R Exercises

1. The goal of this exercise is to get acquainted with different abilities of the R statistical software. It is recommended to use the distributed R tutorial as a guide.

R contains more than 50 datasets and more can be loaded using optional packages. The package VR is depending on the package MASS which contains the dataset survey. This dataset comprises of measurements and answers taken from 237 students of statistics at the university of Adelaide. The following variables are available

Sex gender of student

Wr.Hnd span width in cm (from thumb to pinky) of the writing hand NW.Hnd span width in cm (from thumb to pinky) of the non-writing hand

W.Hnd writing hand

Fold When folding your arms - which one is on top?

Pulse beats per minute

Clap When clapping your hands - which on is on top?

Exer How often do you exercise? Smoke How often do you smoke?

Height body length in cm

M.I Preference of either metric (cm/m) or imperial (feet/inches) units?

Age age in years

> library(MASS) makes the datasets of the MASS package available

PC: Install first the package VR

data() shows a list of all available datasets
 help(survey) gives a description of the dataset survey
 data(survey) makes the dataset survey available

Useful functions to get a first overview of the dataset:

str(survey), summary(survey), table(survey\$Sex), table(survey\$Sex, survey\$Smoke)
The notation survey\$Smoke accesses the variable Smoke in the dataset survey.

> attach(survey) puts the dataset survey on level 2 of the list of available objects. The working directory is on level 1. The variables in the dataset survey can now be accessed directly with their names, i.e. instead of typing survey\$Smoke you may access the variable directly with Smoke.

Dealing with missing values (NA):

> mean(Pulse) result is NA

> mean(Pulse, na.rm=T) the missing values are removed from the calculation of the mean

> na.omit(Pulse) all missing values are removed

> Pulse[!is.na(Pulse)] same as above, but generated by hand

Useful functions for graphics:

```
> hist(Height)
                                                                 histogram
 > boxplot(Height)
                                                                 boxplot
 > boxplot(split(Height, Sex))
                                                                 boxplots of two variables
 > boxplot(Height[Sex=="Female"],Height[Sex=="Male"])
                                                                 boxplots
 > plot(Wr.Hnd,NW.Hnd)
                                                                 scatter plot
 > plot(Sex,Height)
> detach(survey) disconnects the dataset survey from level 2, i.e. variables can no longer by
accessed directly, but only using \$ or [\cdot,\cdot]:
> plot(survey$Wr.Hnd,survey$NW.Hnd) or plot(survey[,2],survey[,3]).
Selecting observations, i.e. only the first 50:
> plot(survey[1:50,2],survey[1:50,3])
```

Do not forget about the online help:

```
> help(survey)
```

> help(plot)

. . .

Now analyse the dataset **survey** using descriptive methods. Therefore produce tables and contingency tables of the categorical variables and calculate location and deviation properties for the continuous variables. Provide suitable graphical representations. Comment on the distributions. Are there any outliers?

Answer the following questions:

- a) Is the span width of the writing hand in general larger than the span width of the non-writing hand?
- **b)** Do the two oldest students smoke?
- c) Which factors might have an influence on the student's pulse?
- d) It is generally believed that the pulse of an individual decreases with increasing age. The function lm fits a linear regression. Investigate the output of the following code:
 - > Agejung <- Age[Age<30]; Pulsejung <- Pulse[Age<30]; plot(Agejung,Pulsejung) Comment on the output. What does the above code do?
 - > lmobj <- lm(Pulsejung \sim Agejung); plot(Agejung,Pulsejung); abline(lmobj)

2. Vectors

What is the output of the following commands? Try to predict the solutions before you type in the commands. We define:

```
commands. We define.
x <- c(5,2,1,4); xx <- c(1,10,15,18); y <- rep(1,5)
z <- c(TRUE,FALSE,TRUE,TRUE); w <- c("Marie","Betty","Peter")
a) sum(x)
    range(x)
    length(y)
    sum(y)
b) c(x,y,13)
c) xx - x
    c(x,12) * y
    1:6 + 1
    1:9 + 1:2</pre>
```

```
d) x <= 2
x <= 2 & z
```

```
e) substring(w,2,4)
   paste(substring(w,1,2),substring(w,5,5),sep="..")
f) cbind(x,xx)
   cbind(2,6:1, rep(c(3,1,4),2), seq(1.1,1.6,by=0.1))
```

3. Sequences of Numbers

Create the following sequences. Use the commands rep and seq.

```
a) 1 2 3 4 5 6 7 8 9
b) "m" "w" "m" "w" "m" "w" "m" "w" "m" "w"
```

- c) 1 2 3 4 1 2 3 4 1 2 3 4
- d) 4 4 4 3 3 3 2 2 2 1 1 1 Hint: Use argument each of the function rep.
- e) 1 2 2 3 3 3 4 4 4 4 5 5 5 5 5
- f) 1 1 3 3 5 5 7 7 9 9 11 11

4. Matrices.

a) Generate the following matrices.

[,1]	[,2]	[,3	3] [,4]						
[1,]	1	101	20	1 3	301						
[2,]	2	102	20	2 3	302						
[3,]	3	103	20	3 3	303						
[4,]	4	104	20)4 3	304						
[5,]	5	105	20	5 3	305						
	[,1]] [,2	2] [,3]	[,4]	[,5]	[,6]	[,7]	[,8]	[,9]	[,10]
[1,]	ļ	5	0	0	0	0	0	0	0	0	0
[2,]	(0	5	0	0	0	0	0	0	0	0
[3,]	(0	0	5	0	0	0	0	0	0	0
[4,]	(0	0	0	5	0	0	0	0	0	0
[5,]	(0	0	0	0	5	0	0	0	0	0
[6,]	(0	0	0	0	0	5	0	0	0	0
[7,]	(0	0	0	0	0	0	5	0	0	0
[8,]	(0	0	0	0	0	0	0	5	0	0
[9,]	(0	0	0	0	0	0	0	0	5	0
[10,]	(0	0	0	0	0	0	0	0	0	5

b) Explore the properties of your generated objects. Which class of R-objects do they belong to? How are they structured?

