor.example<-as.factor(string.vector.example)  
# show it  
factor.example  
  
# show the unique factor levels/ categories  
levels(factor.example)  
  
# Important: if you work with categorical data also see the sidetopic 4.5.1 below  
  
## (4.3.4) Logical data  
logical.vector.example<-double.vector.example>2 # which elements of the numeric vector are greater  
# show it  
logical.vector.example  
# note: ">" is the logical operator for greather than. Other important ones are: >=,<,<=,!=,&,&
```

```

## (4.3.5) Missing Values
# are denominated by NA
missings.example<-c(1,2,3,NA,5,6,7,NA,9,10)

# if we want to operate on the string we might remove the missings in some cases for the operation
mean(missings.example) # may not be the desired result
mean(missings.example,na.rm=TRUE) # works!

## (4.4) Matrices and dataframes
## (4.4.1) Matrices
# first we create a vector ranging from 1 to 12
vector.2<-1:12

# from this we can create a simple matrix where we split up the vector in four columns
matrix.example<-matrix(vector.2, ncol = 4)
# have a look
matrix.example

# we can assign column names to the matrix (e.g. variable names)
colnames(matrix.example)<-c("variable.a","variable.b","variable.c","variable.d")
# Note: matrices work only with numeric values.
# if you need to work with other data types (e.g. categorical data) use dataframes

# we can assign rownames as well (e.g. observations)
rownames(matrix.example)<-c("Peter","Gabriela","Fritz")

# show it
matrix.example

## (4.4.2) Dataframes
# we convert our matrix to a dataframe like this
dataframe.example<-as.data.frame(matrix.example)

# another way to create a data.frame from scratch is to construct vectors and combine them with the
dataframe.example.scratch<-data.frame(
  variable.1=1:10,
  variable.2=c("a","b","c","d","e","f","g","h","i","j"),
  variable.3=rnorm(10)) # rnorm creates randomized normally distributed numbers between 0 and 1, in

# show it
dataframe.example.scratch

# add a new COLUMN the easy way
dataframe.example.scratch$variable.4<-1:10

# show it
dataframe.example.scratch

# add a new COLUMN with cbind (works for multiple columns from e.g. two different matrices)
dataframe.example.scratch<-cbind(
  dataframe.example.scratch,
  variable.5=11:20) # bind a column do the dataframe.

# show it
dataframe.example.scratch

# add a new ROW to the dataframe
dataframe.example.scratch<-rbind(
  dataframe.example.scratch,
  c(11,"a",0.5,11,21)
)

# show it
dataframe.example.scratch

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# Rotate/transpose a dataframe
dataframe.example.transposed<-t(dataframe.example.scratch)

# have a look
dataframe.example.transposed

## (4.4.3) Side Topic: Adding new categories/ Factor Level Creation.
# lets say we want to add a new category inserting data like this:
dataframe.example.scratch<-rbind(
  dataframe.example.scratch,
  c(11, "xyz", 0.5, 11, 21)
)
# doesnt work instead of the new observation xyz a missing value was introduced because the categor
# show the unique factor levels/ categories
levels(dataframe.example.scratch$variable.2)
# add the new category as a new factor level
levels(dataframe.example.scratch$variable.2)<-c(
  levels(dataframe.example.scratch$variable.2),
  "xyz")
# add a new entry to the vector that corresponds to this category
dataframe.example.scratch<-rbind(
  dataframe.example.scratch,
  c(11, "xyz", 0.5, 11, 21)
)
# now it worked
dataframe.example.scratch

## (4.5) Lists
# serve like archives and can help us to store various objects
list.example<-list(double.vector.example, vector.2)

#show it
list.example

# unlist: if we want to use both elements of the list as a single vector we have to unlist it
unlist(list.example)

## (4.6) Useful Functions to get Informations on Objects
str(dataframe.example) # gives information on datatype and other
summary(dataframe.example) # gives some summary statistics
length(vector.2) # shows the total length of a vector
head(dataframe.example) # shows the first rows of a dataframe
View(dataframe.example) # shows data in a seperate window. only Rstudio!

