

```

#####
## L1.2. INSTALL R PACKAGES
#####
install.packages("rgl")
## if does not work:
##   a) Select all repositories in "packages" menu
##   b) if still does not work - use Bioconductor installation
## source("http://bioconductor.org/biocLite.R")
## biocLite("your package")

#####
## L1.3. R INTERFACE
#####
##-----#
## L1.3.1. Typing commands
##-----#
2*2
2^10
sqrt(16)
16^(1/2)

##-----#
## L1.3.2. Calling functions
##-----#
log(100)
log(100, base=10)
log(100, b=10)
log(100, 10)

##-----#
## L1.3.3. Embedded help
##-----#
help("sqrt") # help on "sqrt" function
?sqrt # ...the same...
?round
??round # fuzzy search for "round" in all help topics
apropos("plot") # propose commands with the word "plot" inside name

## Demos
demo() # show available demos
demo("image") # start demo "image"
demo(persp)
demo(plotmath)

#####
## L1.4. VARIABLES and BASIC OPERATIONS
#####

##-----#
## L1.4.1. Variables
##-----#
x = 2
x

```


A

```

A=A-1 ... # add scalar
A

A=A+a ... # add vector
A

t(A) ... # transpose

B=A+t(A) # add a transposed matrix
B

B*B ... # by-element product

B%*%B ... # matrix product

cbind(c(1,2,3,4),c(10,20,30,40))

rbind(c(1,2,3,4),c(10,20,30,40))
##-----
## Data frame
Data = data.frame(A) ... # alternatively:
Data → → → ... # Data=data.frame(matrix(nr=5,nc=5))

## let us add a column to Data
mice = sprintf("Mouse_%d",1:5)
Data = cbind(mice,Data)

## put the names to the variables
names(Data) = c("name","sex","weight","age","survival","code")
Data

## put in the data manually
Data$name=sprintf("Mouse_%d",1:5)
Data$sex=c("Male","Female","Female","Male","Male")
Data$weight=c(21,17,20,22,19)
Data$age=c(160,131,149,187,141)
Data$survival=c(T,F,T,F,T)
Data$code = 1:nrow(Data)
Data

## visualize data as a table
fix(Data)

## see the structure of the objects
str(Data)

## see the head of the objects
head(Data)

## summary on the data
summary(Data)

##-----
## Factors

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## Let's use factors
Data$sex = factor(Data$sex)
summary(Data)

## usefull commands when working with factors:
levels(Data$sex) # returns levels of the factor
nlevels(Data$sex) # returns number of levels
as.character(Data$sex) # transform into strings

# -----
## L1.4.6. Lists
# -----


L=list()
L$Data=Data
L$descr = "A fake experiment with virtual mice"
L$num = nrow(Data)
str(L)

## how to access the fields? Simple!
L$Data
L$"Data"
L$num
## or
L[[1]]
L[[3]]


## clear all
ls()
rm(list=ls())
ls()


#####
# L1.5. DATA IMPORT AND EXPORT
#####

# -----
## L1.5.1. Current folder
# -----


getwd() ## shows current folder

dir() ## shows files in the current folder

setwd("D:/Data/ABS2016") ## sets folder

# -----
## L1.5.2. Scan - reads arbitrary data
# -----


## File from Internet / disk
SomeData = scan("http://edu.sablab.net/data/txt/currency.txt",
                what = character(0))
SomeData

```

```

## · HTML · from · Internet
Google = scan("http://google.com", what = character(0))
Google

## -----
## L1.5.3. · Read · table · (from · Internet · or · local · folder)
## -----
Currency =read.table("http://edu.sablab.net/data/txt/currency.txt",
                      header=T, sep="\t")
str(Currency)

## · let's · ask · to · do · not · transfere · strings · to · factors
Currency =read.table("http://edu.sablab.net/data/txt/currency.txt",
                      header=T, sep="\t", as.is=T)
str(Currency)
head(Currency)
summary(Currency)
fix(Currency)
## · first · plot · :
plot(Currency$EUR)

## -----
## L1.5.4. · "GE" · a · big · dataset: · use · "download.file" · and · "load"
## -----
download.file("http://edu.sablab.net/data/all.Rdata",
              destfile="all.Rdata", mode = "wb")
## · check · the · current · folder · for · ".Rdata" · file
getwd() # show current folder
dir(pattern=".Rdata") # show files in the current folder
load("all.RData") # load the data
ls()
str(GE.matrix)
## · see · the · annotation · for · dimentions
attr(GE.matrix, "dimnames")
rownames(GE.matrix)
colnames(GE.matrix)

## -----
## L1.5.5. · Data · export
## -----
write.table(Currency, file = "curr.txt", sep = "\t",
            eol = "\n", na = "NA", dec = ".",
            row.names = FALSE, quote=FALSE)

save(Currency, file="Currency.Rdata")
save(list=ls(), file="workspace.Rdata")

getwd()
dir()
## · if · you · need · to · set · working · folder, · use
setwd("...put here desired path...")

## · clear · all
rm(list=ls())

#>>>>>>>>>>>>>>>>>>>>>>>
```