Hint: class(), dim(), str(), summary().

5. Lapwings

For various meadows at Zurich Airport we counted the daily number of lapwings on several occasions. For every bird we noted the kind of activity (resting, feeding, flying) as well as the ground conditions (damp, dry, wet). The data was stored in vogel.dat:

	Datum	Zeit	${\tt Feld.Nr}$	Anzahl	Taetigkeit	Boden
1	910903	10.06	1411	22	ru	t
2	910903	10.07	1413	15	ru	t
3	910906	10.01	1411	29	fr	t
4	910906	10.03	1413	44	fr	t
5	910910	15.19	1410	34	ru	n
6	910911	10.00	1413	41	fr	f
7	910912	12.38	1411	10	ru	t
8	911014	15.17	1409	2	fl	t
9	911203	13.05	1413	2	fl	t

It contains the columns date (Datum), time (Zeit), id of meadow (Feld.Nr), count (Anzahl), activity (Taetigkeit) (ru resting, fr feeding, fl flying) and ground condition (n wet, t dry, f damp).

- a) To read the data into R type: d.vogel <- read.table("http://stat.ethz.ch/Teaching/Datasets/NDK/vogel.dat", sep =";", header=TRUE)
- b) Create a new data frame that only contains the meadow id and the counts. How many birds were counted on average?
- c) Create a data frame only with the data of meadow 1413.
- d) Create a vector that contains the number of birds of meadow 1413.
- e) On how many occasions(days) did we observe feeding birds? How many birds were counted while feeding? What were the corresponding observation numbers?
- f) In the data frame change the number of lapwings of the eighth observation (row) to 6. Delete the third and seventh observation from the data frame.

Hint: mean(), sum(), which().

6. Meteo

The data set meteo70.txt contains several measurements of weather variables between 1994 and 2007. The mean daily air temperature is stored in the variable X211.

a) Read the data into R:

fname <- "http://stat.ethz.ch/education/semesters/ss2014/regression/uebungen/meteo70.txt"
d.meteo <- read.table(fname, header=T)</pre>

The missing values are encoded by the value 32767. Change this value to NA and rename the variable X211 in temp.

b) Calculate the mean of temp separately for each day of the week.

Hint: aggregate()

c) Plot the mean of temp for each year. Why is the mean in 2007 so high?

7. Getting to know Data: Iris blossoms

The data set iris contains measurements of the length and the width (in cm) of petals and sepals of three iris species: 1: Setosa, 2: Versicolor and 3: Virginica.

(Source: R. A. Fisher, *The Use of Multiple Measurements in Taxonomic Problems*, Annals of Eugenics, Vol. 7, Part II, 1936, pp. 179-188.)

- a) This data set iris is already part of the standard R installation. Consider the object iris. How is it structured? How many observations (lines) does it contain? How many variables (columns)? Hint: nrow(), ncol(), dim(), str()
- b) To get an overview of the range of values, look at the summary() of the data set. Which information on the data set does it provide?
- c) For the variable Sepal.Length check the results above by using the R-functions min(), max(), mean(), median(), quantile(). If necessary, make use of the help functions ?quantile etc.

8. Missing Values

Statistics needs data. Unfortunately, data often cannot be collected fully. Therefore many data sets contain "gaps", non-existing measurements, so-called NAs (not available). In this exercise you will get to know how R deals with NAs. We work with the data set iris. Make a copy of the iris data set by d.iris <- iris.

a) Assume that we were unable to take the second observation of Petal.Length and Petal.Width, and for the fifth observation, the data for Sepal.Length, Sepal.Width and Petal.Width are missing. Replace these five fields by NA.

Hint: Replace the values by NAs using e.g. d.iris[2, 3:4] <- NA

- b) Consider the first eight observations of the modified data set, to observe how the NAs are displayed by R. The commands class(), nrow(),ncol(), dim(), str() also work for the data set with missing values. What changes in the summary()?
- c) Try to confirm the given values for the variable Sepal.Length using min(), max(), mean(), median(), quantile(). Is there a difference?
- d) There are functions that cannot handle NAs (Result 'NA' or 'Error: missing observations'). There is a trick to make them calculate the correct results: simple functions such as min(), max(), mean(), median(), quantile(), range() etc. can take an argument na.rm. When you set its value to TRUE, the NAs will not be considered in the calculation.
 Try to confirm the values provided by summary() again, using this new argument.
- e) Why should missing values always be coded by NA, and not, for instance, filled with a zero? Explain for the case of the mean() function.
- f) Experiment with missing values in the statistical functions var(), sd(), cor(). Can you explain the behaviour of R?
- g) Select only those observations which have missing values in either Sepal.Length or in Petal.Length.
 Hint: is.na()
- h) The function na.omit() eliminates all observations from the data frame for which any(!) variable contains NAs. Save the result of na.omit(d.iris). How many observations remain? How many remain using na.omit(d.iris[,1:3])?

Notes

Higher-level functions such as t.test() or wilcox.test() have an argument na.action, with which the reaction to NAs can be determined. na.action=na.omit first deletes all lines (observations) with NAs before anything is calculated.