# ----- (5) Indexing Specific Parts of Data -----
## We use the built-in datasets to show how indexing works
# Mean annual flow of the Nile river in cubic meters per second measure at Ashwan from 1871-1970
nile.flow<-as.vector(Nile)
# check out
nile.flow
# Dataframe on height, age and seed of Loblolly Pine Trees
pinetrees<-Loblolly
#check out
head(pinetrees)

## (5.1) Indexing Vectors
# single value
nile.flow[3]
# Change the value e.g. to correct a transcription
nile.flow[3]<-1300

# several values
nile.flow[c(1,3,5)]
# or if they are in sequence
```

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nile.flow[3:5]

## (5.2) indexing matrices and dataframes
# single value e.g. second row, first column
pinetrees[2,1] # matrix notation: first the row than the column

# several values
pinetrees[2:5,1]

# whole column
pinetrees[,1] #note: nothing means all. Seems strange but is like that

# Excluding a column
pinetrees[,-1] #note: nothing means all. Seems strange but is like that

# whole row
pinetrees[1,]

# if you have a dataframe you can index columns as well with the $ sign
pinetrees$height # will return the whole column as a vector

## (5.3) indexing lists
list.index.example<-list(pinetrees,nile.flow)
# requires two brackets [[]]
list.index.example[[2]] # returns the whole vector

# indexing something inside this vector is straightforward
list.index.example[[2]][3] # just put one index after another

## (5.4) Finding specific values (addresses) and logical subscripts
# finding the maximum and minimum
min(nile.flow)
max(nile.flow)

# finding values within a range with the which function
which(nile.flow>800 & nile.flow<1000) # returns the index of the observation in the vector

# do the same but return the value of the observation
nile.flow[nile.flow>800 & nile.flow<1000] # returns the values of the observation

## use indexing to subset dataframes
# example 1: return all rows where the pinetree seed is 301
pinetrees[pinetrees$Seed==301,]

# example 2: return all rows where the pinetree seed is not 301
pinetrees[pinetrees$Seed!=301,]

# example 3: return only the first column (height) where the pinetree seed is not 301
pinetrees[pinetrees$Seed!=301,1]

# example 4: combining return all rows where the pinetree seed is either 301 or 303
pinetrees[pinetrees$Seed==301 | pinetrees$Seed==303,]

# example 5: return all rows of pinetrees where the Seed is equal 301 and the age is equal to 5
pinetrees[pinetrees$Seed==301 & pinetrees$age==5,]

# (5.6) Side Topic: subsetting dataframes using the subset function
# subset based on two criteria, only returning one column
subset(pinetrees,Seed==301 & age==5, select = height)

# ----- (6) Importing, Exporting -----
## (6.1) CSVs
# write csv
write.table(dataframe.example,"output/dataframe.csv",sep=",")

```

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## read csv
dataframe.example.reimported<-read.table("output/dataframe.example.csv", sep=",",header = TRUE)
# note since the first row of the csv contains the variable names we set header = TRUE
View(dataframe.example.reimported) # have a look

## (6.2) xlsx
# write out some data to excel format
write.xlsx(dataframe.example.scratch,
           "output/dataframe.example.scratch.xlsx")

# import the data again
dataframe.example.scratch.reimported<-read.xlsx("output/dataframe.example.scratch.xlsx",
                                                sheetIndex = 1)

## (6.3) SPSS
# activate the library
library("foreign")

## Angry Birds example
# import the Angry Birds data-set
honesty.lab<-read.spss("input/honesty.lab.sav",to.data.frame = TRUE)

## (6.4) Import and Export R objects with lists
# save a single object
save(dataframe.example.scratch,
     file= "output/dataframe.example.scratch.Rdata")

# reimport it again
load(file = "output/dataframe.example.scratch.Rdata")

## Save several objects (and functions) with lists
# Save everything you need first in a list
my.list<-list(dataframe.example.scratch,
              dataframe.example,
              vector.2)

# export the list with saveRDS
saveRDS(my.list,file="output/my.list.Rdata")

# reimport the list
my.list.reimported<-readRDS("output/my.list.Rdata")

# save all objects from your environment (do this at the end of your session)
save.image("rdata/lookup-script-2016.Rdata")

# ----- (7) Functions -----
## (7.1) Functions operating on vectors
# very simple
mean(dataframe.example$variable.b)

## (7.2) Create custom functions
# create with two parameters
function.example<-function(a,b){
  tmp.data<-mean(a)/mean(b)
  return(tmp.data)}

# usage
function.example(pinetrees$height,pinetrees$age)

## (7.3) Apply functions several times
# on different columns in a matrix/dataframe
apply(dataframe.example, 2 , summary) # example with the summary function

# on different rows
apply(dataframe.example, 1 , mean) # example with the mean

```

```

## (7.4) nested functions (calculates from inside to outside)
sqrt(mean(dataframe.example$variable.a))

# you can go on as you like
round(sqrt(mean(dataframe.example$variable.a)),digits=2)

## (7.5) Side Topic: Using masked functions
# sometimes functions from different packages have the same name so they are masked from the global
# In order to use them you will have to write: package::function e.g.
base::sum(1:15)