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## Custom functions
##-----

## Let us write a function to print vectors
printVector = function(x, name=""){
  print(paste("Vector", name, "with", length(x), "elements:"))
  if (length(x)>0)
    for (i in 1:length(x))
      print(paste(name, "[", i, "] =", as.character(x[i])))
}

printVector(Shop$Payment, "Payment")

##-----
## Run script, saved in other files
##-----


source("http://sablab.net/scripts/getFiles.r")

ls()

#####
# L1.7. DATA VISUALIZATION
#####

##-----
## L1.7.1. Plot time-series and smooth
##-----
## get data
Currency =read.table("http://edu.sablab.net/data/txt/currency.txt",
                      header=T, as.is=T)

## initiate window
x11(8,5) # try x11()
## plot the currency behaviour for the last 10 years
plot(Currency$EUR, pch=19, col = "#0000FF11", cex=2)

## let's make it more beautiful
x11(8,5)
?par
plot(Currency$EUR, col="#00FF0033", pch=19,
      main="EUR/USD ratio for 11 years",
      ylab="EUR/USD",
      xlab="Measures (working days)")

## add smoothing. Try different "f"
smooth = lowess(Currency$EUR, f=0.1)
lines(smooth,col="red",lwd=2)
## add 1 level
abline(h=1,col="blue",lty=2)

## (*) add years
year=1999 # an initial year
while (year<=2009){ # loop for all the years up to now

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# take the indexes of the measures for the "year"
idx=grep(paste("^", year, sep=""), Currency$Date)
# calculate the average ratio for the "year"
average=mean(Currency$EUR[idx])
# draw the year separator
abline(v=min(idx), col=1, lty=3)
# draw the average ratio for the "year"
lines(x=c(min(idx), max(idx)), y=c(average, average), col=2)
# write the years
text(median(idx), max(Currency$EUR), sprintf("%d", year), font=2)
# write the average ratio
text(median(idx), average+0.05, sprintf("%.2f", average), col=2,
      font=2, cex=0.8)
year=year+1;
}

```

##-----
L1.7.2. Mouse phenom :)

```

##-----  
## load data
Mice=read.table("http://edu.sablab.net/data/txt/mice.txt",
                 header=T, sep="\t")
str(Mice)

```

```

## initiate window
x11(10,8)

```

```

## plot a factorial data
plot(Mice$Strain, las=2,
      col=rainbow(nlevels(Mice$Strain)), cex.names =0.7)
title("Number of mice from each strain")

```

```

## plot a factorial data as pie
pie(summary(Mice$Sex), col=c("pink", "lightblue"))
title("Gender composition (f:female, m:male)")

```

```

## try to use special command "barplot" as well
## a histogram
hist(Mice$Starting.weight, probability = T,
      main="Histogram and p.d.f. approximation",
      xlab="weight, g")
lines(density(Mice$Starting.weight, width=0.5), lwd=2, col=4)

```

```

## (!) a box-plot of the population on the basis of sex
boxplot(Starting.weight~Sex, data=Mice, col=c("pink", "lightblue"))
title("Weight by sex (f:female, b:male)",
      ylab="weight, g", xlab="sex")

```

##-----
L1.7.3. Show all data frame at once
##-----

```
plot(Mice)
```

```

plot(Mice[, -(1:3)])  

##-----  

## L1.7.4. 3D visualization and custom functions  

##-----  

## see demo  

demo(persp)  

## use RGL library  

library(rgl)  

  

x=Mice$Starting.weight  

y=Mice$Ending.weight  

z=Mice$Fat.weight  

plot3d(x,y,z)  

  

