

```
#####
## 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  
  
y <- 3  
y  
  
x + y -> z  
z  
  
## Variables are case-sensitive  
Z  
  
## Another way to show the data  
print(z)  
cat("x=", x, ", y=", y, ", z=", z, "\n")  
  
## show variables in memory  
ls()  
## remove all variables from memory  
rm(list=ls())  
ls()  
  
##-----  
## L1.4.2. Scalar types of data  
##-----  
  
##-----  
## Numeric (integer, double)  
  
i=5  
i  
i*2  
i/2  
i%/%2 # integer division  
i%%2 # remainder of integer division  
round(1.5)  
  
## (*) for bitwise operation install and use  
## "bitops" package and bitAnd, bitOr, ...  
  
## Double  
r=1.5  
r  
  
l=pi*2*r # let us calculate the circumference for circle with r  
l
```



```
## -----
## NA - Not-Available (missing data)
na = NA
na + 1
100>na
na==na
is.na(na)

## Inf - Infinity (+/- infinite data)
1/0
-1/0
is.infinite(1/0)
is.finite(1/0)

## NaN - Not-A-Number
0/0
0*1/0
is.nan(sqrt(-1))

## -----
## L1.4.4. Vectors
## -----
```

```
## Vector creation
a = c(1, 2, 3, 4, 5)

a
a[1]+a[4]

b=5:9
a+b # (*) try b=5:10. Can you explain the effect?
→ #(ans: "t!tfihs ralucric" :)
```

```
seq(from=1,to=10,by=0.5) #sequence
seq(1,10,0.5)

rep(1:4, 2) # same as rep(1:4, times=2)

rep(1:4, each=2) # not the same

txt = c(st, "Let's try vectors", "bla-bla-bla")
txt

boo = c(T,F,T,F,T)
boo

##!!!!!!!!!!!!!!!
## Extremely important !!
##!!!!!!!!!!!!!!!
```

```
## Vector indexes
```



```
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

## 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("E:/DOCUMENTS/Pedagogics/R-Course_2010/Data") ## 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")

##-----
## 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())

#>>>>>>>>>>>>>>>>>>>
#> please, do Tasks L1.5a, L1.5b
#>>>>>>>>>>>>>>>>>>
idx=Shop$Payment=="Visa"
write.table(Shop[idx,], file = "shop.txt", sep = "\t",
            row.names = FALSE, quote=FALSE)

#####
## L1.6. CONTROL WORKFLOW and CUSTOM FUNCTIONS
#####
Shop = read.table("http://edu.sablab.net/data/txt/shop.txt",
                  header=T, sep="\t")

a=1
b=2

##-----
## IF condition

if (a==b) {

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```
print("a equals to b")
} else {
  print("a is not equal to b")
}

## use if in-a-line
ifelse(a>b, a, b)

##-----
## FOR loop

## print all information for the first client
for (i in 1:ncol(Shop))
  print(Shop[1,i])

##-----
## WHILE loop

## print all information for the first client
i=1;
while (i <= ncol(Shop)){
  print(Shop[1,i])
  i=i+1
}

##-----
## REPEAT loop
i=1
repeat {
  print(i)
  i=i+1
  if (i>10) break
}
## "break" and "next" - help to control flow

##-----
## 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)  
  
## let's make it more beautiful  
x11(8,5)  
?par  
plot(Currency$EUR,col="#00FF00",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=2,lwd=2)  
## add 1 level  
abline(h=1,col=4,lty=2)  
  
## (*) add years  
year=1999 # an initial year  
while (year<=2009){ # loop for all the years up to now  
# 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
windows(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), 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, m: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="kendal")
```

```

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

x=Mice$Weight.change
##-----
## L1.9.1. z-score
##-----
x11()
plot(abs(scale(x)),pch=19,col=4,main="| z |")
abline(h=3,col=2)

Mice$Weight.change[abs(scale(x))>3,]

##-----
## L1.9.2. Iglewicz-Hoaglin method
##-----
x11()
plot(abs(x-median(x))/mad(x),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) {
  x1[x1==outlier(x1)]=NA
}
x11()
plot(abs(x-mean(x1,na.rm=T))/sd(x1,na.rm=T),pch=19,col=2,
     main="Outliers by Grubb's Test")
points(abs(x1-mean(x1,na.rm=T))/sd(x1,na.rm=T),pch=19,col=4)

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

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