# ----- (8) Control Structures -----
## (8.1) loops
# for loops serve for iterations.
for(i in 1:10){print(i+5)}

# Storing loop results
# if we want to store the results of a loop in a new object,
# we will first have to create an empty object to receive the results and then use a clever indexin

# create empty vector
loop.results<-vector()

# loop and store results
for (i in 1:10){loop.results[i]<-i+5}

# show it
loop.results

# General note: avoid loops, if possible because they are slow. have a look at the plyr package for
# other types of loops (e.g. while loops) see e.g. here: https://blog.udemy.com/r-tutorial/

## (8.2) if else statements
# use it to define multiple outcomes of a test
summary(pinetrees)
ifelse(pinetrees$height>50,
      "big",
      "small")

# you can nest it as well to check for multiple conditions
ifelse(pinetrees$height<20,
      "small",
      ifelse(
        pinetrees$height>=20 &
        pinetrees$height<50,
        "medium",
        "big"
      ))
# you can use it to create a new column
pinetrees$height.class<-as.factor(
  ifelse(pinetrees$height<20,
        "small",
        ifelse(
          pinetrees$height>=20 &
          pinetrees$height<50,
          "medium",
          "big"
        ))
  )))

# ----- (9) Plotting -----
# (9.1) Common types of plots
# barplot
barplot(pinetrees$height)
# histogram

```



```

hist(pinetreesh$height)
# boxplot
boxplot(pinetreesh$height)
# scatterplot
plot(pinetreesh$age,
      pinetreesh$height)
# scatterplot with smoothed curve
scatter.smooth(pinetreesh$age,
               pinetreesh$height)

# (9.2) Styling a plot
scatter.smooth(pinetreesh$age,
               pinetreesh$height,
               xlab = "Age of pinetreesh",
               ylab = "Tree height in meters",
               main = "Loblolly Pinetreesh growth and age",
               col = "blue"
              )

# for more information on how to style your plot use the help function of each plot

# (9.3) Plotting with ggplot2 - Example
# NOTE: ggplot two is more complex to learn but later far more powerfull to make sophisticated grap
# easy example: scatterplot with smoothed line
xy.plot.example<-ggplot(data=pinetreesh,
                        aes_string(x="age", y="height"))+ # defines the basics
  geom_point()+ # add a layer with the points
  geom_smooth() # add a layer with the smoothed curve

#plot
xy.plot.example

# styled ggplot
xy.plot.example<-ggplot(data=pinetreesh,
                        aes_string(x="age", y="height"))+ # defines the basics
  geom_point(color="darkgreen", # layer specific styles go into the layer itself
            size=5)+
  geom_smooth()+
  ggtitle("Loblolly Pinetreesh growth and age")+
  ylab("Tree height in meters")+
  xlab("Age of pinetreesh")+
  theme( # general styles relating the plot go into this theme box
    plot.title=element_text(color="red", size=16, vjust=2), # adjust the title
    axis.title.x=element_text(color="blue", size=11, vjust=0.5), # adjust the x-axis title
    axis.title.y=element_text(color="#A86BA7", size=11, vjust=1), # adjust the y-axis title with a
    panel.border = element_rect(colour = "black",fill=NA) # add a boarder to the plot area
  )

#plot
xy.plot.example

# ----- (10) Closing a Session -----
# remove all unneeded objects from your environment e.g. the first created vector from above
rm(result.example) # if you want to remove several objects separete them with commas.

# save your environment
save.image("rdata/lookup-script-2016.Rdata")
# next time you open your project load the data by: load("introduction.data.1.Rdata")
# Insert the load command at the beginning of your script

## exporting your script as a html file for printing
library("highlight")
highlight( "scripts/lookup-script-2016.R",
          output = "scripts/lookup-script-2016.html",

```

Basic R Lookup Script

```
renderer = renderer_html(  
  stylesheet = "/home/jschielein/R/costum_functions/css/costum_1.css") # only needed if  
)  
  
# ----- Additional Remarks -----  
  
## Important packages  
  
# foreign: to import and export several file formats. Comes with basic installation. Just call libr  
# ggplot2: Advanced plotting and mapping tool. Package needs to be installed.  
# plyr: Tools for splitting up datasets and applying functions on several parts of it (automation)  
# and combining the parts again. Package needs to be installed  
# rgdal: to use gdal libraries to work with spatial data. Package needs to be installed  
# raster: most extensive data-set to work with raster data. Package needs to be installed  
# reshape2: For reshaping all kinds of tables e.g. from wide to longformat.  
  
## Other important topics you might consider studying more in depth:  
# merge, union and intersect data  
# rank, sort and order dataframes  
# pattern matching in string with grep
```