## make it more beautiful  

color = as.integer(Mice$Sex)*2  

plot3d(x,y,z,  

       col=color,type="s",radius=1,  

       xlab="Starting weight",  

       ylab="Ending weight",  

       zlab="Fat weight")  

  

#>>>>>>>>>>>>>>>>>>>  

#> please, do Tasks L1.7ab  

#>>>>>>>>>>>>>>>>>>>  

  

#####
#####  

# L1.8. DESCRIPTIVE STATISTICS  

#####  

## clear memory  

rm(list = ls())  

  

## load data  

Mice=read.table("http://edu.sablab.net/data/txt/mice.txt",  

                 header=T,sep="\t")  

str(Mice)  

##-----  

## L1.8.1. Measures of center  

##-----  

summary(Mice)  

  

## mean and median  

mean(Mice$Ending.weight)  

median(Mice$Ending.weight)  

## for mode you should add a library:  

library(modeest)  

mlv(Mice$Ending.weight, method = "shorth")$M  

  

## mean and median if NA values present: add na.rm=T

```

```

mn = mean(Mice$Bleeding.time, na.rm=T)
md = median(Mice$Bleeding.time, na.rm=T)
mo = mlv(Mice$Bleeding.time, method = "shorth", na.rm=T)$M

## let us plot them
x11()
plot(density(Mice$Bleeding.time, na.rm=T), xlim=c(0,200), lwd=2,
      main="Bleeding time")
abline(v = mn, col="red")
abline(v = md, col="blue")
abline(v = mo, col="cyan")
legend(x="topright", c("mean", "median", "mode"),
       col=c("red", "blue", "cyan"), pch=19)

prop.f = sum(Mice$Sex=="f")/nrow(Mice)

#-----
## L1.8.2. Measures of variation
##-----

## quantiles, percentiles and quartiles
quantile(Mice$Bleeding.time, prob=c(0.25,0.5,0.75), na.rm=T)

## standard deviation and variance
sd(Mice$Bleeding.time, na.rm=T)
var(Mice$Bleeding.time, na.rm=T)

## stable measure of variation -- MAD
mad(Mice$Bleeding.time, na.rm=T)
mad(Mice$Bleeding.time, constant = 1, na.rm=T)

#-----
## L1.8.3. Measures of dependency
##-----


## covariation
cov(Mice$Starting.weight, Mice$Ending.weight)

## correlation
cor(Mice$Starting.weight, Mice$Ending.weight)

## coefficient of determination, R2
cor(Mice$Starting.weight, Mice$Ending.weight)^2

## kendal correlation
cor(Mice$Starting.weight, Mice$Ending.weight, method="kendall")
## spearman correlation
cor(Mice$Starting.weight, Mice$Ending.weight, method="spearman")

#>>>>>>>>>>>>>>>>>>>>
#> please, do Task L1.8
#>>>>>>>>>>>>>>>>>>>>

#####
# L1.9. DETECTION OF OUTLIERS
#####

```

```

## clear memory
rm(list = ls())

## load data

#Mice=read.table("d:/data/abs2016/data.txt/mice.txt",header=T,sep="\t")

Mice=read.table("http://edu.sablab.net/data/txt/mice.txt",
                header=T, sep="\t")
str(Mice)

x = Mice$Weight.change
# -----
## L1.9.1. z-score
# -----
x11()
z = scale(x)
plot(z,pch=19,col=4,main="z")

plot(abs(z),pch=19,col=4,main="|z|")
abline(h=3,col=2)

Mice$Weight.change[abs(z)>3]

# -----
## L1.9.2. Iglewicz-Hoaglin method
# -----
x11()
zih=(x-median(x))/mad(x)
#zih=(x-median(x))/median(abs(x-median(x)))
plot(abs(zih),pch=19,col=4,
     main="|z| by Iglewicz-Hoaglin")
abline(h=3.5,col=2)
# -----
## L1.9.3. Grubb's method
# -----
library(outliers)
x1=x
while(grubbs.test(x1)$p.value < 0.05) {
  print(outlier(x1))
  x1[x1==outlier(x1)]=NA
}
x11()
plot(abs(x-mean(x1,na.rm=T))/sd(x1,na.rm=T),pch=19,col="red",
     main="Outliers by Grubb's Test")
points(abs(x1-mean(x1,na.rm=T))/sd(x1,na.rm=T),pch=19,col="blue")

#>>>>>>>>>>>>>>>>>>>>
#> please, do Task L1.9
#>>>>>>>>>>>>>>>>>>>
```